

ANTOLOGIA PALEONTOLOGICA
TOMO 4
ISMAEL FERRUSQUIA VILLAFRANCA

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FERRUSQUIA, V. I. COAD.
ANTOLOGIA PALEONTOLOGICA

SOCIEDAD MEXICANA DE PALEONTOLOGIA, A. C.

**ANTOLOGIA PALEONTOLOGICA - LECTURAS SELECTAS
COMENTADAS SOBRE CONCEPTOS RECIENTES DE
PALEOBIOLOGIA.**

ISMAEL FERRRUSQUIA VILLAFRANCA, COMP.

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Estimado Dr. Ferrusquia-Villafranca:

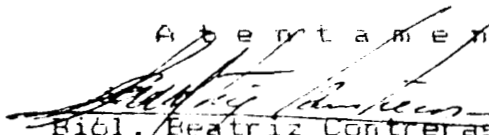
Me es grato saludarlo y hacer de su conocimiento, que el trabajo titulado:

"Antología Paleontológica-Lecturas Selectas Comentadas Sobre Conceptos Recientes de Paleobiología".

En fecha reciente se sometió a la consideración de la Sociedad Mexicana de Paleontología para su publicación, se ha recibido el dictamen aprobatorio del Comité Editorial, quien ha recomendado por el tamaño de la obra (702 cuartillas), se le presente como una "Publicación Especial", ya que no podría incluirse en la Revista; relacionado con lo anterior, cabe destacar que el costo de la publicación excede la capacidad económica actual de la SOMEXPAL, por lo que ya se busca la implementación de mecanismos de financiamiento adicional, que permita a la brevedad posible publicar su interesante trabajo, que estamos seguros será de gran utilidad en la docencia y en la investigación paleontológica y de disciplinas científicas afines.

Sin otro particular, al despedirme hago propicia la ocasión para expresarle mi consideración y aprecio.

Atentamente,


Biól. Beatriz Contreras y Montero
Presidenta

PROLOGO

La formación de recursos humanos de alto nivel, es la labor prioritaria de un país, ya que su riqueza y grandeza son en gran medida una función del número y calidad de gente preparada que posee.

En el caso de México esta actividad es todavía de mayor importancia, por la obligación que se tiene de superar el enorme rezago que muestra, con respecto a otros países, en muchos de los campos de la cultura y la economía, así como la imperiosa necesidad de elevar en términos reales el nivel de vida de todos los mexicanos, apoyándose, claro está, en un desarrollo integral, armonioso y plenamente adecuado a la naturaleza del territorio nacional y a la idiosincracia de su población.

En este sentido resulta muy acertado el esfuerzo que realiza la UAM-Iztapalapa, para alentar a su personal académico a la preparación de textos y obras de índole diversa, que sean coadyugantes eficaces en la docencia a nivel profesional y de posgrado y que subsanen, aunque sea sólo en forma parcial, la necesidad que existe de estas obras en el país.

La presente **ANTOLOGIA PALEONTOLOGICA**, se inserta como una contribución en este noble esfuerzo universitario. El conjunto de trabajos comentados que la integran reflejan los avances recientes en torno al origen y evolución de los seres vivos y su diversificación cronoespacial que los llevó a ocupar todos los ámbitos disponibles integrando así el componente más dinámico del Sistema Tierra, la Biósfera.

La antología pretende ser un vehículo hacia una mejor comprensión de este sistema al que pertenecemos, lo cual sin duda, nos permitiría enfrentar el reto y responsabilidad de conservarlo. En ello va en juego no sólo la supervivencia del hombre como una bioespecie más, sino su misión de hacerlo de una nueva era y una nueva forma de vivir.

La preparación de la obra en si se inició en 1991, cuando el suscrito, investigador del Instituto de Geología, UNAM, y profe-

sor del Departamento de Biología, División CBS, UAM-I, realizó una estancia sabática como docente en la Facultad de Ciencias Biológicas de la Universidad Autónoma del Estado de Morelos, con objeto de proporcionar apoyo al Área disciplinaria de Paleontología, y entre otras cosas, se le encomendó la responsabilidad de preparar una Antología Paleontológica que complementara en lo posible, la formación académica de los estudiantes, poniendo a su alcance, literatura científica especializada y relevante, organizada de manera temática y comentada, para facilitar su comprensión. Esta obra subsanaría parcialmente el gran vacío que al respecto acusa nuestro país, haciendo posible el acceso de material bibliográfico a usuarios potenciales, que difícilmente podrían estar en contacto con él.

La aceptación de tal responsabilidad se hizo de buen grado, ya que como paleontólogo profesional y docente en el área, se tenía plena conciencia de esta necesidad. La realización de la obra constituyó un reto y una experiencia enriquecedora. Pero también implicó un esfuerzo y un tiempo mucho mayores de los considerados, ya que se analizaron unas 5000 referencias bibliográficas, consultándose casi 900 trabajos diversos, seleccionándose de manera preliminar unos 200 e incluyéndose finalmente sólo 60; aun así, la obra consta de 700 cuartillas. El trabajo académico principal se desarrolló en las bibliotecas de los Institutos de Geología (UNAM) y Mexicano del Petróleo, en México, así como en las del Department of Geological Sciences, The University of Texas-Austin, y el Natural History Museum, Los Angeles, California, en Estados Unidos. No sólo se examinó la literatura, sino que se duplicaron o adquirieron los trabajos más relevantes, muchos de ellos en ese país, por no estar disponibles en México. Posteriormente, la duplicación xerográfica definitiva, la redacción de la obra y la preparación de la bibliografía respectiva, se realizó en el Instituto de Geología.

Con la elaboración de la versión xerográfica de la obra, concluyó la responsabilidad del suscrito ante la Universidad Autónoma del Estado de Morelos, cuyas autoridades ante la imposibilidad de publicarlo, dejaron al autor en libertad de buscar los mecanismos para hacerlo, ya que a todas luces los resultados del esfuerzo realizado, merecían ser compartidos con la comunidad académica y quedar a disposición de los estudiantes y maestros en las áreas de Biología y de Geología. De la misma opinión fueron los colegas a quienes se les mostró el trabajo, incluidas las autorida-

des de la División de Ciencias Biológicas y de la salud, UAM-I, quienes recomendaron presentarla en el CONCURSO DE LA ELABORACION DE LIBROS DE TEXTO Y MATERIAL DIDACTICO DE APOYO A LA DOCENCIA que ofrece esta división cuya convocatoria se publicó el 22 de marzo de 1993.

La adecuación de la obra a los requisitos y lineamientos de la convocatoria, conllevó un esfuerzo adicional enorme consistente en su transcripción completa a un formato de de procesador de palabras. Al efecto el suscrito solicitó y recibió el entusiasta apoyo de sus alumnos de biogeografía, Trimestre 93I, quienes lo ejecutaron en las instalaciones de la UAM-I.

AGRADECIMIENTOS

Una obra como la presente, implica la colaboración y el apoyo de diversas instituciones y personas; a todas ellas el autor les queda muy reconocido. En la fase inicial, la Universidad Nacional Autónoma de México, al través de las instancias del caso -Instituto de Geología y H. Consejo Técnico de la Investigación Científica- autorizó la estancia sabática; la Facultad de Ciencias Biológicas y la Dirección de Investigación y Postgrado de la Universidad Autónoma del Estado de Morelos, amablemente extendieron la invitación para realizar la estancia, ejecutaron las gestiones académico-administrativas correspondientes, y durante ella, tuvieron con el suscrito únicamente amabilidades, brindándole todo su apoyo; por ello, este se encuentra muy agradecido, especialmente con los profesores Biól. Luciano Vargas Mendoza y M. en C. Gerardo Avila García, Directores respectivos de esas instituciones. La subsecretaría de Educación Superior e Investigación Científica de la SEP, y el Consejo Nacional de Ciencia y Tecnología, proporcionaron el apoyo económico que posibilitó la estancia.

El trabajo académico conducente a la preparación de la presente Antología, se benefició ampliamente con las discusiones sostenidas con los Doctores James Sprinkle, Keith Young, Jhon A. Wilson y Ernst L. Lundelius, de la University of Texas-Austin; Edward C. Wilson y Craig C. Black, del Natural History Museum, Los Angeles; y Robert Douglas, de la University of Southern, California, quienes compartieron sus experiencias y conocimientos, así como permitieron el acceso a sus bibliotecas particulares, ofreciendo además valiosas sugerencias.

ADVERTENCIA

Aunque la presente Antología se encuentra terminada y se dispone de prototipos, su duplicación xerográfica preliminar múltiple, y desde luego, su eventual publicación, requieren la consecución de los permisos pertinentes, que deberán otorgar en su caso, los tenedores de los derechos respectivos -copy right-. Por tanto, deberán realizarse las gestiones necesarias.

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8.5 (a)

The earliest known ants: an analysis of the Cretaceous species and inference concerning their social organization. *

E. O. Wilson

Abstract. The known Cretaceous formicoids are better interpreted from morphological evidence as forming single subfamily, the Sphecomyrminae, and even a single genus, Sphecomyrma, rather than multiple families and genera. The females appear to have been differentiated as queen and worker castes belonging to same colonial species instead of winged and wingless solitary females belonging to different species. The former conclusion is supported by the fact that the abdomens of workers of modern ant species and extinct Miocene ant species are smaller relative to the rest of the body than is case for modern wingless solitary wasps. The wingless Cretaceous formicoids conform to the proportions of ant workers rather than to those of wasps (Figs. 1-2) and hence are reasonably interpreted to have lived in colonies.

The Cretaceous formicoids are nevertheless anatomically primitive with reference to modern ants and share some key traits nonsocial aculeate wasps. They were distributed widely over Laurasia and appear to have been much less abundant than modern ants.

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Introduction

This article attempts to resolve a problem in systematics that bears significantly on the origin and early evolution of the ants and hence the antiquity of advanced social behavior in insects

generally. Studies of four mid-Cretaceous amber ant subfamily Sphecomyrminae across a wide portion of present-day North America: Sphecomyrma freyi from New Jersey (Wilson et al. 1967 a,b) and S. canadensis from Alberta (Wilson 1985a) Dlussky (1975,1983) has described an important additional collection of ant like forms from the Upper Cretaceous of the Taymyr Peninsula (extreme north-central Siberia), southern Kazakh S.S.R., and the Magadan region of extreme eastern Siberia. He erected 10 new genera to accommodate this material. In his more recent article, he also created a new family, the Armaniidae, to accommodate some of the genera while elevating the Sphecomyrminae to family rank (hence Sphecomyrmidae) to receive others. This classification is summarized in Table 1.

Dlussky's taxonomic interpretation, which is based careful and descriptions of the new material, has sweeping consequences for our conception of the ants. First, it presents a picture of an extensive radiation of ants-like forms by the early part of the Upper Cretaceous- in other words, by no later than 80 ma B.P. Second, Dlussky suggested that the Sphecomyrmidae are not true ants or even precursors of the Formicidae but a closely related side branch, while the Armaniidae are the true ancestors of the ants. He doubted are the true ancestors of the ants. He doubted that either the Sphecomyrmidae or the were eusocial; in other words he questioned whether they possessed the most advanced mode of social organization in which distinct queen and worker castes from overlapping adult generations and care for the developing young (see Wilson 1971).

Because of the light that the early fossil can shed on the origin of the ants and their distinctive social systems, which are matters hitherto largely unexplored due to the exclusively eusocial status of modern ant species, I decided to reexamine closely the morphological and biogeographic evidence provided by the Cretaceous material. I have arrived at a wholly different conclusion from that of Dlussky. In essence, the differences among the fossil cannot support the separation of the two new families, the Sphecomyrmidae and Armaniidae, from the Formicidae. It is difficult to justify even the recognition of any genus other

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than *Sphecomyrma* on the basis of the morphological evidence. The most parsimonious explanation of the data is that the winged females and males from the Soviet deposits are queen and males of eusocial colonies of which the wingless *Sphecomyrma* and wingless Soviet species are the workers.

In proposing this view, I wish to emphasize something earlier by Dr. Dlussky and myself. There is only one exception, the supposed divided condition of the hind trochanter in the Armaniidae, to be discussed later. Instead, the disparity is due to a difference of interpretation of the facts with reference to higher classification and phylogeny. This is all to the good. By such contrasts the basic issues can be better clarified and the gathering of new data stimulated.

Materials and Methods

All of the characters known to vary among the Cretaceous species were first broken into character states. Each genus and species was then redescribed character by character in matrix form to allow close comparison of taxa. The holotypes of two of the best preserved and taxonomically most important Soviet species, *Armania robusta* and *Pseudarmanis rasnitsyni*, were studied closely in comparison with the descriptions. I also worked with all of the four known specimens of the North American formicoids, placed in *Sphecomyrma* (Wilson 1985a).

In addition, measurements were of a wide range of contemporary queen and worker ants and the winged and wingless females of other, nonsocial aculeate families deposited in the Museum of Comparative Zoology, in order to detect consistent differences in body proportions between social and nonsocial females. A single representative specimen was taken from the series available in each species, and the species in turn were selected to provide a large amount of phyletic diversity. This information was needed to infer the level of social evolution of the earliest fossil. Further measurements were utilized from the monograph of Miocene ants of North America by Carpenter (1930). All of these fossils belong to extinct species but surviving subfamilies and in some cases surviving genera. They were considered useful because as rock fossils their

shapes were likely to have been distorted in the same manner as some of the Soviet specimens, which had been similarly preserved.

In particular, it was important to take into account the possible increase in length of the abdomen, which tends to occur when this softest of body parts is crushed laterally or dorsoventrally.

Head length was adopted as a reliable as an index of body size as a whole (see Wilson 1971). The size of the abdomen is critical as an indicator of reproductive as apposed to nonreproductive status in females, because it is the main part of the body containing the ovaries, organs that are proportionately large in full social hymenopterans.

The abdomen in turn was defined in two ways for the present study. First, a "functional" definition marks the abdomen as the posteriormost discrete body part, commonly called the gaster by ant specialists—the ovary-bearing portion behind the one or two segments of the waist. In addition, a strictly homologous definition of the abdomen (or, more precisely, posteriormost major body tagma) was used: all of the true abdominal segments of the waist in ants and a few aculeate wasps. The first, functional definition is intuitively the better because the waist has by definition been reduced to a relatively thin, largely muscled portion that increases the mobility of in order to evaluate the situations as fully as possible.

The results were evaluated with reference to the following criteria derived from standard systematic practice on mode faunas of insects and other animals. Individual taxa, whether species, genera, or representative of higher taxa, should be distinguished and made only if they differ by character states. The states can be relatively minor in the case of species, but should be more substantial in the case of genera (e.g., in ants they include the number of antennal and palpal segments, presence or absence of antennal scrobes; and they preferably should exist in multiples). In the case of families, traits should be truly major, as for example the presence or absence of the petiole, presence or absence of principal exocrine glands, and the pattern of wing folding—again preferably occurring in multiples. Dlussky (1975, 1983) appears to have used lighter criteria of the kind more

commonly accepted in paleontology, in which it is recognized that fossils separated by large geographic distances and stretches of geological time are more likely to belong to different genera or higher categories. Add to this the fact that characters are often obscured due to imperfections in fossilization, and reliance is therefore placed on minor character states.

Because of the importance of the evolutionary issues involved, I believe the criteria employed in assessing the early ant fossils must be the stricter ones used in neontology. That is, it is preferable not to recognize taxa unless the character states separating them can be seen and are of approximately the same magnitude used in recognizing contemporary taxa of same rank.

Results

The result of the character-state analysis for all of the Cretaceous formicoid genera are summarized in Table 2 and 3. An inspection shows that no single character state or combination of states can be used to separate a distinct family, the Armaniidae, if neontological standards are applied. Moreover, almost none of the genera can be unambiguously supported from the existing evidence. The only exception is *Cretomyrma*, possessing protuberance on the propodeum (in *C. unicornis*) more compact gaster, and proportionately longer legs; a second, weaker possibility is *Dolichomyrma*, with an elongate head and slight constriction of gaster (Dlussky 1975, 1983, and new details provide the author in litt.). It cannot be denied that still other genera might be represented by the Soviet fossils; we are only sure that few can be defined by neontological standards with existing data. Hence the synonymy suggested in Table 4 is for the moment the appropriate nomenclature arrangement.

Figure 1 and 2 show that the measurable, relatively undistorted wingless females among the North American and Soviet fossils, in other words *Shecomyrma freyi* and *Dolichomyrma longiceps*, have small abdomes in proportion to the remainder of the body, by both the functional and strictly homologous definitions of the abdomen. In this respect they resemble the worker caste of

modern ants more than they do the wingless females of modern nonsocial aculeate wasps. Moreover, the winged female type of *Pseudarmania rasnitsyni*, a relatively undistorted specimen, is within the range of the queens of modern ants (as well as the females of nonsocial aculeate wasps), as shown in Fig. 3. In sum, the wingless females among the Cretaceous fossils are best interpreted as worker ants rather than wingless reproductive aculeate wasps belonging to solitary species while the winged females are probably the queens.

Several additional observations support this conclusion. In his description, Dlussky (1983) states that a diagnostic trait of the Armaniidae is the approximately equal length of the scape and second funicular segments. Although this is quite correct, another and more revealing way of putting the matter is to say that the scape is quite short relative to the funiculus, while the second funicular segment is long relative to the first and third funicular segments. It turns out that these are precisely the *Sphecomyrminae*. Hence both the winged and wingless Cretaceous females share the same another reason for associating them closely.

Dlussky (1983) gives as another diagnostic trait of the Armaniidae the possession of a second, free trochanter on the middle and hind legs. This condition is weakly indicated in the drawing of *Pseudarmania aberrans*, but is wholly lacking in the drawing of all of the other armaniids and by direct examination I confirmed that it is indeed absent in the holotypes *Armania robusta* (the type genus species of the family) and *Pseudarmania rasnitsyni*.

In the winged females of Armaniidae the petiole are more broadly attached posteriorly to the gaster than in the *sphecomyrmine* wingless females. But this is not a subfamilial or even species-level character. It is a common difference between the queens and workers belonging to the same species among ants, and taxonomic character to separate higher formicoid taxa.

Discussion

The most parsimonious interpretation of the Cretaceous formicoid fossil neatly joining the facts

we know, is that they all belong to the subfamily Sphecomyrminae of the family Formicidae, or true ants. Furthermore, so long as contrary evidence is lacking the Cretaceous fossil should all be placed provisionally in the genera *Sphecomyrma* and *Cretomyrma*. It is entirely possible that other genera, and even taxa in additional subfamilies or still higher categories, existed in Cenomanian to Santonian times and might be represented by the existing times and might be represented by the existing fossils, but until supporting evidence emerges, the conservative taxonomic arrangement suggested here (Table 4) is both more accurate and heuristic.

This interpretation among the Cretaceous fossils are most reasonably interpreted to be queen, worker, and male formicoids, respectively, in other words, what we would call ants as opposed to wasps. This hypothesis is more clearly depicted by juxtaposing the best preserved representatives of the three phases as though they are members of the same colony (see Fig. 4). The hypothesis receives considerable support from the size differences between the best-preserved winged fossils and the best-preserved wingless ones, consistent with their being queens and workers. It receives additional support from the fact that the proportionate size of the abdomen in the Cretaceous ants is closer to modern ants than to modern aculeate wasps (Figs. 1,2).

Dlussky (1983) made two inferences connecting anatomy to behavior inclining him to the hypothesis that the Cretaceous formicoids were not eusocial. The first is that the tips of antennal funiculi are too far removed from the mandibles to allow the precise coordination required for social behaviour: "The antennae in these insects did not permit them to control the manipulation of small objects, so that they could not have the manipulation of small objects, so that they could not have transported their brood or entered into trophallaxis with their larvae—that is, they could not have been true social insects." This supposition is surely incorrect. The eusocial vespid wasps have similarly proportioned antennae yet experience no difficulty in transporting all prey objects and placing them on the larvae. They also engage in trophallaxis, or liquid food exchange. The flexibility of the funiculi contribute to these skills, and the twists and

curves of the Cretaceous ant antennae suggest that their funiculi were likewise flexible. To this may be added the fact that some modern ants, such as the primitive *Amblyopone pallipes* (Traniello 1982) and more advanced *Pogonomyrmex badius* (Wilson 1971) do not engage in trophallaxis. Hence this form of food exchange was not essential for the evolution of eusocial behavior in ants.

Dlussky also inferred that the short, wasplike mandibles of the Cretaceous formicoids "indicates that these insects did not build true nests, and could have used only pre-existing hollows." But this overlooks the fact that some primitively eusocial wasps and bees use similar mandibles to build quite elaborate nests, mostly from carton and wax. A few, such as the halictid bees, excavate soil in a very antlike fashion.

To summarize, nothing in the observable anatomy of the Cretaceous formicoids precludes their having possessed a eusocial organization, characterized by brood care, overlap of adult generations, and division of labor between reproductive and nonreproductive castes. Although direct evidence either way is lacking on the matter, these insects could also have constructed nests in the soil, rotting wood, or arboreal cavities.

The low accumulation rate of Cretaceous formicoids to date indicates that they occurred in low densities compared with modern ants. Only two individuals (*Sphecomyrma canadensis*) have been found so far among thousands of insects in amber from Alberta Province, Canada (Carpenter et al. 1939; J.F. McAlpine, pers. comm.). Formicoids constituted just 13 of the 1200 insect impressions in the Magadan collection and 5 of the 526 impressions in the Kazakhstan collection, in other words about 1% in both cases (Dlussky 1983). These figures contrast sharply with Oligocene and Miocene deposits. In the Florissant and other shales of North America (Carpenter 1930), as well as the Baltic amber of northern Europe (Wheeler 1914) and amber of the Dominican Republic (Wilson 1985b), the ants are among the most abundant insect specimens. Thus the adaptive radiation that took place in the late Cretaceous or early Tertiary, yielding at least three of the dominant modern

subfamilies (Myrmicinae, Dolichoderinae, Formicinae) by mid-Eocene times (Wilson 1985a) was accompanied by a marked increase in abundance.

The ants are seen to have paralleled the mammals by achieving dramatic increases in diversity and abundance around the close of the Mesozoic Era. How they accomplished this breakthrough and managed to sustain a dominant position in the insect world to the present time is a matter of unusual interest, which additional paleontological studies should help to illuminate.

ACKNOWLEDGEMENTS.

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TABLE I

O. E. WILSON

Taxon	Sex	Locality	Formation and age before present (million years a.p.)
Superfamily Formicoidea			
Family Sphecomyrminae Dlussky 1983			
<i>Sphecomyrma</i> Wilson and Brown 1967	Female (wingless)	New Jersey, U.S.A.; Alberta, Canada	New Jersey: Santonian (80)
<i>Cretomyrma</i> Dlussky 1975	Female (wingless)	Taymyr Peninsula	Santonian (80)
<i>Paleomyrmex</i> Dlussky 1975	Male	Taymyr Peninsula	Santonian (80)
Family Armaniidae Dlussky 1983			
<i>Archaeopone</i> Dlussky 1975	Male	Southern Kazakh S.S.R.	Turonian (90)
<i>Armania</i> Dlussky 1983	Female (winged)	Magadan	Cenomanian (100)
<i>Armaniella</i> Dlussky 1983	Female (winged)	Magadan	Cenomanian (100)
<i>Dolichomyrma</i> Dlussky 1975	Female (wingless)	Southern Kazakh S.S.R.	Turonian (90)
<i>Poneropterus</i> Dlussky 1983	Male	Magadan	Cenomanian (100)
<i>Pseudarmania</i> Dlussky 1983	Female (wingless)	Magadan	Cenomanian (100)
Incertae Sedis (unplaced to family)			
<i>Cretopone</i> Dlussky 1975	Female (wingless?)	Southern Kazakh S.S.R.	Turonian (90)
<i>Petropone</i> Dlussky 1975	Female (wingless?)	Southern Kazakh S.S.R.	Turonian (90)

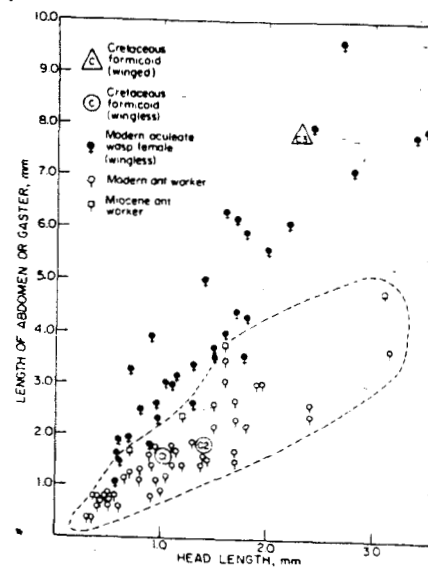
TABLE 3

Character	Species (Male)			
	<i>Palaeomyrmex zherichini</i>	<i>Archaeopone kyzylharica</i>	<i>Archaeopone layardi</i>	<i>Poneropterus sphecoideus</i>
Ratio, length of 3d antennal segment to 2d segment	2-3	?	5	3
Genitalia covered by terminal abdominal tergite, or not	Not covered	Not covered	Not covered	Covered
Petiole trapezoidal (or nearly cubical) versus tapered anteriorly (delimiting a node)	Trapezoidal	Trapezoidal	Trapezoidal	Tapered
Trochanter	1-jointed	?	?	?
Source of data	Dlussky (1975)	Dlussky (1975)	Dlussky (1983)	Dlussky (1983)

TABLE 4

- Superfamily Formicoidea Latreille 1802
Family Formicidae Latreille 1802
- Formicidae Latreille 1802, *Hist. Nat. Gen. Part., Crust. & Ins.* 3:352. Type genus: *Formica*.
- Sphecomyrminae Dlussky 1983, *Paleontol. Zhurn.* 1983, no. 3, p. 65. Type genus: *Sphecomyrma* Wilson and Brown 1967. NEW SYNONYMY.
- Armaniidae Dlussky 1983, *Paleontol. Zhurn.* 1983, no. 3, p. 66. Type genus: *Armania* Dlussky 1983. NEW SYNONYMY (tentative).
- Subfamily Sphecomyrminae Wilson and Brown 1967
Sphecomyrma Wilson and Brown 1967
- Sphecomyrma* Wilson and Brown 1967, *Psyche* 74:8. Type species: *S. freyi* Wilson and Brown 1967.
- Dolichomyrma* Dlussky 1975, *Trans. Paleontol. Inst.* 147:121. Type species: *D. longiceps* Dlussky. NEW SYNONYMY (tentative).
- Palaeomyrmex* Dlussky 1975, *Trans. Paleontol. Inst.* 147:118. Type species: *P. zherichini* Dlussky. NEW SYNONYMY (tentative).
- Archaeopone* Dlussky 1975, *Trans. Paleontol. Inst.* 147:120. Type species: *A. kyzylharica* Dlussky. NEW SYNONYMY (tentative).
- Armania* Dlussky 1983, *Paleontol. Zhurn.* 1983, no. 3, p. 67. Type species: *A. robusta* Dlussky. NEW SYNONYMY (tentative).
- Pseudarmania* Dlussky 1983, *Paleontol. Zhurn.* 1983, no. 3, p. 69. Type species: *P. rasnitsyni* Dlussky. NEW SYNONYMY (tentative).
- Armaniella* Dlussky 1983, *Paleontol. Zhurn.* 1983, no. 3, p. 71. Type species: *A. curiosa* Dlussky. NEW SYNONYMY (tentative).
- Poneropterus* Dlussky 1983, *Paleontol. Zhurn.* 1983, no. 3, p. 73. Type species: *P. sphecoideus* Dlussky. NEW SYNONYMY (tentative).
- Cretomyrma* Dlussky 1975
Cretomyrma Dlussky 1975, *Trans. Paleontol. Inst.* 147:115. Type species: *C. arnoldii* Dlussky.
- Incertae Sedis
Petropone Dlussky 1975, *Trans. Paleontol. Inst.* 147:119. Type species: *P. petiolata* Dlussky.
Cretopone Dlussky 1975, *Trans. Paleontol. Inst.* 147:119. Type species: *C. magna* Dlussky.

FIGURA I



PAUL

FIGURA 1

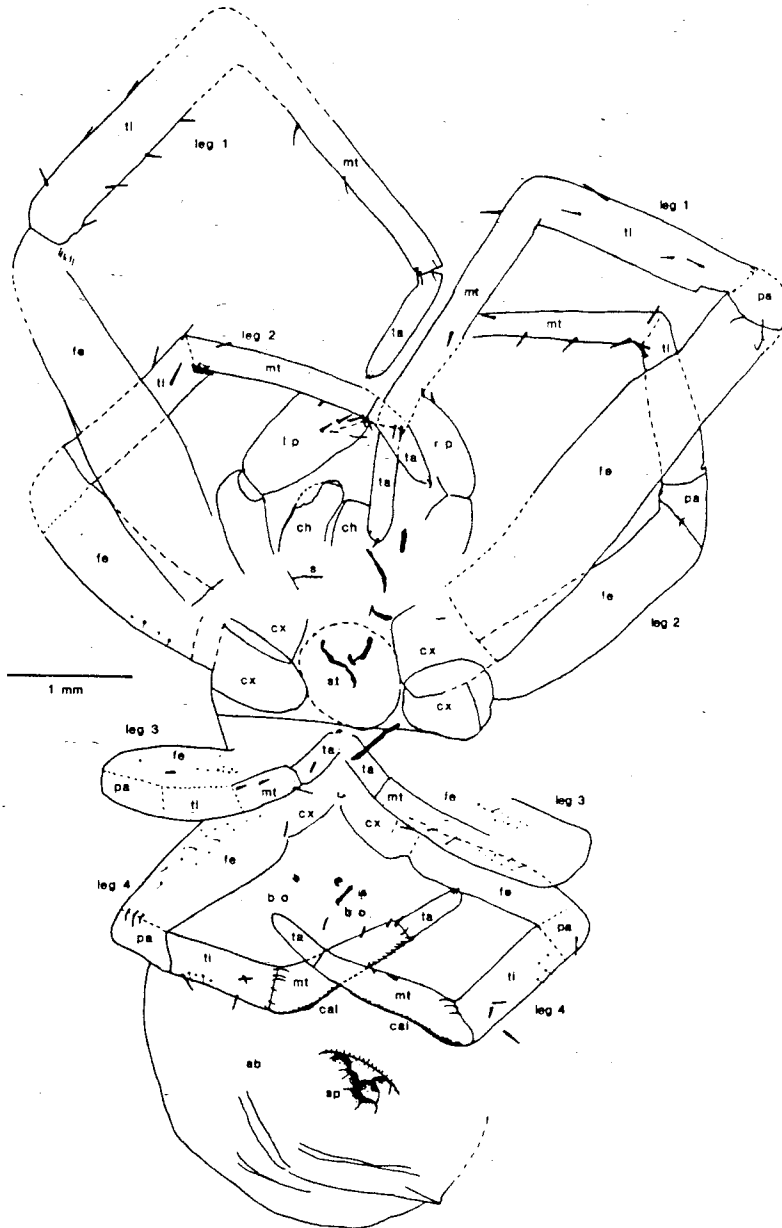


PLATE I

PLATE I

SELDEN, *Palaeodorus*

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WILSON, O. E.

PLATE

2

PLATE 2



SELDEN, *Palaeouloborus*, *Philoponella*

LOWER CRETACEOUS SPIDERS FROM THE SIERRA DE MONTSECH, NORTH-EAST SPAIN

by PAUL A. SELDEN

Abstract. Four specimens of spiders (Chelicerata: Araneae) from Lower Cretaceous (Berriasian- mValanginian) lithographic lime stones of the Sierra de Montsech, Lerida Province, north-east Spain, are described, as *Crataceous vilaltae* gen. et sp nov. All belong to the infraorder Araneomorphae. *Palaeuloborus* is the oldest representative of the superfamily Deinopoides, *Crataceus* is referred to the superfamily Araneoides and *Macryphantes* is the oldest record of the superfamily Araneoidea, family Tetragnathidae (metine-tetragnathine-nephiline group). All three spiders were web weavers; *Macryphantes* and *Palaeuloborus* wove orb webs, and may have used a wrap attack to prey on the abundant contemporaneous insect life preserved in the Montse deposit.

SPIDERS are rare rocks of Mesozoic age. Only two specimens, *Juraneus rasnitsyni* Eskov, 1984, and *Jurarchea zherikhini* Eskov, 1987, from the Jurassic of the USSR, have been adequately described. Three specimens from the Lower Cretaceous of New South Wales (Jell and Duncan 1986) and twelve from the Trias of France (Gall 1971) were not identified to a taxonomic rank below that of order. Spiders reported from Canadian Cretaceous amber (McAlpine and Martin 1969), the Jurassic and Cretaceous of Siberia and Mongolia (reported in Eskov 1984), and Cretaceous of Botswana (Waters 1989) await description. The four specimens described here are sufficiently well preserved for their taxonomic affinities to be determined with some precision, and thus they represent only the third find of Mesozoic spiders to be described and

named. The fossil spiders described here are placed in extant superfamilies or families but closer assignment is considered unwise pending the of current studies on living and fossil members of these groups.

GEOLOGICAL SETTING

Locality. The Sierra de Montsech lies in the eastern Pyrenees, between Balaguer and Temp in Lepida Province, north-east Spain (see Scharer and Janicke (1970) for details and location map). Three quarries in the vicinity of the abandoned village of Rubies, in the eastern part of the sierra, yield remarkable fossils. La Pedrera de Meia and La Cabrua have been worked for many years, and El Reguer is currently under investigation.

Stratigraphy The 50m succession of limestones exposed in the quarries the Calcaires lithographiques a Plantes et Vertebres de la Pedrera de Rubies, is a facies development of the 100m Calcaires a Charophytes du Montsech (Peybernes 1976). The deposit has been determined as late Berriasian to early Valangian in age on the evidence of ostracodes (Peybernes and Oertli 1972; Brenner, Goldmacher and Schroeder 1974; Wahalley and Jarzembowski 1985), a conclusion consistent with evidence from palynology (Barale et al. 1984; R. Porter, personal communication 1988).

Sedimentology The Calcaires a Charophytes du Montsech is a sequence of mostly pale, fine grained thinly-bedded limestones. Sedimentary structures include laminations, fine grading, minor deformation, and roll marks; trace fossils include arthropod trackways (Scharer and Janicke 1970). The environment of deposition deduced from the sedimentological and palaeontological evidence (see below) is that of coastal lagoons within a large area of algal flats separating marine conditions to the north from the Ebro continent to the south (Barale et al. 1984). This spider-bearing sequence the Lithographic Limestones, represents a particularly tranquil lacustrine depositional episode (Lacasa and Martinez 1986).

History of the fauna and flora. The exceptionally preserved biota of the Montsech lithographic limestone has received attention from paleontologists since the beginning of this century. In the

last few years, renewed interest in the deposit has brought new material to light, including the specimens described here, and avian fossils of phylogenetic importance from a locality of similar age and lithology in the neighbouring province of Cuenca (Sanz et al. 1988). Lists of fauna and flora were given by Barale et al. (1984), with the most recent summary, especially of the insects, provided by Lacasa and Martinez (1986). All the indigenous microflora is of terrestrial or non-marine aquatic provenances; the macroflora includes a wide range of gymnosperms, progymnosperms, a few ferns and horsetails, and other, unclassified, plants. Animals include ostracodes, few decapod crustaceans, larval and mature insects belonging to eight extant orders, a wide variety of fish, a few frogs and reptiles, and some birds remain. The spiders were first mentioned by Lacasa (1985, p. 2228) and a preliminary report of the results presented here was given by Selden (1989).

MATERIAL AND METHODS

Preservation. The spiders are preserved on thin slabs of pale buff-grey limestone. Grains are not visible in the rock, and the hackly fracture and vitreous appearance under high-power microscopy suggest crystallization from a lime mud. Calcified cracks cross some specimens. The spiders are preserved as pieces of cuticle on the bedding surface. The cuticle is brittle and brown; thicker parts are deep brown and the thinnest cuticle pale buff. The cuticle has not been chemically analysed; it is presumed still to be organic, but probably not of the original composition. The best-preserved parts are visible through a thin layer of translucent limestone, but their morphological details are hazy due to the presence of the overlying matrix. In such instances, 22-4% hydrochloric acid are used, sparingly and with care, to remove the matrix and thus to reveal fine structural details. The spiders are in varying state of original decay: for example in LC 1753 AP the podomeres, of leg 1 in particular, are crowded with subcircular objects along the central parts of the shaft. These objects are interpreted as the decayed remnants of muscle. LC 1754 AP is very poorly preserved: mainly a mould remains, and this is

interpreted as a specimen in which decay has progressed further.

Both part and counterpart of specimens LC 1753 AP and LC 1754 AP are preserved, but only the part of specimens LC 1150 IEI and LP 1755 AP was collected. Specimens LC 1753 AP B, LC 1754 AP A, LC 1150 IEI and 1755 AP represent lower slabs preserving mainly ventral features of the specimens; LC 1753 AP A and LC 1754 AP B are upper slabs preserving mainly ventral features of the specimens; LC 1753 AP A and LC 1754 AP B are upper slabs with dorsal features. However, splitting the rock has not resulted in perfect separation of dorsal and ventral, and due to the mode of preservation within the limestone (described above), most of both dorsal and ventral parts are preserved on LC 1753 AP B, and LC 1150 IEI. LC 1754 AP is mainly an external mould with little cuticle remaining, but on LP 1755 AP ventral parts body and both dorsal and ventral sides of the distal parts of the legs are preserved. On all specimens, the legs and abdomen (when present) are crushed flat. The carapace has sufficient convexity to produce relief in the fossil, so that on LP 1755 AP, for example, the carapace shape can be determined by the relief of the sternum and coxae are outlined by seta and cuticle. The mallepals of LC 1753 AP, LC 1754 AP, and LC 1500 IEI appear to occupy depressions in the matrix. This is probably because they were bulbous structures in life therefore created a concavity in the sediment into which the palp collapsed during burial.

The spiders were studied under a Wild M 7S stereomicroscope, with the specimens immersed in ethanol or glycerine to enhance their contrast against the pale rock background. Camera lucida drawings were photographs were under same conditions. In addition, a Zeiss photomicroscope was used, in reflected light mode with oil immersion objectives, to view and photographic details at higher magnifications. The computer programs MacClade, version 2.1 (Maddison and Maddison 1987) proved very useful for exploring relationships.

Terminology Setal terminology is somewhat problematic, since different authors have the same terms in different ways. Further more, there is

a completa gradation in size between setae (small and may be or long thick or thin), bristles, and spine (large). Macroseta is a term use, for example by Opell (1979), to describe a large seta, which is smooth apart from few minute accessory spine which are no longer than the mean thickness of the seta (see Lehtinen 1967, fig. 10a; Kullmann 1972, fig. 7). Two fairly distinct setal types are commonly called plumose (e.g. by Forters and Wilton 1973; Forster and Blest 1979). The first, which is here called plumose, is generally thicker than the serrate seta, and has helical ridges bearing small accessory spine which are much shorter than the width of the seta (see Lehtinen 1967, fig. 8; Kullmann 1972, fig. 8). The second, which is here called feathery, is fine, smooth, and has long accessory branches which are much longer than width of seta (see Lehtinen 1967, fig. 9) Many other types of seta and spine exist; there may be complete gradations between them, and the extent of their phylogenetic importance unknown.

Terminology of sclerites of male palps differs between workers in different group of spiders due to a lack of understanding of the homologies between the sclerites. Thus the task of recognizing palpal sclerites in fossil spiders is problematical. Useful descriptions of the constituent parts of male palpal organs are found in Cromstack (1948), Levi (1961), Merrett (1963) and Millidge (1977). In leg formulae (e.g. 1243), the leg lengths are ranked in order longest (first) to shortest (last). Abbreviations used in the text and text-figures are as follows:

ab abdomen
 - l labium
 s serrula
 bo book-lug operculum
 lp left palp
 st sternum
 ca cephalic area
 m maxilla
 t tegulum
 cal callamistrum
 ma median apophysis

ta tarsus
 ch chelicera
 mt metatarsus
 ti tibia
 cx coxa
 pa patella
 ti tibial apophysis
 e embolus
 pc paracymbium
 tr trochanter
 f fovea
 pe pedicel
 fe femur
 rp right palp

Provenance and depository. Three of fossil specimens, LC 1150 IEI, LC 1753 AP, and LC 1754 AP, come from the quarry of La Cabrera, the fourth, LP 1755 AP, is from the locality of La Pedrera de Meia. Exact stratigraphical provenance is not known, but both of these localities are in the same 50m sequence of lithographic limestones, the Calcaires lithographiques a Plantes et Vertebres de la Pedrera de Rubies, described above. The specimens are deposited in the Institut d'Estudis Ilerdencs, Lerida.

Preserved specimens of extant spiders were studied for comparative purposes, and in addition to those in the author's collection of mainly British species, the following specimens were examined. Uloboridae: *Uloides flavidus*, female, Funchal Madeira, M. J. Jones Collection No. 119. M-*Archester* Meseum; *Phloponella* sp., male and female, Lake Naivasha. mature and immature males, Nairobi, Kenya, J. Murphy Collection NOs 1302, 1363. Deinopidae: *Deinopsis guianensis*, females British Guiana, British Museum, (Natural History) (BM(NH)) No. 1939.3.24.228; *Deinopsis stauntoni*, female Durban, South Africa, BN (NH) No.1903.8.20.1: *Deinopsis* sp., female and immature, Kilifi, Kenya, J. Murphy Collection; *Menniscus camelus*,

females (types), Durban, South Africa BM (NH) No. 1903.7.10.22; *Mennus* sp., male, Kitale, and females Nairobi, Kenya, J. Murphy Collection *Avella angulata*, Gayndah, *Aebutina binotata*, Aguas Negras near Tarapu, Napo, Ecuador, British Museum (Natural History) Arachnid Collection.

SYSTEMATIC PALEONTOLOGY

Order ARANNEAE Clearck, 1757

Suborder OPHISTHOTHELAЕ Pocock 1892

Infraorder ARANNEOMORPHAE Smith, 1902

Remarks. The phylogenetic scheme for the higher classification of spiders which is currently widely accepted originated with Pocock (1892). In this scheme, the liphistiomorphae (Mesothe lae) is the sister group to the Opisthotelae (Mygamorphae + Araneomorphae). The morphology of liphistiomorphs distinctive (Platnick and Gertsch 1976, fig. 5) The fossil spiders described here synapomorphies: sclerite of the first abdominal segment (Platnick and Gertsch 1976), and labium as wide as or wider than the sternum (Raven 1985) Thus they are referred to the Ophisthotellae.

Mygalomorph synapomorph include the following external morphological feature: cheliceral fang which operates in the sagittal plane (orthognathous), complete loss of expression of the anterior median spinnerets, three or four articles in the posterior lateral spinnerets (Raven 1985). The alternative character states typify the Araneomorphae, and it is to this latter group that the Montesech spiders are referred. In particular, the chelicerae of *Crataaraneus* gen. nov. is more complex than the mygalomorph palp; and *Palaeouloborus* gen. nov. possesses a calamistrum and a cribellum; the latter is homologue posterior median spinneret. Neither labial or maxillary cuspules nor three- or four segmented posterior lateral spinner are known in any of the Montesech spiders, and each of them shows greater affinity to araneomorph families than to mygalomorphos.

Assignment of the Cretaceous spiders to lower taxonomic categories presents greater difficulty.

A competent araneologist can place most living spiders seen in the field into a family with a high degree of certainty. Even in the absence of field characters, such as web type and habitat, spider familial keys are relatively straightforward, only requiring the use of a binocular microscope and normally no dissection. Spider families are diagnosed on unique combinations of morphological characters such as number and position of eyes, spinner pattern, and number of tarsal claws. However, research has shown that many of the characters on which the families are based are plesiomorphic at higher taxonomic levels, convergent with other group, or are less characteristics (Lehtinen 1978). A classic example of this is the recognition of the cribellum as a synapomorphy for Araneomorphae: it prevails in lower categories ie merely the expression of the plesiomorphic state, and its absence illustrates convergence of the apomorphy in many araneomorph families. Whilst unique combinations of plesiomorphies may work as familial descriptions in practice, they cannot reflect true relationship between families. Of necessity, therefore, recent cladistic analysers have sought new or previously overlooked characters as apomorphies. These characters are commonly behavioural or anatomical, and may be poorly known in groups outside those under particular study. More important in the present work, such characters are most unlikely to be preserved in the fossil record.

An additional concern encountered when working rare fossils which have Recent relatives arises from the possibility that the fossils may be ancestors of Recent species. Should the fossils be classifying fossils may be ancestors of Recent species. Should the fossils be classified in the same as their Recent relatives or in a separate scheme? Such problems have been discussed by Crowson (1970), Farris (1981) and others. A useful device for classifying fossils with their Recent relatives is the plesion (Patterson and Rosen 1977): This presupposes, however, that the Recent classification is soundly based shared derived characters.

In spider systematics this is not yet true. In this paper, the new taxa are taxonomically classified as

close as possible to their presumed position in the Recent classification. Inevitably this entails placement within taxa yet to be defined by synapomorphies or defined on shared derived characters which cannot be seen easily in fossils. Because these are among the first Mesozoic spiders to be described, and because of the paucity of diagnostic features in the fossils, and the present lack of knowledge of synapomorphies in living families, the spiders are assigned to superfamilies, and in one case to family, but to no extant lower taxa. (Note that among Cretaceous insect, classification to modern genera is not unusual). More specimens of Mesozoic spiders which are becoming available for study may help to elucidate further the taxonomic positions of the specimens described here, and assist in unravelling the complexities of spider phylogeny and evolution. The consensus spider classification scheme given in Shear (1986b) is followed here in general, but with discussion concerning the status of groups assigned to the Araneidae following recent work by Coddington (1989a,b, 1990).

Superfamily DEINOPOIDEA Koch, 1851

Remarks .. This superfamily consists of the two cribellate families Uloboridae and Deinopidae, which weave orb webs (most Uloboridae) or spin modified orb webs which are then thrown reticularly-like at their prey (Deinopidae). Much more is known about uloborids than deinopids, though work on the latter is currently in progress. Although these two families have been considered closely related for nearly a century (Simon 1892), Coddington (1986, p. 359), with reference to his cladogram (p. 358), remarked that: the monophyly of the uloborid-deinopid lineages is based on only three characters, primarily because so little is known about deinopids. The three characters Coddington accepted as deinopoid synapomorphies are: puffed cribellate silk, pseudoserrate plumose hairs, and fourth-tarsal macrosetae; however, he qualified this by suggesting that the first two characters may actually be synapomorphies for all orb-web weavers (Deinopoidea and Araneoidea) (Coddington 1986, pp. 327, 359). Recent work by Coddington (1989a,b, 1990) has

resolved the apparent trichotomy between Araneoidea, Uloboridae, and Deinopidae (reported in Platnick 1986); Deinopidae and Araneoidea being seen as sister group of orb-web weavers: the Orbicularie Walckenaer, 1802.

Palaeouloborus gen. nov., described below, possesses three tarsal claws accessory claws typical of web weavers and lacks the tarsal adaptations of the superfamilies Thomisoidea, Philodromoidea, Lycosoidea, Clubionoidea, Salticoidea and Pholcoidea. The fossil genus also lacks the synapomorphies of Scytodoidea (Lehtinen 1986), Dysderoidea (Forster and Platnick 1985), Palpimanoidea (Forster and Platnick 1984), Hersilioidae, Hypochiloidea (Platnick 1977), Eresoidea, and Agelenoidea. Dictynoidea are cribellates, and the deinopoid families have, at one time or another, been referred to this superfamily, and to the family Dictynidae in particular. However, dictynids are generally small spiders, with short legs of approximately equal length, and lack femoral trichobothria (see below); thus they are quite unlike *Palaeoulobus*.

The characters which are most useful in placing *Palaeouloborus* are: femoral trichobothria on legs 2,3, and 4, leg 1 more than five times the length of the carapace and more than twice the length of leg 3, all leg tarsi with large accessory claws and apparently non-pectinate paired claws, presence of calamistrum and cribellum, and presence of plumose hairs.

Femoral trichobothria occur in only two groups of araneomorph spiders: the metine-tetragnathine lineage of the superfamily Araneoidea, and the family Uloboridae (Opell 1979; Lehtinen 1980; Levi 1981). In both of these groups there is great disparity in length between the elongate anterior legs (1 and 2) and the short third pair (Pl. 22, fig. 9). a feature also found in some other Araneoidea (Argiopinae Levi 1983). *Palaeouloborus* is cribellate, and since only the Deinopoidea, but not the Araneoidea, are cribellate, this genus must be referred to the former superfamily. The calamistrum of *Palaeouloborus* is situated in a curved depression on the superior surface of fourth metatarsus. Such a curvature occurs in uloborids (Pl. 2, fig. 10), to a much lesser degree in deinopids (

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Shear 1986a), and also in *Aebutina* a poorly known genus tentatively referred to the Dictyidae (Only females and immatures of this genus are known). Additionally.

Palaeouloborus bears plumose setae, which are found in Deinopoidea but not Araaneioidea but not Araneoidea. The characteristic tarsal macrosetae of deinopoids (see below) appear to be absent from *Palaeouloborus*.

Palaeouloborus is referred to be the superfamily Deinopoidea, on the evidence given above. The fossil clearly resembles members of the family Uloboridae more closely than Deinopidae; the latter legs all of a similar length, web-throwing and lacks the femoral trichobothria typical of the Uloboridae and the fossil genus.

The position of *Palaeouloborus* within the Deinopoidea is now discussed. In a recent revision of the Uloboridae, Opell (1979) gave the following characters as diagnostic of the family: (1) lack of poison glands, (2) cribellate orb-web weavers, (3) femoral trichobothria, and (4) row of macrosetae (short spines) on metatarsus and tarsus of leg 4. The tarsal spine cannot be a synapomorphy for the Uloboridae since they also occur in Deinopidae (see below). Coddington (1986) added characters of the silk-glands described by Kooor (1977), and some behavioural traits, to the list of uloborid synapomorphies, but omitted femoral trichobothria. Coddington's cladograms (1986, p. 358; 1989b, fig. 108) show the metinetetragnathines, which also have femoral trichobothria, far from the dichotomy of Araneoidea with Deinopoidea which presumes that femoral trichobothria are a convergent phenomenon in uloborids and metinetetragnathines. The alternative hypothesis (that they are a synapomorphy for all orb-weavers) would require their loss in many separate lines.

The row tarsal and metatarsal macrosetae of uloborids was considered to be a synapomorphy of the family by Opell (1979), but in a later paper, Opell (1982) mentioned finding a poorly developed row on leg 4 of deinopids as well. My own observations confirm that macrosetae are present on the inferior surface of the distal half of the metatarsus and the tarsus of leg 4, and to a lesser extent on leg 3, in the deinopids *Deinopis*,

Menneus and *Avella*. These macrosetae are rather similar to the comb of serrate bristles seen in the Theridiidae in a similar position on the legs. However, they differ from theridiid bristles in being plumose, not serrate. As in the Theridiids, they may need to be searched for, since blend into the general hirsuteness of the tarsus. The macrosetae are not greatly different from the curved bristles normally present on the inferior surface of the distal end of the tarsus. Also, as in theridiids, they are not strictly confined to mt4 and ta4, also being present on leg 3, and they vary from species to species. In all deinopids I studied they were quite unlike the comb of short, upstanding spines of the uloborid *Zosis geniculatus* illustrated by Opell (1979, pl. 1, figs. A and C). Rather they resemble the row of macrosetae of the uloborid *Hyptiotes cavatus* figured by Opell (1982, pl. 1, fig. C.) The similarity between theridiids and deinopids in this feature is presumably due to convergence in their prey-wrapping strategies rather than synapomorphy. Clearly a comb of macrosetae on the fourth leg is a derived character of uloborids and deinopids which is not present in *Palaeouloborus*.

The legs of uloborids bear fine feathery setae amongst setae; neither deinopids nor *Palaeouloborus* bear them.

Palaeouloborus cannot be included in the family Uloboridae because it has neither feathery setae nor fourth tarsal macrosetae. The fossil lacks the many specializations of deinopids, and in addition the fourth tarsal macrosetal comb is absent. Rather than redefining the family Uloboridae to accommodate the fossil genus, it is left here within the superfamily Deinopodea, closer to the Uloboridae than the Deinopidae, but not placed in either family. This placement indicates that the loss of a fourth tarsal macrosetal comb is autapomorphic for the fossil genus.

Diagnosis. Deinopoid with ovate carapace bearing marked break of separating anterior of carapace from sloping posterior area; leg 1 more than five length of carapace and more than twice length of the 3; many trichobothria on superior? retrolateral surface of femur of leg 2 and superior? prolateral surfaces of femora of legs 3 and 4; paired tarsal claws small, without teeth, median

claw long, without teeth, pair of large accessory claws; superior surface of metatarsus of leg 4 in gentle S-shape proximally convex and the concave, straighten out about half-way along podomere, bearing calamistrum which becomes row of bristles towards distal end of podomere; plumose present. Row of macrosetae absent from tarsus.

Palaeouloborus lacasae sp. nov.

plate 1; plate 2, fig 1-5, 7, 8, 10; text fig. 1

Derivation *of name After Antonio Lacasa - Ruiz Palaeontologist at the Institut d' Estudis Ilerdencs. Lerida Spain.

Types specimen Holotype LP 1755 AP, from the quarry of La Pedrera de Meia Sierra de Montsech, north-east Spain, and held in the collections of the Institut d' Estudis Ilerdencs, Spain. Diagnosis As for the genus.

Description The carapace shape is determined from fragments of cuticle from the relief of the matrix. These scraps of cuticle in the carapace region mainly represent coxae and sternum. However on the left side between legs 2 and 3, cuticle of the carapace edge can be seen lying in a depression by pressure of the edge into the matrix. Faint traces of cuticle from the posterior edges of the carapace can also be discerned between the tarsi of legs 3. These cuticle remnants and the faint deep depression by the carapace margin suggests an ovate carapace with truncated anterior margin. The carapace is 1.73 mm long, and 1.50 mm wide. It is widest just posterior to its midpoint. There does not appear a well defined cephalic area, nor an obvious fovea. There is a marked transverse break transverse break of slope just posterior to the widest part of the carapace and separating the raised foveal region from the backwardly sloping part of the carapace. This slope is found in some living spiders (e.g., the uloborid *Phyllonella*, Pl. 2, fig 8), and accommodates an abdomen which extended forwards beyond the pedicel. Eyes not seen, no obvious tubercles.

The chelicerae are large, 0.67 mm long, and 0.33 mm wide and somewhat forwardly directed, with parallel sides. The fangs cannot be seen, since the anterior edges of the chelicerae are obscured by overlying matrix.

The palps are clothed with setae, and short spines occur on the superior surface of the tarsus. The tarsus is oval, indicating that it was tumid in life. The distal end is not seen, so the presence of a claw cannot be confirmed. The specimens is therefore not a mature male, and could be an immature or a female. Very little of the basal parts of the palp can be seen, but superimposed on the anterior part of the carapace area is a dark line which, under high magnifications, is seen to be serrate (Pl 2, fig. 1). This interpreted as the serrula of the left maxilla: cuticle is absent is absent where the serrula of the maxilla would have been preserved.

The leg formula is 1243. The coxae measure approximately 0.58 mm long and the trochanters 0.18 mm. Lengths of the more distal podomeres, in mm are as follows: leg 1: fe 3.27, pa 0.77, ti 1.64, mt 2.50, ta 0.96, total 9.14; leg 2: fe 1.11, pa 0.48, ti 1.35, mt 1.44, ta 0.77, total 6.15; leg 3: fe 1.35, pa 0.48, ti 1.06, mt 0.48, ta 0.48, total 3.85; leg 4: fe 1.06, pa 0.48, ti 0.87, mt 1.025, ta 0.552, total 4.76.

The legs are clothed with setae of the plumose type. Under high magnification, they are seen to bear a sculpture of striations arranged in a helical in a pattern, with abundant minute serrae (Pl. 2, fig 7). No feathery setae can be seen on the legs. The femora are only sparsely setose, setae and bristles becoming more abundant on distal parts of the legs. Groups of trichobothria are certainly present on the femora of legs 2-4 and on the tibia of leg 4. These trichobothria are not feathered (Pl. 2, fig. 2). Isolated trichobothria are more difficult to see, and their presence elsewhere on the legs cannot be confirmed. The leg spines are not large, and because so few are seen, their presence elsewhere on the legs cannot be confirmed. The leg spines are not large because only part of their presence elsewhere on the legs cannot be confirmed. The leg spines are not large, and because only part of the specimen is preserved, the numbers of spines given below are not the maximum number which may be present on the legs. All tarsi are spineless and bear two small, non-pectinate paired claws, a large median claw which appears to be non-pectinate or if pectinate then with only minute teeth and two large accessory claws (Pl. 2, fig. 3). Femora bear few setae, mainly on the inferior surface (where they are curved) and on the super-

rior surface, especially proximally and distally. Trichobothria may be present on fel (two possible trichobothrial bases can be seen fel on the right side) but cannot be confirmed. Fel and pal are spineless.

Till bears four inferior spines and three superior spines along the shaft of the podomere, one proximal lateral spine, one lateral subdistal spine, and one superior subdistal spine. Mtl bears three inferior and two lateral spines down the shaft of the ppopodomere. Proximally, mtl bears one superior one lateral, and one inferior spine, and two inferior distal spines are present. Fe2 is spineless and bears few setae, principally on the inferior and superior surfaces. Both second femora are poorly preserved but a few trichobothrial bases can be seen proximally, apparently on the retrolateral surface. Pa2 is spineless. Ti2 bears bristles proximally and along the inferior surface. Ti2 bears two inferior distal spines, one lateral subdistal spine subdistal spine, and one superior distal spine. More spines are probably present on ti2 but on both sides specimen and bears bristles, especially along the inferior surface. Mt2 bears superior and inferior proximal spine, two median inferior spine and one lateral and two inferior distal spines. Fe3 bears few setae, one tiny spine laterally, and many trichobothria (Probably 20-40 in life) over a large area of the prolateral surface. Pa3 is spineless. Ti# is poorly preserved on both sides of the specimen, but bears many bristles. Mt3 bears two short inferior spines and one lateral distal spine. Fe4 is spineless, bears strong, curved bristles distally, and many trichobothria (as many as on fe3) over a large area of the superior? prolateral surface. Pa4 bears large bristles laterally. Ti4 bears one superior proximal spine, and prolateral, retrolateral and superior spines subdistally. Ti4 bears about four trichobothria on the proximal; superior surface, and long curved bristles distally. The superior surface of mt4 follows a gentle S-shape, proximally convex and then concave, before straightening out about half-way along the podomere. The superior surface of mt4 bears a calamistrum composed of curved setae, apparently in one row, running from the proximal end of the podomere for about two-thirds of its length, where it passes indistinctly into a row of curved

bristles which continues to the distal end of the podomere (Pl 2, figs. 5 and 8). Mt4 bears two short inferior median spines, at least two very short spines inferodistally, and large bristles distally.

The abdomen measures 3.67 mm long, and 2.89mm wide. It is ovate, wrinkled posteriorly, and compressed to the right, indicating that it was globose in life. The abdomen is sparsely setose, the setae becoming thicker, but not longer, posteriorly, where they show plumose under high magnification (Pl.2, fig. 7) A pair of subtriangular areas of darker cuticle each with a small dark patch anteriorly, is situated at the anterior end of the abdomen. These are interpreted as book-lung opercula. Three pairs of spinnerets are visible in the posterior half of the abdomen. They are compressed to the right. Their position indicates that they were not terminal, but ventral, in position in life. A recurved line immediately anterior to the anterior pair of spinnerets represents the cribellum. Little detail can be discerned the cribellar plate is not preserved (this is presumably on the counterpart) only the fold of cuticle anterior to the plate. (In living spiders the cribellum is commonly invaginated into a fold in front of spinnerets when not in use, see Pl. 2, fig. 6) Along

this fold there are numerous short, blunt setae of setae a types not seen elsewhere on the spider (Pl. 2, fig. 4).

Superfamily ARANEOIDEA latreille 1806

Remarks Cretaneusgen. nov. and Macryphantes gen. nov., describe below, are assigned to this superfamily. Both genera lack a calamistrum and cribellum. This does not, by itself, exclude them from cribellate superfamily because mature males of many living cribellate genera commonly abandon web weaving with the concomitant degeneration of the cribellum and calamistrum, and the preserved specimens of both genera are mature males. However, evidence is presented below that Crataraneus and Macryphantes show greater resemblance to araneoids than to any cribellate group. Tarsal adaptations (two claws, scopulae, Tarsomeres, etc.) for cursorial, saltatorial, and

other specialized locomotory habits, as found in the superfamilies rejected *Cretaraneus*.

Cretaraneus palp appears to be relatively simple for an araneoid, palps are variable within spider families. Therefore *Cretaraneus*, though apparently lacking a comb of setae on the fourth tarsus, cannot be excluded with confidence from the Therididae.

Stridulatory ridges are commonly present on the chelicerae of male linyphiids but are obscure in some species, and the labium is usually reborded. Neither stridulatory ridge nor a reborded labium is seen in *Cretaraneus*. A link between *Cretaraneus* and linyphiids is provided by the raised cephalic, a feature common in linyphiid mals. However, a presumed lateral condyle on the chelicera and accessory tarsal claws (serrate bristle) are features not found in the Linyphiidae, but characteristic of the Araneidae.

The rotation is characteristic of the Araneidae.

The rotation of the male palpal sclerites in some Araneidae mentioned by Levi (1983) cannot be confirmed in *Cretaraneus* because of the rotation produced during fossilization, although the general appearance of the palp in fossil genus is not reminiscent of the araneid palp. The apparent presence of a median apophysis in the palps of the fossil may provide information on its placement, but at present this character is unresolved (Coddington 1989b). The wide labium, presumed cheliceral condyle, and accessory tarsal claws of *Cretaraneus* are characteristic of most araneoids (Levi 1980). The rather large, forwardly directed chelicerae, and relatively simple palp of *Cretaraneus* suggest the metine-tetragnathine lineage, but the fossil lacks other characteristics of some linyphiids also possess enlarged chelicerae. Male palps with a superficially simple appearance are characteristic of the genus *Nephila* (Schultze 1983), but *Cretaraneus* bears few other similarities to the genus.

The Jurassic spider *Juraraneus rasnitsyni* Eskov, 1984 was placed as a monospecific family, the Juraraneidae. Eskov, (1984) defined the family on a unique combination of araneoid characters and could find no apomorphies for the family.

Cretaraneus resembles *Juraraneus* in many ways, including overall size shape; leg (but not tarsal) lengths; approximate shapes of sternum, labium and maxillae; general shape of chelicerae. *Cretaraneus* differs from *Juraraneus* in lacking the irregular group of denticles on the inner margin of the chelicera, and the palpal characters interpreted by Eskov (1984) as a large, hook-like paracymbium, a large, pointed median apophysis, and a long straight conductor (= embolus?).

The family Archaeidae is included in the Araneidae by many arachnologists but, in a radical revision of archaeids and some other small families (for example Mimetidae, previously always placed in Araneidae), Forster and Platnick (1984) removed them to the Palpimanidae. They also created the monogeneric families Prachaneidae and Holarchaeidae for genera previously included in the Archaeidae. Forster and Platnick (1984, p. 99) proposed two synapomorphies for the superfamily Palpimanoidea: cheliceral peg-teeth (modified setae as opposed to cuticular teeth), and an elevated cheliceral gland mound. They also mentioned that peg-teeth are found in some unrelated thomisoid and scytodoid genera, as convergent phenomena and have been secondarily lost in members of six families assigned by them to the palpimanoids. *Cretaraneus* possesses true teeth on the chelicerae, but mentioned here because of some similarities with the Pararchaeidae.

The Pararchaeidae differ from the other palpimanoids in having serrate, rather than plumose, hairs, and show similarity with *Cretaraneus* in the enlarged chelicerae with a prominent keel and large bristles on the tarsus. A cheliceral keel is also found in other groups, for example the Leptonetidae (Gestrich 1974). The male palp of pararchaeids has a strongly developed embolus and a large tegular plate (Forster and Platnick 1984, p. 70), feature also apparent in the palp of *Cretaraneus*. It is possible, therefore, that some relationship exists between *Cretaraneus* and the Pararchaeidae, which may or may not be an araneoid family. An archaeid spider has been described from the Jurassic (Eskov 1987).

Since it seems impossible to refer *Cretaraneus* to an araneoid family, there are two available options. First, a new be difficult, give the lack of specialized features displayed by fossil, and would not provide any additional phylogenetic information. The second option, and the one chosen here, is leave the genus unplaced within the superfamily Araneoidea. It is possible that future discoveries will reveal that *Juraneus* and *Cretaraneus* belong to the same group of Mesozoic araneoids.

Genus CRETARANEUS gen. nov.

Derivation of name creta, chalk, and araneus, a spider.

Type and only known species *Cretaraneus vilaltae* sp. nov.

Diagnosis. Araneoid spider with subelliptical carapace bearing raised cephalis and no fovea; subtriangular aternum; small, subtriangular labium; serrate setae converging all parts of body.

Chelicerae relatively large (0.4 x length of carapace), forwardly directed (at least in adult male), with inner and outer row of denticles (not peg-teeth), and mesal ridge; male palp with long embolus, and small, proximal paracymbium; leg relatively equal in length, about three times the length of carapaces; femora, tibiae and metatarsi with spines; tarsi with pectinate paired claws, small median claw, and associated serrate bristle; not true trichobothria; globose abdomen.

Derivation of the name. Ramon Vilalta-Oliva. President of the Institut d'Estudis Ilerdenses, Llerida.

Type specimen. Holotype and only known specimen. LC 1150 IEI. complete specimen on single piece of limestone from of La Caubus. Sierra de Montsec, north-east Spain; held in collections of Institut d'Estudis Ilerdenses, Spain.

Diagnosis. As for the genus.

Description. A well-preserved spider, and smallest of the specimen known from Montsec. The carapace cuticle is preserved and is golden brown in color. The carapace is 1.73 mm long and 1.37 mm wide, and pyriform: its greatest

width occurs at four-fifth of the length behind the anterior margin from greatest width.

Carapace is covered in setae. At the anterior of the carapace, these forwardly directed setae and long curved bristles. At the anterior of the carapace two circular structures are interpreted as median eyes. Just posterior. At the anterior margin of the carapace, a deep, drop-shaped depression, with its blunt end anteriormost, marks the fovea. The shape is confirmed by specimen LC 1754 AP, in which the carapace is 3.25 mm wide.

A pair of deep depressions just to the anterior median eyes are surrounded by dark cuticle. These structures are interpreted as the proximal ends of the chelicerae which are directed ventally; the chelicerae presumably disappear into the matrix beneath the specimen. The sternum appears to be circular; its shape is suggested by the arrangement of the coxae, as seen on both LC 1753 AP and LC 1754 AP. The coxae of legs 4 meet in the midline, and on both the holotype and paratype some remnants of the anterior edge of the aternum can be discerned.

The palp on the right side of LC 1753 AP B (the left side of LC 1753 AP A) is preserved (Pl. 4, fig. 3), and confirms that the specimen is an adult male. The palpal tarsus is subcircular in outline with a mesal bulge. This bulge represents either the mesal part of a circular cymbium or a mesally situated crescentic cymbium. The bulge extends beyond the oval dark oval band which is interpreted as the embolus (this structure could be the conductor or some other sclerite functioning as a guide for the embolus). The embolus is coiled in a horizontal viewed from above. On the proximalateral side, a thick, reniform lobe is interpreted as a median apophysis. A dark area can be seen on the proximal edge of the cymbium within the embolus; this is inferior in position; this dark area belongs to a sclerite of the palp. A smaller dark patch occurs just anterior to the midpoint of the palp, on the inferior surface; this represents part of another sclerite. Numerous long bristles run right laterally from the base of the tarsus, and some curved bristles are present on its superior surface.

Larger, curved spinew are present around the base of the tarsus; the appear to originate proximal to the tarsus, that is, on tibia, which is otherwise poorly preserved.

The leg length formula is 1243. The legs are very unequal in length, leg I being more than twice the length of leg 3. The coxae are approximately equidimensional, each about 0.58 mm long. The trochanters are not easily visible, but each measures about 0.25 mm in length. Lengths of the more distal podomeres and total leg lengths, in mm, are as follows: leg 1: fe 4.33, pa 1.35, ti 3.00, mt 4.55, ta 1.70, total 15.76; leg 2: fe 3.17, pa 1.00, ti 1.64, mt 3.00, ta 1.15, total 10.79; leg 3: fe 2.01, pa 0.39, ti 1.15, mt 1.64, ta 0.85, total 6.78; leg 4: fe 3.00, pa 0.81, ti 2.04, mt 1.98, ta not preserved (presumed to be C. 0.95 by comparison with ratio of mt4 to ta4 seen in LC 1754 AP, see below), total c. 9.61. LC 1754 AP is poorly preserved, and slightly larger than LC 1753 AP, and its leg measurements are as follows (in mm but with a lesser degree of certainty than in LC 1753 AP): coxae 0.9, trochanters 0.3; leg 1: fe 4.7, pa 1.4, ti 3.1, mt 5.0, ta 1.8, total 17.2; leg 2: fe 3.8; leg 3: fe 2.3, pa 0.6, ti 1.4, mt 1.3, ta 0.9, total 7.7; leg 4: fe 2.8, pa 0.9, ti 1.9, mt 2.1, ta 1.8, total 17.2; leg 2: fe 3.17, pa 1.00, ti 1.64, mt 1.64, ta 0.85, total 6.87; leg 4: fe 3.00, pa 0.81, ti 2.04, mt 1.98, ta not preserved (presumed to be c. 0.95 by comparison with ratio of mt4 to ta4 seen in LC 1754 AP, see below), total c. 9.61. LC 1754 AP is poorly preserved, and slightly larger than LC 1753 AP and its leg measurements are as follows (in mm but with a lesser degree of certainty larger than LC 1753 AP): coxae 0.9 trochanters 0.3; leg 1: fe 4.7, pa 1.4, ti 3.1, mt 5.0, ta 1.8, total 17.2; leg fe 3.8; leg 3 fe 2.3, pa 0.6, ti 1.4, mt 1.3, ta 0.9, total 7.7, leg 4: fe 2.8, pa 0.9, mt 2.1, ta 1.0, total 9.9.

All legs are thickly clothed with long appear amoth. but high magnification reveals minute accessory spines, especially distally, thus they are the serrate type. No plumose or feathery hairs can be seen. Spines are mostly large and numerous, occurring on all podomeres except the coxae and trochanters. Some spines show a helical pattern of longitudinal lines. Most spine are quite large (mean length 0.48 mm) and straight (Pl. 4, fig 6), but spines end at the distal end of podomeres are commonly curved and/or short. Fel bears five

pairs of spines along the superior surface and an inferoanterior distal spine (this may be one of a pair. Fel also has many very short, fine hairs on the inferior surface of the distal of the podomere. Pal bears a prominent posterior spine, and antero- and posteroinferior distal spines. Ti1 bears a whorl of about the spine proximally, two pairs of spines inferiorly and two single spines superior along the inferior surface, including and distal pair. Fe 2 bears four pairs of superior spines along the shaft, and posterior and anteroinferior distal spine. Pa2 bears one lateral and antero- and posteroinferior distal spiners. Ti2 bears ten long spines along the shaft of the podomere, and two short distal spine. Mt2 bears a pair proximally, a whorl of four spines and then five more along the shaft, with a whorl of five spine distally. Ta2 on the right side of LC 1753 AP B is particularly well preserved (Pl. 4, fig. 5 and 7), and shows curved paired claws each with six teeth, an equally long curved median claw, and numerous serrate accessory claws (i.e. hypertrophied bristles, as seen in living Araneidae see Foelix 1970a). Ta2 bears two small spine on its inferior surface. Along the superior prolateral side of the proximal three-quarters of fe3 are about 24 trichobothria arranged mainly in two rows (Pl. 4, fig. !). The trichobothrial hairs are not feathered. This podomere bears many fine, curved hairs inferiorly, two median superior spine. Pa3 appears to be spineless. Ti3 bears long, stiff seta, and one median and two subdistal apines. Pa appears to be spineless. Ti3 bears long, stiff setae, and one median and two subdistal spines. Mt 3 has three inferior and one superior proximal spines, one superior, antero- and posterolateral, and two short inferior and one lateral and three curved inferior spines and a whorl of five distal spines. Ta3 bears two small spines on its inferior surface, like those which occur on. Ta2. Fe4 bears at least one superior prolateral row of about sixteen trichobothria, fine, curved hairs inferiorly, three superior, two inferior, and two posterior subdistal spines. Pa4 bears one distal spine. Mt4 bears antero- and two inferior and posterolateral spines proximally, two inferior median spines, one small curved superior median spine, two small curved distal spines, and two small and one large inferior distal spines. The numerous setae on the superior surface of mt4 are

gently curved, giving the appearance of weak calamistrum (Pl. 4, fig. 6). However, the high density of setae may be an artefact of compression, since a similar density of curved setae are no different in structure from any others. Thus mt4 does not bear a calamistrum.

No trace of the abdomen is preserved on LC 1753 AP, but it is preserved on LC 1754 AP; it is oval, and measures 4.80 mm long and 4.20 mm wide. The greatest width is in the anterior half. The abdomen is elongated, and forms a compact group in a subterminal position on the abdomen.

MODES OF LIFE

Palaouloborus can be compared most closely to the *Uloboridae*. *Uloborids* are orb-web weavers which use a characteristic 'wrap attack' to subdue prey (Robinson 1975). In *Uloborus* and *hyptiotes* (Nielsen 1932 the tarsal claw is relatively large, the paired claws are fine and bear few (Hyptiotes) or no teeth (*Uloborus*), and in both genera there are large, accessory claws. A similar pattern of tarsal claws occurs in *Palaouloborus*, and it is unlike that found in araneids, in which the paired claws are large and pectinate (see, for example, Levi 1978). Nielsen (1932, pp. 26-28) described the method of silk handling by *Hyptiotes* using this claw pattern. Whilst accessory claws are widespread among web-spinning spiders, and are used for silk handling they are particularly well developed in orb-web weavers; the pattern in the *uloborids* studied seems distinctive, and these genera are orb-web weavers.

Trichobothria occur on the femora of *Palaouloborus* and *uloborids*. The function of these is not known, but it is interesting that among living araneomorph spiders they are only in tetragnathines and *uloborids*, both of which are orb-web weavers. Femoral trichobothria are absent from some adult *Pachygnatha*, tetragnathines which make no web (Levi 1980). Many orb-web weavers have no femoral trichobothria, but their presence in tetragnathines *uloborids* appears to be linked with the habit. Another behavioral similarity between tetragnathines and *uloborids* is in resting postures (Levi 1980). Tetragnathines and many metines rest with their long legs 1 and 2 stretched out

forwards, the fourth legs outstretched behind, and short third legs pointing backwards and where necessary gripping the twig on which the spider is resting. Opell and Eberhard (1983) distinguished four types of resting posture in *uloborids*; in tree, legs 1 and 2 are stretched forwards in some manner, whereas in the legs 1 and 2 are half folded with the femora projecting at right angles to the long axis of the body. Opell and Eberhard (1983) remarked on the close similarities of resting postures between *uloborids* and araneids. The femoral trichobothria are generally on the superior or retrolateral sides of femoral 1 and 2 on the prolateral sides of femoral 3 and 4 (Opell 1979). Thus they point laterally when the animal is in the normal resting posture. It is likely that the presence of femoral trichobothria is linked with the *uloborid* and tetragnathine-metine resting postures. These behavioural characters may be due to convergence, but could conceivably be synapomorphies for all orb-web weavers (Shear 1986a). The great similarity in leg lengths, femoral trichobothrial pattern, and tarsal claws between *Palaouloborus* and the *Uloboridae* suggests that the web-building and resting behaviour of the fossil genus resembled that of typical member of the living family.

The wrap-attack prey capture in *uloborids* was described by Opell (1979) and Lubin (1986). In it, the spider hangs by the first and second pairs of legs whilst throwing silk over prey using the fourth leg-pair. After further entanglement of the prey in these threads, the spider approaches closer to the prey and, holding it now with the second and third leg-pairs wraps it more tightly with silk combed by the row of macrosetae on the fourth legs. A wrap attack is found in a number of spider groups, such as the Theridiidae and Nesticidae (both of which have a comb of setae on the fourth legs). Metinae, Tetragnathinae, Araneidae, Oecobiidae, Hersiliidae, and Pholcidae (Coddington 1986). The wrap attack of *uloborids* differs from that of other spider families because *uloborids* lack poison glands and the prey is killed by digestive enzymes during (Opell 1979). Since a wrap attack is found in families which do not have a comb of setae on the legs, such a comb is not essential for this method of prey capture. Wrap attack is therefore

a possible method of prey capture in *Palaeouloborus*.

Macryphantes compares most closely in general appearance with large long legged araneoids such as the argiopine araneids, tetragnathines, and nephilines. Argiopines are similar in general appearance and habits to uloborids (both are orb-web weaving wrap-attack predators) but they lack femoral trichobothria. *Macryphantes* has an araneoid pattern of basal claws, and bears femoral trichobothria. Therefore, it is suggested that *Macryphantes* wove an orb web, rested in a posture like that of uloborids or metines-tetragnathines and may have used a wrap-attack method of prey capture. *Cretaraneus* has few positive features which indicate its possible mode of life. The fossil genus is small, short-legged, and bears pectinate, paired claws and serrate accessory claws, which indicate that it is a web-legged, and bears pectinate, paired claws and serrate accessory claws, which indicate that it is a web-weaving spider. Small, short-legged araneoids, such as Cretaceous and most theridiids and linyphiids, are weavers of sheet webs in litter, undergrowth, or bushes; such webs catch pedestrian or small flying prey. It is likely that *Cretaraneus* occupied a similar ecological niche.

A great variety of orb webs are woven by uloborid and araneoid spiders, each designed to capture a specific type of prey. They are put up for short periods or longer, day or night, and in open or secluded situations (Riechert and Gillespie 1986; Stowe 1986; Lubin 1986). They vary from massive, collective structures to minimalist devices hardly recognizable as orb web derivatives. It is impossible to suggest what type of prey

Macryphantes and *Palaeouloborus* captured with their orb webs. There was a wide diversity of insect in the Montsech area during the early Cretaceous. Which suggests that prey specialization may have been practised by orb-web weavers at that time.

CONCLUSIONS

Described here the oldest known representatives of the spider superfamily Deinopoidea family Tetragnathidae, and the second oldest record of the superfamily Araneoidea in the fossil

record. The Deinopoidea and the Araneoidea both contain weavers of orb webs of remarkable similarity. Indeed, there is continuing debate about whether the orb webs of remarkable similarity. Indeed, there is continuing debate about whether the orb webs of remarkable similarity. Indeed, there is continuing debate about whether the orb web evolved only once, in the common ancestor of the Deinopoidea and Araneoidea, or is a convergent phenomenon in these two groups. Shear (1986a) comprehensively reviewed the evidence for and against these conflicting hypotheses, and araneoids in the Lower Cretaceous indicates that, whichever hypothesis is favoured, both groups of orb-web weavers were in existence at time, and suggests that the orb web originated earlier in the Mesozoic than before.

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AP. Explanatory drawing for Pl. 1. See Terminology for explanation of Abbreviations.

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TEXT FIG 1 *Paleouloborus lacasa* gen. et sp. nov. holotype, LP 1755 AP. Explanatory Drawing for Pl. 1. See Terminology for explanation of abbreviations.

EXPLANATION OF PLATE 1

Paleouloborus lacasa gen. et. sp. nov., holotype; Lithographic Limestone Lower Cretaceous; Sierra de Montsech, Lerida Province, Spain LP 1755 AAP, whole specimen. lower slab, under ethanol, 21. See text- fig 1 for explanation.

Figs 1-5, 7, 8. *Paleouloborus lacasa* gen et sp. nov., holotype; Lithographic Limestone, Lower Cretaceous: Sierra de Montsech, Lerida Province, Spain, LP 1755 AP, oil immersion. 1. Serrula of left maxilla, bristle at left end, 240. 2. Femoral trichobothria, x 150. 3., Claws on distal end of tarsus of left leg 4; note long median claws and accessory claws, x 100. 4. Spinnerets: anterior pair (at top), median pair, and posterior pair (Part); recurved line of short setae (Seen at left) in front of anterior spinnerets mark approximate position of cribellum; compare with fig 6, x85.5 Proximal part of metatarsus of right leg 4 showing

curvature of superior surface calamistrum note long, curved bristle at distal end of tibia (bottom left) compare with fig. 10x 60.7 Plumose seta, x 625.8 Distal end of metatarsus of leg 4 showing calamistrum (overlying tarsus of 4 right leg 4 on right) terminating in of curved bristle to left x 85.

Figs 6, 9, 10. *ilopponella* sp., Lake Naivasha, Kenya; M J Murphy Coll. No. 1363. under ethanol. 6. Ventral view of posterior end of abdomen of immature male, Showing spinnerets and cribellum in front, x 36.9 Left lateral view of immature male, x 13. 10 Metatarsus of right leg 4 of mature female, showing curvature of superior surface calamistrum, x 50.

TEXT-FIG 2 *Cretaraneus vilaltae* gen. et. sp. nov. holotype, LC 1150 IEI, Explanatory drawing for text fig 3 See Terminology for explanation of abbreviations

TEXT-FIG 3. *Cretaceous vilaltae* gen. et sp. nov., holotype Lithographic Limestone Lower Cretaceous: Sierra de Montsech, Lerida Province Spain LC 1150 IEI. See text-fig 2 for explanation x 13.

TEXT FIG. 4 *Cretaceous vilaltae* gen. et. sp. nov., holotype Lithographic Limestone Lower Cretaceous: Sierra de Montsech: Lerida Province Spain LC 1150 IEI, a tarsal claws on left leg 2 note large pectinate paired claws, small, uncinata, median claw (bottom centre) and curved serrate bristles, especially inferiorly x 2200, b, typical spine, on tibia of left palpal tarsus: see f for explanation. x .88 e, tarsal claws on right leg 4. x 200 f, explanatory drawing of d, g. explanatory drawing of see Terminology for explanation abbreviations.

TEXT-FIG 5 *Macrophantès cowdeni* gen. et sp. nov.- holotype, LC 1753 AP B. Explanatory drawing for Pl. 1, Fig 1 See Terminology For explanation of abbreviations.

EXPLANATION OF PLATE 3

Figs. 1 and 2 *Macrophantès cowdeni* gen. et. et sp nov., holotype: Lithographic Limestone, Lower Cretaceous; Sierra de Montsech Lerida Province, Spain under ethanol. 1, LC 1753 AP B,

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lower slab, x 7, see text fig 5 for explanation 2, LC 1753 AP A, upper slab, x 7.

TEXT- FIG 6. *Macrophantes cowdeni* gen. et sp. nov., paratype, LC 1754 AP A. Explanatory drawing for Pl. 4, fig 4. See Terminology for explanation of abbreviations.

Figs 1-7 *Macrophantes cowdeni* gen. et sp. nov., holotype: Lime stone, Lower Cretaceous; Sierra de Montsech, Lerida Province, Spain 11, 3, 5-7, LC 1753 AP B, holotype, oil immersion 1, Trichobothria setae and spines on prolateral surface of femur of right leg 3, x 85.3 Right palP, SHOWING APOPHYSIS ON RIGHT SIDE, X 47.5 Distal end of tarsus of right leg 2, showing paired pectinate claws, long median claw, and curved, serrate bristles (accessory claws), x 320.6 Spines and setae on shaft of matatarsus of left leg 4, superior surface to top, x 85.7 Distal half of tarsus of right leg 2 showing terminal claws and short spine on inferior surface of podomere, x 130.2 LC 1754 AP B, paratype upper slab, x 4.54 LC 1754 AP A, PARATYPE, LOWER SLAB, X 3.6; See text-fig. 6 for explanation. Both under ethanol.

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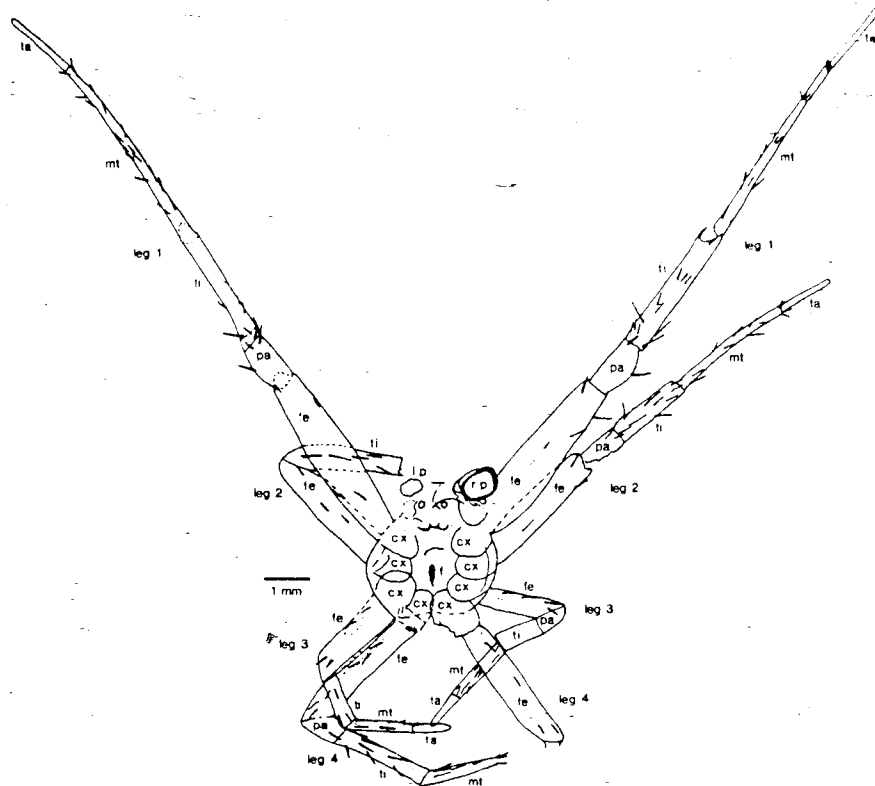
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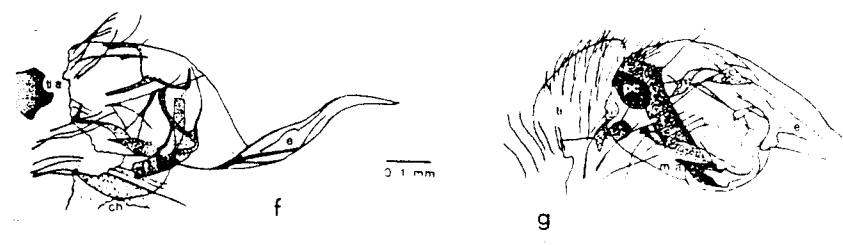
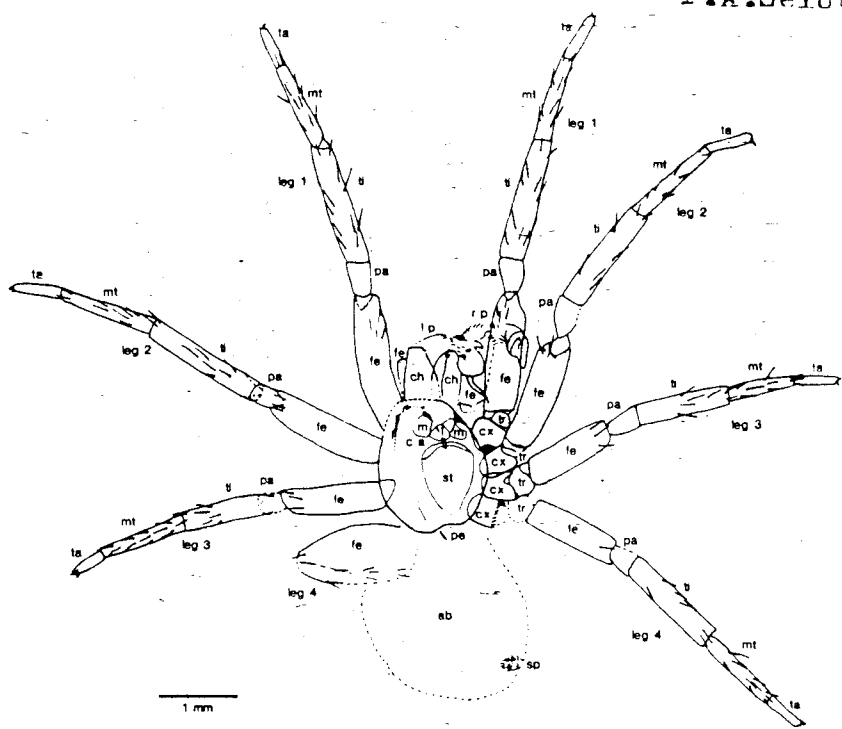
P.A. Selden



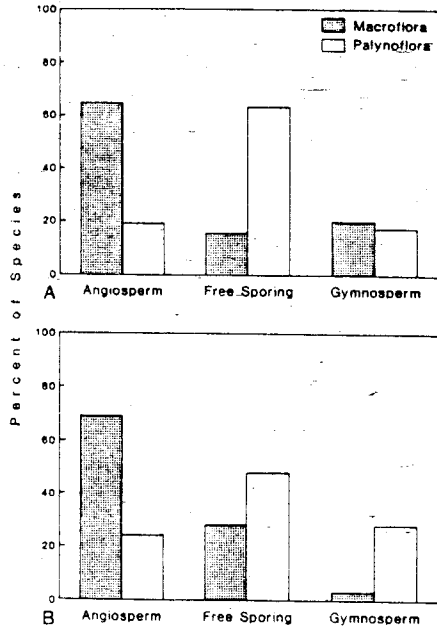
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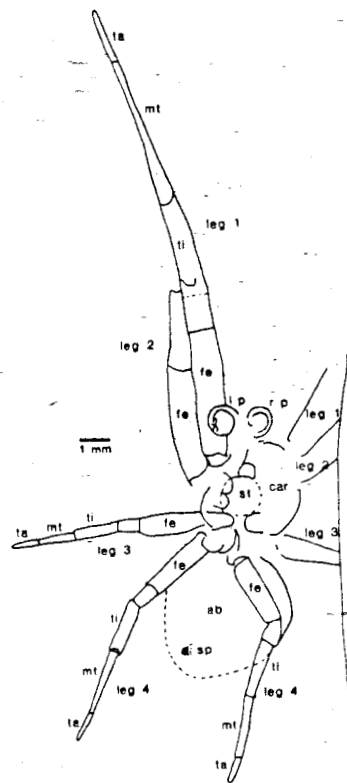


P.A. Selden



HAU	BAR	APT	ALB	CEN	TUR		
						MORPHOLOGY	LEAVES
						Simple ¹	
						Pinnate ²	
						Palmate ³	
						MARGIN	
						Entire ⁴	
						Chlor. teeth ⁵	
						VENATION	
						Pinnate ⁶	
						Parallel ⁷	
						STOMATA	
						Anomocytic ⁸	
						Paracytic ⁹	
						SEX DISTRIBUTION	FLORAL ORGANIZATION
						Hermaphrodite ¹⁰	
						Unisexual ¹¹	
						ARRANG. OF PARTS	
						Acyc.-Hemicyc. ¹²	
						Cyclic ¹³	
						INSERT. OF PARTS	
						Hypogynous ¹⁴	
						Epigynous ¹⁵	
						CARPEL NO./ARRANG.	GYNOECEIUM
						Many apocarpous ¹⁶	
						Five apocarpous ¹⁷	
						Three apocarpous ¹⁸	
						Six syncarpous ¹⁹	
						Five syncarpous ²⁰	
						Three syncarpous ²¹	
						One ²²	
						OVULE TYPE	
						Anatropous ²³	
						Orthotropous ²⁴	
						STAMEN NO./ARRANG.	ANDROECIUM
						Numerous free ²⁵	
						Nine free ²⁶	
						Five free ²⁷	
						Three fused ²⁸	
						ANTHER ATTACHMENT	
						Basifixed ²⁹	
						Dorsifixed ³⁰	
						ANTHER DEHISCENCE	
						Simple ³¹	
						Valvate ³²	
						POLLEN APERTURES	POLLEN
						Monosulcate ³³	
						Tricolpate ³⁴	
						Tricolporate ³⁵	
						Triporate ³⁶	
						EXINE SCULPTURE	
						Pallate ³⁷	
						Reticulate ³⁸	
						EXINE STRUCTURE	
						Granular ³⁹	
						Acol. semi-tect ⁴⁰	
						Col. semi-tect ⁴¹	

P. A. Selden



8.6 MACROFAGOS VERTEBRADOS

8.6 (a)

PALAEOCLOGY, TAPHONOMY, AND DATING OF PERMO-TRIASSIC * REPTILES FROM ELGIN NORTH-EAST SCOTLAND.

Michael J. Benton and Alick D. Walker.

ABSTRACT Three Permo-Triassic reptile and one footprint fauna are known from the area of Elgin, north-east Scotland. The footprints are probably Permian in age and they indicate an assemblage of mammal-like reptiles. The mammal-like reptiles, *Pareiasaurus* and *Procolophon* from the Cutties Hill Sandstone Formation (new name) are probably latest Permian in age. They were found at the base of an aeolian unit, just above pebbly sheet-flood deposits. The Lossiemouth Sandstone Formation is dated as Lower Norian on the basis of its varied reptile fauna. This fauna shows affinities with northern and southern hemisphere faunas of its time, and contains Britain's oldest dinosaur, the coelurosaur *Saltopos*. The Lossiemouth reptiles (c. 130 individual animals from four or more localities) were apparently preserved at the bases of barchan dunes and many of them just above flood plain deposits. Most of the skeletons are well preserved in articulation and only a few show disturbance, possibly by scavenging. The fauna consisted of *Hyperodapedon* and *Stagonolepis*, two moderate-sized herbivores (25% each), a selection of small omnivores:

Leptopleuron (23%), *Brachyrhinodon* (9%), *Scleromochlus* (5%), and *Erpetosuchus* (2%), and medium-large carnivore, *Ornithosuchus* (10%) and small canivore *Saltopos* (1%), some of the smaller animals show adaptations to living on sand. The last reptile fauna, from the Rhaetic of Linksfield, consists of odd bones of marine reptiles.

VERTEBRATE

Palaeontologists who study Mesozoic reptiles often concentrate on individual genera and do not record information on faunas. This paper presents information on the Elgin reptile faunas that we have gathered by personal observation of the specimens and the geology, and by studying unpublished documents and the scattered published literature.

Two main faunas of fossil reptiles are known from the neighbourhood of Elgin, north-east Scotland. The earlier, from the Cutties Hill Sandstone Formation (defined below; probably latest Permian) consists of two genera of mammal-like reptiles, a *Pareiasaurus* and a *Procolophon* and the later, from the Lossiemouth Sandstone Formation (late Triassic) comprises at least four genera of thecodontians, a rhynchosaur, a *Sphenodontid*, a *Procolophon*, and a coelurosaur dinosaur. A third fauna from the Rhaetic of Linksfield contains a few marine reptile bones. The Hopeman Sandstone Formation (in the restricted sense proposed in this paper, see below) has also yielded a fauna consisting of two or three kinds of footprints. Both main faunas are unusual in showing close affinities with those of southern continents as well as with those of the rest of western Europe and North America. Both faunas are also unusual in that the remains are preserved in aeolian deposits, clearly not the natural habitat of the majority of the animals.

The aims of this paper are fivefold: to review the composition of the faunas to consider the sedimentology and environments of the reptile beds; to discuss the taphonomy of the remains; to assess the ecology of the late Triassic reptiles; and to review the relationships and age of the faunas. Most of the data have never been published before, and it is hoped that this paper will provide a useful summary of the Elgin reptiles and their

environments. The toponomic observations are original and should be of value for comparison with other aeolian deposits. Most attention will be devoted to the late Triassic fauna because of the larger number of remains, and its greater importance in Triassic faunal development.

Repository abbreviations used in this work are: BMNH, British Museum (natural History); EM, Elgin Museum; GSE, Geological Survey Museum (Edinburgh), GSM, Geological Survey Museum (London); MM, Manchester Museum; NUGD, Newcastle University, Geology Department; RSM, Royal Scottish Museum, Edinburgh.

THE FAUNAS.

Footprints from the Hopeman Sandstone Formation.

At least two kinds of footprint have been identified from the Hopeman Sandstone Formation. Type A (text-fig 1A) (roughly circular, impressions 30-40mm long, stride length 110-120 mm, width of track way 80-90 mm, no sign of toe marks) is presented by a slab collected in 1850. This was the first fossil from Elgin recognized as reptilian, and the track way was initially interpreted as having been formed by a tortoise (Brickenden 1850, 1852).

Numerous slabs of larger footprints (Type B, text-fig. 1B) were collected afterwards (Beckles 1859; Huxley 1859b; Hickling 1909). The fore- and hind-foot were clearly different. The print of the forefoot is semicircular, about 40 mm long and 60 mm wide, with traces of four or five claws at the front. The print of the hind-foot is larger, about 90 mm long and 80 mm wide, with traces of five claws at the front. The prints overlap in pairs, and stride length was 300-400 mm. These were named *Helichnus megachelarius* Huxley 1877. Larger specimens (print 170 mm long, 140 mm wide) were figured by Huxley (1877, pl 15, fig 6) and some similar tracks may still be seen on a large slab at Clashach Quarry (NJ 163702). These large tracks measure 150-250 mm long and 100-150 mm wide, and the stride length is 700-800 mm (text-fig 1c). A possible third kind of track was described from the coast by Huxley (1877) and Hickling (1909, pl 2 fig. 7. 8) and from Cutties Hillock (Watson and

Hickling 1914). This is supposedly distinguished by having broader toes than types A or B, but the generally smudged preservation of most specimens makes such a distinction inadvisable. The Hopeman footprints were probably formed by two or more mammal-like reptile genera, possibly *Dicynodonts* (Haubold 1971).

Reptiles from the Cutties Hillock Sandstone Formation.

Reptiles bones were collected from a quarry on Cutties Hillock in 1884, and some nearly complete skeleton were object in 1881 and 1885 (Judd 1885, 1886a, b, Traquair 1886). Three genera described by E.T. Newton (1893); *Gordinia*, *Geikia* and *Elginia*.

Gordinia and *Geikia* are *Dicynodonts*, a group of herbivorous mammals-like most of which had no teeth except for a tusk *Gordinia* (text-fig 2A) is represented by remains of the skulls and skeletons of eight to thirteen individuals, and four species were originally described, although they are perhaps all synonymous (Walker 1973). *Gordinia* had a heavy broad skull, 100-180 mm long, and it was clearly provided with powerful jaw muscles. The relationships of *Gordinia* are uncertain; Cluver and King (1983, p 268) state possibly related to *Kingoria* or *Dicynodon*. The single specimen of *Geikia* has no teeth at all and the snout is box-like. Rowe (1980) has redescribed the specimen Africa and Zambia. He also placed *Dicynodon locusticeps* from Tanzania in the genus *Geikia* and in the new Family *Aulacephalodontidae*, and Cruickshank and Keyser (1984) have confirmed this assignment of *Geikia*.

Elginia was a *pareiasaur* with a remarkably spinescent skull (text-fig. 2c). The 210mm long holotype skull is broad and covered with rough pits and spines of various sizes. The teeth are leaf-like in shape, and *Elginia* was probably a herbivore. Other *pareiasaur* remains include vertebrae and a sacrum probably belonging to the holotype as well as and undescribed partial skeleton and skull.

A fourth reptile from Cutties hillock is represented by a specimen that was described by Newton (1893, pp 461-462, pl 33 fig 5) as a tail of?

Gordonia. This specimen consist of seven dorsal vertebrae, the blades of two scapulae and the blade of an ilium of a procolophonid (Walker 1973). Finally, remains of an un-named dycynodont were collected from nearby Knock of Alves (or York Tower) Quarry in 1953 by A.D.W.

Reptiles from the Lossiemouth Sandstone Formation (table 1).

A slab containing scutes of the thecodontian reptile *Stagonolepis* (text-fig 3A) was the first fossil found in the Elgin Permo- Triassic, but it was initially interpreted as a large ganoid fish (Agassiz 1984). later finds of limb-bones led to its interpretation as a reptile (Huxley 1859a,b), and in particular as an ancestral crocodile (Huxley 1875, 1877). More recent study (Walker 1961) has shown that it was an aetosaur- a probable herbivore. *Stagonolepis* is represented by skull and skeletal remains of thirty or more individuals which up to 2-7 m long. The snout had a curious blunt end, probably for digging, and the teeth were peg-like, *Stagonolepis* was shaped rather like a crocodile and had an extensive armour around its body.

The first Elgin reptile to be recognized as such was *Leptopleuron lacertinum* Owen 1851 (objective synonym, *Telerpeton elginense* Mantell 1852), based on a small skeleton collected at Spynie (Benton 1983c). Further finds (Huxley 1867; Boulenger 1904; Huene 1912a, 1920) showed it to be a procolophonid, a small omnivorous or herbivorous reptile with a triangular, spiked skull (text-fig. 3E). The skull was about 52 mm long, and the body may have been up to 400 mm in total length.

The third animal to be obtained from the Lossiemouth Sandstone Formation was named *Hyperodapedon* by Huxley (1859a). It was interpreted as a rhynchosaur (Huxley 1869, 1887; Burckhardt 1900; Boulenger 1903; Huene 1929; Benton 1983d, 1984), a group of diapsid reptiles that were abundant in many faunas worldwide in the middle and late Triassic. its closest relative is *H. huxleyi* from the Maleri formation of India. *Hyperodapedon* was a bulky 1-3 m long quadruped with strong limbs- the hind-foot had large claws and was probably adapted for scratch digging. The skull was the most remarkable feature

(text-fig. 3B). It was 100-200 mm long and very broad at the back-three was an anterior premaxillary beak and the teeth were arranged in multiple rows on the maxilla. the dentary had a sharp edge and it cut into a groove in the maxillary toothplate, providing a strong shearing bite.

Remains of the medium sized carnivore *Ornithosuchus* (text-fig 3C) were described by Huxley (1877), Newton (1894), Boulenger (1903), Broom (1913), Huene (1914), and Walker (1964). This animal was probably partly quadrupedal and partly bipedal, and it had sharp dagger-like teeth. Several specimens are known and these indicate a range in skull length of 50 to 450 mm and in body length of 0.5-3.5 m. *Ornithosuchus* was regarded by Walker (1964) as an ancestral carnosaur, but comparison with *Riojasuchus* (Bonaparte 1969) suggests that it had a crocodile-reserved type of ankle joint (Cruick shank 1979) rather than the mesotarsal joint typical of dinosaurs, and was thus a thecodontian closely related to some South American forms (Bonaparte 1969).

A second small carnivore (or insectivore) is *Erpetosuchus* which was collected from a block built into a wall in Lossiemouth in the early 1890's (Newton 1894). The specimens includes a skull (text-fig 3D), the fore-limbs and shoulder girdle, and the armour plates, which indicate a total body length of about 600 mm. It seemed to have some crocodilian characters (Walker 1968) but these were later considered to be the result of convergence. *Erpetosuchus* is another advanced pseudosuchian thecodont with a narrow 75 mm long skull which has a huge antorbital fenestra and a broad square posterior skull roof (Walker 1970, pp 364, 367- 368; Krebs 1976, pp 87-89).

Extensive collecting by William Taylor, a local naturalist, between 1890 and 1920, produced abundant remains of these animals from Lossiemouth and Spynie, as well as the holotypes of three new genera: *Scleromochlus*, *Saltopus*, and *Brachyrhinodon*.

Scleromochlus was based on several skeleton which an animal with a short body, short forelimbs, but long hindlimbs and long tail (Woodward 1907; Huene 1914). The skull (text-fig 3G) is relative ly thecodontian whose skull specialization suggest a relationship to the

aetosaurs (Walker 1970, p 361; Krebs 1976, p 90). One specimen (BMNH R3146) shows transverse bands of thin dorsal scutes, figured by Woodward (1970) as abdominal ribs.

Saltopus (text-fig. 6A), represented by only one skeleton, and unfortunately lacking the skull, is Elgin's only dinosaur (Huene 1910a). The vertebrate and limbs are rather poorly preserved in outline, and the total body length was probably about 600 mm. It is a coelurosaur (Walker 1970, p 359), and may be a podokesaurid (Huene 1914; Steel 1970) or a procompsognathid (Romer 1966).

One small undescribed thecodontian is represented by a somewhat obscure skeleton in Manchester Museum (MM L8270; Watson 1909a). This consists of a disarticulated skull, some of the skeleton, and some scutes. It is probably a new form, but is too poor to be described.

The record of a lungfish tooth-plate from Spynie, identified as *Ceratodus* (Traquair 1895, p 280) is an error. This record is attributed by Taylor (1920) to Huxley, but he gives no references and we have been unable to find that Huxley mentions it. The earliest reference appears to be that of Judd (1866a). The error was caused by confusions between the Triassic locality Spynie and New Spynie (now called Quarrywood), which lies on Upper Old Red Sandstone 4 km to the west. The tooth-plate is probably of an epipluron (T.S. Westoll, pers. comm.) and the locality is most likely Leggat Quarry (NJ 176635), 850 m to the south-west of Quarrywood, as suggested by Taylor (1901, p 48).

Reptiles from the Rhaetic of Linksfield

A large glacial erratic at Linksfield, near Elgin (NJ 222641) has yielded some reptile remains, as well as fish. The sediments were originally determined as Pliocene or Wealden (Duff 1842), but they were later dated as Rhaetic (Moore 1860, Jones 1863, Anderson 1964) on the basis of the fish and ostracods. Duff (1842, pl 4.5) figured spines, scales and teeth of such fish as *Hybodus*, *Lepidotes*, *Sphenochus*, and *Acrodus*, as well as a plesiosaur vertebra and teeth, and femur of a chelonian reptile which Seeley (1891) later described as the femur of a crocodile *Saurodesmus rodertsoni*.

STRATIGRAPHICAL TERMINOLOGY.

The Permo-Triassic of the Elgin area outcrops in two belts, one along the coast between Burghead and Lossiemouth, and the other in the region of Elgin itself (text-fig 4). The beds near Elgin occur in small fault-blocks within petrologically similar Upper Old Sandstone.

The nomenclature of the Hopeman and Cutties Hillock sandstones.

Formal lithostratigraphical terms for the main units of the Permo-Triassic of the Elgin area were introduced by Warrington et al (1980). The Hopeman Sandstone Formation was proposed to include predominantly aeolian sandstone some 60 m thick cropping out along the coast between Cummingston and Cove Sea Skerries Light house, and also for a separate strip of similar rocks, 30-45 m thick, lying on top of Quarry Wood ridge and in the Knock of Alves area to the west of Elgin. Although there is some evidence which suggests that the sandstones of these two regions are broadly equivalent, and while it is true that the coastal sandstones are much better exposed (although the base is not seen). The most important of these is that the name Cutties Hillock for a world-famous fossil vertebrate locality and set of beds has disappeared. To be replaced by an unfamiliar name. Furthermore, apart from a small scrap of bone (Peacock et al 1968, p 59) the coastal sandstone have yielded nothing but reptilian tracks, whereas the reptilian fauna entirely from the inland localities. Thus the accurate determination of the age of this formation has to be obtained from a study of the reptiles, none of which occurs in the type area.

The tracks from the coastal sandstones are believed to have been made by dycinodont reptiles, but this is a group which ranges in time from the middle of the Permian until late in the Triassic. Hickling (1909) considered that the closest comparisons of these tracks were to be made with those from Magnesian Limestone and is thus early Upper Permian in age, while the tracks from Penrith are late Lower Permian. In contrast, the Cutties Hillock fauna is probably very late Permian in age (see below). Tracks were also seen at Cutties Hillock Quarry itself in 1878 before the

reptiles were discovered (Peacock et al 1968, p 73) and have occasionally been seen since. A single small print from a nearby quarry, the position of which is now uncertain, was figured by Watson and Hickling in 1914. However, these tracks can surely only be used as evidence of a broad time-equivalence between the sandstone of the two areas.

While we do not agree with the conclusions of Glennie and Buller (1983) concerning the subdivisions of the Hopeman Sandstone on the coast, these differences of interpretations serve to underline the necessity for a separate formal lithostratigraphical unit to include the reptile-bearing beds, to avoid confusion, and until such time as definite correlations shall have been established. It is proposed, therefore, that these inland beds be termed the Cutties Hillock Sandstone Formation.

Peacock et al (1968) preferred to use the name Sandstone of Cutties Hillock (Quarry Wood) instead of Cutties Hillock Sandstone because Cutties Hillock does not occur on Ordnance Survey maps (it does actually appear on a map by Gordon (1982)). However, this is not a strong objection since the position of the quarry is given by Judd (1886a, pp 397-398) and is well established by local tradition. The term Cutties Hillock Sandstone Formation has in fact already been used by Smith et al (1974, p 25), although not formally defined.

Origin and correct form of the name cutties hillock

Although various suggestions have been made, including shorth hillock and witches hillock, it seems most likely that the name derives from a Gaelic original. It does not seem to have appeared in print until after the discovery of the reptiles in 1884. Mackie, an important worker on the local geology and discoverer of the Rhynie Chert, always used the form Cuttieshillock in several papers (e.g. 1897) and in articles in local newspapers. Other local writers tended to do the same; furthermore, the old labels on the specimens from this locality in Elgin Museum also use this form. Maxwell (1894) mentions, among other similar names Kitty shalloch in Galloway and Cuttyshallow in Ayrshire. In the last

instance he gives the derivation as the Gaelic *ceide sealghe*, pronounced to sound something like *ked-dyshalluh* and meaning hill-brow of the huting. It appears. Therefore, that originally the name had nothing to do with the words cutty or hillock, but when it was written down, authors tended to split it into two words, transferring the 's' to first of these and sometimes adding a singular or plural apostrophe as seemed appropriate.

Cutties hillock Sandstone Formation .

Type area. Quarries and natural exposures of Quarry Wood ridge, the Knock of alves and Carden Hill west of Elgin. The formation is seen to rest unconformably on Rosebrae Beds of the Upper Old Red Sandstone in Rosebrae Quarry (NJ 473633) and York Tower Quarry (NJ 162629). The reptile fauna comes mainly from Cutties Hillock Quarry (NJ 185638).

OCCURRENCE OF THE REPTILES.

Sedimentology of the Hopeman Sandstone Formation(s.s.)

The Hopeman Sandstone Formation of the coast between Cummington and Covelea Skerries and Halliman Skerries, some 60 m thick, shows evidence of largely aeolian deposition; large-scale cross bedding is common, and the sandstone is generally composed of well-rounded grains of quartz and feldspar, often of high sphericity, with only a little mica (Peacock et al 1968, p 59). However, the action of water is indicated in places by lenses of coarse sandstone and well-rounded pebbles with only a little mica (Peacock et al, 1968, p 59). However, the action of water is indicated in places by lenses of coarse sandstone and well-rounded pebbles with small-scale cross-bedding, as well as contorted beds (Peacock 1966). Williams (1973) identified four phases of dunes; sea dunes at the base, followed by three phases of beach dunes. Each of the first three phases is topped by contorted beds and sheet flood or playa lake deposit. The fossil tracks have been obtained principally from Masonshaugh Quarry, Cummington town (NJ 125692). Occasional footprints have been observed in Greenbrae Quarry (NJ

137692) and Clashach Quarry (NJ 163 7902), and Peacock et al (1968, p. 59) report an unidentifiable bone fragment from Greenbrae.

TAPHONOMY OF THE HOPEMAN SANDSTONE FORMATION FOOTPRINTS.

The footprints of the Hopeman Sandstone Formation may be preserved on low-angle dune foreset, but this has only been observed in a few in situ occurrences. The slabs collected in the nineteenth century may include some from horizontal bedding planes. However, there is usually a mound of sand behind each print (Brickenden 1852; Huene 1913; Watson and Hickling 1914), and this would suggest that the animals were moving uphill. These mounds are seen also behind the large footprints at Clashach (text-fig. 1 c).

Sedimentology of the Cutties Hillock Sandstone Formation

The Cutties Hillock Sandstone Formation is between 30 and 45m thick. It may be divided into two units (Peacock et al, 1968; William 1973): a lower phase consisting of a series of pebbly sandstone that lie discordantly on the Old Red Sandstone, and an upper phase which consists of about 30 m of large-scale light brown and yellow cross-bedded sandstone. The lower pebbly beds have been interpreted as sheet flood deposits, but the presence of dreikanter pebbles with rounded edges suggests that they were exposed to wind erosion before being reworked by water (Mackie 1902; Watson 1909b; Watson and Hickling 1914; Williams 1973) The upper phase shows well-rounded quartz grains and unidirectional forests which indicate fossil barchan dunes. The reptiles *Elginia*, *Gordonia*, and *Geikia* came from Cutties Hillock Millstone Quarry (NJ 185638) and an isolated footprint and other trackways were found nearby (Linn 1886, Huene 1913; Watson and Hickling 1914). An unidentified bone in Forbes Museum was found in Crown head Quarry (NJ 183630) on the south side of Quarry Wood hill and is of interest as the only bone recorded from this part of the outcrop. A slab in Elgin Museum showing footprints with a tail-drag on top of ripple-marks probably came from Robbies Quarry,

the position of which is uncertain but it was probably one of the Crownhead group of quarries.

Taphonomy of the Cutties hillock Sandstone Formations reptiles.

The reptiles from the Cutties Hillock Sandstone Formation appear to have been collected from the base of the upper phase, just above the pebbly sandstone. Judd (1886a, pp 400-401) noted that 20 feet of the reptiliferous Sandstone was to be seen above the pebbly layers, and that the remains of five reptiles all came from one horizon and that a sixth came from the bed below. Phillips (1886) confirmed this Gordon (1892, p 242) referred to a portion of this conglomerate containing reptilian remains. Newton (1893, pp 462, 466) also noted that the specimens of *Gordonia juddiana* also pebbles in the specimen of *Gordonia duffiana*. These blocks (E:M 1978. 559.1.2) show quartz pebbles up to 20 mm and up to 7 mm in diameter respectively.

Individual bones may be distorted. The bones are represented by cavities in the rock from which black material containing iron, manganese, and cobalt (Newton 1893, p 435) The cavities may be compressed with opposite impressions almost touching. Limb bones may be particularly flattened, and the ends may be hard to interpret either because of poor preservation and compression, or because the ends were largely cartilaginous in life. The centra of the dicynodont vertebrate are usually missing or very poorly preserved, although neural arches and ribs may be easy to cast. By contrast, the centra of *Elginia* are well preserved. Skulls are often vertically compressed (Newton 1893; Walker 1973; Rowe 1980), and in *G. duffiana* most of the squamosals and the right half of the occiput are missing, which presumably indicates post-mortem, pre-fossilization damage.

Sedimentology of the Lossiemouth Sandstone Formation

He is distributed in several small fault-bounded blocks at Lossiemouth, Spynie, and Findrassie (Peacock et al 1968, pp 67- 69; Williams 1973) The

thickness of the unit varies from 7 to 30 m. It is underlain by strata which have been interpreted as thin representative of the Burghead Sandstone Formation. Peacock et al (1968) suggested that the Burghead Sandstones (mainly point bars in their type area, according to Williams (1973)) interdigitate with the Lossiemouth as a reduced lateral equivalent of the Burghead Sandstone Formation. However, this interpretation is hard to prove because there is a large geographic gap between the outcrops of typical Burghead Sandstones in their type area. The Lossiemouth Sandstone Formation is overlain by the Cherty Rock (sandy limestone and chert).

The sandstone may be finely laminated, but more usually they show large-scale cross-beds on well-weathered surfaces. These features, as well as the absence of pebbles, the rarity of micaceous and heavy minerals, strongly suggest aeolian depositions.

Field analysis of the cross-beds indicates a prevailing south westerly wind forming barchan dunes. These have been interpreted as reworked sand from the underlying Burghead Sandstone Formation, which he regards as composed of flood-plain sandstones at Lossiemouth. Dunes up to 20 m high migrated across the flood plain during arid periods, and the aeolian deposition was terminated by large-scale flooding and the deposition of silicified and calcareous sandstone (Sago Pudding Sandstone Formation).

The reptiles were obtained from Lossiemouth East Quarry (NJ 236707), Lossiemouth West Quarry (NJ 231704), Spynie (NJ 223657), and others, and Findrassie (NJ 202650). Taylor (1920) reported a specimen of *Leptopleuron* (EM 1920.5) from a glacial erratic of Triassic beds on the Hill of Meft, north-west of Urquhart (NJ 268642).

Taphonomy of the Lossiemouth Sandstone Formation reptiles

The skeletons of *Stagonolepis* and *Hyperodapedon* were apparently found just above the base of the Lossiemouth Sandstone Formation. Murchison (1859, p 428) stated that the bones found then were collected in the lowest part of

the freestones being quarried at Lossiemouth. Gordon (1859, p 46) confirmed this, stating that the lowest beds at Lossiemouth were red clay, succeeded by yellowish soft sandstone and then harder sandstone. The red clay may be equivalent to that reported by Peacock et al (1968, p. 65) as micaceous siltstone, the yellowish soft sandstone may be the Burghead Beds equivalent and the harder sandstone is probably the Lossiemouth Sandstone Formation. The bones were found immediately under this hard siliceous sandstone in a quarry half-way to the new harbour from Rockhouse, and in the face of the wall of rock that overhangs the houses fronting the old harbour... This refers to the east end of Lossiemouth East Quarry (NJ 237707). Judd (1873, p 137) stated that the reptiles were found 100 ft, below the top of the sandstones, which would imply about the base of the Lossiemouth Sandstone Formation, if its complete thickness is taken into account. Judd (1886a, pp 397, 403) added that the reptile remains all came from a single band of soft rock. Further, Gordon (1892, p 245) suggested that if any excavation were to be made for fossils alone, it should be, in the first place, in the Lossiemouth quarries, and in the platform left by the quarrymen.

They did not go further down, because the bed was softish and rubbly, but it was at this horizon more than elsewhere in the quarry that most fossils were found.

At Spynie the reptiles also appear to have been found low in the Lossiemouth Sandstone Formation. The type specimen of *L. laceratinum* was found by Mr. William Young at the bottom of a shaft which had been sunk through 51 feet of sandstone down to a soft rubbly bed (Duff in Murchison 1859, p 435) Gordon (1859, pp 45-46) added that the specimen was extracted from the living rock, deep in a quarry opened on the west end of the hill and Martin (c. 1860) stated that the specimen was found low down, in the bottom of the quarry. The quarry has been identified as an old pit at NJ 2206 6557 (Peacock et al 1968, p 68). *Hyperodapedon* apparently came from another pit nearby: the most westerly of the Spynie quarries (NJ 2192 6555) (Gordon in Huxley 1877; Linn 1886; Peacock et al 1968, p. 68) These openings were probably abandoned over 100 years ago,

judging by the large trees growing in them, and *Ornithosuchus*, collected in 1891, may have come from the large quarry still in operation (NJ 2225 6565). Thios was the sites of two fine skulls of *Hyperodapedon* collected in about 1948 (now in RSM).

The first finds of *Stagonolepis* from Findrassie were made near the east entrance to Findrassie House, and among the debris of a pit opened up for road material (Gordon 1859, p 44). The East Longde of the Fidrassie Estate is situated at NJ2074 6545, and the pit could be one of the remaning Findrassie quarries which lie south and south-west of the entrance (Peacock et al. 1968, p 69), or it could have been filled (Walker 1961, p 106). On the original geological survey of the Elgin area, Linn (1886) recorded that *Stagonolepis* was found in the more westerly of a line of three quarries (NJ 2015 6495). Peacock et al (1968, p 137) suggest a more easterly pit at NJ 205651 as the source of the reptiles. Most of the Findrassie specimens figured by Huxley (1877) are in the form of well preserved moulds. However, some specimens in Elgin Museum labelled Findrassie look different and have bone preserved-this may indicate a different locality. There are occasional pebbles in the matrix, and the early specimens at least occurred at the base of the reptiliferous sandstone, just above the Upper Olde Red Sandstone (Gordon 1859; Walker 1961).

Outline sketches of the more complete skeletons of *Hyperodapedon* (text-fig. 5) show that most were fossilized flattened in a horizontal plane, although it is hard to say whether they are lying on their bellies or on their backs. Only the type specimen lies partially on its side (text-fig 5). The vertebral column is usually unbroken and the ribs and gastralia retain their association. The limbs are often in a natural resting pose, with the forelimbs flexed and pointing forwards, the hindlimbs are often in a natural resting pose, with the forelimbs flexed and pointing forwards, the hindlimbs pointing forwards or backwards. The shoulder girdle and pelvis may retain their original positions but they usually collapse. The skull is often present undamaged and in close articulation with the vertebral column. In these specimens there is no evidence of disarticulation by water currents, wind, or moving sand. The animals generally

appear to have died naturally and to have been covered by sand fairly rapidly. One or two isolated bones, such as tooth-bearing elements (EM 1926.6; BMNHR3151, R4780; MM L8272), show that some skeletons were broken up before fossilization.

The remains of the other medium-sized reptiles, *Stagolepis* and *Ornithosuchus*, are also generally preserved with vertebrae, limbs, and skull in articulation. Even the double row of dermal scutes above the vertebrae of *Ornithosuchus* often occurs in its original position. However, scutes of *Stagonolepis* are very often found isolated, and small bones may be slightly sattered in some cases (e.g. foot of *Ornithosuchus*; Walker 1964, p 95). The type specimen of *O. woodwardi* (BMNH R 2409-24190) is preserved with the skeleton lying on its side and the skull in a horizontal plane. The head is turned back close to the sacrum presumably by drying out of the neck ligaments after death. The *Ornithosuchus* skull, BMNH R3143, is preserved on its side. However, it should be stressed that there are no complete specimens available of *Stagonolepis* and *Ornithosuchus*. In fact, only a few specimens of *Stagonolepis* show more than a small group of associated bones, and it is clearly not possible to say a great deal about the attitude and completeness of the specimens. The main reasons for the lack of complete specimens were that there was usually no one with sufficient professional interest around at the time of discovery to make sure that everything was picked up, and it was clearly difficult to collect large fossils which were preserved as moulds in a virtually structureless sediments. This is particularly true of *Stagonolepis* which also had hundreds of scutes around its body; these would have made careful collection even more difficult. Many acutes were doubtless removed from their proper association because they made attractive specimens on their own. Only specimens of *Stagonolepis* and *Ornithosuchus* from Findrassie show signs of transport. They are disarticulated and slightly damaged, and Walker (1961, p 106) noted that pebbles are occasionally present in the Findrassie matrix. These facts may indicate some reworking by water.

The small reptiles of the Lossiemouth Sandstone Formation are often preserved in an undisturbed well-articulated state and flattened in a horizontal plane (text-fig. 6). The skull is in place, the back bone may be straight or curved, and the limbs are in natural squatting pose, as with the specimens of *Hyperodapedon*. However, there are about ten partial skull of *Leptopleuron*-isolated mandibles, maxillae, parts of the cheek region. In one specimens the skull roof behind the orbits has been lost, and probably also the upper part of the braincase. One *Brachyrhinodon* has also lost the skull roof. The type specimens of *Erpetosuchus* (BMNH R 3139) has the jaws wide open and the forelimbs in a standing pose. Two individuals of *Scleromochlus* are preserved together on one slab (BMNH R 3146) with the head of one lying partly over the anterior trunk of the other.

In detail most bones are excellently preserved and they show little sign of crushed or distorted, even in association with other unaffected elements. A right ilium (BMNH R 4788) and a posterior caudal vertebra (GSM 90884) of *Stagonolepis* became distorted without fracturing, and this led to problems of identification (Walker 1961, pp 106-107). A skull of *Ornithosuchus* (BMNH R 4782; RMS 1967. 10.1A) are rather compressed (text-fig. 7a, B) and the skull of BMNH R 4782 has also collapsed slightly. The compression of all of these bones has probably been caused by the weight of superincumbent sediment, unusual in a sandstone matrix. Delicate skulls, like those of *scleromochlus*, may be completely flattened. Various specimens of *Hyperodapedon* (e.g. BMNH R 4782; RSM 1967. 10.1A, 2; GSM 90932/5) show fine cracks or larger fractures on the surface of the bone (text-fig. 7c) which may be connected with the same process. The cracks have not opened in any way, and they are probably not sun cracks.

Signs of predation or scavenging appear to be rare, but this hard to assess because of collection failure in many cases. Walker (1964, pp. 129-131) described a specimen of *Ornithosuchus* (BMNH R 3562) in which an articulated skeleton of a large individual has been crushed and broken in certain areas. The anterior scutes of the left side are damaged, the transverse processes of the as-

sociated vertebrae are broken off short, the anterior edge of the left scapula, which lies immediately below these vertebrae, is also broken and incomplete. The humerus is broken at the same point and the ribs of the left side are missing. The body has been considerably disturbed in the region of the left shoulder and side, but the rest of the skeleton is not affected, so that the damage was almost certainly caused by a predator or scavenger.

Two isolated skulls of *Hyperodapedon* (RSM GY 1984. 20.1.2) show similar localized damage. In one the parietal is broken off just behind the attachment of the epipterygoids, and small displaced bone chips are preserved, and the top of the braincase is slightly crushed. This damage occurred before fossilization and probably at, or after, death since the breaks are clean, but there are no tooth or claw marks on the bone. The damage was probably not caused by physical processes since fine sclerotic plates are preserved, only slightly displaced, in the orbit. In the other specimens the braincase has been completely removed, and it is represented by only two fractured, displaced bone slivers. The atlas, axis, and two anterior cervical vertebrae are preserved in articulation, but pushed in between the quadrates. Ossified hyoid elements are preserved in place, so the disarticulation is unlikely to have been caused by sedimentary processes.

The bone material is either absent (some Fiddisie specimens), or very soft and sometimes partly replaced by iron oxide (goethite) and fluorite (Spynie, Lossiemouth). When original material is preserved the structure of bones and teeth may be extremely clear, and the cavities highlighted by the replacement sandstone. Positive preparation in the study of some specimens of *Stagonolepis* and *Ornithosuchus*. Normally, casts have been made from the natural moulds left as cavities in the rock, and various methods that involve flexible synthetic rubbers have been developed in order to preserve the rock mould and produce a high-fidelity copy or the bone (Walker 1961, 1964, 1973; Benton and Walker 1981).

ECOLOGY OF THE LOSSIEMOUTH SANDSTONE FAUNA (text-fig. 8)

The composition of the fauna of the Lossiemouth Sandstone Formation and the average sizes of its constituents are summarized in Table 2. The absence of plants, invertebrates, and fish, and the relatively small numbers of specimens make it futile to draw up food-chains and calculate biomass and productivity.

Stagonolepis and Hyperodapedon dominate the faunas as medium-sized herbivores that must have fed in relatively well-vegetated and watered areas distinct from the dune-fields in which they are preserved. Stagonolepis had strong dorsal and ventral armour and a heavy tail. It may have grubbed for plants or invertebrates with its shovel-like snout, and masticated them with its small pointed teeth by means of a slicing jaw action. The structure of the forelimb and shoulder girdle show that Stagonolepis normally walked on all fours, and the forelimb may have been used for digging also. For fast movement it may have been able to hold its limbs under the body for short periods, as in crocodiles. Stagonolepis was a thoroughly terrestrial animal (Walker 1961).

Hyperodapedon was a shorter more bulky terrestrial reptile which had powerful limbs. The massive laterally flattened claws of the foot and the construction of the hindlimb strongly suggest that it

was used for scratch-digging. Hyperodapedon may have fed on buried roots and tubers and other lowlevel vegetation which it raked up with its premaxillary beak and manipulated with a large tongue.

The jaws operated like a penknife, with the dentary cutting into a deep groove on the maxilla. There were multiple rows of teeth that were not replaced from below. There were massive spaces for adductor muscles at the back of the skull, and Hyperodapedon probably probably cut up tough plant material with a powerful precision shear bite (Benton 1983b, d, 1984)

Stagonolepis and Hyperodapedon were probably preyed upon, or their carcasses scavenged, by large Ornithosuchus, Ornithosuchus may have

been capable of bipedal locomotion for short spells, as well as quadrupedal locomotion. The long, curved teeth were clearly those of a carnivore, and the spinose scutes along the back may have been necessary for protection against cannibalistic attack (Walker 1964, 1977).

Erpetosuchus was a rare small carnivore and/or insectivore. The dentition was peculiar, with long sharp recurved teeth at the front of the jaws and toothless longitudinal ridges behind which may have been used for crushing the prey. There was also an incipient secondary palate, which may have been connected with the need to masticate the food.

Saltopus Elgin's only dinosaur, was a small presumably carnivorous coelurosaur. It is more primitive than Huene (1910a) indicated, having three rather than four sacral vertebrae, and the anterior blade of the ilium is not very long. Further, Saltopus shows no particular adaptations to hopping, as he suggested. Huene also suggested that it had elongate cervical vertebrae, but these are not preserved: the anterior portion and the skull of the only skeleton is missing.

Saltopus, small individuals of Ornithosuchus, and possibly Erpetosuchus, probably fed on the small herbivores of the Lossiemouth Sandstone Formation; Leptopleuron, Brachyrhinodon, and Scleromochlus. Leptopleuron may be seen as a reptilian rodent. It had transversely broad back teeth consisting of two cusps linked by a narrow cross ridge, that were probably used for grinding up and chopping tough plants sliced off by its sharp, chisel-like front teeth. The teeth were set well forward in the jaws, the lower jaw was deep with a strong (coronoid) process, and the temporal area of the skull was spiny which may have acted to deter predators, to break up the animals against a background of spiny vegetation, and/or to form part of an interspecific combat display. The size and form of Leptopleuron and its inferred habitat, are reminiscent of two desert-dwelling lizards; the Coast horned lizard, Phrynosoma cornutum from North America, and the Moloch, Moloch horridus from Australia.

Brachyrhinodon a tiny sphenodontid with acrodont teeth on the jaw margins and on the palate, and a very short snout, was probably also

a herbivore that could chop up tough vegetation with ease. It was much smaller than the living *Sphenodon* (*Brachyrhinodon* had a skull length of 20-30mm; *Sphenodon* has a 70 mm skull), and its habits need not have been the same.

Scleromochlus is equipped with long hindlimbs that have been interpreted (Woodward 1907; Huence 1914) as adaptations to jumping, and it may have sought out foot on the dunes in which it is preserved. Jumping animals (saltators) have lengthened hindlimbs, the distal segments in particular, several tarsal, or metatarsal, bones may lengthen, the tail is long (for balance), and the dorsal vertebral column may be shortened and the pelvis strengthened in order to withstand the impact of landing (Hildebrand 1974, pp. 512-515). *Scleromochlus* displays all of these features and in addition, it has a deep intercondylar fossa (patellar groove) at the distal end of the femur (Huene 1914, p 10), a feature seen in jumping mammals to accommodate a strong quadriceps femoris tendon. *Scleromochlus* may be compared with the jerboa and kangaroo rat which are similarly adapted and live in hot, sandy deserts. The suggestion by Huene (1914) that *Scleromochlus* climbed trees and made bold leaps from a branch or even glided, is not likely since the feet hardly appear to be adapted for grasping branches. Likewise, the suggestion by Wilfarth (1949, pp 7-9) that *Scleromochlus* lived underwater and used its powerful hindlimbs to propel itself to the surface for air, seems even less probable. *Scleromochlus* also appears to have some specific adaptations for living on sand, like the North America desert-living iguanid lizards *Uma* and *Dipsosaurus* the nares are nearly closed by lateral flanges, there is posterior flanges from the squamosal and quadratojugal protecting the tympanic region, the lower jaws is countersunk, and the metatarsus is flattened.

The sedimentological and taphonomic data suggest that most of the Lossiemouth Sandstone Formation animals normally lived in well-vegetated areas, possibly around interdune pools or close to water outside the dune-field. During arid periods, large sand dunes migrated across the floodplain, and many of the animals may have been overwhelmed by the sand. Others died from natural causes within the area of dunes. Nevertheless, it is

striking that no fossils have been preserved in the underlying water-laid beds, of animals like fishes, labyrinthodont amphibians, or phytosaurs.

The cadavers were normally buried rather rapidly by drifting sand dunes, which is suggested by their relatively well-articulated state. There may have been some scavenging of the carcasses by *Ornithosuchus* and the smaller carnivores.

AGE VARIATION AND SEXUAL DIMORPHISM.

Stagonolepis, *Ornithosuchus*, *Hyperodapedon*, and *Leptopleuron* are represented by enough specimens to permit some observations on variation in size and proportions.

The material of *Stagonolepis* segregates into two size groups, each containing similar numbers of specimens. Individuals within each group are remarkably uniform in size, and all measurements differ by 25-30% between the groups. Proportions are similar in both groups and morphological differences slight. These groups have been interpreted as male and female animals (Walker 1961). Romer and Prince (1940) found two size-groups in material of *Dimetrodon limbatus* and they also explained these in terms of sexual dimorphism. Among living reptiles, male crocodiles and alligators tend to be larger than females of the same age (Guggisberg 1972, p.127). Male lizard and tuataras also tend to be larger than females, but snakes often show exactly the opposite relationship (Goin and Goin 1971, pp. 115-116; Porter 1972, p.309). Both situations may be found in chelonians.

Ornithosuchus displays a broad range of size, with estimated skull length ranging from 125 to 210 mm. with the majority in the range 170-180mm. One small fragment (RSM 1966.43.2) represents the tip of a juvenile dentary, but the other specimens cannot be segregated into distinct size-groups (Benton 1983d). Thus, as with *Ornithosuchus*, the size distributions corresponds to age alone. As is general among living reptiles, age size-classes may be apparent among juveniles, but they are usually blurred in older animals because of individual differences in growth rate (Gibbons 1976).

However, the material of *Hyperodapedon* may be divided into two classes according to proportions. For example, if two skull of identical length are compared, certain important elements (tooth-plate, occipital condyle, glenoid and quadrate facets) have the same dimensions, but bone bars and sheets between skull opening may differ markedly. Forms may be classified as gracile or robust on the basis of measurements of interorbital width, thickness of postorbital, or jugal, which differ by 20-70%. There is no relationship between these proportions and overall skull-size. It has been suggested (Benton 1983d) that the robust forms may be male, and the gracile forms female by analogy with certain living and fossil reptiles. The extant lizard *Lacerta jacksoni* shows sexual differences in the shape of the snout, nares, frontals, and occiput (Degen 1911). Proposed males and females of the dinosaur *P. Orotoceratops* achieve the same maximum size, but differ in certain secondary sexual characters (frill, horn), as well as in the nasal height of the skull, size of the parietal fenestra, size of nares, height of the coronoids process, and other features (Dadson 1976). Similarly, different skull patterns have been ascribed to sexual dimorphism in *Dimetrodon limbatus* (Romer and Prince 1940), *Seymouria baylorensis* (Vaughn 1966), and *Tetragonias* (Cruickshank 1967).

DATING THE FAUNAS

The Hopeman Sandstone Formation

The footprints from the Hopeman Sandstone Formation (s.s.) are regarded as those of mammal-like reptiles, probably dicynodonts, and they closely resemble tracks from strata close to the Lower/Upper Permian boundary of Dumfriesshire, Penrith, and Mansfield (Hickling 1909; Watson 1909b; Watson and Hickling 1914; Haubold 1971; Smith et al 1974). As noted already (p215), Glennie and Buller (1983) divided the Hopeman Sandstone Formation into two units, assuming that the contorted sandstones are confined to one horizon at about the same topographical level in the cliffs. However, since Williams (1973) recognized three well-separated horizons of contorted strata in upward succession

from east to west, this assumption is open to question. Furthermore, the Clarkly Hill bore hole (Peacock et al 1968, p 130) encountered pebbly sandstone at the base of the Hopeman Sandstone, many of the pebbles being faceted. This pebbly sandstone would naturally be taken to be the equivalent of the dreinkanter bed at the base of the Cutties Hillock Sandstone Formation, whereas Glennie and Buller correlate the upper of their two units with this formation. In addition Glennie and Buller (p.57) imply that the reptilian tracks occur in the lower of their two units, again if anything suggesting a correlation of this unit with the Cutties Hillock beds. The latter, however, are of very late Permian age. In fact, if we interpret Glennie and Buller correctly, the tracks would come predominantly from the upper of their two units, since they were most common in Masonsshaugh, Greenbrae, and Clashach quarries. However, the evidence of the footprints cannot be regarded as very strong (see above), and the uncertainty over the succession within the Hopeman Sandstone Formation simply emphasizes the necessity for separate Cutties Hillock Sandstone Formation in order to avoid confusion.

The Cutties Hillock Sandstone Formation

The reptiles from the Cutties Hillock Sandstone Formation were initially assumed to be of the same age as those from the Lossie mouth Sandstone Formation (Judd 1885, 1886a). However, Newton (1893) hinted that the faunas were distinct, and Taylor (1894) suggest a Permian age. Independently, Huene (1902) and Boulenger (1904) made the same suggestion. Walker (1973) discussed the age of the Cutties Hillock reptiles in detail, comparing *Elginia* and *Geykia* with animals from South Africa and Tanzania. He concluded that the Elgin formation was to be placed very close to the Permo-Triassic boundary, as had Watson and Hickling before him (1914), tentatively suggesting that it might lie at the very base of the Triassic, equivalent to a position low in the Lys trossaurus Zone of South Africa. The main evidence for this suggestion was that both *Elginia* and *Geikia* are more advanced than their closest relatives from the late Permian of South Africa and Russia, particular stress being laid on the

specialization of the skull of *Elginia*. This view was not only based on the high degree of the spinescence, as stated by Rowe (1980), but reference was made to other skull characteristics. Rowe (1980) considered that the relationships of *Geikia* point to an uppermost Permian (i.e. *Daptophthalmus* zone) horizon for the Cutties Hill Lock Sandstone Formation. Using different evidence from the late Permian of Tanzania, is close to the ancestry of *Geikia*, the cryptodontid dicynodonts, all come from the late Permian of South Africa or Zambia. On the other hand, pareiasaurs less specialized in the skull that *Elginia* persist into the higher part of the *Daptophthalmus* zone (Kitching 1977) and there seems no reason why this group should have become extinct world-wide at the Permian-Triassic boundary. On balance, it seems best to take a conservative course, and regard the Cutties Hill Lock Sandstone Formation as lying at the extreme summit of the Permian, but bearing in mind that *Elginia* may represent a relict line which lingered on rather later in Scotland than elsewhere.

The Lossiemouth Sandstone Formation

The dating of the Lossiemouth Sandstone Formation has an even more involved history. The Elgin sandstone were all initially considered to be Devonian in age, until finds of reptile foot prints and bones in the period between 1850 and 1860 convinced most palaeologists that some of them must be regarded as younger. Local geologists long considered that the Lossiemouth sandstone were Devonian because they wanted to have the oldest reptiles in the world (e.g. Phillips 1886; Gordon 1892). Also, certain geologists (e.g. Sir R.I. Murchison, Charles Lyell) had their own reason for preferring to assign the reptiles to the Old Red Sandstone (Benton 1983 c.). However, Huxley (1867) argued convincingly for a Triassic age, and Murchison (1867, p. 267) accepted an Upper Triassic assignment.

Huene (1980) correlated the Lossiemouth sandstone with the German Lettenkohle (Late Ladinian) on the assumption that *Hyperodapedon* also occurred in the Lower Keuper sandstone of the English Midlands together with

amphibians and plants typical of the German formation.

There is now strong evidence for a lower Norian (Upper Triassic) assignment for the Lossiemouth Sandstone Formation. Walker (1961) pointed out that *Stagonolepis* is very closely similar to *Aetosaurus* from the German Stubensandstein (Middle Norian; Anderson and Cruickshank 1978; Tucker and Benton 1982). *Aetosaurs* occur also in the German Blasensandstein (Ebrachosaurus Lower Norian), the Dockum Group of Texas and the Chinle Formation of Indian (unnamed: L. Norian) the Ischigualasto Formation and Los Colorados Formation of Argentina (*Aetosauroides* and *Neo-aetosauroides*, respectively; L. Norian and U. Norian /Rhaetian), and the New Haven Sandstone of Connecticut (*Stegomus*: M. Norian). *Aetosauroides* is slightly more primitive than *Stagonolepis* (Walker, in discussions to Warrington 1970, p. 218). *Ornithosuchus* is most closely related to *Riojasuchus* from the Los Colorados Formation of Argentina (Norian; Bonaparte 1978). *H. gordonii* is remarkably similar to *H. huxleyi* from the Maleri Formation of India, and *Scaphonyx* from the Santa Maria Formation of Brazil and Ischigualasto Formation of Argentina (all late Carnian or early Norian). The Elgin procolophonid *Leptopleuron* appears to be very close to *Hypsognathus* from the Newark Group of New Jersey (latest Triassic or earliest Jurassic; Olsen and Galton 1977). *Brachyrhinodon* is most like *Polysphenodon* from the Gipskeuper (Carnian) of East Germany (Walker 1966). The other Elgin reptiles do not appear to have had relatives elsewhere, as far as we know.

The close relationship of *PyHyperodapedon* from Elgin and India is important. The Maleri Formation has also yielded a primitive phytosaur close to *Francosuchus* from the German Blasensandstein (Lower Norian). The Maleri Formation fauna also contains specimens of the Labyrinthodont *Metoposaurus* which is restricted to horizons in Germany ranging from the Schilfsandstein to the Blasensandstein (Upper Carnian-Lower Norian). The Maleri Formation is very probably lower Norian in age, and the Lossiemouth Sandstone Formation also. Unfortunately the age of the Lossiemouth beds cannot

be confirmed independently by means of other fossil groups.

The Lossiemouth Sandstone fauna shows some affinity with other northern hemisphere faunas of the late Triassic. It shares aetosaurs, Sphenodontids, procolophonoids and, coelurosaur with the North America and German faunas. However, the Elgin beds lack the metoposaur amphibians and phytosaurs that dominate all of these faunas. The other elements of the Lossiemouth Sandstone fauna show affinities with India and the southern continents; similar late Triassic rhynchosaurs are known from India and South America (a few scraps from North America), the closest relative of *Ornithosuchus* comes from South America, the South America and Indian faunas also have aetosaurs.

It is hard to find palaeogeographic reasons for these apparent distributional anomalies. The solution may depend on the environments in which the animals lived. The sediments in which the German, North America, and Indian faunas are found are largely water-laid, and metoposaurs and phytosaurs are clearly aquatic or semi-aquatic animals. However, the sediments of the South America formations are also largely water-laid, and amphibians are rare and phytosaurs are absent. The distinction between these two kinds of faunas is probably environmental, however, and they have been distinguished as a Metoposaur/Phytosaur Empire (Germany, North America) and a Rhynchosaur/Diademodontid Empire (South America, Africa, India) (Tucker and Benton 1982; Benton 1983a). An interesting problem is the virtual absence of rhynchosaurs in North America, and their complete absence in Germany which is hard to explain in view of their overwhelming abundance elsewhere.

SUMMARY

1.- Fossil reptile remains are known from four horizons in the region of Elgin, North-east Scotland: the Hopeman Sandstone Formation, the Cutties Hillock Sandstone Formation, the Lossiemouth Sandstone Formation, and the Rhaetic.

2.- The remains from the Hopeman Sandstone Formation of the coastal region are footprints of

several kinds. These indicate a range of two or three genera of mammal-like reptiles as the producers, and they resemble other mid-to late-Permian trackways from elsewhere.

3.- The reptiles from the Cutties Hillock Sandstone Formation include two dicynodonts, *Geikia* and *Gordonia*, a pareiasaur, *Elginia*, and an un-named procolophonid. These reptiles indicate a date in the late Permian near the Permian-Triassic boundary.

4.- The reptiles from the Lossiemouth Sandstone Formation include the rhynchosaur *Hyperodapedon*, the thecodontians *Stagonolepis*, *Ornithosuchus*, *Erpetosuchus*, and *Scleromochlus*, the procolophonid *Leptopleuron*, the sphenodontian *Brachyrhinodon*, and the early dinosaur *Saltopus*. These reptiles individually suggest a late Triassic age, and jointly a lower Norian age.

5.- The reptiles from the Rhaetic of Linksfield include plesiosaurs and crocodiles, represented by odd teeth, vertebrae, and limb bones.

6.- The Cutties Hillock Sandstone Formation is formally defined here, and it is distinguished from the Hopeman Sandstone Formation. There is little evidence for a direct correlation of the two, and the new name is required to show the distinctness of the two formations.

7.- The reptiles of the Cutties Hillock Sandstone Formation and the Lossiemouth Sandstone Formation occur in aeolian sediments. The skeletons are preserved fairly completely, and with only occasional disturbance (by predators). The bones are sometimes preserved, and sometimes completely lost, so that casts can be made. There is good evidence that the skeletons occurred low in their respective formations, at the base of large aeolian dunes.

8.- The Lossiemouth fauna includes medium-sized herbivores that must have fed in well-watered areas, as well as smaller lizard-shaped animals that show adaptations for running around in the dunes. For some genera there are enough specimens to show age variations and sexual dimorphism.

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Text-fig 1.- Reptile footprints from the Hopeman Sandstone Formation, near A small prints, after Brinckenden (1952; B, medium prints. *Chelichmus megacheirus* after Huxley (1877, pl. 14); c, large prints, field photograph. A, B, from Mason-shaugh Quarry; c from Clashach Quarry, scales are shown for A and B; c, shows tracks which are 100-150mm wide and have a stride length of 700-800mm.

Text-fig. 2. The reptiles of Cutties Hillock Sandstone Formation of Cutties Hillock Quarry, near Elgin. Skull of A, *Gordonia*; b *Geikia*; and c, *Elginia*, all drawn to the same scale. (A, Newton 1893; B, after Newton 1893 and Rowe 1980; C, after Newton 1893 and original.)

Tabla 1.- Classification of the reptiles of the Lossiemouth Sandstone Formation.

Text.-fig. 3. The reptiles of the lossiemouth sandstone Formation, near elgin (late triassic; early Norian). Skull of A, *stagonolepis*, B, *Hyperodapedon*; c; *Ornithosuchus*; D, *Erpetosuchus*; E, *Leptopleuron*, F, *Brachyrhinodon*, and G, *scleromochlus*, drawn to three different scales (A-C, D, E-G). The skull of *Saltopus* is not known, (A, after Walker 1961; B, after Benton 1983 d; C, after Walker 1964; D, original; E, original; F, after Huene 1910b, G, original).

Text-fig 4 The distribution of the Permo-Triassic beds around Elgin, north-east Scotland. The formations are indicated by shading, and the main reptile and footprint localities are named.

Text-fig. 5 skeleton of *Hyperodapedon* as preserved to show typical position of fossilization. Vertebrae and ribs are shaded black, and all other elements are shown in outline.

Text-fig. 6 Skeleton of some smaller reptiles from the lossiemouth sandstone formation to show the positions of fossilization. Vertebrae ribs are

shaded black, and all other elements are shown in outline.

Text-fig. 7 Compression and pre fossilization damage to bones of *Hyperodapedon* from the Lossiemouth sandstone Formation, A,, lateral view of distorted partial left scapula, compared with B, lateral view of left scapula, restored from cast, both from animals of similar size, C, crushed distal end of a femur, showing pre-fossilization damage.

Tabla 2.- The Lossiemouth Sandstone Formation fauna, composition. Information from Walker (1961, 1964) Benton (1977, 1983 d), and original.

Text-fig. 8 A scene at elgin, north-east Scotland, in Lossiemouth Sandstone Formation times showing reconstruction of the reptiles in an imaginary scene with typical late Triassic plants. Three *Hyperodapedon* feed on seed ferns in the foreground. Behind them on *Ornithosuchus* runs towards the armoured *stagonolepis* which is looking over its shoulder. Behind *Stagonolepis* two *Erpetosuchus* feed on a small carcass. On the rocks in the left the side of the pond. In and around the pond there are horsetails, cycads, and ferns, and there are tall lycopods in the distance. Based on a colour painting by Jenny Halstead in Benton (1983b).

M.J.Benton and A. D. Walker.

Figura 1

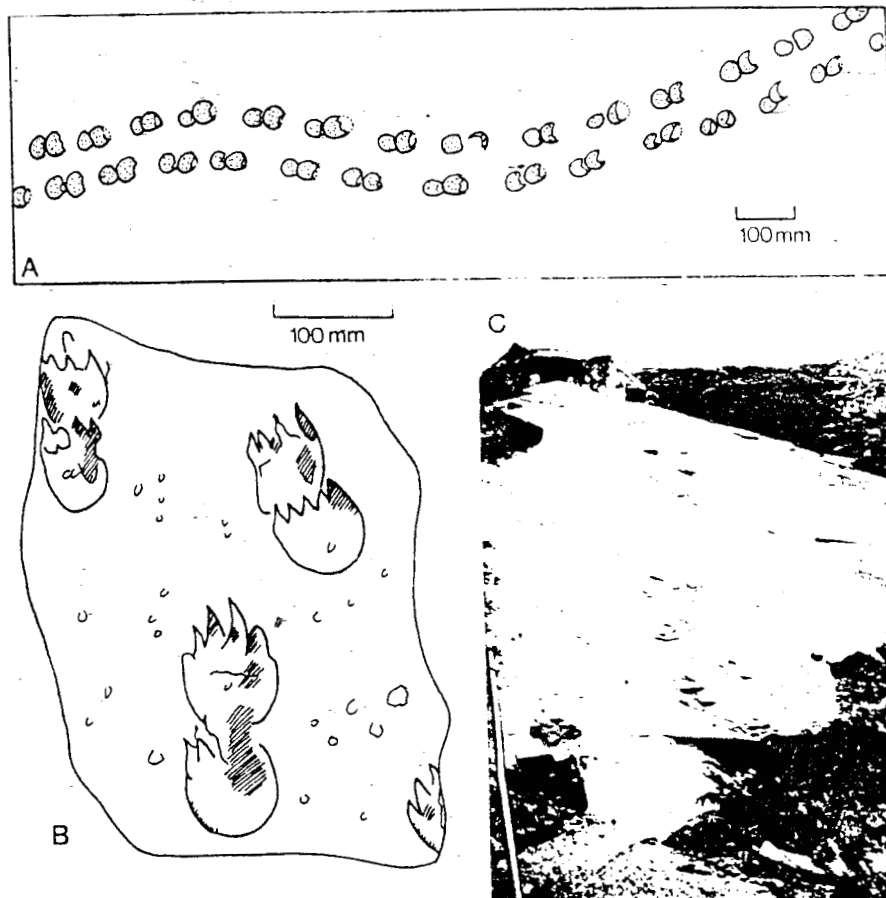


Figura 2

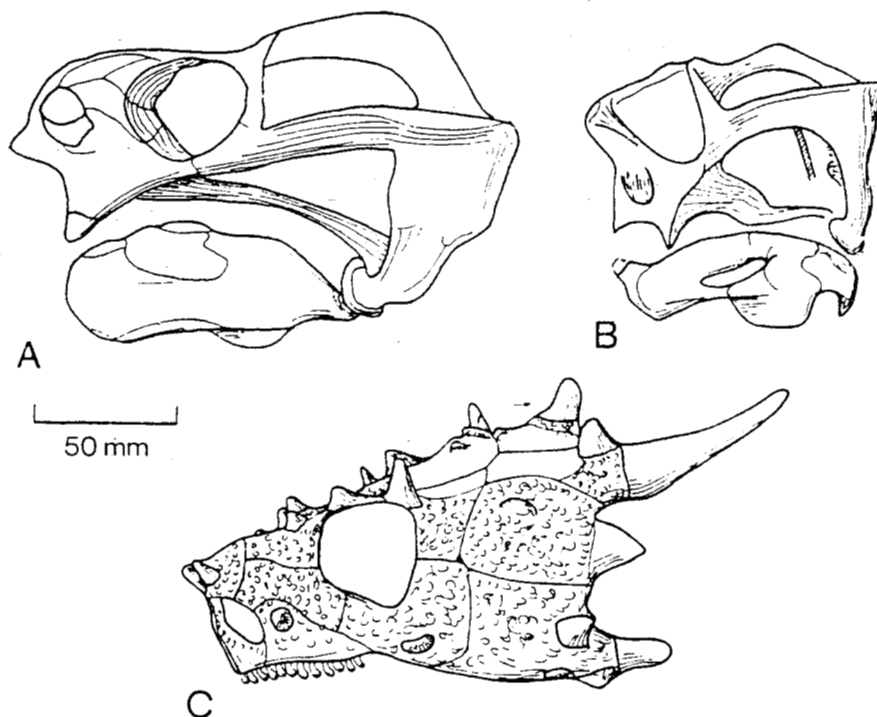


Tabla 1.

SUBCLASS ANAPSIDA

Family Procolophonidae Cope 1889

Leptopleuron lacertinum Owen 1851 (objective junior synonym, *Telerpeton elginense* Mantell 1852)

SUBCLASS DIAPSIDA

Cohort Archosauromorpha Huene 1946

Superorder Rhynchosauria Osborn 1903

Family Rhynchosauridae Huxley 1887

Hyperodapedon gordonii Huxley 1859 (subjective junior synonym, *Stenometopon taylori* Boulenger 1903)

Superorder Archosauria Cope 1869

Order Thecodontia Owen 1859

Family Stagonolepididae Lydekker 1887

Stagonolepis robertsoni Agassiz 1844

Family Ornithosuchidae Huene 1908

Ornithosuchus longidens (Huxley 1877) (subjective synonyms, *Dasygnathus longidens* Huxley 1877;*O. woodwardi* Newton 1894; *O. taylori* Broom 1913)

Family Erpetosuchidae Watson 1917

Erpetosuchus granti Newton 1894

Family Scleromochliidae Huene 1914

Scleromochlus taylori Woodward 1907

Order Sauria Seeley 1888

Family *Brachyrhinodon* Huene 1910*Brachyrhinodon taylori* Huene 1910

Cohort Lepidosauromorpha Benton 1983

Superorder Lepidosauria Haeckel 1866

Order Sphenodontia Williston 1925

Family Sphenodontidae Cope 1870(?)

Brachyrhinodon taylori Huene 1910

Figura 3

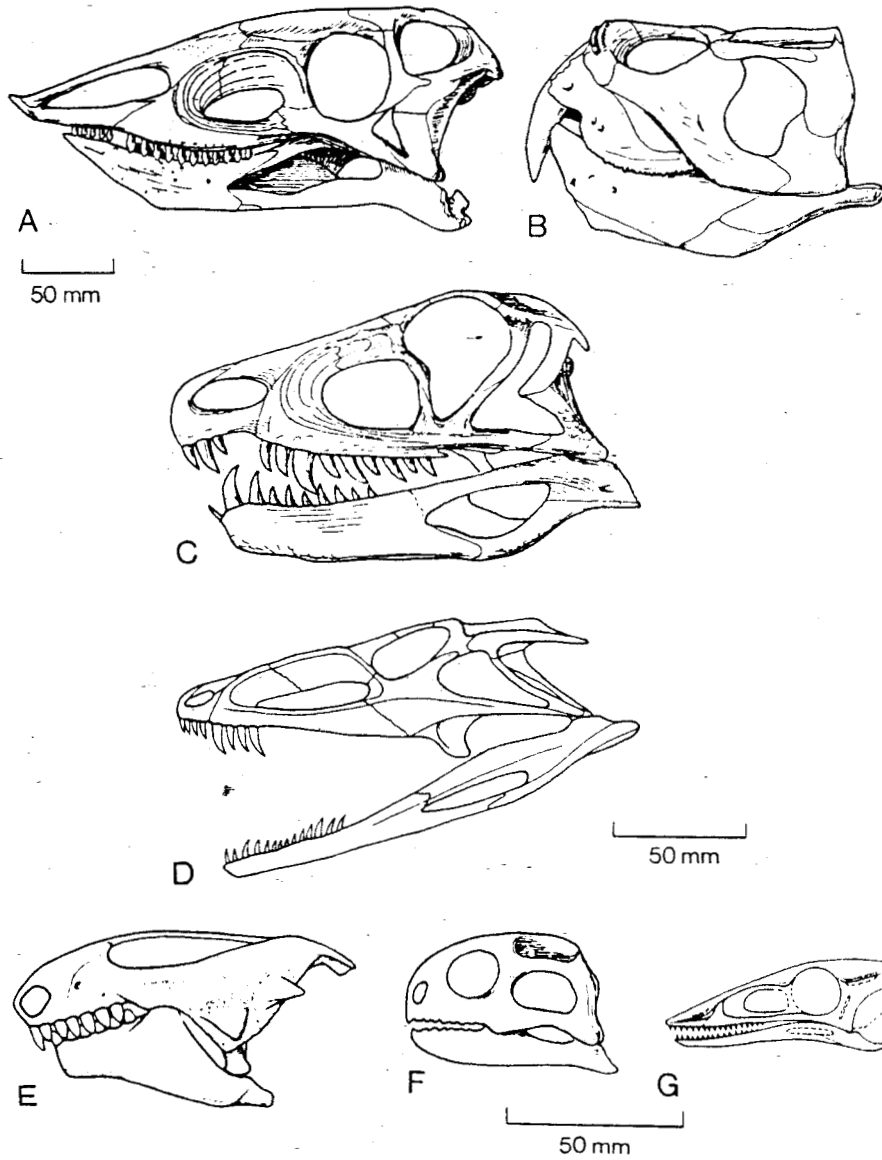


Figura 4

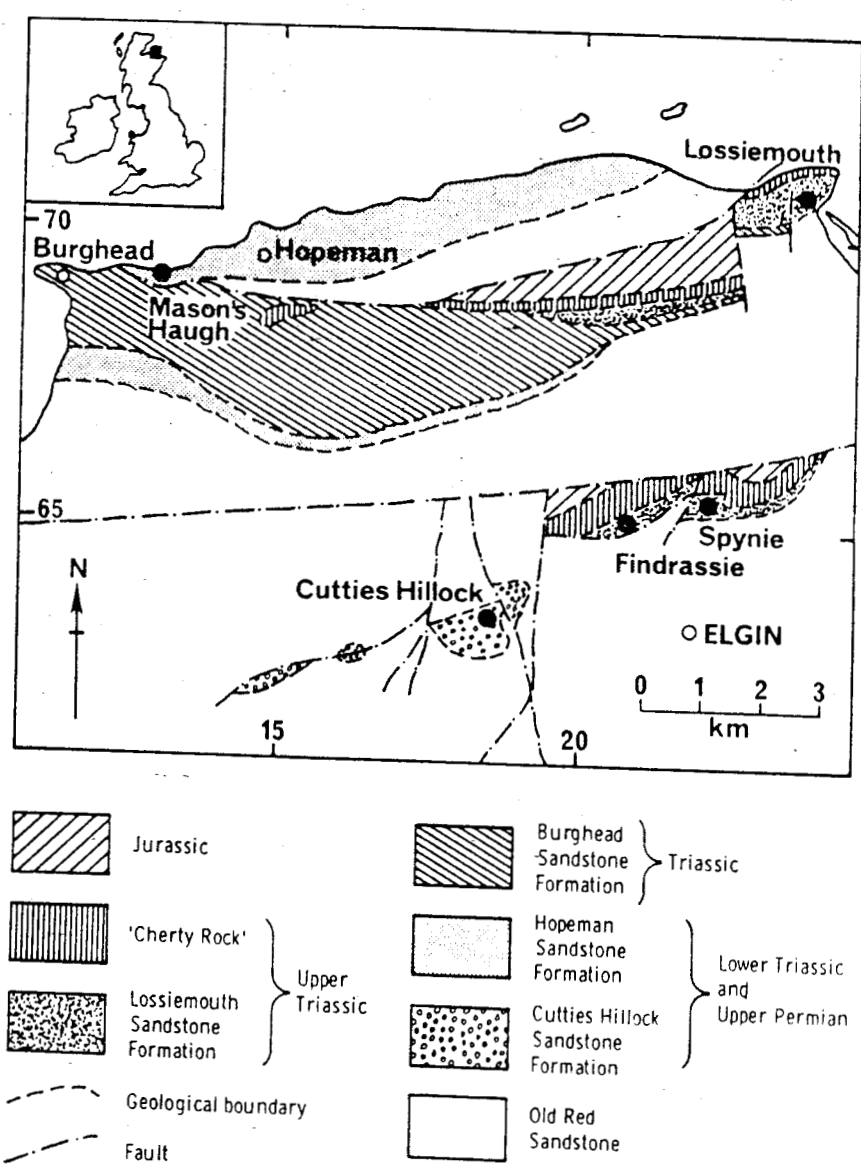


Tabla 2

	Body length	Approx. number of individuals
Herbivores		
<i>Stagonolepis</i>	2.1-2.7 m	30(+)
<i>Hyperodapedon</i>	1.0-1.5 m	35
Carnivores		
<i>Ornithosuchus</i>	1.0-3.7 m	12
<i>Saitopus</i>	600 mm	1
Carnivore, insectivore		
<i>Erpetosuchus</i>	2700 mm	2/3
Small omnivores		
<i>Leptopleuron</i>	110-250 mm	230
<i>Brachyrhinodon</i>	150 mm	211
<i>Scleromochlus</i>	200 mm	7
		<hr/> 128

The classification of the Permo-Triassic rocks around Elgin, using the new terminology (Warrington *et al.* 1980), and as recommended here, is:

Cherty Rock (?Norian-Rhaetian)

Lossiemouth Sandstone Formation (Carnian-Norian)

(Synonyms: Reptiliferous Sandstone (pars) (Symonds 1860; Harkness 1864; Judd 1873, 1886a, b; Gordon 1892; Traquair 1895; Mackie 1897; Boulenger 1903); Elgin Sandstone (Newton 1894); *Stagonolepis* Beds (Boulenger 1904; Huene 1910a); *Stagonolepis* Sandstone (Watson 1909b; Huene 1910b, 1912a, b, 1913, 1914); Sandstones of Lossiemouth, Spynie, and Findrassie (Westoll 1951); Sandstones of Spynie, Lossiemouth, and Findrassie (Peacock *et al.* 1968)).

Burghead Sandstone Formation (?Anisian-Carnian)

(Synonyms: Burghead Sandstones (Westoll 1951); Burghead Beds (Peacock *et al.* 1968)).

Hopeman Sandstone Formation (Late Permian)

(Synonyms: Sandstones of Cummingstone (Huxley 1859b, 1877); Cummingstone Beds (Hickling 1909; Watson 1909b; Watson and Hickling 1914); Reptiliferous Sandstone (pars) (Symonds 1860; Harkness 1864; Judd 1873, 1886a, b; Traquair 1895); Sandstones of Cutties Hillock and Hopeman-Cummingstone (pars) (Westoll 1951); Sandstones of Cuttie Hillock (Quarry Wood) and Hopeman (pars) (Peacock *et al.* 1968)).

Cutties Hillock Sandstone Formation (Late Permian)

Reptiliferous Sandstone (pars) (Judd 1886a, b; Traquair 1895; Mackie 1897); Elgin Sandstone (Gordon 1892; Newton 1893); Cutties Hillock Beds (Watson 1909b; *Gordonia* Beds (Boulenger 1904); *Gordonia* Sandstone (Huene 1913); Sandstones of Cutties Hillock and Hopeman-Cummingstone (pars) (Westoll 1951); Sandstones of Cutties Hillock (Quarry Wood) and Hopeman (pars) (Peacock *et al.* 1968); Cuttie's Hillock Sandstone (Walker 1973)).

Figura 5

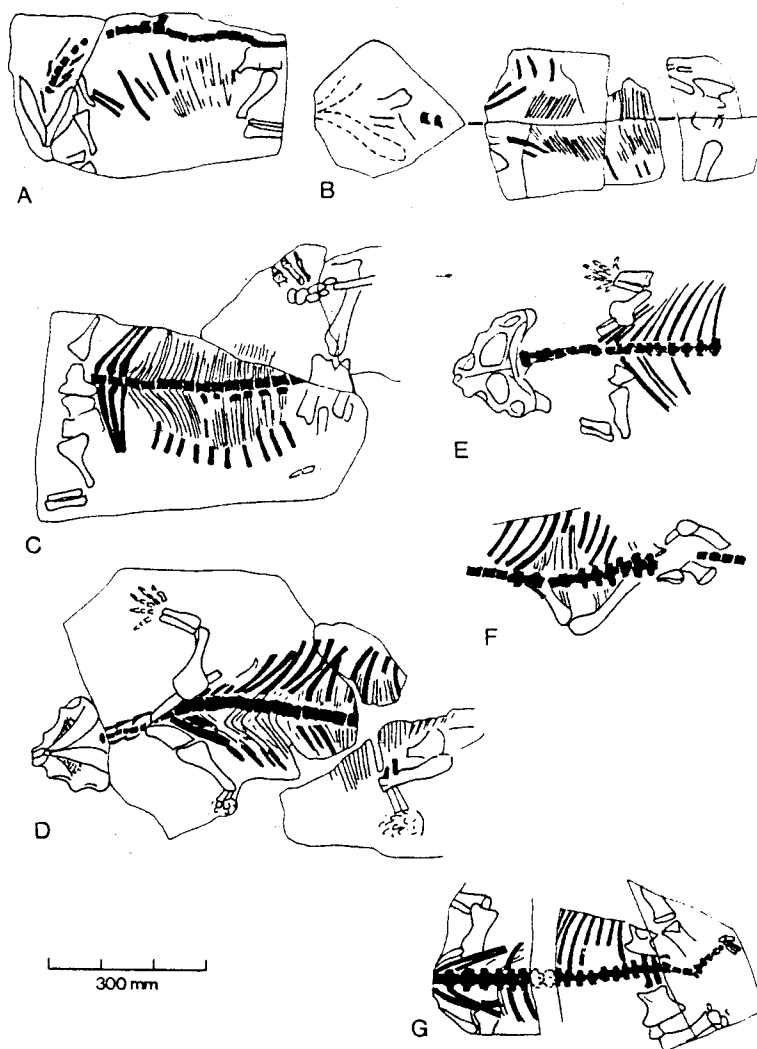


Figura 6

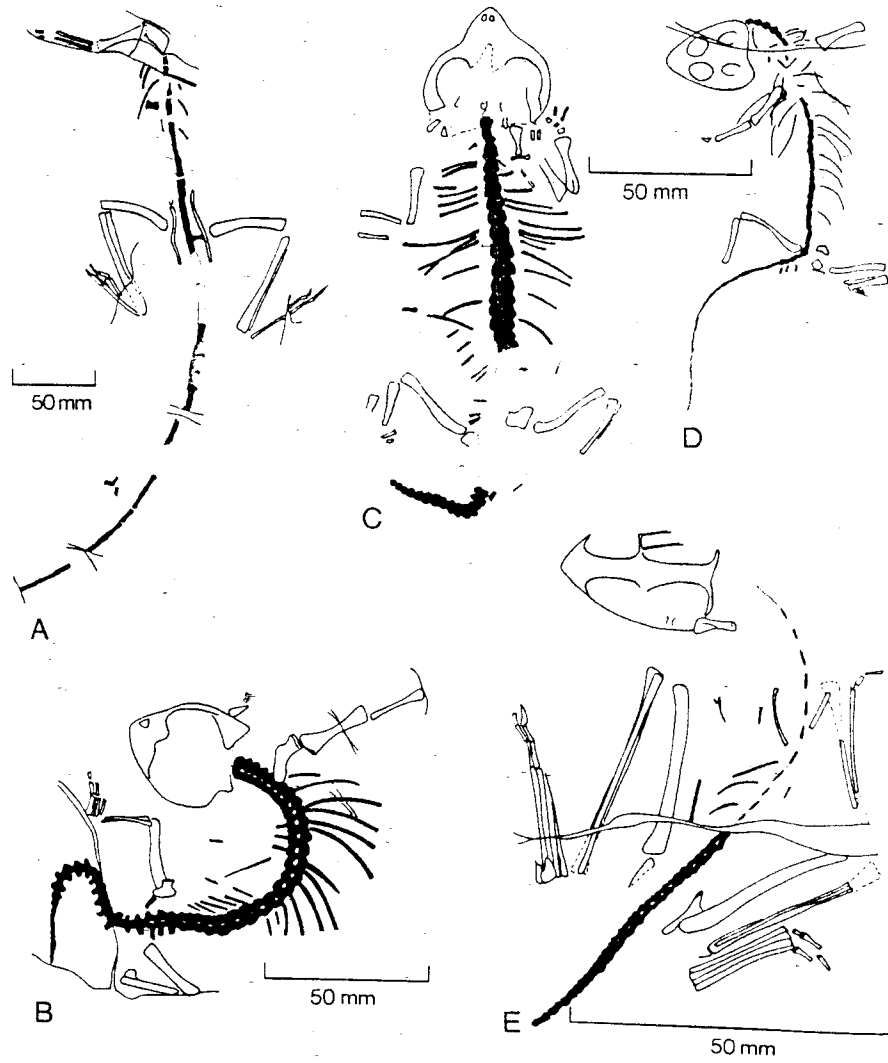


Figura 7

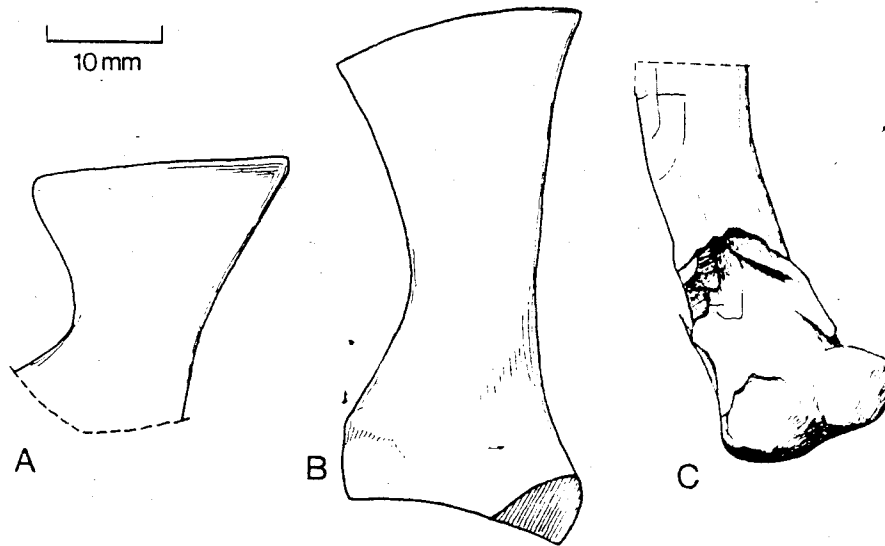
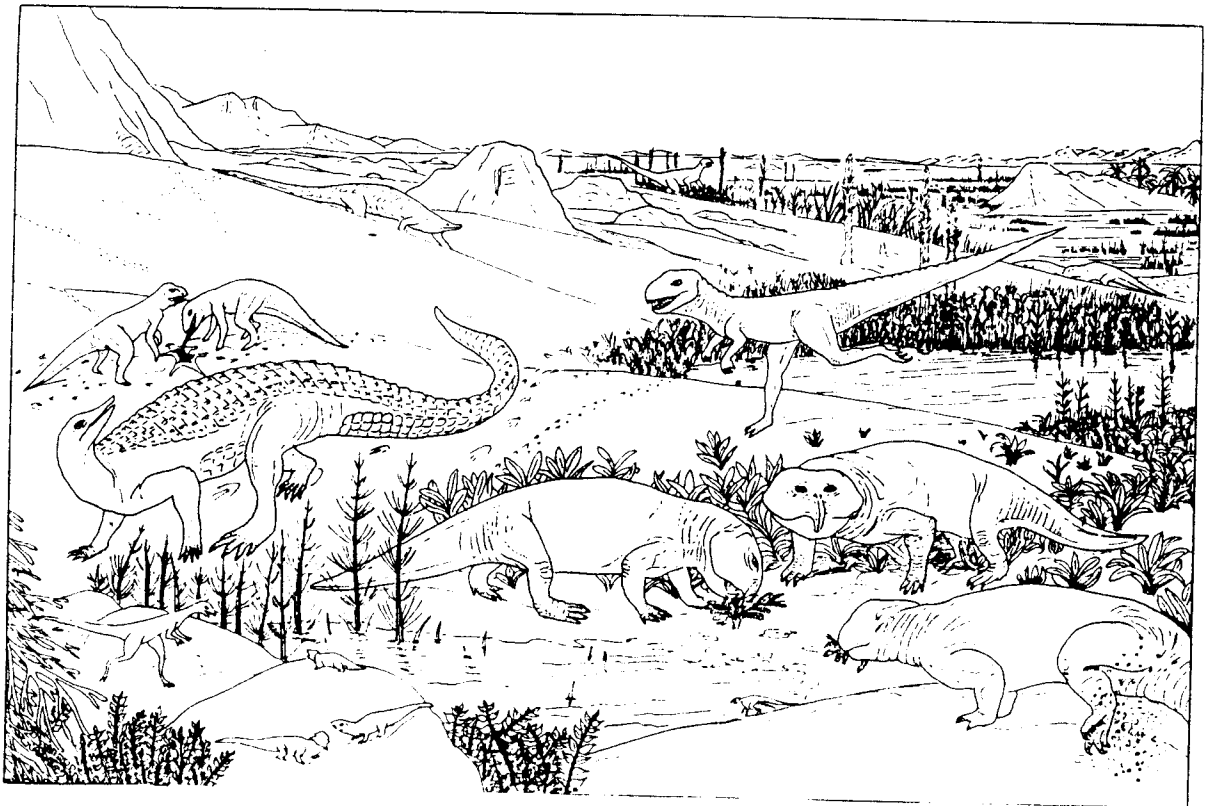


Figura 8



in PADIAN, K. & CHURE, D.J., CONVENERs, 1989. THE AGE OF DINO SAURS, SHORT-COURSES IN PALEONTOLOGY NUMBER 2: KNOXVILLE, TENN., THE PALEONTOLOGICAL SOCIETY, p. 7-21

8.6 (b)

THE ORIGIN of DINOSAURS and the BEGINNING of the AGE of DINOSAURS

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INTRODUCCION

What are dinosaurs? what fauette characterize them? Who were their closest relatives? What were the times like when the dinosaurs first evolved? How did their appearance change the balance of diversity in terrestrial ecosystems? During the past decade we have come to know much more about these topics than ever before. What follows is only a brief review. For more extensive information, readers are referred to the chapters in Padian (1986a), particularly the Introduction, on questions of the Late Triassic- Early Jurassic transition; and to Gauthier (1984-1986) on the characteristics of dinosaurs and their relations. Because this is meant to be a general account, like the other papers in this book, it is impossible to cite every worker, discovery, or relevant paper in the field, and some personal judgments must be exercised.

THE DINOSAURIA

The first remains of dinosaurs to be recognized as fossils of extinct reptiles were found in England in the 1820s (for good accounts, see Desmond 1975 and Charing, 1979). *Megalosaurus* was a giant carnivore; *Iguanodon*, vegetarian with leaf-shaped teeth and a curious spike for a thumb. Neither animal was preserved and early restorations perhaps understandably made both

cretares quadrupedal (after all, who had ever seen a bipedal reptile?): *Iguanodon*'s spike was a first stuck on its nose. In the following decades several new taxa were added to the problematic giant land-going reptiles of southern England, as well as new discoveries of the unrelated marine forms, the ichthyosaurs and plesiosaurs, and the flying pterosaurs. In 1841 Richard Owen, the consummate genius of Victorian zoology, created a new group of reptiles, the Dinosauria, to encompass the half-dozen forms and to set them apart from other reptiles by virtue of their erect limbs, their pelvises that incorporated at least five sacral vertebrae, and their terrestrial habits. In so doing, Owen also drove another nail into the coffin of the transmutational evolutionists by denying evolutionary progress.

In a brilliant article, Desmond (1979) argued that Owen's purpose was to show that ancient reptiles were more sophisticated in form and physiology than their living counterparts. Ironically, he created a grade of evolutionarily advanced reptiles in order to deny evolution.

The concept of Dinosauria has shifted over the years, and questions of dinosaur monophyly and membership have persisted. It had a common origin and can be diagnosed by a suite of features. Gauthier (1984, 1986) listed some of these shared by the common ancestor of all known dinosaurs (Figures 1 and 2). The vertebral column was highly regionalized, and there were at least three sacral vertebrae. Digits 4 and 5 on the hand were reduced in size and number of phalanges, and the distal end of the first metacarpal (the base of the thumb) is offset so that digit 1 is somewhat opposable to the others. The hip bones have reduced their contact around the socket (acetabulum), so that the socket is at least partly open. The upper rim of the socket is buttressed, and there is a deep channel on the lower part of the pelvis behind the socket that housed the caudofemoralis brevis muscle. This tibia (shin bone) has a crescent-shaped knee crest and is widened at its distal end. Tibia is much more robust than the fibula, the other lower leg bone, much as in humans; and the proximal ankle bones that they contact (the astragalus and calcaneum, respectively) are similarly disparate in size (in crocodiles, lizards, and turtles, they are

more nearly equal because there is more rotation in the lower leg during walking). The astragalus has large ascending process against the tibia. The two proximal ankle bones together form a double roller joint (as on the drumstick of the chicken) for the main articulation of the hinge of the ankle. The metatarsals (sole bones) are long and closely appressed; the foot is functionally tridactyl, and the stance is digitigrade. This is what the first dinosaurs were like.

Some of these characters are shared by the closest relatives dinosaurs, including pterosaurs and the small archosaur *Lagosuchus* (Figure 3B; Bonaparte, 1975), but the unique combination of all these characters is known only in the dinosaurs. Pterosaurs, of course, are highly modified for flight, so it is difficult to know whether they passed through any or all of the dinosaurian characteristics before becoming so modified. *Lagosuchus* and other related forms of the South American Triassic are not yet well enough known to say just how many dinosaurian characters they shared. Clearly all these groups had a common ancestor in the Middle Triassic and pursued closely related evolutionary paths. The available evidence indicates that their common ancestor was a small, agile, lightly built, bipedal archosaurian reptile, predaceous rather than herbivorous, that used its forelimbs (now freed from locomotion) for a variety of functions. This seems to have been the great invention of the dinosaurs. Many reptiles of today, including crocodiles and lizards, can achieve an erect stance when they want to. In its 'high walk,' for example, the crocodile pulls its hind limbs underneath the body and proceeds, instead of sprawling. These reptiles are versatile in this respect.

But dinosaurs had no choice: their hip and joints tell us that they had to stand fully erect, limbs tucked under and moving under the body. This suggests two things: first, that they had the energetic capability to be habitually erect; and second, that they were using their hands for other things, as their offset thumbs attest. Just what those uses were is not entirely clear; seizing prey and manipulating objects come to mind. But one could reasonably infer that some selective force must have contributed to the switch from the energetically less costly and more versatile stance

retained by other reptiles. It also must have played an important role in respiration, and thereby in the metabolism necessary to sustain activity, as Carrier (1987) has elegantly shown.

I will return to the relationships of dinosaurs to other Mesozoic reptiles after explaining a bit about the time period that witnessed their origin.

THE AGE OF DINOSAURS

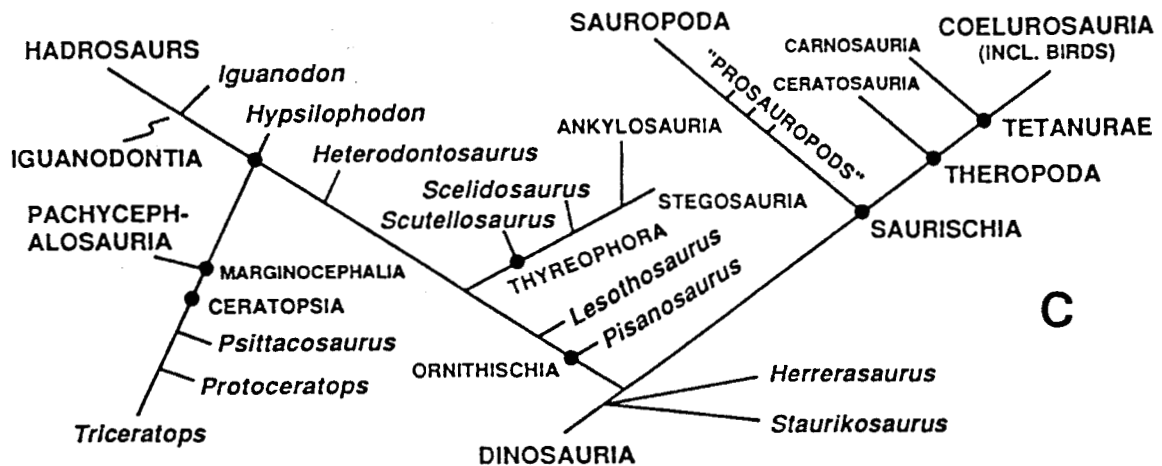
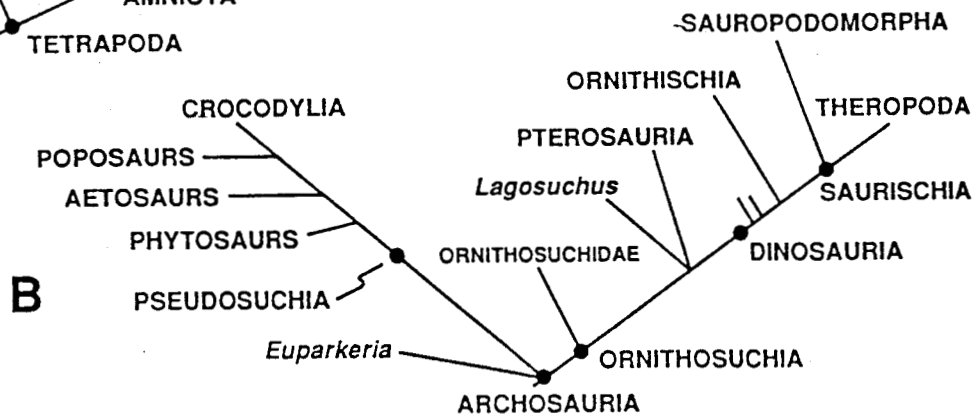
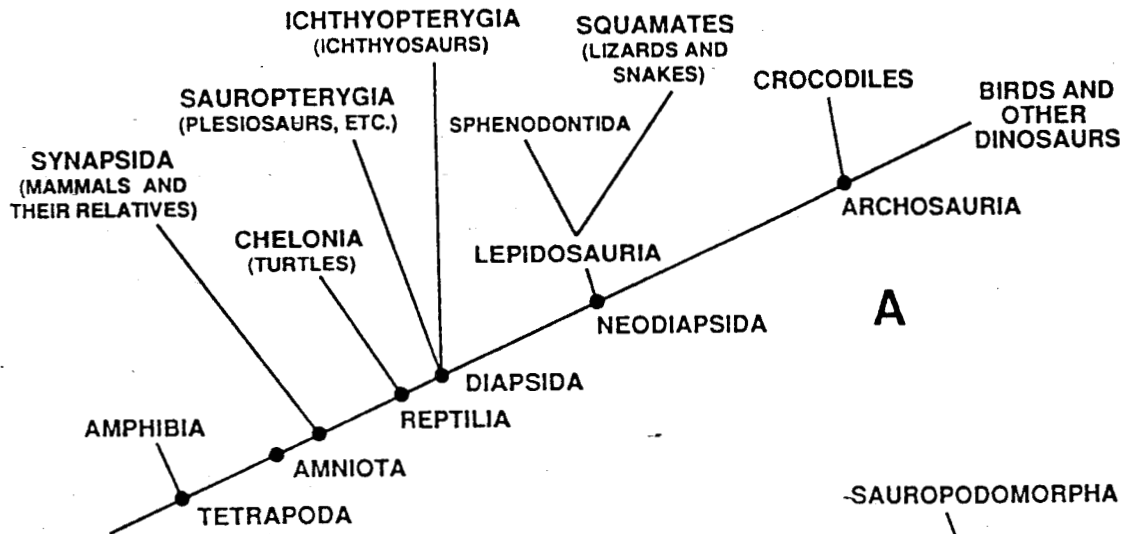
The first known dinosaurs appeared in the latest Middle Triassic or earliest Late Triassic of South America, and are known from only a few incomplete skeletons (*Staurikosaurus*, *Herrerasaurus*, *Ischisaurus*, *Pisanosaurus*, and *Trialestes*; see Gluy 1982 for illustration of some of these). *Pisanosaurus* appears to be an ornithischian dinosaur but the rest are neither ornithischians nor saurischians proper, as Gauthier (1984) showed. There are a couple of apocryphal Middle Triassic records of dinosaur footprints (Haubold 1986), and certainly there must have been dinosaurs at some point in the Middle Triassic, but it is really in the Carnian and Norian that the first good records of dinosaurs occur, and Africa (Padian 1986b). The "Age of Dinosaurs," therefore, does not precisely coincide with the Mesozoic Era, because the dinosaurs got a late start.

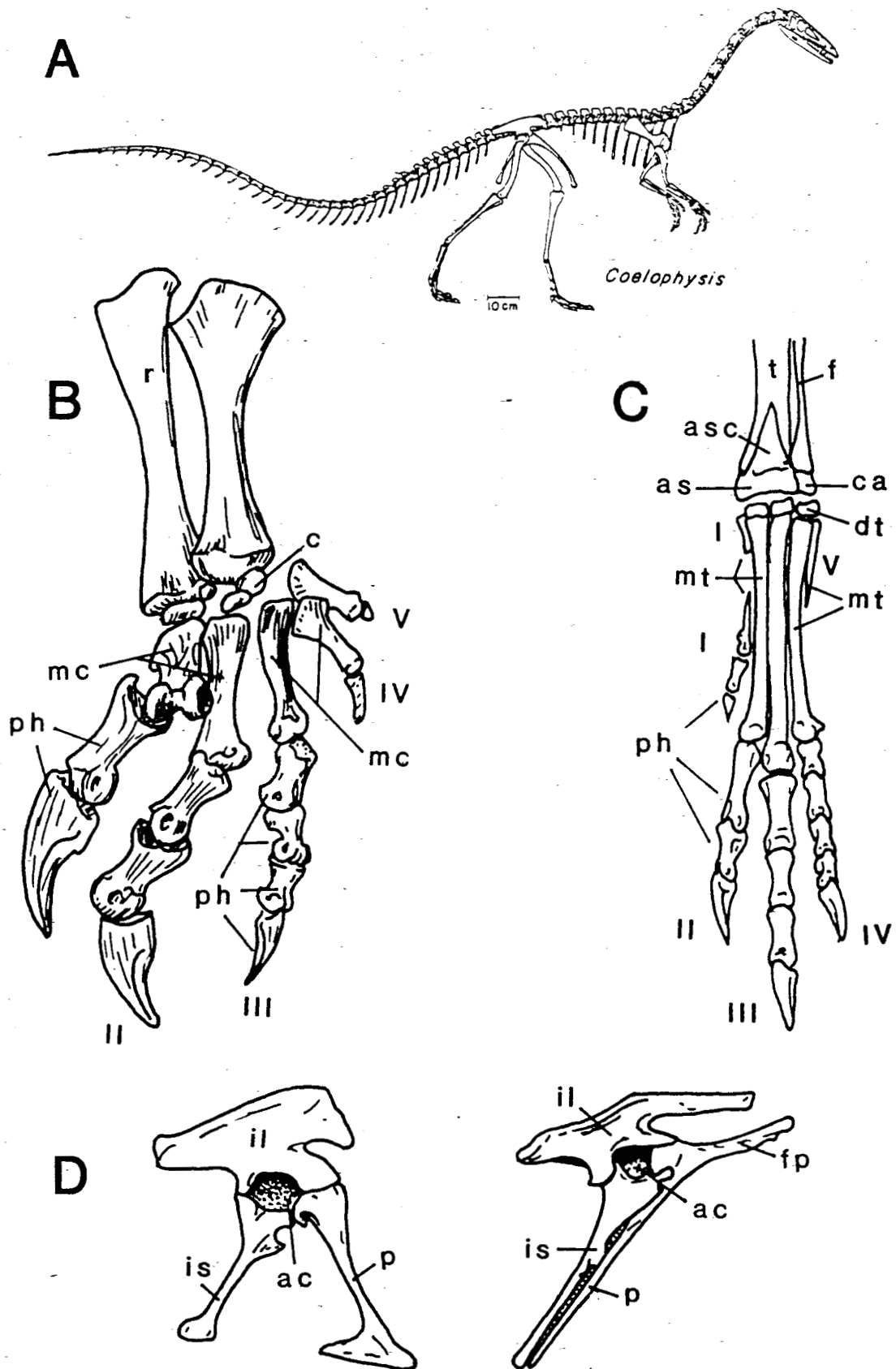
Moreover they did not disappear by the end of the Cretaceous. As the chapter in this book by Gauthier and Padian demonstrates, the birds are descended from small carnivorous dinosaurs, and so by the criterion of morphology are formally to be considered dinosaurs (see also Gauthier, 1984, 1986; Gauthier and Padian, 1985).

The Triassic was an exceptionally busy time in vertebrate history (Padian 1986a, pp. 3-4; see also Padian and Clemens, 1985). More recognized taxonomic orders of tetrapods lived at one point or another during the Triassic than at any other time in vertebrate history, with the exception of the Tertiary (the explosion of mammals and the addition of the taxonomically inflated birds does this). This is because Triassic terrestrial life really consists of three major faunal elements: (1) group that survived from earlier periods; (2) group that lived only during the Triassic, and (3) group that

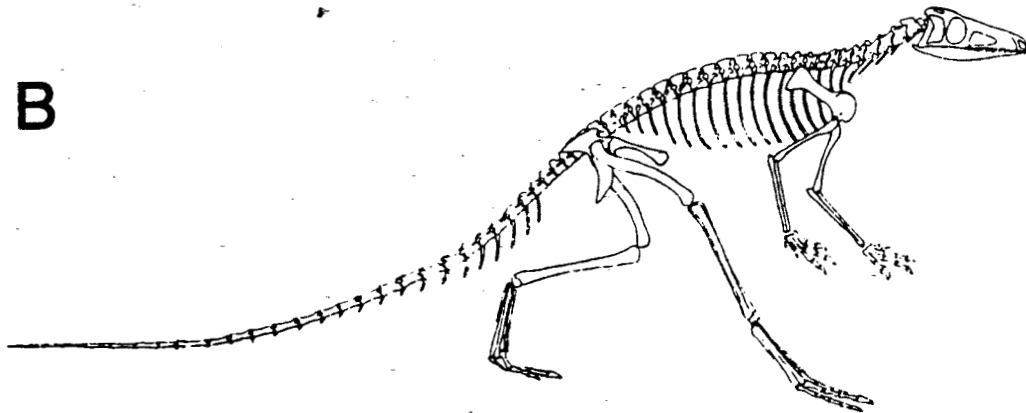
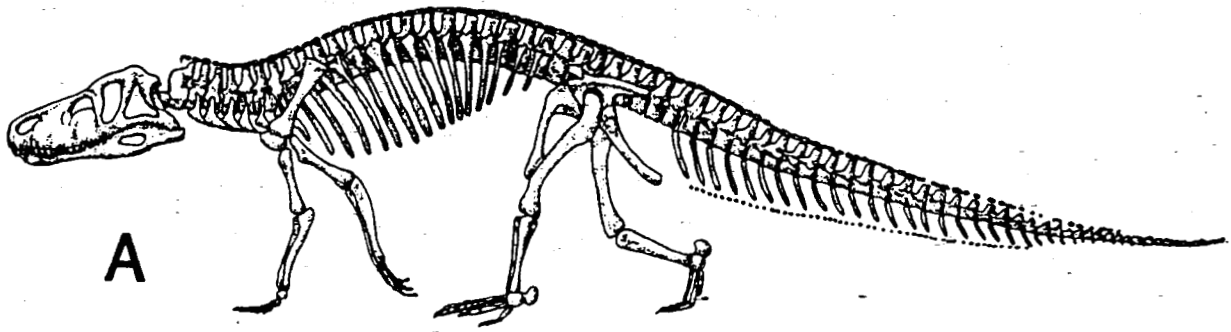
appeared at the end of the Triassic but reached their greatest diversities in later periods.

The first group which I call "Paleozoic holdovers" comprises mostly large, flat-skulled amphibians that prowled the water ways, plus the therapsid relatives of mammals (often misnamed "mammal-like reptiles" see below). These groups were on the wane and had nearly all disappeared by the end of the Triassic. The second group includes tetrapods, mostly reptiles, that arose and then died out during the Triassic, in most cases the Late Triassic. Many of their names are not generally familiar but they include metroposaurs, phytosaurs, eosaurs, rauisuchids, poposaurs, lagosuchids, triphosaurs, rhynchosaurs, and tanystropheids [Some of these will be encountered later; skeletons and restorations of many of them can be found in Colbert (1972-1985), Charig et al. (1976) Carroll (1987), and Long and Houk (1989).] The reasons suggested for their demise range from competition with the dinosaurs (not readily demonstrable, and anyway dinosaurs were few in number and diversity back then) to ecological catastrophe (difficult to see what could have affected them that would have spared the dinosaurs) to simple luck of the draw (possible, but a non explanation in that respect).





Kevin P.



In: PADIAN, K. & CHURE, D.J., CONVENER, 1989.
THE AGE OF DINOSAURS, SHORT COURSES IN
PALEONTOLOGY NUMBER 2: KNOXVILLE, TENN.,
THE PALEONTOLOGICAL SOCIETY, p. 12-133.

The Origin of birds and the Evolution of Flight

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INTRODUCTION

One of the most salient advances in vertebrate paleontology in recent years has been the settling of the question of the origin of birds, a problem that has vexed evolutionary biologists since well before Darwin. To be sure, the consensus is not unanimous, and many details of this branch of the phylogenetic tree are yet to be worked out, but we now have a much clearer picture of this problem than we had a decade ago. Less settled, but equally stimulating, has been the controversy over the origin of flight in birds and other flying vertebrates. Was there a gliding stage? Did flight begin from the ground up or from the trees down? Were birds initially arboreal? What selective pressures drove the ancestors of birds to take advantage of the aerial opportunity?

The overarching interest in such questions, it is that they get at the heart of the study of "macroevolution," in its fullest sense. Macroevolution can mean the origin of "major groups" of taxa, such as the birds. It can also mean the origin of "major adaptations" as they are assembled, piece by piece, through the phylogenetic sequence. Fortunately, new evidence on these questions and new methods of analysis have come to predominate scientific work of the past decade. They have brought us to the point where the origin of birds, once the most problematic of evolutionary "missing links," is now probably the best resolved major evolutionary transition known in all of paleontology.

New evidence has come mostly from the discovery of new theropod dinosaurs of the Mesozoica, including many very closely related to the first birds. Among new methods of analysis, certainly the use of phylogenetic analysis (cladistics) has had the biggest impact, for two major reasons. First, it insists on setting out taxa that are diagnosed only by shared derived characters, and linked with other taxa only on the basis of such characters. Second, it forms a phylogenetic framework against which other questions of the early history of birds and their adaptations must be measured. After all, there is no limit to the number of hypotheses one can generate about an evolutionary event, but there is only one true history. Working with the best supported phylogenetic hypothesis constrains to some degree many of the limitless possibilities of adaptive and other historical explanations. It also provides the best opportunity for evolutionary understandings that are based on a concordance of independent lines of evidence.

The various questions surrounding the origin of birds and flight are reviewed in the volume of proceedings of the first International Archaeopteryx conference, held in Eichstätt, West Germany, in 1984 (Heech, Ostrom, Viohl, and Wellnhofer, 1985). [Copies of this book are still available, and at a generous discount, from the the Freunde des Jura-Museum, Willibaldsburg, D-8078 Eichstätt.] This book is a very good compendium of recent work on these questions, with extensive and historical reviews; for the sake of economy we will tend to these later papers, to which readers may turn for more background on the questions. Another source of information is a volume of proceedings of a AAAS symposium held at the California Academy of Sciences in June, 1984 (Padian 1986; copies available from the Publications office of the Academy).

In the following pages we provide a review of some of the major advances in the study of the origin of birds and the evolution of flight and some comments on other related questions.

THE ORIGIN OF BIRDS

By the late 1860s, T. H. Huxley had deduced that dinosaurs were the closest relatives of the birds

(Huxley, 1868, 1870), on the basis of extensive examination of large living ratite birds and comparison to Mesozoic theropod dinosaurs (review in Desmond, 1975; Gauthier, 1986). He presented his conclusions to the Geological Society of London in 1869, providing a list of 35 characters to support this relationship. In the audience was H. G. Seeley, an accomplished if maverick paleontologist who had made his reputation the unlikely combination of South African dicynodont synapsids and British pterosaurs; he would later (1888) sunder Owen's (1841) concept of Dinosauria by proposing the two component orders Ornithischia and Saurischia, based mainly on hip structure.

Seeley, who had a great range of largely self taught knowledge, was known for expressing the odd and heterodox view that was prone to put in fly in the ointment. On this date his reaction to Huxley's lecture was no exception. Seeley thought it possible that the peculiar structure of the hinder limbs of the Dinosauria was due to the functions they performed rather than to any actual affinity with birds. That is to say, why could not the similarities be convergences, inasmuch as these dinosaurs and birds both walk on their hind limbs? With these words, Seeley effectively neutralized Huxley's hypothesis of relationships and his list of 35 characters, without providing a single character of his own nor any alternative hypothesis of relationship. Today we would take a different tack. We would say that convergence has to be demonstrated by proving an alternative hypothesis that more parsimoniously explains all the available evidence. This has been the primary problem with all hypotheses of relationships of birds (Gauthier, 1984, 1986; Gauthier and Padian, 1985). It is less question of evidence than of method.

Regardless of the merits of Huxley's or Seeley's arguments, the fact is that Huxley's idea was bypassed by history. Perhaps this should not surprise us. In Victorian Era it was common for anttransmutationists such as Richard Owen to invoke convergence as an explanation of why groups that supposed to be taxonomically distinct had certain morphological similarities: these were seen as necessary to function or development. It is indeed difficult to counter this point of view, un-

less one insists that convergence be demonstrated, not merely asserted. Although we like to think that the transmutationists won the battle over evolution in the 19th Century, in fact Owen's archetypal transcendentalism has had much more influence on biological education. If you find this hard to accept, pick up your child's life science textbook and see whether its presentation of biology accords more with Darwin's sweeping evolutionary view or with Owen's essentially static, topological comparative approach.

In the following decades, candidates for the closest relatives of birds included pterosaurs, ornithischian dinosaurs, mammals, thecodontian archosaurs, crocodylomorphs, and theropod dinosaurs (not to mention a variety of less likely candidates). Of these groups, pterosaurs are indeed birdlike in many respects (Padian, 1983, 1985) but their wings are formed of completely different morphological structures, as are those of bats. Ornithischian dinosaurs, like birds, have a pubis that has migrated backwards to lie alongside the ischium, but few other birdlike features that are convergent or shared or shared by most archosaurs. A bird-mammal link has been recently resuscitated by Gardiner (1982), but effectively falsified by Gauthier, Kluge, and Rowe (1988a), among other respondents.

Heilmann (1926), in a book of seminal influence on all later workers, reviewed evidence in favor of these various possible relationships, using anatomy, development, integument, stratigraphy, and many other lines of evidence. He, too, approached the conclusion that small carnivorous dinosaurs were the most relatives of birds. But he drew back from this conclusion because these Mesozoic theropods apparently lacked clavicles, the bones that represent the furcula (wishbone) of birds. Heilmann reasoned from a rather literal interpretation of Dollo's law that structures once lost in evolution could not be regained, and so he rejected the theropod ancestry of birds. Instead, he argued, the origins of birds must be from a more distant archosaurian relative, and he pointed to the thecodontians as the most likely source of candidates. This is really a failsafe argument: while rejecting all specific candidates for ancestry, one can throw an evolu-

tionary origin farther back into the obscured lower branches of the phylogenetic tree. And like the argument from the asserted convergence, Heilmann's won favor with biologists. Until nearly the present day it has been the standard textbook story.

Since the early 1970s, three different taxa have been advanced as the principal candidates as the closest relatives of, and ultimately the phylogenetic source of, Archaeopteryx and the rest of the birds. These are (1) crocodylomorphs; (2) thecodontians; and (3) theropod dinosaurs.

The crocodylomorph hypothesis.-- The principal exponent of this view has been Dr. Alick Walker, who based his reasoning on a series of extremely careful and detailed studies of the skull (especially the braincase and ear region) of an early crocodylomorph from the Triassic-Jurassic boundary sediments of South Africa, *Sphenosuchus*. There is no doubt that the early crocodylomorphs were much more gracile and lightly built than their extant counterparts, and they show a number of adaptations for an active lifestyle reminiscent of similar features in birds and dinosaurs (see discussion of early crocodylomorphs in Chapters 2 and 14). Unfortunately, until very recently there was very little available skull material of small carnivorous dinosaurs that could also be compared to the skulls of Archaeopteryx (which is not rich in such details) and extant birds. So a full comparison could not be made.

Walker was followed in his view by L.D. Martin and K. Whetstone (see e.g., Whetstone and Martin 1979, 1981) who argued on the basis of characters in the teeth, tarsus, and braincase that birds and crocodiles shared a very close common ancestor to the exclusion of other groups. However, no specific taxon for closest relationship was proposed, and the majority of hypothesized synapomorphies were eventually shown to be convergences or more generally distributed than these authors suggested (Gauthier, 1986: 3-4). In 1985, Walker criticized Whetstone's (1983) interpretation of the braincase of Archaeopteryx, and in the light of new evidence rejected his previous view of a close crocodile-bird relationship in

favor of the theropod hypothesis (Walker, 1985; Gauthier, 1986).

The thecodontian hypothesis. Following Heilmann's reasoning, the principal advocates of this view in recent years have been S.F. Tarsitano and M.K. Hecht (e.g., 1980). Again, without pointing to any specific thecodontian, they have made the case that the birdlike characters of all known Mesozoic theropods are too specialized or too late in time to qualify these dinosaurs as the stem group from which birds evolved. And indeed there are several small, lightly, agile-looking Triassic archosaurs from which one could derive birds by further transmutation of the skeleton. (However, we would agree with P.E. Olsen [pers. comm.] that one would really have to go through a theropod to do so.) The problem is that every statement of the "thecodontian" hypothesis has had to rely on an "unknown thecodont" (Tarsitano and Hecht, 1980). Certainly, complete knowledge of all fossil forms will never be available, and we must be mindful of the incompleteness of our knowledge. On the other hand, an appeal to that ignorance as the incompleteness of our knowledge. On the hand, an appeal to that ignorance as the basis of a phylogenetic or anti-phylogenetic statement appear to restrict severely any attempt to reconstruct evolutionary history. An additional difficulty for the "thecodontian" hypothesis is the concept of "thecodontians" itself has come under heavy criticism in recent years. As Gauthier and Padian (1985) stressed, "Thecodontia has been a convenient term for archosaurs that are not crocodiles, pterosaurs, dinosaurs, or birds, but its use has obscured relationships more than clarified them. In phylogenetic analysis, this admittedly paraphyletic grouping is useless." Instead, Gauthier (1980; summarized in Gauthier and Padian, 1985) used morphological synapomorphies to show that the supposedly "basal archosaurs" usually called thecodontians could be neatly divided into those that were closer to crocodiles (*Pseudosuchia*; *Crocodylotarsi* of Benton and Clark, 1988), closer to birds (*Ornithosuchia*; *Ornithotarsi* of Benton and Clark, 1988), closer to birds common ancestry of birds and crocodiles (*Archosauria sensu stricto*).

The "thecodontian" hypothesis thus becomes problematic. With the dissolution of this as-

semblage, it is no longer possible to pick and choose birdlike characters from among a range of taxonomically separated, primitive archosaurs. The "Thecodontia", a group of "Flying Dutchmen" of uncertain phylogenetic relationships that do not belong to any well-defined monophyletic taxon, is not a natural taxon, and its use should be abandoned (Gauthier and Padian, 1985; Padian, 1989b; see Chapter 2). Without any specific hypothesis of relationship, the "thecodontian" hypothesis appears to have no material basis, and reduces to a denial of other hypotheses.

The theropod hypothesis.—Huxley's original idea was based on a far smaller range of dinosaurian material than is available to us today, and he drew his phylogenetic inferences from the whole spectrum of dinosaurs known to him then. In the early 1970s, the small Early Cretaceous coelurosaur *Deinonychus* and its relative provided J.H. Ostrom with a more specific insight into the origin of birds. In a series of papers, Ostrom (1969, 1973, 1975a,b, 1976) used detailed osteological comparisons to show unique features shared only by coelurosaurs and birds, and to demonstrate possible morphological transitions that occurred between these basal coelurosaurs and *Archaeopteryx* and the bird. Gauthier (1984, 1986; reviewed in Gauthier and Padian, 1985; see also Rowe, this volume) used cladistic analysis to arrange Ostrom's hypothesized synapomorphies and dozens of others in a fuller sequence that placed birds squarely within the coelurosaurian, theropod, saurischian dinosaurs.

The conclusion reached by these studies is birds are dinosaurs, not just descended from them, by the principles of group membership that are required by the monophyly criterion of phylogenetic systematics. This realization is an especially effective one to bring to students, who delight that they eat dinosaur at Thanksgiving, have dinosaur in their backyards, and occasionally go out for dinosaur McNuggets. The dinosaurs are not extinct; living dinosaurs are restricted to a fragment of their former morphological diversity, but their extant representatives comprise thousands of living species. Furthermore, following the same principles of group membership discussed above, and reviewing the basic dichotomy of amniotes into

Synapsids (mammals et al.) and Reptiles (see Chapter 2), birds belong to the Reptilia. The living species of birds and other reptiles, in fact, outnumber mammals. So, despite allegations that the Cenozoic is the Age of species diversity the reptiles still have the edge.

Who are the coelurosaurs?—As Gauthier (1986) has redefined them, Coelosauria includes Aves as well as all other tetanurine theropods that are more closely related to living birds than they are to carnosaurs ($Tetanurae = Carnosauria + Coelosauria$). As used here, Coelosauria is a monophyletic taxon, including birds. Coelosaurs have not always been treated in this way, for that name has traditionally been applied to all small theropods aside from birds (e.g., Ostrom, 1969). Recent analyses have demonstrated the inadequacy of dividing theropods on the basis of size alone. For example, it is now clear that the huge carnosaurs are more closely related to birds than are several of the small theropods formerly referred to Coelosauria (Gauthier, 1984, 1986). Moreover, the usually small coelosaurs contain both large forms, such as *Dromornis* (a ratite estimated to weigh some 500 kg) and *Deinonychus* (a giant ornithomimid known only from its eight foot long arms), as well as small forms, such as the pigeon-sized *Archaeopteryx* and the diminutive hummingbirds.

Gaps in the fossil record complicate our understanding of the initial radiation of Coelosauria. The oldest coelosaurs currently known are from the Late Jurassic. Some of the species, such as *Ornitholestes hermanni* and *Elaphosaurus bambergeri*, are known from single, partial skeletons, and others, such as *Coelurus fragilis*, have been reported only from incomplete skeletons. Ironically, the most completely known early coelosaur is *Archaeopteryx lithographica*, and it is already a bird. That is unfortunate because it implies that coelosaurs that are even less closely related to living birds must have diverged still earlier in the Jurassic. Terrestrial sediments of mid-Jurassic age are very rare, and as a consequence we have yet to discover a truly archaic coelosaur. The situation is further complicated because many groups of coelosaurs that diverged before the origin of birds, such as ornithomimids, caenagnathids, dromaeosaurids, and troodontids, are best

known from their highly modified Late Cretaceous representatives, which may be removed from the ancestral coelosaur by as much as 70 million years (see Currie, this volume).

Despite these complications, it is relatively easy to distinguish coelosaurs from other theropod dinosaurs (Gauthier, 1986). For example, the quadrate bone, which suspends the jaw on the skull, is set close to the braincase in birds, with an inner attachment to the prootic bone and an outer articulation with the opisthotic (braincase). The head of the quadrate in other coelosaurs is not so modified, but it is intermediate in form in that the quadrate attaches near the base of the opisthotic, rather than near the outer edge of that element as in other theropods. But the most obvious of the diagnostic characters of Coelosauria are several modifications in the neck, forelimbs, and hindlimbs that are normally thought of as peculiarly avian. In contrast to other theropods, coelosaurs have very long arms and hands, and their long neck bones join another in a distinctively avian fashion, with flexed zygapophyses and incipiently heterocoelus centra. Their hindlimbs are equally diagnostic in having an enlarged ascending process of the ankle that firmly binds it to the shank, and in having compressed the metatarsal bones connecting the toes to the ankle (see Rowe, this volume).

The Manuraptoran Coelosaurs.-- Manuraptora (Gauthier, 1986) is a subgroup of coelosaurs that is even more markedly birdlike. It includes such well-known coelosaurian groups as Caenagnathidae, Dromaeosauridae, Troodontidae, and, of course birds (Norman, 1985; Currie this volume). Manuraptorans can be distinguished from such coelosaurs as ornithomimids by several evolutionary novelties, the most noteworthy of which involve the forelimbs and shoulder girdles. It is clear that the earliest manuraptorans were unable to fly, yet they had most of the modifications that are essential to flight in living birds (Ostrom, 1969).

The hand in living manuraptorans (bird) is reduced and nearly fingerless, with the remaining elements largely fused to one another. But the elements of the hands of early manuraptorans, including early birds, are virtually identical, down

to the proportions of the individual bones comprising each of three (Ostrom, 1976). The prey-seizing hands that arose in the ancestral theropod became finely tuned, raptorial organs in manuraptorans. These huge, grasping hands have long, slender fingers; the shorter thumb converges slightly on the longer outer fingers during grasping, and the third finger has a slender and bowed at its base whose joint surfaces may have enabled that finger to oppose the others as well (Russell and Seguin, 1982). The fingertips are supplied with very large, sharply pointed, trenchantly margined, deeply recurved claws reminiscent of on the prehensile feet of hawks and owls. The wrist is formed by a half-moon-shaped bone that enables the hand to move in a broad arc upon the forelimb (Ostrom, 1969, 1976; Gauthier and Padian, 1985). The forelimbs in manuraptorans are very long, and they are operated by comparatively powerful muscles arising from a prominent breastbone. The clavicles are fused, forming the "wishbone" so characteristic of birds, in order to brace the shoulder against the forces imposed by struggling prey (Barsbold, 1983). And the lower part of the shoulder girdle is modified in such a way as to indicate that a chest muscle that formerly swung the arm down is now beginning to lift it up, thus is now beginning to lift it up, thus providing greater power on the recovery stroke when seizing prey. Such muscle scars as remain on the forelimbs and girdles suggest that the arms could be folded against the body at rest in much the same manner seen in living birds (Ostrom, 1974). Moreover, the entire forelimb and shoulder apparatus operated as in living birds: the arms swung down and forward, then up and back, thus describing the classic figure-eight motion that characterizes the flight stroke in birds and all other animals that fly, whether through the air or in the water (Gauthier and Padian, 1985; Padian 1985; Rayner, 1988). The manuraptoran foot, with its hyperextendable and sickle-clawed second toe, is equally diagnostic (Ostrom, 1969). Living carinid birds have similarly modified feet, with sickle claws that are used for defense as well as for killing prey. More or less prominently developed raptorial second toes are present in Archaeopteryx, Ornitholestes, dromaeosaurids, and

troodontids, and perhaps in caenagnathids. Ornithurine birds (all those post-Archaeopteryx birds with reduced tails) generally lack such a specialized toe, so its presence in some later birds is probably secondary. Manuraptoran feet are often portrayed as didactyl, with the raptorial claw held aloft so as not to blunt its tip during locomotion. However, *Cariama* walks on all its toes and the tip of its deeply arched sickle claw touches the ground, so it seems reasonable to assume the earlier manuraptorans did likewise.

In manuraptorans, the tail is shorter than in other theropods, and it is stiffened by a combination of modified vertebral articulations and (in forms such as *Deinonychus*) a network of stiffened zygapophyseal extensions similar to the ossified tendons of ornithischians (see papers by Norman and Sereno, this volume). Ostrom (1969) suggested that this tail acted as a dynamic stabilizer for the animal when slashing at prey with its feet; in running, it would also provide greater maneuverability, and this function seems to have been retained in birds (in which the function became more aerial than terrestrial).

Although the origin of birds from among coelurosaurs has become a fairly settled issue in the systematic community, there is still much to be known about the specific relationships of the early manuraptoran groups, and which of these is the closest to birds. One problem has been that not all skeletal parts of all manuraptoran forms are known, so detailed comparisons are difficult. As new forms constantly turn up (see Currie this volume), the weight of evidence shifts. Part of the reason is that the group is subject to extensive convergences (e.g., the auditory recess in ornithurines and troodontids, the reversed pubes in Avialae and Dromaeosauridae, the bulbous parasphenoid in Troodontidae and Ornithomimidae, the large sickle-claws in Dromaeosauridae and Troodontidae, and the extensive pneumatization of the skull in troodontids, caenagnathids, and Ornithurae). Moreover, there are problematic age-related characters (such as ossified uncinate and ventral ribs, the calcified sternum, fused metatarsal bones) that make it difficult to tell whether the features shown by an individual specimen are characteristic of its taxonomic group or its ontogenetic stage. For the

present, it is perhaps best to take Currie's conservative, open-minded view of the interrelationships of these theropods (this volume), because the state of our knowledge is rapidly changing with new evidence and new analyses.

THE EARLY EVOLUTION OF BIRDS

Definition of birds.-- Birds have feathers and no other organisms do. This unites birds as a group. On the basis of feathers, *Archaeopteryx* is usually classified as a bird. Over the years there has been dispute about whether it could fly or not, but it was clearly doing something aerial with its has no features that debar it was clearly doing something aerial with its feathers, and it has no features that debar it from ancestry or very close relationship to later-birds. Of course, it is possible that we might find other, non-flying coelurosaurs with feathers, which may have evolved initially for functions other than flight (see papers in Hecht et al., 1985). This raises the question of how to define birds. Let us say that we found a *deinonychosaur*, very close to *Archaeopteryx*, with contour or downy feathers, but no flight feathers, but couldn't fly, and presumably none of its ancestors could. How would we classify this animal? Gauthier (1984, 1986) suggested that the term *Aves* be restricted to the living groups of birds and all the descendants of their common ancestor. This suggestion would provide stability to the definition of *Aves*, regardless of any future fossil discoveries that would fill in the evolutionary gaps and muddy the clean break between birds and other reptiles. However, recognizing that *Archaeopteryx* represents a decisive step toward living birds, and has several crucial avian features such as flight feathers, Gauthier proposed the taxon *Aviales* to include it and all more derived birds. In consequence, *Archaeopteryx* (and other primitive birds like *Hesperornis* and *Ichthyornis*) is a "bird". But not a member of *Aves*. This provides both recognition and stability to the taxon.

Archaeopteryx.-- The six known specimens of *Archaeopteryx lithographica* (Wellnhofer, 1974, 1988a, b; Ostrom, 1975a, b, 1976) represent the oldest winged, or avialan coelurosaurs. They are still the earliest known bird specimens. Molnar (1985) has ably reviewed other early birds or

would-be birds; for further comment see Jensen and Padian (1989) and Padian (1988b), and other papers in Hecht et al. (1985).

For over a century Archaeopteryx has been seen as the "missing link" between reptiles and birds, though until recently it was far from clear which reptiles were being linked. The appearance of Archaeopteryx in the fossil record has been essentially equated with the origins of birds and of flight. However, given the advances of recent years, it is now apparent that Archaeopteryx has far more obvious link with other Mesozoic, non-flying theropods than it does with the living groups of birds. This statement may seem ironic, but it turns out that nearly every characteristic of Archaeopteryx is shared by non-flying Mesozoic theropods, and very few link it to later birds. It is perhaps for this reason that the Eichstatt specimen of Archaeopteryx, which has only the faintest of feather impressions, was initially classified as a juvenile specimen of Compsognathus. Only when Ostrom (1972a) recognized the feather impressions on the Teyler specimen, a discovery that changed its taxonomic assignment instantly from Pterodactylus to Archaeopteryx, did curators in museums all over the world with holdings from the Solnhofen limestones re-examine their collections assiduously -- with the result of the announcement of the "new" Eichstatt specimen, so beautifully described by Wellnhofer (1974).

The cited papers by Ostrom, Wellnhofer, Gauthier and Padian have laid out in detail the evidence that Archaeopteryx retains characters that mostly have a distribution among theropod dinosaurs that indicates a much more ancient origin -- for examples, the fused clavicles, the tridactyl hand reduced first finger and bowed third, the halfmoon-shaped carpal, the calcified breastbones, the long arms and legs, the backwards-pointing pubis, the shortened, stiffened tail, and the reversed hallux on the tridactyl foot. Most of these characters have been explained by evolutionists as adaptations for flight or arboreal life. These modifications may facilitate flight, but it is now clear that their persistence in non-flying theropods requires a more general explanation than the particular role(s) that they currently play in avian biology (Padian, 1985; Gauthier and Padian, 1985; Gauthier, 1986). Most of these char-

acters have been introduced previously, and further explanations of their roles in avian evolution can be gleaned easily from the references just cited and from the papers in Hecht et al. (1985).

Birds after Archaeopteryx.-- In addition to the being the earliest known bird, Archaeopteryx is the one known from the most complete material until one gets to Hesperornis and its flightless, dividing relatives from the Late Cretaceous (Marsh, 1880). In the Late Jurassic and Early Cretaceous there are many fragmentary bird bones, as well as impressions of feathers, on which taxa have been based. Molnar (1985) has thoroughly reviewed most of these taxa and some other would-be middle Mesozoic birds. To his reviewed we append only a few updated comments. Marsh's Laopteryx is probably a pterodactylid (Ostrom, 1986). Palaeopteryx (Jensen, 1981) is a nomen nudum (Jensen and Padian, 1989), and most of the "Dry Mesa" (Uncompahgre) material originally considered avian or avian-like is pterodactylid, with a few probable maniraptoran pieces. Cossosaurus is an aquatic archosauriform (see discussion in Evans, 1988) and has nothing to do with birds.

Olson (1985) has reviewed many occurrences of Cretaceous birds, particularly those supposed to represent extant families, and has rejected every case. Cretaceous birds, it seems, had specializations that converged on many of toes seen in living birds (for example, the loon-like diver Hesperornis and the ternlike Ichtiornis), but in Olson's view the supposed phylogenetic affinities to living bird families are wishful thinking. Particularly important early bird discoveries of the past decade include C.A. Walker's (1981) report of the very unusual Enantiornithes from the Cretaceous of South America, and the report by Sanz et al. (1988) of Early Cretaceous ornithurine birds and feathers from Spain. The latter bird has strong coracoids braced to the sternum, a wishbone with a hypocleidium, a long, fused foot, an expanded pelvis, and a reduced tail with pygostyle.

The presence of a palaeognath bird, Gobipteryx, in the Late Cretaceous sediments of Mongolia (Elzanowski, 1981) indicates that the primary avian clades, Palaeognathae and Neognathae, had already diverged from their last common an-

cestor by the close of the Mesozoic Era (Cracraft, 1986). It follows further that the full suite of evolutionary novelties underlying the level of flight performance fundamental to extant birds was already in place more than 75 million years ago. Even at that early date, the ancestor of Aves could be distinguished from the ancestor of Archosauria by at least 213 synapomorphic modifications in the skeleton alone (Gauthier, 1986). Systematists looking to link birds with other groups, matters were only made worse by the extinction of the few remaining nonavian archosaur group till extant at the end of the Cretaceous. The disappearance of intermediate forms, groups still extant at the end of the Cretaceous.

The disappearance of intermediate forms, together with an astonishing diversification of birds and the ensuing Cenozoic Era, has further accentuated the differences between birds and their nearest living relatives, the crocodiles. We still have large gaps in the early history of birds, especially in the Jurassic and Cretaceous, and because of problems with preservation, we may never have a very complete record of some episodes in early avian history. Using what we have, however, the fossil record is surprisingly good for answering some central questions about the origins of birds and flight.

Before moving to a consideration of flight, the recent announcement of a Triassic bird, to be named "Protavis" by its discoverer, Dr. Sankar Chatterjee, deserves mention. To date this discovery is not only an announcement, and no scientific report on it has appeared. We will not consider here the various claims and counterclaims made about it in the press to date. It is worth noting that occasionally a Triassic or other pre-Jurassic fossil emerges and is reported to knock Archaeopteryx off its perch, and "Protoavis" is the most recent candidate. But the "Protoavis" question, so far reduce to one major issue. We already have a well already have a well documented phylogeny of theropod evolution.

Presumably if it is correctly interpreted, "Protoavis" will fit somewhere on this evolutionary tree. But it does not promise, on basis of claims made so far, to alter this tree in any major way. If the claims are correct, then birds would simply

have a more ancient geologic record than usually thought. Skeptics and supporters alike now need to wait a published report before further comment; already the creationist press has seized upon Dr. Chatterjee's preliminary claims to mock the competence of paleontologists and their inferences about the fossil record.

Final question is Archaeopteryx a forgery, as Hoyle and Wickramasinghe (Hoyle et al. 1985) have claimed? Obviously not, as Rietschel (1985) and Charig et al. (1986) have shown; these claims, of course, is to replace conventional evolutionary theory. The motivation for these claims, of course, is to replace conventional evolutionary theory, which Hoyle and Wickramasinghe reject, with their own notion of "germs from space" periodically bringing new viral material to infect the DNA of Earth's organisms, thus causing bursts of evolution. But then, these workers have also testified that insects are smarter than we are they're just not letting on.

CAPITULO 9 ACTUALIZACION BIOTICA: LA DOMINANCIA DE LOS MAMIFEROS.

9.1 PREAMBULO

Este capítulo se dedica al grupo que durante todo el Cenozoico, y desde luego, en la actualidad, ha dominado el ambiente terrestre, diversificándose ampliamente y explotándolo con una amplitud y diversidad que no tienen paralelo en toda la historia de la Tierra. Se decidió no incluir información sobre productores, esto es sobre plantas, ya que las angiospermas que han dominado la escena desde el Cretácico fueron ya consideradas.

En el primer trabajo, Farish Enkins resume la información sobre el origen del grupo, a partir de terápsidos muy avanzados, discutiendo las modificaciones craneales y postcraneales, así como algunas de la anatomía blanda, que son discernibles en el registro fósil en esta fase de transición, y les da una interpretación morfológico-funcional, básicamente mediante un avance gradual, pero continuo en el control de la homotermia como clave de sobrevivencia en un ambiente arbóreo confinado. Discute también el surgimiento de linajes independientes de mamíferos, y su radiación adaptativa temprana, conducente al surgimiento de numerosas líneas paralelas. Cabe destacar que durante dos terceras partes de la duración geológica del grupo desde el Tríasico Temprano al Reciente los mamíferos constituyeron un grupo subordinado, "marginado" por los dinosaurios en el Mesozoico, y es al extinguirse estos, que los mamíferos rápidamente adquieren esa dominancia ecológica, que tienen al presente.

En el segundo trabajo Dave Krause describe la evolución mamíferiana durante el Paleoceno, primera de las épocas o subdivisiones formales del Cenozoico, y que testifica la radiación adaptativa de los "nuevos amos" del dominio terrestre, que les permitió ocupar ampliamente el ecoespacio dejado por los dinosaurios. Esta radiación produjo una mastofauna arcáica, esto

es, constituida por organismos diferenciados desde el Cretácico y que acusaron cambios mínimos en el Paleoceno, o bien, organismos que aparecen en esta época, pero cuyo linaje se extingue en el Terciario Temprano. La diversidad alcanzada por esta fauna es considerable, e incluye formas herbívoras, carnívoras, insectívoras y otras; modalidades de locomoción también variados, cursores, corredores, fosoriales, etc. hábitos terrestres, arbóreos; y tamaños desde unos pocos gramos, hasta tal vez 152kg. Todo ello evidencia la rápida ocupación de nichos ecológicos asombrosamente variados, que desarrollaron estos mamíferos, ocurrida en un lapso geológico breve, tal vez de 5 Ma.

En el tercer trabajo, Dave relata algunos aspectos de la historia evolutiva de los mamíferos modernos, usando como apoyo o tema central, lo que él designa como la "sabanización" de los continentes, es decir el surgimiento y extensión del bioma de sabanas (praderas, pampa, estampa, son designaciones de variedades de este bioma), resultado de la tendencia general Cenozoica hacia condiciones de sequedad y enfriamiento creciente. Este fenómeno tuvo repercusiones muy profundas en la evolución de los mamíferos, ya que diversos grupos desarrollaron adaptaciones que les permitían "prosperar" en estas condiciones y aprovechar a la vegetación dominada por gramíneas o a los consumidores primarios, como alimento principal, tal es el caso de los caballos, los elefantes y los roedores cricétidos. Webb también discute aspectos biogeográficos de estos mamíferos, relacionados con procesos de intercambio faunístico; así como patrones de reemplazamiento faunístico.

En el último trabajo, Catherine Badgley resume la información paleontológica disponible sobre la ancestría de nuestra propia

especie, enfatizando los aspectos filogenéticos, así como el escenario geográfico ambiental en que ocurrió el surgimiento del hombre, es decir el marco paleoecológico de la evolución humana.

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9.3

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9.3 (a)

A Survey of Mammalian Origins.

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INTRODUCTION

The origin of mammals, often treated as a discrete but obscure event that took place sometime between the Late Triassic and Early Jurassic, can also be viewed as the product of many transitions—from the early synapsid radiations during the Pennsylvanian and Early Permian through the emergence of placentals and marsupials in the Cretaceous (see Hopson, 1969). A fundamental dichotomy appeared early in the evolution of amniotes; sauropsids (represented today by living reptiles and birds) constituted one lineage, synapsids (which includes the mammalian descendants) the other. Thus, mammalian ancestry may be traced to pelycosaurs that first appear in the fossil record as part of the earliest known reptilian fauna (Carroll, 1982). And mammalian bony and dental structures continued to undergo substantial modification throughout the Mesozoic, long after the appearance of forms technically classified as Mammalia. Given these phylogenetic changes, the suggestion that there occurred a point when mammals "originated" seems simplistic. Yet the major evolutionary stages may still be evaluated with the purpose of identifying the inception of features and functions basic the radiation of modern mammals. Accordingly, this

survey summarizes our current understanding of the evolution of mammals with full acknowledgment that an account of mammalian origins has neither a definitive beginning nor a climactic end.

SYNAPSID EVOLUTION: GENERAL TRENDS.

Several features of the skull of a representative primitive synapsid (the pelycosaur *Dimetrodon* 1; Figure 1A) represent the beginning of two trends in mammalian development: the evolution of the masticatory complex and the middle ear (see Hopson and Barghusen, in press). Typical features of these primitive mammalian predecessors are an expanded maxilla (mx, Figure 1), enlarged caniniform teeth (presaging the dental differentiation of therapsids, and a single temporal fenestra (tf, Figure 1) related to the jaw adductor muscles. Although not all synapsids appear to have been carnivorous, it is clear that a predaceous habit and specialization of the jaws, teeth and associated muscles represented adaptations fundamental to the beginnings of mammalian history. Mammal-like reptiles, especially, elaborated an array of tooth forms and skull shapes virtually unequalled by any tetrapod group save mammals (Figure 2). The two most obvious aspects of the trend are the increased size of the dentary (dn, Figure 1) in the lower jaw, and the expansion of the temporal fenestra to accommodate an enlarged mass of jaw adductors (tf, Figure 1B-D). In synapsids, the postdentary bones established early a distinctive configuration (including a flange, or reflected lamina, of the angular, and a ventrally oriented retroarticular process of the articular) that was to undergo progressive modification in relation to the development of the mammalian middle ear (see MIDDLE EAR, below). Figure 1 Stages in the evolution of the mammalian skull. A, The pelycosaur *Dimetrodon limbatus* 1, after Romer and Price, 1940, plates 6 and 15. B, The gorgonopsid *Lycaenops ornatus* 1, modified after Colbert, 1948, Figure 1.C, The cynodont *Thrinaxodon liorhinus* 1, modified after Hopson, 1969, Figure 8. D. the mammal *Sinoconodon*

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changchiawaensis1, from Crompton and Sun, in press.

ADVANCED MAMMAL-LIKE REPTILES: CYNODONTS.

The latest radiation of mammal-like reptiles, beginning in the Late Permian, gave rise to the cynodonts which, considered together, represent transitional stages in the establishment of definitive mammalian features (Figure 3). Among these the procynosuchids are the most primitive. Procynosuchus 1 exhibits numerous characters, many of which are related to the masticatory apparatus, that were further elaborated among later cynodonts (Kemp, 1979; Hopson and Barghusen, 1968); the zygomatic arches are bowed laterally and the coronoid is centered in the temporal fosas (rather than close to a Zygomatic arch); and the postcanine teeth have anterior and posterior accessory cusps, as well as cuspules along the lingual (internal) margin of the crown.

In various cynodont families (e.g., Thrinaxodontidae, Probainognathidae and Chiniquodontidae) are found further refinements of cranial features that established the basic plan inherited by the earliest mammals (Hopson and Barghusen, in press). Differentiation of the teeth into incisors, canines, and post canine (molariform) teeth is clearly evident. The number of upper and lower incisor teeth are lost. The structure of the postcanine molariforms, in which there were three or four cusps aligned anteroposteriorly and a central primary cusp (Figure 6A), established the primitive pattern for mammals. However, in most cynodonts (excluding the so-called gomphodonts in the families Diademodontidae and Traversodontidae, and the Tritylodontidae) the reptilian mode of tooth replacement persisted (postcanine teeth were replaced continuously throughout life in an alternating sequence) and the roots were single (not multiple, as in mammals). Modifications of the skull and jaw related to mastication are apparent in several lineages: the braincase consolidated as an enclosed structure with the epipterygoid suturally joining the prootic, and thus provided for an extensive origin of the temporalis muscle (Hopson, 1969). A secondary

palate completely separated the oral and nasal cavities. This structure has been interpreted as evidence that cynodonts undertook some oral processing of food, although the original function of the secondary palate may have been as a structural cross-brace (A. W. Crompton, personal communication). It is worth noting in this context that among carnivorous cynodonts (those with trir- or quadricuspid postcanine teeth of the pattern typical of mammalian occlusion (tooth-to-tooth contact did not develop, it is therefore unlikely that "mastication" of the mammalian type was present. Occlusion did develop, however, in certain herbivorous cynodonts, such as the traversodontids and tritylodontids, clearly indicating that oral processing of food was important in some forms.

In their postcranial skeleton (Figure 4) cynodonts exhibited a number of features intermediate between "reptilian" and mammalian grades of organization, and in this sense were truly transitional forms (Jenkins, 1971; Kemp, 1980). The cynodont atlas-axis complex retained a reptilian arrangement of its constituent ossicles, but in the pairing of the occipital condyles, the differentiation of a dens (a neomorphic, anterior outgrowth of the atlas body) and the reduction of the prezygapophyses of the axis, showed incipient specializations to permit atlanto-occipital flexion and extension as well as atlanto-axial rotation (Jenkins, 1969). The unique specializations of mammals at these two joints for mobility of the head were already underway among cynodonts. Similarly, the remainder of the vertebral column exhibited certain modifications toward a mammalian level of organization but without attaining the distinctive features that uniquely characterize mammals. Both cervical and lumbar ribs were retained; only a subtle gradational change occurred between the thoracic and lumbar series in the shapes of neural spines and articular processes. It is unlikely that cynodonts flexed and extended the vertebral column in a sagittal plane as do mammals; the principal movements were probably lateral undulation, as in reptiles, facilitated in some forms by an enlarged iliocostalis system of muscles inserting on imbricating costal plates.

The shoulder girdle of cynodonts was comprised of a trough-shaped scapula, and anterior and posterior coracoids; a robust clavicle articulated with the anterior margin of the scapula (the homologue of the scapular spine/ acromion of mammals). This primitive arrangement was inherited virtually unchanged by the early mammals and persists today among monotremes in an only slightly modified form. Propulsive movement of the humerus appear to have involved a significant amount of elevation long axis rotation, a primitive characteristic, was retained. The cynodont hindlimb and pelvic girdle also showed a mosaic of features that represented an intermediate stage in the development of a mammalian pattern. Among these were an ilium with a substantially elongated and broadened anterior portion (figure 4), a socket-like acetabulum and a bulbous, inflected femoral head. The femur, held at angles of about 55 grads to a sagittal plane, was posturally intermediate between the more sagittal orientation of early mammals. Cynodonts possessed a well-developed calcareal tuberosity for insertion of the triceps suare, indicating that at this stage the major pedal plantarflexors acted directly on the calcaneus. Cynodonts likewise exhibit the beginning of the mammalian condition in which the astragalus is superimposed on the calcaneus. In mammals the astragalus has no "plantar contact" but lies entirely between the tibia above and the calcaneus below. In cynodonts this condition is incipient, for the astragalus still has a partial plantar contact along its medial half; the lateral half of the astragalus was supported by the calcaneus through a rudimentary sustentaculum tali. Figura 4. Reconstruction of the cynodont *Thrinaxodon liorhinus*. Figura 5. Reconstruction of a Liassic morganucodontid. From Jenkins and Parrington, 1976, Figura 18.

THE EARLIEST KNOWN MAMMALS.

Given the nature of the reptile-mammal transition, it is not surprising that there have been varied opinions on defining the Class Mammalia (for a review, see Crompton and Jenkins, 1973). The most widely accepted criterion is the presence of dentary condyle on the jaw articulat-

ing with a glenoid in the squamosal, although many other features may be cited (Crompton and Sun, in press).

Fossil remains of the earliest known mammals are now known from a number of localities throughout the world (Clemens et al., 1979). Of particular importance are the fissure deposits in Wales and southwest England, and the bone beds of continental Europe (Germany, Switzerland). More recently, a fauna of early mammals has been discovered in north-eastern France (Sigogneau-Russell, 1976, 1983). In Africa, isolated but important mammalian specimens have been recovered from the Stomberg Group in Lesotho (Crompton 1974). From the Lufeng beds of the Yunnan Province of China are known a number of important specimens that are (Sigogneau-Russell and Sun, 1981; Crompton and Sun, in press; Zhang, 1984). In North America, an early mammal fauna has been discovered in the Kayenta Formation of northeastern Arizona (Jenkins et al., 1983). Following the biostratigraphic analyses of Olsen and Galton (1977) and Clemens (1980), the preponderance of evidence favors an Early Jurassic (Liassic) age assignment for these faunas, rather than Late Triassic (Rhaetian) or "Rhaeto-Liassic" as was equivocally stipulated in earlier papers.

Nearly all the earliest known mammals were an order of magnitude smaller than any of the Middle Triassic cynodonts. In head-body length, they perhaps reached ten centimeters and a body weight estimated between 20 to 30 gr. Only *Sinocodon* 1 appears to have been larger; its skull length is about 5 cm, compared to 3 cm in morganucodontids (Crompton and Sun, in press).

MORGANUCODONTIDS.

Among the different kinds of early mammals known from Liassic deposits, the family Morganucodontidae is known in greatest detail by virtue of the many isolated skeletal elements or *Morganucodon* 1, and a nearly complete skeleton each of the related forms *Megazostrodon* 1. On dental evidence, the morganucodontids appear to have been allied to certain "nontherian" families of mammals of the

later Mesozoic, such as the tricinodontids and docodontids. With the possible exception of the monotremes, this group of early mammals left no living descendants. Nevertheless, they clearly had attained a mammalian grade of organization, and in terms of our present knowledge of the fossil record offer the best view of the adaptations of early mammals. The following summarizes their most important features. Figure 6. Representative postcanine teeth of *Thrinaxodon* 1 (from Crompton, 1963, Figure 4) and molars of *Morganucodon* 1 and *Kuehneotherium* 1 (from Hopson and Crompton, 1969, Figure 7). Anterior ends of the teeth are to the left.

The jaw joint was transitional in structure, comprising a dentary-squamosal contact as well as primitive, reptilian quadrate-articular contact. Post dentary bones persisted in close relation to the lower jaw (Figure 8; see MIDDLE EAR, below). Teeth were well differentiated into incisors, canines, premolars and molars, and the postcanine teeth were double rooted. During occlusion, the lower jaw moved dorsomedially and a consistent pattern of wear facets formed as the outer surface of the lower molars sheared against the inner surface of the upper, and was further elaborated among later "nontherian" as well as therian mammals. The typical morganucodontid cusp pattern (with a primary central cusp (A,a), an anterior secondary cusp (B,b), and two posterior secondary cusps (C,c;D,d), all aligned anteroposteriorly) is clearly reflective of a cynodont heritage (Figure 6B). Other features include the ossification of the braincase and the development of the internal ear: an anterior lamina of the prootic established a lateral wall to the cavum epiptericum (lodging the trigeminal ganglion) and provided a foramen for the maxillary and mandibular divisions of the trigeminal nerve; a bony "floor" enclosed the cavum epiptericum, and separated two foramina for the exit of the anterior (greater petrosal) and posterior (primary, or Hyomandibular) of the facial nerve; and the cochlear region became relatively enlarged (Kermack et al., 1973, 1981; Crompton and Jenkins, 1979; Crompton and Sun, in press).

The structure of the postcranial skeleton in morganucodontids gives evidence of the early

establishment of many typically mammalian features (Jenkins and Parrington, 1976). The specializations of the atlas-axis complex begun among cynodonts (i.e., double occipital condyles, a dens, reduction of the atlanto-axial zygapophyses) were further elaborated to the point that these early mammals apparently employed as much flexion-extension at the atlanto-occipital joint and rotation at the atlanto-axial joint as do modern forms. Clear differentiation of the cervical vertebrae in *Megazostrodon* 1, which included narrow laminae and pedicles, broadly spaced zygapophyses, and rod-like, posteriorly throughout later mammalian evolution. The enlargement of the cervical vertebral foramina, evidence of a spinal cord intumescence corresponding to the origin of the branchial plexus, has been interpreted as evidence of greater neuromuscular control of the forelimb. Evidence of a cervico-thoracic flexure indicates an upright posture to the head and neck as in modern forms (figure 5). In *Morganucodon* 1a series of three mid-dorsal vertebrae exhibit "anticlinal" and "diaphragmatic" characteristics indicating a sharp transition between the thoracic and lumbar series. This differentiation is related to musculoskeletal specialization for axial flexion and extension typical of the locomotor movements and posture of mammals. The pelvic girdle, with a narrow, rod-like ilium directed anterodorsally, a large obturator foramen, and a reduced pubis, was fully advanced to a mammalian grade of organization (Figure 5). Aspects of the limbs and particularly the joints are consistent with an interpretation of well-developed mobility. The hallux appears to have been somewhat divergent, and extension independent of the other digits were possible. This feature, together with sharp, trenchant claws, supports the interpretation that these early mammals were active foragers with climbing abilities similar to those of many small mammals today.

As in the skull, where primitive features (such as the postdentary bone) persist despite the development of structure that established the mammalian plan, the postcranial skeleton also is a mosaic. The morganucodontid shoulder girdle in particular is little modified from that in

cynodonts (cf. Figure 4,5); the scapula remains a trough-shaped blade, both caracoids persist, and the glenoid is a semilunar, posterio-ly directed concavity. The only major difference is that the anterior coracoid has been reduced in size. The humero-ulnar joint is composed of a condyle on the distal end of the humerus and a corresponding cup on the ulna; this condition, inherited from cynodonts, is retained by monotremes and other nontherian mammals, and contrast with the trochlear joint typical of more advanced forms (Jenkins, 1973).

KUEHNEOTHERIIDS

Kuehneotherium (Kermack et al., 1968) and Woutersia (Sigogneau-Russell, 1983) represent an entirely different family of early mammals that is known only on the basis of isolated teeth and jaws. Despite this, the group appears to occupy a singularly important position in mammalian ancestry on the basis of the pattern of the cusps and the occlusal relations between molar teeth. Unlike the condition in other early mammals in which the cusps are aligned anteroposteriorly, in kuehneotheriids the secondary cusps are arranged in a triangular pattern (Figure 6C). Cusps B and C of upper molars are buccally positioned (toward the tongue). This pattern of "reversed triangles" presages the more complex interlocking mechanism of tribosphenic molars that developed later in mammalian evolution (Kernack et al. 1968). Although the shear facets created by this type of occlusion were relatively simple in comparison to those in later mammals, the occlusal pattern (including the talonid-like function of cusp d and the fact that cusp E is positioned as a primitive parastyle) has the fundamental features of a primitive therian dentition (Crompton, 1971). However, as in morganucodontids, the quadrate and articular remained in the primitive configuration (Figure 7E), and thus kuehneotheriids also had a double jaw joint.

THE DIVERSITY OF EARLY MAMMALS

Until recently, representatives of the families Morganucodontidae and Kuehneotheriidae were

the primary fossil evidence on which to base interpretations of the earliest known stage in mammalian evolution. A third family, the Haramiyidae, had long been known from European deposits; represented only by rare, isolated teeth with paired roots and with multiple cusps arranged around the margins of a broad crown, these fossils provided more of a puzzle than any solution to interpreting mammalian origins (although Hahn, 1973, has advanced credible arguments for a relationship to multituberculates). On this limited evidence, Crompton and Jenkins (1973, 1979) interpreted morganucodontids and kuehneotheriids as representing a subtle but important dichotomy between closely related forms leading to "nontherian" and therian mammals, respectively. Recent discoveries, however, have shown that these early mammalian faunas were more diverse than previously thought, and it is now clear that the earliest known mammals include representatives of families other than haramiyids, morganucodontids and kuehneotheriids (Clemens, 1980; Jenkins et al., 1983). These findings not only create new systematic problems, but they also provide evidence that substantial diversification of mammals occurred before the "earliest known" faunas of Liassic age. More importantly, however, morphological studies of the new specimens alter previous conception of how mammalian structural and functional features arose. These new findings can be summarized as follows:

Dinnetherium. Jenkins et al. (1983) described molar wear facets in this genus (from the Kayenta Formation of northeastern Arizona) that are indicative of medial translation and rotation of the mandible during the occlusal phase. These movements appear to have been more highly developed than in any other contemporaneous mammal. The structure of the jaw is also unusual in possessing an angular process (a neomorphic outgrowth of the dentary) that expanded laterally over the postdentary bones and created a new insertion area for a superficial masseter muscle (Figure 7). Oron and Crompton (in press) have shown that in the modern insectivore *Tenrec* and in other mammals the combined activity of the tem-

poralis (acting dorsomedially) and superficial masseter (acting dorsolaterally) are instrumental in effecting medial translation and rotation of the jaw. The occurrence of a neomorphic angle in *Dinnetherium* 1 is evidence that a superficial masseter was positioned and functioned as in modern mammals, an interpretation consistent with the reconstruction of jaw movements based on wear facets. This finding challenges the previous supposition that the therian angle arose by posterior migration of the pseudangular process that took place as certain postdentary bones (i.e., the articular and angular) were incorporated into the middle ear. This would imply that the evolution of the mammalian masticatory apparatus and middle ear were not so tightly coupled either masticatory apparatus and middle ear were not so tightly coupled either structurally or functionally as previously had been supposed.

Figure 7. The lower jaw of *Dinnetherium nezourum* showing the pseudangular (psang) and angular (ang) processes.

Sinoconodon. Crompton and Sun (in press) provide an analysis of new and better preserved material of this important genus originally described by Patterson and Olson (1961). There is no question that *Sinoconodon* 1 (Figure 1D) is a mammal by the conventional definition, for it clearly possesses a dentary-squamosal joint; the dentary condyle is large and bulbous, and the apposing glenoid is deeply concave (as opposed to the morganucodontid condition in which it is very shallow). The dentary itself is large and robust, and the postdentary bones appear to have been smaller than in other early mammals. But Crompton and Sun also point out that, in addition to these distinctly mammalian features, *Sinoconodon* 1 retains some characteristics of a cynodont such as *Thrinaxodon* 1. Among these are the massive, angular nature of the symphyseal region of the mandible, and the fact that the condyle lies below the level of the postcanine alveoli (cf. Figure 1C,D). Postcanine teeth do not appear to have been clearly differentiated into premolars and molars by structural differences and, even more surprising, precise occlusal relations between upper and lower postcanine teeth are

absent. It is even possible that these teeth did not actually come into occlusal contact during jaw closure. Thus it appears that a load-bearing dentary-squamosal joint does not necessarily entail occlusion, although in every known case where precise, unilateral occlusal relations occur the dentary-squamosal joint is well developed. The different occlusal patterns known among other early mammals (Crompton, 1974; Jenkins et al., 1983) may well have arisen subsequent to a "*Sinoconodon* 1 stage" in which the dentary-squamosal joint was present but occlusion was not yet established.

Gobiconodon. This late surviving representative of the family Amphilestidae, known from the Early Cretaceous Cloverly formation of western North America and also from deposits of comparable age in Mongolia, would not otherwise be pertinent to this account of mammalian origins except for one unusual feature. The two Cloverly specimens, both of which appear to be adult on the basis of epiphyseal fusion, clearly show "molar" teeth in the process of being replaced. The long accepted definition of a molar is that it is not preceded by a deciduous ("milk") tooth a relation initially established on the basis of observations of living mammals and the heretofore found consistent with the fossil record. Diphyodonty and the nondeciduous nature of molars have been widely accepted as unique mammalian characteristics. Hopson (1973) has argued cogently that the rapid growth rate of juvenile mammals and their dependence upon maternal milk for nourishment is related to this type of tooth replacement. As in the case *Sinoconodon* 1, *Gobiconodon* 1 upsets our conception of a standard. By definition, its molariform teeth cannot be "molars" although in all other structural and functional respects they functioned as mammalian molars. Hallautherium. Clemens (1980) erected this new genus on the basis of several isolated teeth from Hallau, Switzerland. Although the general proportions and alignment of the cusps are comparable to those in morganucodontids certain details, including an enlarged posterobuccal basin, are so divergent from any pattern previously known that *Hallautherium* 1 is appropriately considered "*Mammalia incertae*

sedis." it is clear, however, that Hallautherium 1 adds to our growing appreciation of dental diversity among the earliest known mammals.

THE MIDDLE EAR

The auditory apparatus of mammals is characterized by an articulated chain of three bones: the malleus, incus and stapes conduct vibrations from the tympanum, a membrane sensitive to airborne sound, to the inner ear which lodges that part of the VIIIth (vestibulo-cochlear) nerve associated with the Organ of Corti. On embryological evidence, it was established well over a century ago that certain structures of the reptilian jaw and jaw joint were homologous with bones in the mammalian middle ear (Figure 8B, C). More recently, major phylogenetic aspects of this transformation have been clarified, principally through a landmark study by Allin (1975). Allin traced the evolution of the synapsid middle ear to its beginnings among pelycosaurs, and demonstrated convincingly that auditory structure in this lineage was fundamentally different from that in other reptiles. The reflected lamina (rl) of the angular (an) is a persistent and dominant feature (Figure 1). In primitive forms, the reflected lamina was probably associated in part with an oral-pharyngeal diverticulum (the recessus mandibularis), and in therapsids developed as a principal support for a membrane (tympanum) with true auditory function (Figure 8A). The residual homologue of the angular, the ectotympanic bone, retains this same function in mammals (Figure 8B). The retro-articular process of the articular, lying posteriorly and medially to the tympanum, persists as the manubrium of the malleus: and the primitive reptilian jaw joint between the articular and quadrate, although reduced to miniscule size, serves to link the mammalian incus and malleus.

Figure 8-A, the posterior half of the skull in a cynodont (*Thrinaxodon*1) shows that the jaw joint (between the articular in the lower jaw and the quadrate lodged in the squamosal) also was part of the auditory apparatus. Vibrations of the tympanum, largely supported by the reflexed

lamina of the angular, were conducted via the articular and quadrate to the stapes. B, the posterior half of the skull in a Virginia opossum (*Didelphis*) shows the typical mammalian jaw joint formed by a dentary-squamosal articulation. The tympanum and middle ear ossicles, much reduced in size, lie behind the jaw joint but retain the same relations as in cynodonts. C, the lower jaw joint of a fetal mammal (medial view) reveals that the middle ear ossicles develop in position and form comparable to those in cynodonts (see inset, B). D, reconstruction of the dentition and jaws of the Triassic mammal *Morganucodon*1; E, the lower jaw is shown in medial view. In this and other early mammals the auditory apparatus persisted in its association with the lower jaw although the quadrate and articular were relieved of their function as the primary jaw joint. F, in *Morganucodon* 1a distinctive pattern of wear facets on the molars was developed as a result of occlusal contact. In order to display these relations, the internal surface of the upper molar (which wears against the external surface of the lower) is represented as if transparent. The relative positions of upper and lower molars varied among genera of early mammals, and thus produced generically distinctive facet patterns. (From Crompton and Jenkins, 1979, Figure 3-2).

Figure 8 (See legend on opposite page).

NEEONTOLOGICAL PERSPECTIVES ON PALEONTOLOGICAL PROBLEMS

Anatomical and physiological studies of modern mammals and reptiles have provided complementary approaches to questions concerning the early evolution of mammals. A few examples may be cited:

How this mammalian homeothermy arise? Crompton et al. (1978), on the basis of energetic studies of a monotreme, a marsupial and several insectivore species, posed the hypothesis that mammalian temperature regulation developed in two stages. At first, a body temperature of 28 - 30 °C allowed early mammals to occupy a "nocturnal insectivore" niche was largely unexploited. Subsequently, higher metabolic rates enabled mammals to maintain higher body

temperature (38 - 40 C, comparable to those of most modern mammals) and thus permitted diurnal activity. The first stage was modelled on the finding that three insectivores (tenrec, setifer and hedgehog) possess a relatively low, reptile-like metabolic rate. From additional studies of other mammals, Oron et al, (1981) concluded that the unusual energetic regime found in the three insectivores may not represent a primitive condition, but rather "a specialization for the conservation of energy" Nonetheless, it is clear that homeothermy is possible in mammals at lower metabolic rates than are present in most forms today (Taylor, 1980).

To what degree do mammalian and reptilian limb muscles differ? Jenkins and Goslow (1983) compared the results of a cineradiographic and electromyographic analysis of the shoulder muscles in the Savannah monitor lizard with those from an earlier study on the Virginia opossum (Jenkins and Weijs, 1979). A large number of putatively homologous muscles in two species were found to be comparable in attachments, in activity patterns with respect to phases of the step cycle, and in apparent actions; these muscles, designated as "functional equivalents," were interpreted as a representation of the pattern inherited from the common tetrapod ancestor of living mammals and lizards. The muscles that were found not to be equivalent in attachment, activity or action were considered "Functionally divergent"; such divergence could be interpreted as a specialization peculiar to either mammals or lizards, but in any case provides a basis on which to stipulate the functional, rather than simply anatomical, differences that separate mammals and reptiles. A major difference lies in the proportion of muscles used in the propulsive phase (limb in contact with the substrate) versus the swing phase (limb not in contact with the substrate). The opossum uses only about a quarter of its shoulder muscles for swing phase activity, the remainder being active during propulsion; monitors, in contrast, activate half of their total shoulder musculature during the swing phase. Parallel studies of the pelvic musculature now in progress are revealing a similar pattern: a mammal employs more of its musculature during the propulsive phase than does a reptile.

How did the mammalian tibio-astragalar joint evolve? A radiographic study of tarsal relations and movements in a variety of climbing mammals (Jenkins and McClearn, in press) has provided the basis for reinterpreting the origin of mammalian ankle structure. Two primary patterns have been identified among living mammals: in didelphids and phalangerids, the proximal surface of the astragalus bears two sulci to receive apposing tibial condyles (the extinct multituberculates have a nearly identical arrangement), whereas in other marsupials (such as macropodids) and in eutherians the astragalus is of a distinctive trochlear shape. This structural diversity is explicable in terms of parallel evolution from the tibio-astragalar joint of advanced mammal-like reptiles and the earliest known mammals in which the bulbous, hemispheroidal proximal surface of the astragalus bears two indistinct facets for the tibia. These facets are represented in didelphis and multituberculates as sulci, and in macrodids and eutherians as the proximal and medial surface of the trochlea. Monotremes alone among living mammals appear to have retained the basic features of the primitive mammalian condition, and in echidnas it occurs with slight modification as a ball and socket joint.

ACKNOWLEDGMENTS.

My attempt to summarize our knowledge of mammalian origins is quite obviously a compendium of the observations and ideas of many workers. I should like especially to acknowledge the major contributions made by A. W. Crompton and James A. Hopson, and to thank them for their criticisms and suggestions in preparing this presentation. Edgar F. Allin reviewed the account of the MIDDLE EAR, a subject to which he has substantially contributed. Figures 2 and 3, modified from Hopson (1969), indicate relationships that were specifically suggested by J. A. Hopson based on his paper (in press) with H. R. Barghusen. The reconstruction of *Sinoconodon* 1 (Figure 1D) was generously contributed by A. W. Crompton and Sun Ai-Lin in advance of their publication. Figure 8 is reproduced with permission of the University of California Press. I am also very grateful to Lillian L.W. Maloney for her special assistance, to L. Laszlo Meszoly for rendering Figures 1-4, 6 and 7, and to Deedra McClearn for editorial review.

FIGURA 1

Jenkins A.Jr.

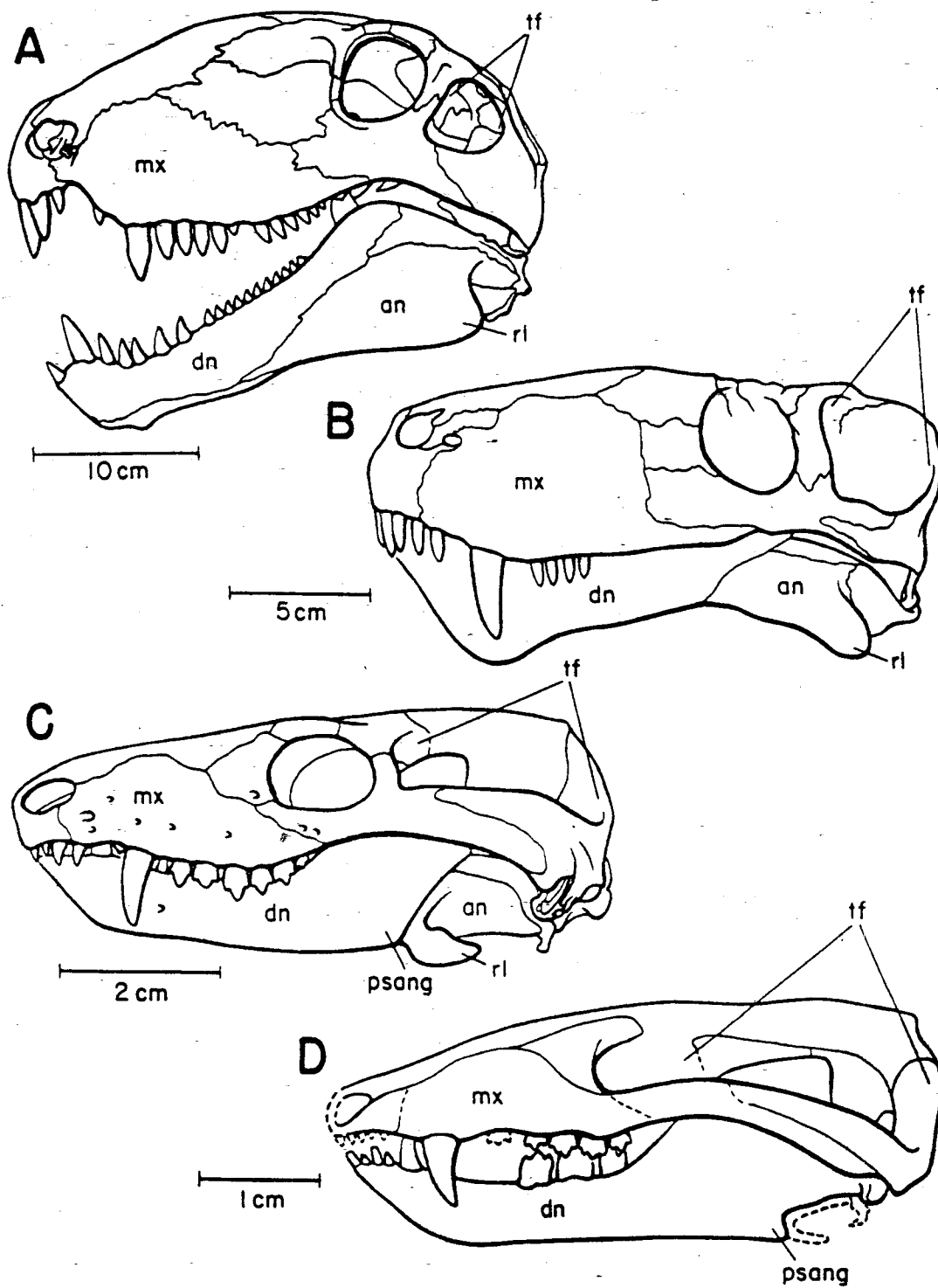


FIGURA 2

Jenkins A. Jr.

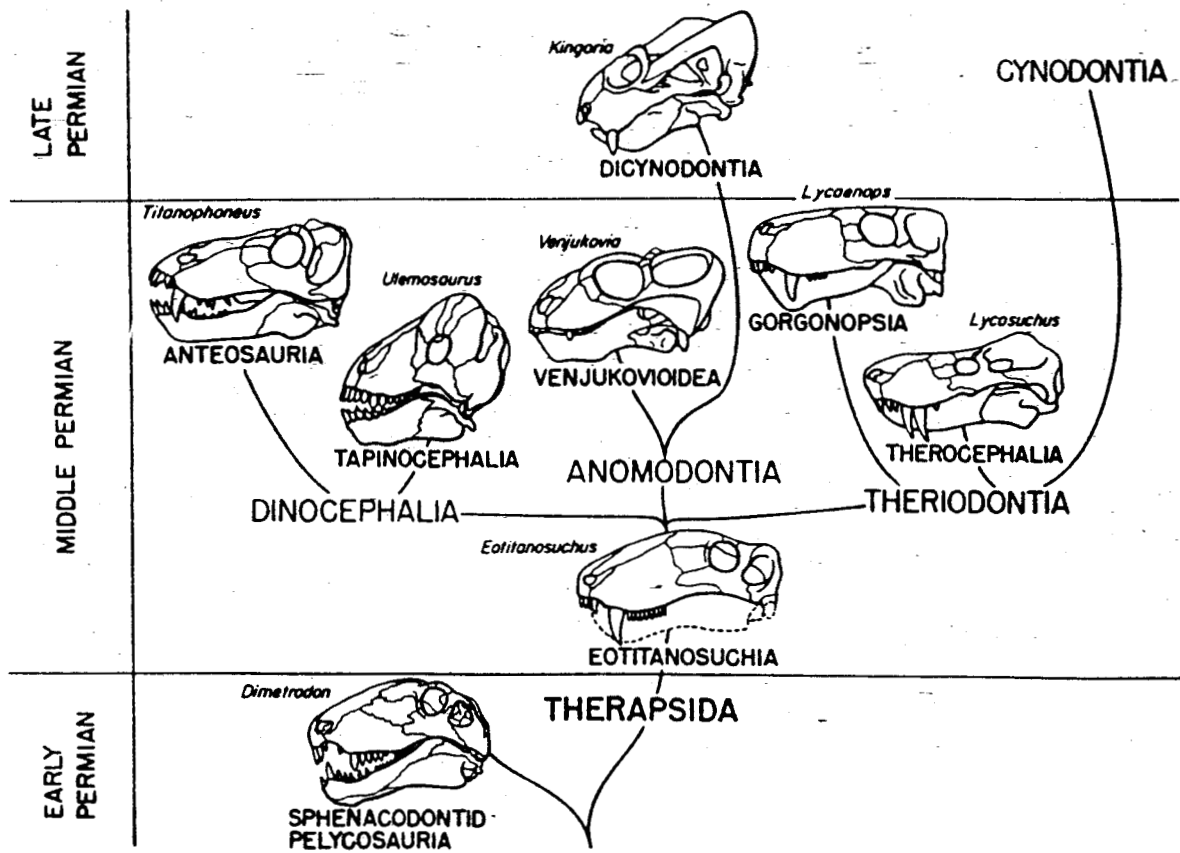
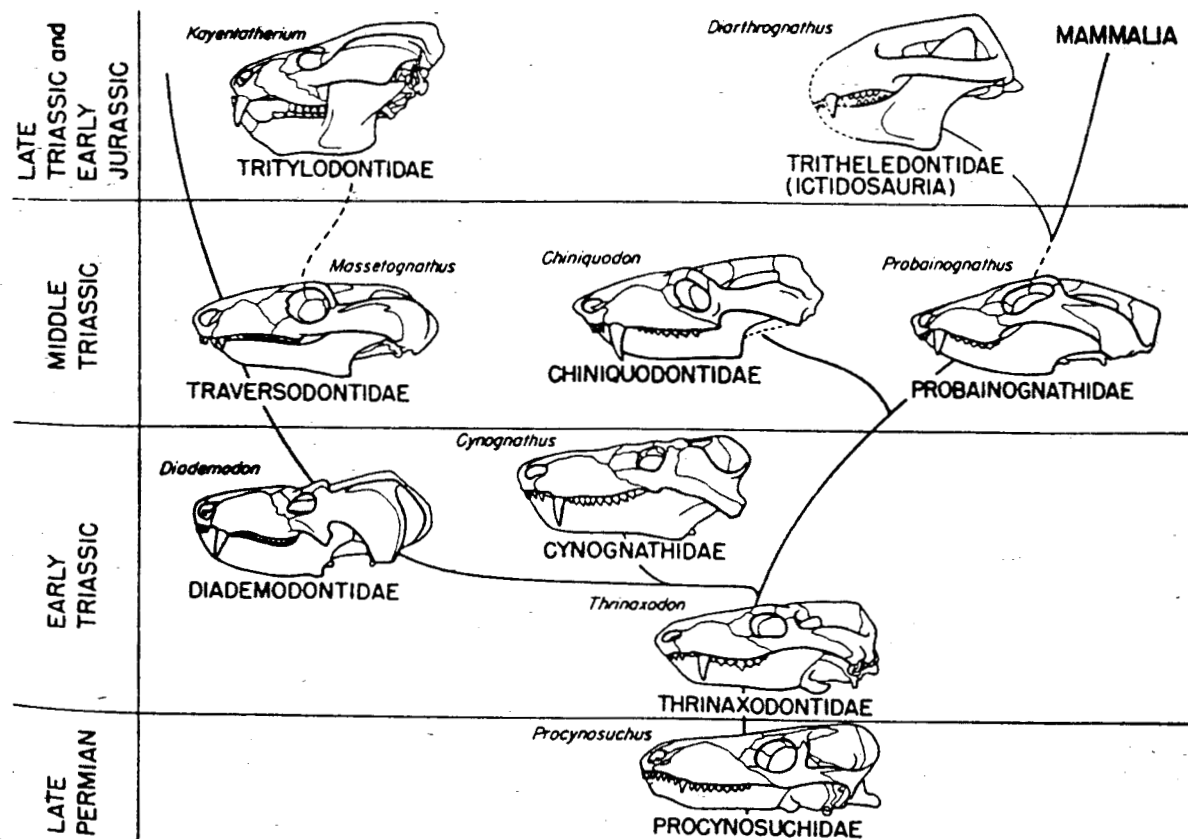


FIGURA 3



Jenkins A. Jr.

FIGURE 4

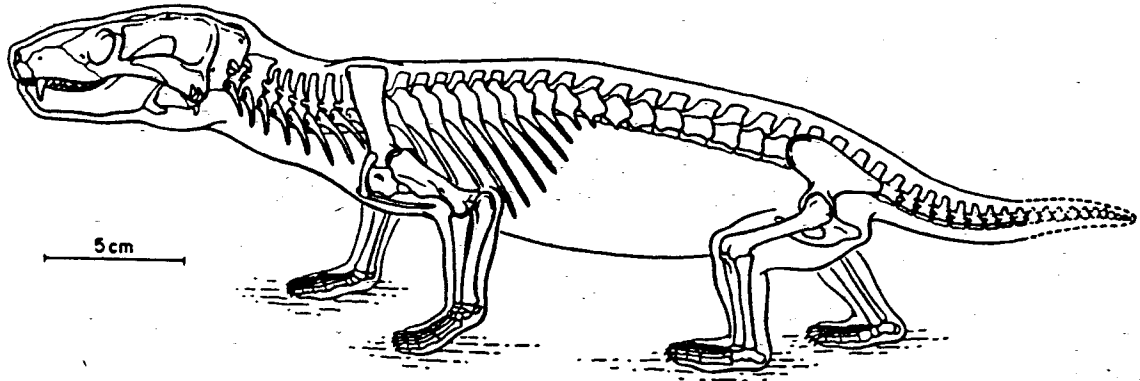


FIGURE 5

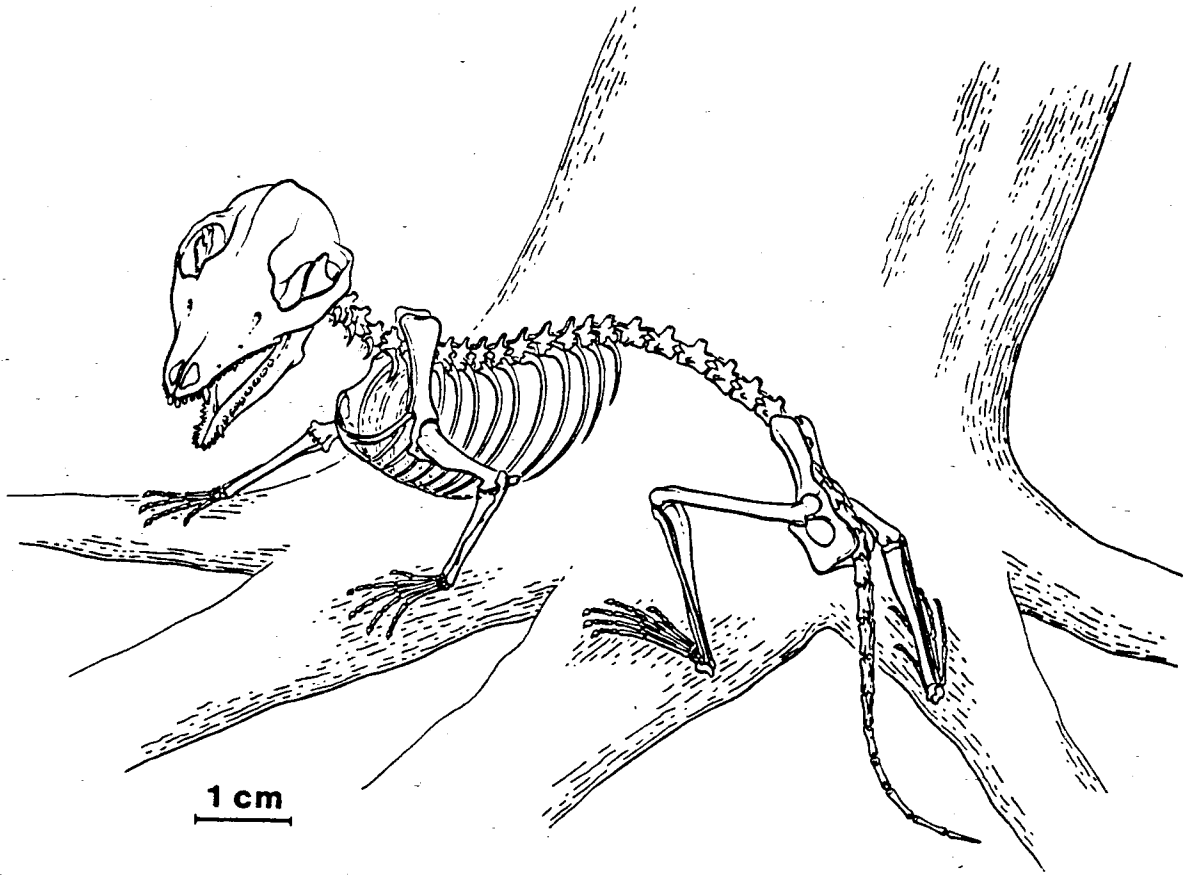
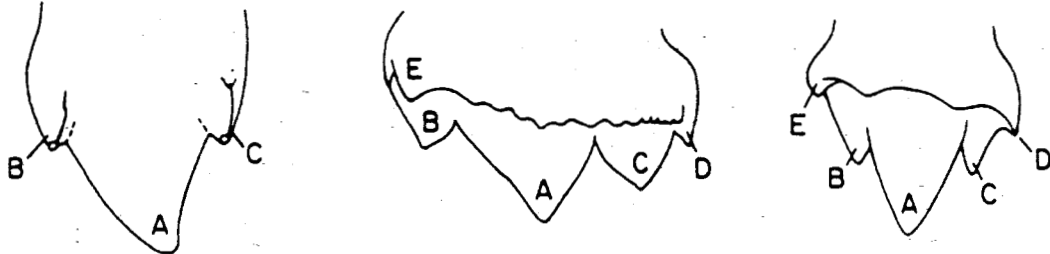
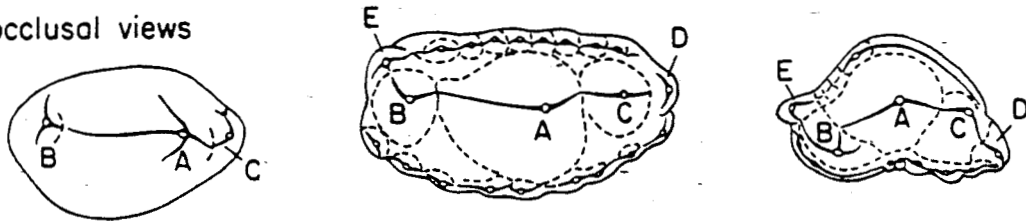


FIGURE 6

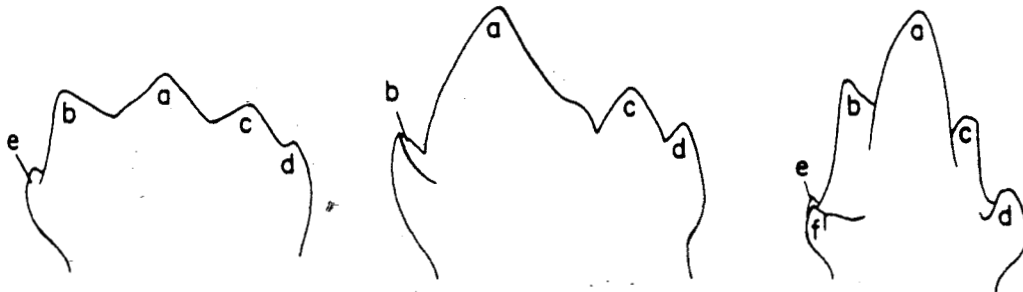
Upper teeth, internal (lingual) views



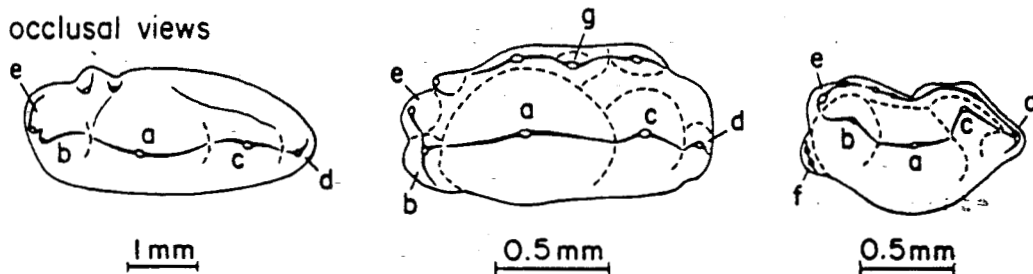
occlusal views



Lower teeth, external (buccal) views



occlusal views



1mm

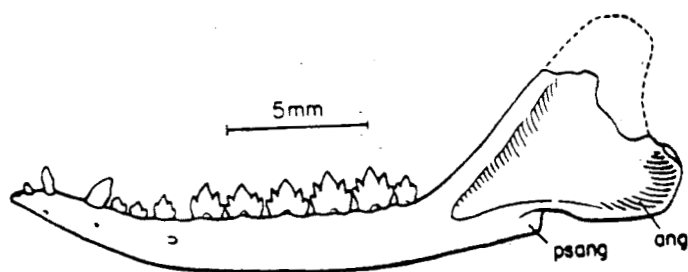
0.5mm

0.5mm

A *Thrinaxodon* B *Morganucodon* C *Kuehneotherium*

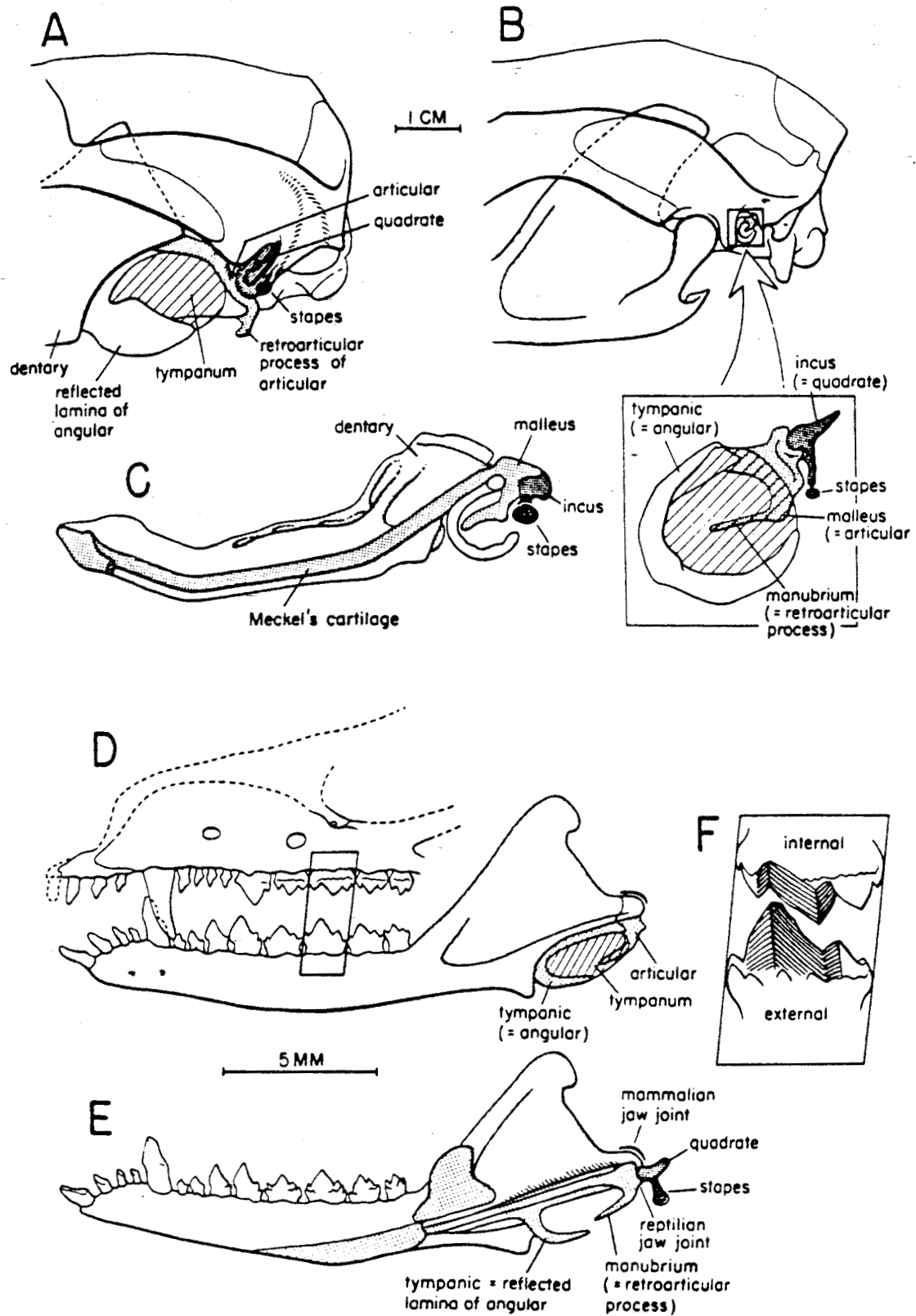
Jenkins, A. Jr.

FIGURE 7



Jenkins, A. Jr.

FIGURE 3



IN GINGERICH, P.D. 7 BADGLEY, C., ORGANIZERS, 1984. MAMMALS, NOTES FOR A SHORT COURSE: KNOXVILLE, TENN., UNIV. TENNESSEE DEPT. GEOL. SCE. STUDIES IN GEOLOGY 8, AND THE PALEONTOLOGICAL SOCIETY, p 87-109.

9.4 (a)

Mammalian Evolution in the Paleocene: Beginning of an Era.

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INTRODUCTION

The Paleocene (approximately 65 to 55 million year ago) is the first epoch of the Cenozoic Era; it is also the beginning of an era in which mammals underwent a major adaptative radiation and became dominant land animals. It was a fascinating and exciting era of transition in mammalian evolution between the dark ages of the Cretaceous, when mammals were few and small and lived in the shadow of the great, "ruling reptiles", and the explosive ages of the Eocene, when mammals were diverse and in full blossom. In the Late Cretaceous there were approximately eight families of marsupials and placentals; by the early Eocene over 70 had come into being. Were mammals from the Paleocene totally unknown, creationists would surely have plenty of grist for their missing-link mills. But paleocene mammals are now known from five continents and they give evidence of archaic groups of mammals that were unleashed from the constraining suppression of dinosaurs. Paleocene mammals evolved rapidly and included "evolutionary experiments" that failed soon thereafter but also groups that were at the base of enormously successful adaptative radiations later in the Cenozoic. In fact, more than one-third of the known orders of the Cenozoic had their beginnings, or origins, in the Paleocene. Origins are captivating. At one time or

another every one of us has pondered the origin of the universe, the origin of life, the origin of humankind, etc. Without knowledge of Paleocene mammals, many basic questions concerning the origin and evolution of mammalian adaptations would not be answerable.

Although a great deal has been learned about Paleocene mammals since G. G. Simpson's excellent review in 1937 (c), less is still known about mammalian evolution during the Paleocene than for any of the later epochs of the Cenozoic. The prime reasons for this discrepancy in availability of information are 1) Paleocene mammals are older and therefore have had a greater chance of being destroyed by various geological agents; 2) Paleocene mammals are generally smaller and therefore more difficult to find and collect (and to prepare and study once collected); and 3) historically, fewer paleontologists have studied Paleocene mammals.

In the limited space available, I will attempt to provide a brief overview of geographic, climatic, and temporal setting in which Paleocene mammals lived and died, as well as synopsis of some of the better known fossil mammal localities on each continent. This will be followed by a limited survey of some of the more important archaic groups of Paleocene mammals. More progressive groups (such as rodents, bats, artiodactyls, perissodactyls, and primates of modern aspect ("euprimates")) will be treated by K. D. Rose in the following chapter even though the earliest representatives of some of these groups may have had their first appearance in the Paleocene boundary and at the Paleocene-Eocene boundary. For general information on mammalian taxa that lived during the Paleocene the reader is referred to Kurten (1971), Romer (1966, 1968), and Savage and Russell (1983).

PALEOCENE GEOGRAPHY AND CLIMATE

During the Paleocene the positions of the continents and the size and distribution of eperic seas were different from those of today. A useful distillation of the relationships between areas of land and sea during the Paleocene, and its relevance to mammalian biogeography is provided by Savage and Russell (1983:50). The

salient aspects of these relationships are illustrated in Figure 1 and can be summarized as follows:

1) North America and western Europe were contiguous and provided a variably developed avenue for mammalian dispersal until possibly as late as the early Oligocene (McKenna, 1983). Terrestrial Paleocene mammalian faunas of North America and western Europe show many similarities (but not as many as during the early Eocene).

2) Partial or complete epicontinental seaways appear to have separated northwestern Europe (Great Britain and part of Scandinavia) from central and southeastern Europe and European USSR from Asian USSR. The latter seaway, the Obik Sea, projected southwards from the Arctic Basin and may have connected with the Tethyan Sea, which trended in a northwest-southeast direction and separated Africa from Europe and Asia.

Figure 1 -- Positions of the continents and distribution of epicontinental seas approximately 60 millions years ago (modified from Savage and Russell, 1983),

3) A high latitude land connection through Beringia (between northeastern Asia and northwestern North America) was developed in the Late Cretaceous and persisted in to the Paleocene.

4) South America was separated from both Africa and Central America.

5) The Indo-Pakistan region was still an island heading northward in the present Indian Ocean.

6) Australia was not yet an island continent and was still joined to Antarctica, which may or may not have been contiguous with the southern tip of South America.

7) Large flooding seas constricted the aerial extent of several continents. In North America, for example, large parts of both the east and west coasts of the United States were flooded and one seaway extended northward from the Gulf of Mexico to near the present-day Canadian border.

Paleoclimatic evidence from several areas in North America suggest that temperature decreased toward the end of the Cretaceous and that these relatively cool temperatures prevailed into the early Paleocene. During the Paleocene climates fluctuated: mean annual temperatures and equability were relatively low during the early and late Paleocene and high during the middle Paleocene and early Eocene (Sloan, 1969; Wolf and Poore, 1982). Much of the evidence for these changes derives from information on the composition and diversity of both the fauna (Rose, 1981 a, b) and flora (Hickey, 1980) in the Bighorn Basin of Wyoming. The relatively high humidity apparent during the Paleocene is shown by the morphology of fossil leaves from North America (Wolf, 1978), western Europe (Dorf, 1964), and Australia (Kemp, 1978). The temperature fluctuations postulated above have been corroborated by evidence from oxygen isotopes in the North Sea (Buchardt, 1978) and calcareous plankton in the Atlantic Ocean (Haug et al., 1977).

BIOSTRATIGRAPHY AND IMPORTANT PALEOCENE MAMMAL LOCALITIES.

The 10 million year duration of the Paleocene epoch can be, and has been, subdivided on the basis of its mammalian faunas. It was, however, first defined by a paleobotanist, W. P. Schimper, in 1874 on the basis of floras from the Paris Basin, France. The boundaries of Schimper's Paleocene have been modified in the light of later work and a history of these modifications can be gleaned from the works of Gingerich (1975) and Savage and Rosell (1983). The estimated ages for the lower and upper boundaries of the Paleocene are 65 and 55 million years ago, respectively (Berggren et al., 1978).

Paleocene mammals are best known from North America but they are also now well represented in collections from Europe, Asia, and South America. Only one locality in Africa has so far yielded Paleocene mammals and none have yet been recovered from the continents of Australia and Antarctica. Some of the more important Paleocene mammal localities from each continent are listed below and their positions are indicated

on Figure 2. Figure 3 provides a tentative intercontinental correlation of the paleocene on the basis of its contained mammalian faunas.

Figure 2--Map the world showing localities that have yielded Paleocene mammals.

Figure 3-- Tentative intercontinental correlation of Paleocene ages. Shaded areas reflected uncertainties concerning precise correlations with North American Land-Mammal Ages.

North America1.-- (Major references: Russell, 1967; Sloan, 1969).

Paleocene mammals are known from more specimens, more species, more localities and more levels in North America than on any other continent. This has, in turn, resulted in a relatively precise biostratigraphic zonation of the Paleocene. The North American Paleocene has been divided into four Land-Mammal Ages: the Puercan, Torrejonian, and Tiffanian correspond roughly to early, middle, and late Paleocene. The fourth Land-Mammal Age, the Clarkforkian, straddles the Paleocene-Eocene boundary. The Dragonian Land-Mammal Age, proposed by Wood et al. (1941), is now generally recognized as early Torrejonian (Tomida, 1981) and the Mantuan, recently proposed by Van Valen (1978), is considered to be early Puercan (Gingerich et al., 1980; Archibald, 1982).

Each of the North American Land-Mammal Ages has been further subdivided into a series of zones based on mammalian species that make their first appearance in each zone. The most recent species-level biostratigraphic framework for the Paleocene has been summarized by Gingerich (1983 b), who subdivided the Puercan in three zones, the Torrejonian into four, the Tiffanian into five, and the Clarkforkian in three. The Paleocene-Eocene boundary is drawn between the first and second zones of the Clarkforkian (Rose, 1980, 1981 a).

Paleocene mammals were first discovered in North America in 1879 by David Baldwin, who then collected for the renowned O.C. Marsh, in the San Juan Basin of New Mexico and Colorado (Simons, 1963). This Basin contains the type localities for the Puercan, Torrejonian, and Tiffanian Land-Mammal Ages. Fossil mam-

mals from the San Juan Basin have been described in a large number of papers, the most important being Matthew's classic overview of the Puercan and Torrejonian collections. The late Paleocene faunal fauna was described by Simpson (1935). A survey of recent work on Paleocene mammals from the San Juan Basin is contained in Lucas et al. (1981). Paleocene mammals are also known from numerous localities in several other large intermontane basins scattered along and to the east of the Rocky Mountains in western North America. Sediments within these structural depressions were deposited during a major period of mountain-building in latest Mesozoic and earliest Cenozoic time. This is the so-called Laramide orogeny. Broad regional uplift of the western Interior during the late Cenozoic resulted in the rejuvenation of streams and large-scale erosion that has culminated in the highly dissected topography in the area today. Although important and exciting field work is being carried out in many areas, two basins require special mention: the Bighorn Basin of northwestern Wyoming and the Crazy Mountain Basin of south-central Montana. In addition to work in the San Juan Basin, collection from these two basins are largely responsible for our concepts of Paleocene mammalian evolution in North America. The Bighorn Basin, in the Pocatello Bench - Clark Fork Basin area, contains a demonstrable stratigraphic succession containing Puercan, Torrejonian, Tiffanian, and Clarkforkian mammals (Gingerich et al., 1980). The Crazy Mountain Basin is noteworthy because it contains some of the largest and best documented samples of Torrejonian and Tiffanian mammals. Recent work in the Crazy Mountain Basin and Tiffanian mammals. Recent work in the Crazy Mountain Basin (Krause and Gingerich, 1983b; Gingerich et al., 1983) is attempting to build on the early studies by Simpson (1936, 1937 a, b) and others.

Europe.-- (Major Reference: Russell, 1964, 1975).

In 1841 de Blainville described the bear-like arctocynid condylarth *Artocyon primaveus* and he thus became the first person to ever describe a Paleocene mammal. *Artocyon primaveus* is a member of the Cernay fauna, which is known from

two areas in the northeastern part of the Paris Basin: Cernay-les-reims and Mont de Berru. The Cernay fauna is Thanetian (late Paleocene) in age and is correlative with the late Tiffanian of North America (Gingerich, 1976b). A slightly early fauna, from near the town of Walbeck in Germany, is a peculiar, unbalanced assemblage that was deposited in a fissure fill. The Walbeck fauna is probably equivalent to the early or middle Tiffanian of North America (Gingerich 1976 b). The recent discovery of the peculiar plesiadapiform primate *Saxonella*, previously only known from Walbeck, in a middle Tiffanian fauna from Alberta, Canada, lends additional support to this correlation (Fox, 1984). A third locality, at Manat in the south of France, has yielded only three species of mammals and is thought to be equivalent to the earliest of the Tiffanian zones in North America (Gingerich, 1976 b). Finally, a very important Paleocene fauna was recently discovered at the Montain (middle Paleocene) type locality near Hainin (southwest of Brussels) in the Mons Basin of Belgium (Godifriaux and Thaler, 1972). The sample from the Hainin locality represents the oldest known Paleocene mammalian fauna from Europe. Only the multituberculates, marsupials, and condylarths from the Hain fauna have been described in detail (Vianey-Liaud, 1979; Crochet and Sigé, 1983; Sudre and Russell, 1982). The multituberculates and condylarths are dominated by genera that are unknown from elsewhere in Europe, North America, or Asia.

Approximately 50% of the genera of Europe were shared with North America during the early Eocene. During the late Paleocene less than 10% of the genera were shared but, nonetheless, they still provide strong evidence of a land connection between the two continents.

Asia1.--(Major Reference: Chow and Zheng, 1980; Li and Tong, 1983; South China "Redbeds" Research Group, IVPP, 1977).

The discovery of Paleocene mammals in the Xinjiang Region of the People's Republic of China in 1960 signalled the beginning of one of the most important recent advances in vertebrate paleontology (Crow, 1960). Since 1960, new and very productive localities yielding Paleocene

mammals have been discovered in numerous basins in Xin-jiang, Guandong, Jiang-xi, Hu-nan, An-hui, He-nan, and Shaan-xi provinces and in the Nomogen area of Inner Mongolia (Chow and Zheng, 1980; Li and Tong, 1983). More than 125 species of Paleocene faunas have been tentatively divided into two groups; those from the early and middle Paleocene (Shanghuan Provincial Age) and those from the late Paleocene (Nungshaanian Provincial Age). The ubiquitous pantodont *Bemalambda* is restricted to the Shanghuan while *Archaeolambda* is characteristic of the Nungshanian.

A series of faunas from the Inner Mongolian Gashato Formation, discovered by American Museum of Natural History expeditions in the 1920's and formerly regarded as late Paleocene in age (e.g., Szalay and McKenna, 1971), are now thought to be earliest Eocene, correlative with the North American Clarkforkian Land-Mammal age (Dashzeveg and McKenna, 1977; Gingerich and Rose, 1977; Rose, 1981 a).

South America.--(major References: McKenna, 1980a; Patterson and Pascual, 1968; Simson, 1948, 1967, 1978, 1980).

Paleocene mammals have long been known from two major areas in South America: 1) The Itaborai locality in Brazil, and 2) east-central Chubut province in Patagonia. The faunas from these areas were generally considered to belong to the late Paleocene Riochican Land-Mammal Age. Rodrigues Francisco and Sounza Cunha (1978), however, recently suggested an early Paleocene age for the Itaborai fauna and Marshall et al. (1981) now believe that the Riochican spans the middle and late Paleocene. Mammals of probable Paleocene age have recently been discovered in northwestern Argentina, in the provinces of Salta, Jujuy (Marshall et al., 1981), and Tucuman (Soria and Pwell, 1981), but these faunas are incompletely described. Conservatively, the mammals from all of the known South American Paleocene localities are here treated as Riochican in age. The Riochican mammals consist of marsupials, xenarthrans, and several different kinds of ungulates. The ordinal diversity of Paleocene mammals from South America (7) is considerably less than from North

America (15, or even Asia (11). The mammalian fauna from Itaborai is dominated by marsupials, that from Patagonia by ungulates. Peculiarly, the known Riochican faunas exhibit a strong imbalance of mammalian carnivores and herbivores, only three small marsupial carnivores are known from Itaborai and none from Patagonia.

Africa.--Paleocene mammals are known from only one locality in Africa (Cappetta et al., 1978). The locality is in the Quarzazate Basin of Morocco and is thought to be early Paleocene in age. The few specimens from there are isolated teeth of paleoryctid proteutherians, creodonts, and carnivorans.

THE CAST OF CHARACTERS

The familial and ordinal distribution of Paleocene mammals in space and time is summarized in Table 1. Even the briefest synopsis of every family, or every order, of Paleocene mammals is impossible in this forum. My intent, Therefore, is to provide a conspectus of only some of the most diverse and spectacular orders represented in the Paleocene, although a brief discussion of the unexpected recent discovery of a possible xenarthran in China is appended to this section. Table 2 lists the most diverse (species rich) orders of Paleocene mammals within each recognized temporal subdivision on each continent. It is immediately apparent that, archaic orders predominated. Two types of mammalian orders can be characterized as "archaic": 1) those that were represented in the Cretaceous and did not change significantly during the Paleocene, and 2) those that are quite highly specialized, evolved just prior to or during the Paleocene, but became extinct soon thereafter.

Condylarths were the most, or one of the most, diverse Paleocene groups of mammals on every continent. Multituberculates, proteutherians, and, toward the latter half of the epoch, plesiadapiform primates achieved high species diversities in North America and Europe. The Paleocene mammalian fauna of Asia was dominated by anagalids and pantodonts while marsupials and notoungulates comprised the majority of species on South America. Each of these major Paleocene groups will be dis-

cussed individually below; groups that are represented by excellent cranial or postcranial material are illustrated in Figures 4-10.

TABLE 1--Mammalian higher taxa the Paleocene of North America, Europe, Asia, and South America. Subdivisions of the Paleocene are abbreviated as follows: Pu, Puercan; To, Torrejonian; Ti, Tiffanian; Cr, Clarkforkian; D-M, Dano-Montian; T, Thanetian; E-M, Early-middle Paleocene; L, Late Paleocene; R, Riochican. Temporal distributions of taxa are modified from lists provided by Savage and Russell (1983) and Li and Ting (1983).

TABLE 2 -- The most diverse orders of Paleocene mammals. Only the three most diverse orders in each temporal subdivision on each continent are listed. The percentages indicate the ratio of the number of species per order to total number of species in each temporal subdivision times 100. Also listed is the total number of known species in each temporal subdivision on each continent.

Order Multituberculata 1.- General references: Clemens and Kielan-Jaworowska, 1979; Sloan, 1979].

Multituberculata, whose closest living relatives are the monotremes, comprise one of the oldest, longest-lived orders of mammals. They are known from the Late Jurassic to the early Oligocene and were among the most diverse and abundant mammals in the Paleocene. Their geographic distribution is confined to North, America, Europe, and Asia and is therefore entirely Holarctic. Multituberculates are currently divided into three suborders: Plagiaulacoidea, Ptilodontoidea and Taeniolabidoidea. Plagiaulacoids are generally considered to include, or to be representative of, the ancestral stocks from which later multituberculates evolved. Only ptilodontoids and taeniolabidooids are present in Paleocene faunas.

Ptilodontoids were small (shrew-to rat-sized), whereas taeniolabidoidea, although also having shrew-sized members, included forms as large as modern beavers (Figure 4). Like most fossil mammals, multituberculates are best known from dental remains. Ptilodontoids possessed a

pair of long, slender procumbent lower incisors, the crowns of which are completely covered with enamel, and an enlarged, blade-like fourth lower premolar that functioned as a slicing wedge (Krause, 1982a). Taeniolabidoids, on the other hand, are united as a group by the common possession of an enlarged pair of lower incisors bearing a ventrolabially restricted band of enamel. This morphology produced through wear, a self-sharpening tool for gnawing, as in the incisors of rodents. Taeniolabidoid lower, fourth premolars were generally reduced although a few genera (such as *Eucosmodon* and *Neoliotomus*) retained large, ptilodontoid-like blades. All multituberculates have distinctive molars with many cusps arranged in mesiodistal rows -- hence the name Multituberculata.

Figure 4.- Cranial and dental morphology of *Ptilodus* (low. left, modified from Krause, 1982a) and *Taeniolabis* 1 (upper left, modified from Matthew, 1937). On right, reconstruction of multituberculate skeleton based primarily on the skeleton of *Ptilobus* 1 (from Krause and Jenkins, 1983; reproduced with permission from Museum of Comparative Zoology, Harvard University).

The ptilodontid *Ptilobus* known from Puercan Through Clarkforkian levels in western North America, is the most completely known of any multituberculate. Excellent skull and skeletal material of this rat-sized genus has been recovered. *Ptilobus* 1 appears to have been an omnivore, possibly specializing in a diet of small body size suggests that these forms were less herbivorous than previously thought; they almost certainly supplemented their diet with protein-rich foods such as insects, worms, seeds or nuts and were thus quite omnivorous (Krause, 1982a). An analysis of dental function in *Ptilobus* 1 has revealed that ptilodontoids had two distinct masticatory cycles: a slicing-crushing cycle in which the posterior premolars were used to open and initially break down large, hard food items and a grinding cycle in which the mandible was retracted while the molars, with their reciprocally concavo-convex cusps, were held in tight occlusion (Krause, 1982a). This pattern of occlusion, particularly the posteriorly-

directed power stroke of the grinding cycle, appears to be unique among mammals. Taeniolaboids, with their rodent-like incisors, reduced premolars, and enlarged molars de-emphasized the slicing-crushing cycle and placed a greater emphasis on gnawing and grinding.

The postcranial skeleton of North American multituberculates also reveals a number of unusual and even unique specialization including the presence of a post-obturator foramen, an enlarged prepollex and parafibula, a large, lateral flange proximally on the fibula, and a deep excavation posteriorly beneath the proximal articular surface of the tibia (Krause and Jenkins, 1983). The recently-discovered skeleton of *Ptilodus* 1 exhibits several adaptations indicating that it, and all other North American multituberculates for which postcranial material is known, were arboreal. These adaptations include a distinctive tarsal construction for a range of pedal mobility characteristic of mammals that descend trees headfirst, a hallux that could be abducted and adducted for prehension in a plane independent of the other digits, and a long, robust, and prehensile tail (Jenkins and Krause, 1983).

Why did this great group of mammals, which was undoubtedly one of the most successful, long-lived taxa in all of mammalian history, go extinct? It is impossible to specify the exact cause of such an event but strong inverse correlations of abundance and diversity between North American multituberculates and rodents suggest possible diffuse competition between the two groups (Van Valen and Sloan, 1966). Multituberculates may also have competed earlier with plesiadapiform primates and condylarths. Reconstruction of diets, body sizes, activity patterns, habitats, and locomotion in multituberculates indicates that the resource space occupied by the group may have overlapped broadly with that of rodents (Krause, 1981, 1982b). The most rapid decline in multituberculates occurred at the Tiffanian-Clarkforkian boundary, the precise time at which rodents are thought to have immigrated to North America from Asia. [Order Marsupialia [General references: Crochet, 1977; Marshall, 1982a; Paterson and Pascual, 1968; Paula Couto, 1952; Simpson, 1980].

Almost every continent has been suggested as the area of origin for marsupials but the most defensible include North America, South America, Australia, and Antarctica (Marshall, 1980b). Although Australia is known as the "Land of Marsupials," the earliest record of marsupials on that continent is in late Oligocene. Records of marsupials on North and South America extend back to the Late Cretaceous and in Antarctica to the late Eocene (Woodburne and Zinsmeister, 1982). Despite their early diversity in the Cretaceous of North America (at least 9 genera), only one genus (*Paradectes* 1; Family Didelphidae) survived into the Paleocene. Marsupials became extinct in North America in the middle Miocene although *Didelphis* 1, the Virginia opossum, re-invaded from South America in the Pleistocene and is thriving today. In the Paleocene, marsupials achieved their greatest diversity in South America where they are represented by 14 genera of didelphids (more than in the Recent fauna), one genus of Borhyaenidae, and three genera of Polydolopidae (Marshall, 1982a). Didelphids include the opossums and are regarded as the stem family of the Marsupialia; its most primitive members lie near the ancestry of all other marsupial groups. Didelphids of the Paleocene, as now, were mostly omnivores although the smaller species undoubtedly fed largely on insects. Borhyaenids are a group of "dog-like" marsupials that included short-legged carnivorous and omnivorous forms. Its most spectacular member is the saber-toothed marsupial *Thylacosmilus* 1 of the Miocene and Polydolopids such as the best-known Paleocene genus *Epidolops* 1 (Figure 5), have pincer-like incisors, bunodont, multi-cusped molars, and enlarged, laterally compressed posterior lower premolars with serrate edges reminiscent of those in ptilodontoid multituberculates.

Figure 5 -- Cranial and dental morphology of *Epidolops* 1. Left, ventral view of skull (modified from Paula Couto, 1952). Right, lateral view of left dentary (modified from Marshall, 1980b). "Insectivores" [General references: Butler, 1972; Novacek, 1982]

The general notion that most early mammals were tiny, primitive insectivores is incorrect.

The archaic ungulate group, the Condylarthra, had already assumed a position of dominance by the early Paleocene and other groups were also in abundance. Nonetheless, insectivores were present and, if anything, are under-represented in the Paleocene fossil record because of their small size. Screen-washing techniques utilized in the past 20 years have, of course, rectified this situation to some extent.

Mammals generally referred to as insectivores have long been placed in a waste-basket taxon, the Insectivora. The waste-basket has been partly emptied by removing a group now referred to by some as Lipotyphla and by others as Insectivora sensu stricto. Lipotyphla of the Paleocene include a hedgehog-like group (the Erinaceoidea) and a shrew-like group (the Soricoidea). The waste-basket, however, still contains various groups of placental mammals that have diverged little from the ancestral stock. This new waste-basket and primarily early Cenozoic taxon, the Proteutheria, was coined by Romer (1966) and elevated to ordinal status by Butler (1972). McKenna (1975a) attempted to dissolve the Proteutheria and to re-assess its component families to various eutherian "super-groups." The Proteutheria, for obvious reasons, is a group that particularly defies brief characterization. Suffice it to say that, during the Paleocene, there were at least six families present, none of which were particularly diverse or abundant. Some of the more important families include the relatively large and aquatic Pantolestidae and two superficially rodent-like groups, the Apatemyidae and Mixodectidae.

Order Primates - Suborder Plesiadapiformes (Major references: Gingerich, 1976b; Kay and Cartmill, 1977; Rose and Fleagle, 1981).

The earliest primate, *Purgatorius*, belongs to the diverse radiation of primates placed in the Suborder Plesiadapiformes, representatives of which are known from both North America and Europe. *Purgatorius* may have been present in the latest Cretaceous (Van Valen and Sloan, 1965) but is best known from the early Paleocene (Puerca) of Montana (Clemens, 1974). *Purgatorius* is so generalized in its dental morphology that it is difficult to determine its

relationships to later primates. However, in the middle and late Paleocene, plesiadapiform primates became very common and were represented by a number of families with divergent specialization. These include microsypids, plesiadapids, carpolestids, paramomyids, picrodontids, and saxonellids. Like most Paleocene mammals, plesiadapiforms are known primarily from jaws and teeth, although relatively complete skulls are known for a few genera (e.g., *Palaechthon*, *Plesiadapis*, *Zanycteris*) and partial postcranial skeleton are known for the closely related genera *Plesiadapis* and *Nannodectes*.

Plesiadapis, the best known of the Plesiadapiformes, had a rodent-like skull with a small braincase, long snout, laterally-directed eyes, and a distinct diastema between the enlarged lower incisors and the cheek teeth (Figure 6). The skull lacked a post-orbital bar, a very primitive condition among primates. The partial postcranial skeleton of *Plesiadapis* includes most of the limbs (Figure 7). *Plesiadapis* was a heavily-built animal but no consensus has yet been reached on whether it was an arboreal or terrestrial form (Szalay et al., 1975; Gingerich, 1976b). The Plesiadapidae included probable omnivores, hervirores, and granivores (Gingerich, 1976b). Figure 6-- Cranial and dental morphology of representatives of four families of plesiadapiform primates. Top left, *Palaechthon* 1 (Family Microsypidae, modified from Szalay and Delson, 1979). Top right, *Plesiadapis* (Family Plesiadapidae, modified from Gingerich, 1976b). Bottom left, *Ignacius* (Family Paromomyidae, modified from Szalay and Delson, 1979). Bottom right, *Carpodaptes* 1 (Family Carpolestidae, modified from Szalay and Delson, 1979). Stippled areas indicate parts that are still unknown. Scale bars each represent 1cm. Figure 7-- Skeletal reconstruction of *Plesiadapis* 1. Solid black areas indicate parts of the skeleton that are still unknown. (Modified from Tattersall, 1970).

The Microsypidae are the most primitive family of Plesiadapiformes. The earlier, small microsypids, such as *Palaechthon*, were predominantly terrestrial insect-eaters that used primarily tactile, auditory, and olfactory cues to locate their prey (Kay and Cartmill, 1977). The Paromomyidae include small primates with long

snouts, long slender lower central incisors, and a pointed lower fourth premolar (Simpson, 1955). Carpolestids and saxonellids are more specialized Plesiadapiformes. Both groups consist of small, mouse-sized animals that had enlarged, transversely compressed, and multi-tuberculate-like posterior premolars (Rose, 1975). Picrodontids, another highly specialized group, have peculiarly-expanded first molars that are reminiscent of those in some bats (Szalay, 1968). These resemblances led to early suggestions of bat relationships but later workers have instead regarded the similarities as convergent and simply indicative of a bat-like diet of fruit and nectar. The relationship of plesiadapiform primates to other mammals and, in fact, to other primates is obscure. Plesiadapiformes were probably derived from some insectivoran sometime in the Late Cretaceous, but the ancestral insectivoran stock has so far been elusive (Novacek et al., 1983). Plesiadapiform primates are allied with later forms primarily on the basis of detailed resemblances in the cheek teeth, although some postcranial characters have also been used. The pattern of carotid circulation in plesiadapiformes, however, has recently been found to be non-homologous with that in later primates and re-establishes the uncertainty concerning the phylogenetic relationships of this early radiation (MacPhee et al., 1983). [Order Condylarthra -- [General references: Cifelli, 1983; Matthew, 1937; Simpson, 1937a].

The Condylarthra is an extinct radiation of primitive mammals that arose during the Late Cretaceous and flourished during the Paleocene. The order is thought to have been involved, either directly or indirectly, in the ancestry of as many as 18 other orders of mammals including all living hoofed mammals, or ungulates, "subungulates" (hyraxes, proboscideans, sirenians, desmostylans), whales, and tubullidentates (armadillos). In reality, however, few clear-cut connections to later groups have been documented. Condylarths exhibit a tremendous diversity in morphology and undoubtedly occupied a wide range of adaptive zones.

The composition of the Condylarthra has changed, and will indubitably change, numerous times. For instance, arctocyonids and mesonychids (the groups dealt with in greatest detail here) were once placed in the Suborder Creodonta of the Order Carnivora along with three other families: Miacidae, Oxyaenidae, and Hyaenodontidae. With additional study, vertical classifications have resulted in the elevation of the Creodonta to separate ordinal status but including only the Oxyaenidae and Hyaenodontidae. The Miacidae have been allied with "true" carnivores in the Order Carnivora and arctocyonids and mesonychids are now included within the Condylarthra by some and given separate ordinal status (Arctocyonia and Acreodi, respectively) by others. Likewise, some of the earliest whales were originally described as condylarths. The changing concept of the Condylarthra is indicative of a broader, more interesting phenomenon concerning "archaic" mammals. With increasing efforts to trace modern ungulate or ungulate-like clades to their "roots" in the Paleocene, various taxa are removed from the Condylarthra is therefore a grade because available evidence does not permit its subdivision. For the present purposes, I have adopted a conservative classification and thus at least 10 families of condylarths are identified as having been present in the Paleocene.

The earliest condylarths are the Arctocyonidae, known from the Late Cretaceous of North America (Sloan and Van Valen, 1965) and probably South America. They were one of the most diverse families of the Paleocene and were probably at the base of the subsequent condylarth radiation (Van Valen, 1969). Early Paleocene arctocyonids were of small to medium size. They generally had long, low skulls with prominent canines and low-crowned, primitively tribosphenic molars. Salient features in the postcranial skeleton include short limbs, clawed feet, and long tails. Larger forms, the size of small bears, evolved later in the Paleocene and are present in both North America and Europe. Typical of these later forms is Arctocyon (Figure 8), which, like most bears, probably had an omnivorous diet. Figure 8--

Cranial and dental morphology of Arctocyon (left, modified from Russell, 1964) and Dissacus (right, modified from Matthew, 1937).

Mesonychids were more common in the Eocene but at least one genus, Dissacus, attained a broad holarctic distribution in the late Paleocene. Dissacus had simple, triangular molars and high-crowned shearing lowers (Figure 8). Unlike arctocyonids, mesonychids had hooves but they were of unusual construction in that terminal phalanges that supported them were narrow and fissured. Mesonychids were probably carnivorous or piscivorous (Szalay, 1969b) and probably gave rise to whales (Gingerich et al., 1983).

The remaining families of Paleocene condylarths were primarily herbivorous forms. Included here are such divergent forms as the small, generalized hyopsodontids, the peripitychids, with their characteristic wrinkling of molar enamel, meniscotheriids, with had molarized premolars and selenodont cheek teeth, and the only family represented in South America, the Didolodontidae. Space restrictions preclude a detailed treatment of each of the herbivorous condylarth families. Order Pantodonta-- [General references: Chow et al., 1977; Simons, 1960].

The Pantodonta include the largest of Paleocene mammals. They were generally large, heavy-bodied herbivores with small brains, short, stout limbs, and pentadactyl feet. They had an entirely holarctic distribution and presumably filled several of the ecological niches vacated by dinosaurs at the end of the Cretaceous. Pantodonts are present from the early Eocene of North America and Europe and survived into the early Oligocene in Asia, but their greatest radiation was in the Paleocene of North America and Asia. Pantodonts have not yet been discovered from the moderate to large size. The smallest form is the recently described Cyriacotherium, which was probably about the size of a small sheep (Rose and Krause, 1982). Two of the best-known North American genera are Pantolambda and Titanoides (Figure 9). Pantolambda was slightly larger than Cyriacotherium and had simple, conical canines and W-shaped crests on the molars. Its digits terminated in small hooves.

Titanoides, a rhinoceros-sized animal, is one of the few examples among ungulates that possessed claws rather than hooves. It was also unusual in the development of large, sabre-like upper canines, presumably an adaptation for digging up roots. The Asian radiation included, in addition to several large forms, a host of small pantodonts that were as small as a rat. Until recent collecting in the People's Republic of China, the magnitude of this radiation of pantodonts that were as small as largely unknown. One of the Chinese forms, *Bemalambda*, is known from both cranial and postcranial material and is thus among the best-known Paleocene mammals from Asia (Chow et al., 1977). It appears to be even more primitive than the earliest North American form, *Pantolambda*, but already shows the characteristic w-shaped molar crests. It, like *Barylambda* from North America (Gingerich and Childress, 1983), was sexually dimorphic. Figure 9-- Cranial and dental morphology of *Pantolambda* (left, modified from Matthew, 1937) and *Titanoides* (right, modified from Simons, 1960). [Order Anagalida-- [General references: Szalay and McKenna, 1971; Li and Ting, 1983].

The Anagalida is comprised of a large radiation of Late Cretaceous and early Tertiary Asian mammals that was first given ordinal status in 1971 by Szalay and McKenna, despite the fact that members of the order were known since the 1920's. Early workers regarded anagalidans as tree shrews but knowledge of the relationships and diversity of the Anagalida has increased tremendously in the past few years with the discovery of many new species in the Paleocene of China. In their recent tally of anagalid taxa, Li and Ting (1983) list five families (*Anagalidae*, *Pseudictopidae*, *Zalambdalestidae*, *Eurymylidae*, and *Mimotonidae*), 23 genera, and 33 species. A great deal remains to be learned of the Anagalida and of their relationships to other mammals. Symptomatic of this gap in our knowledge is the general confusion concerning the composition of the order. Kielan-Jaworowska et al. (1979), for instance, would not include the *Zalambdalestidae* in the order, while Sych (1971) erected a separate order (*Mixodontia*) for the *Eurymylidae*. Savahe and Russell (1983) did not

recognize the *Mimotonidae* as a family distinct from the *Eurymylidae*, and Gingerich (1981e) removed the *Didymoconidae*, a family originally regarded by Szalay and McKenna (1971) as possibly belonging to the Anagalida, to the *Condylarthra*.

The destitution of anagalidans tends to include molariform posterior premolars, hypsodont and easily worn upper cheek teeth, and procumbent incisors. The postcranium was apparently lagomorphlike. Agomorph characteristics have long been noted in the Anagalida and the group is now regarded as having been near the ancestry of both the agomorphs and *Rocentia* (Hartenberger, 1980; McKenna, 1982).

Order Notoungulata [General references: Patterson and Pascual, 1968; Simpson, 1980].

Notoungulates were a diverse group of South American ungulates that reached their acme in the Oligocene but survived into the Pleistocene. They also achieved a modest diversity in the late Paleocene and early Eocene of Asia and are represented by rare specimens of a single genus in the late Paleocene and early Eocene of North America (in the Bighorn Basin of Wyoming only). This geographic distribution has led to much speculation about centers of origin for the group. The most recent view is based on the reidentification by Marshall et al. (1983) of *Perutherium* (from late Cretaceous beds at aguna Umayo in Peru), previously thought to be a condylarth, as a notoungulate. Because of the early record of *Perutherium*, is South American, Marshall et al. argue that notoungulates originated on that continent and subsequently migrated to North America and then Asia. *Perutherium*1, however, is known from only isolated lower molar fragments and is so primitive that allocation to the Notoungulata may be premature. It has, in fact, previously been referred to three different condylarth families and Hoffstetter (1981), followed by Cifelli (1983), even questioned its status as a eutherian.

There are, conservatively 14 families of notoungulates, only seven of which occur in the Paleocene. Of these, only the *Arctostylopidae* occur in North America and Asia; all other notoungulate orders are restricted to South

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America. Notoungulates exhibit a characteristic pattern of crests, or lophs, on both the upper and lower molars. Also typical of crests, or lophs, on both the upper and lower molars. Also typical of notoungulates is the peculiar osseous construction of the middle ear cavity, the most significant feature of which is that the epitympanic recess is greatly enlarged into a sinus. [Order Xenarthra 1-- [General reference: McKenna, 1980; Patterson and Pascual, 1968; Simpson, 1978, 1980].

The evolutionary history of xenarthrans, the order of mammals to which armadillos, sloths, glyptodonts and anteaters belong, is largely confined to South America. Toward the end of the Cenozoic, however, several forms migrated northward as far as Alaska, and possible xenarthrans possess a suite of purportedly primitive features (e.g., presence of septomaxillary bones, a monotreme-like stapes, complete ossification of ribs, poorly differentiated vagina and uterus, and poor thermoregulation) that has led some workers to believe that they were the earliest group to diverge from the Eutheria. The polarity of most of these characters, however, has yet to be documented (McKenna, 1975a, 1980a; Novacek, 1982).

Relevant to this discussion is a group of fossil, early Cenozoic mammals from North America, the Palaeonodonta. Palaeonodonts, including the families Epoicotheriidae and Metacheiromyidae, have long been regarded as primitive members or ancestors of the Xenarthra. Emry (1970), however, argued that palaeonodonts were related to pangolins (Order Pholidota) and not Xenarthrans. The recent discovery of morphologically primitive palaeonodonts from the late Paleocene of the Bighorn Basin, Wyoming, however, suggests that palaeonodonts may have been ancestral to both pholidotans and xenarthrans (Rose, 1978, 1979; but see Storch, 1978b). Palaeonodonts, in turn may have been derived from pantolestid or pantolestid-like proteutherians (Rose, 1978).

One of the most extraordinary Paleocene mammals discovered in recent years is *Ernanodon*, a possible xenarthran from the late Paleocene of China (Ting, 1979). *Ernanodon* was a col-

lie-sized animal that had very powerful jaws and a dentition consisting of vestigial incisors, large canines, and reduced, peg-like cheek teeth. Its postcranial skeleton is very robust and powerfully-built and the forefeet had large claws. These features suggest that *Ernanodon* represents an early, "experiment" in ant- or termite-eating (Radinsky and Ting, 1984). The discovery of *Ernanodon* in China is intriguing from a biogeographical point of view. The evolutionary history of undoubted xenarthrans is almost entirely restricted to South America (beginning in the Riochican) and if *Ernanodon* is indeed a xenarthran its presence in China becomes very difficult to explain. Figure 10 -- Cranial and dental morphology of *Ernanodon* (modified from Radinsky and Ting, 1984).

FAUNAL TURNOVER

One of the most pronounced intervals of early Cenozoic faunal turnover occurred near the Paleocene-Eocene boundary in North America (Rose, 1980, 1981, b). Other major turnovers surely occurred, in North America and elsewhere, but none is as well documented as that near the beginning of the Eocene. The change in faunal composition took place in two stages, both presumably a result of major immigrations. The first coincides with the Tiffanian-Clarkforkian boundary when rodents and tilodonts, as well as the genera *Haplomylus* (a hyopsodontid condylarth) and *Coryphodon* (a pantodont) all appear simultaneously. The second stage, at the Clarkforkian-Wasatchian boundary, is heralded by the first appearance in North America of artiodactyls, perissodactyls, primates of the families Adapidae and Omomyidae, and probably hyaenodontid creodonts and meniscotheriid condylarths. Correspondingly, major groups that were previously diverse and abundant during most of the Paleocene are drastically reduced or become extinct altogether. Thus, for example, multituberculates decline sharply at the end of the Tiffanian, as do phenacodontid condylarths and plesiadapid and carpolestid primates at the end of the Clarkforkian.

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The reasons for these turnovers must remain speculative but they were probably owing to a combination of changing climates and competition between native and immigrant taxa. Where the immigrant taxa came from is also inferential but paleogeographic considerations as well as new information on the Paleocene fossil records on other continents show that the first wave of newcomers probably came from Asia (Chow et al., 1977; Hartenberger, 1980) whereas the second came from Europe (Godinot, 1981). [By contrast, earlier transitions in faunal composition, such as at the Torrejonian-Tiffanian boundary, appear to have been a result of intracontinental evolution and dispersal. Concerning the Torrejonian-Tiffanian turnover, Sloan (1969) and Gingerich (1976b) noted that, in many respects, mammals of the middle Paleocene and early Eocene were more similar to one another than those of the late Paleocene and early Eocene. They observed that there were a large number of early Eocene taxa that had more suitable structural ancestors in the middle Paleocene than in the immediately preceding late Paleocene. They therefore hypothesized that, during the relatively cool late Paleocene, the lineages of mammals represented at Torrejonian horizons in the Western Interior migrated southern stock then gave rise to the late Paleocene ancestors of early Eocene forms that migrated northward again as climates ameliorated. The character of the late Paleocene - early Eocene faunal transition has now been documented through an intensive field program carried out by P. D. Gingerich and colleagues. No similar work, however, has been done on the middle-late Paleocene transition in western North America and the intriguing Sloan-Gingerich hypothesis therefore remains to be adequately tested.

OVERVIEW

Dinosaurs met their demise at the end of the Cretaceous and their role in terrestrial ecosystems appears to have been quickly occupied by mammals. The Mesozoic-Cenozoic boundary therefore marks the geologically sudden transition from the "Age of Dinosaurs" to the "Age

of Mammals." Whether or not dinosaurs were the direct ecological vicars of some Cenozoic mammals, there is no doubt that Paleocene mammals were vastly different from those of the Mesozoic. Nonetheless, mammals of the Paleocene are typically characterized as "archaic" and they certainly were relative to those of the Eocene, when many orders with living representatives made their first appearance in the fossil record. The primitive nature of Paleocene mammals, as well as their generally fragmentary condition, makes them particularly difficult to study. More so than in the later Cenozoic, they are largely represented only by dental remains. Also, since many of them had not yet diverged a great dental from their ancestral stocks, the most minute of differences become significant. Although an overstatement, there is some truth to Romer's (1966; 29) proclamation: "Were we living in the late Cretaceous we would probably include all placentals then existing in a single family, and even in the Paleocene most forms could be reasonably placed in a single order."

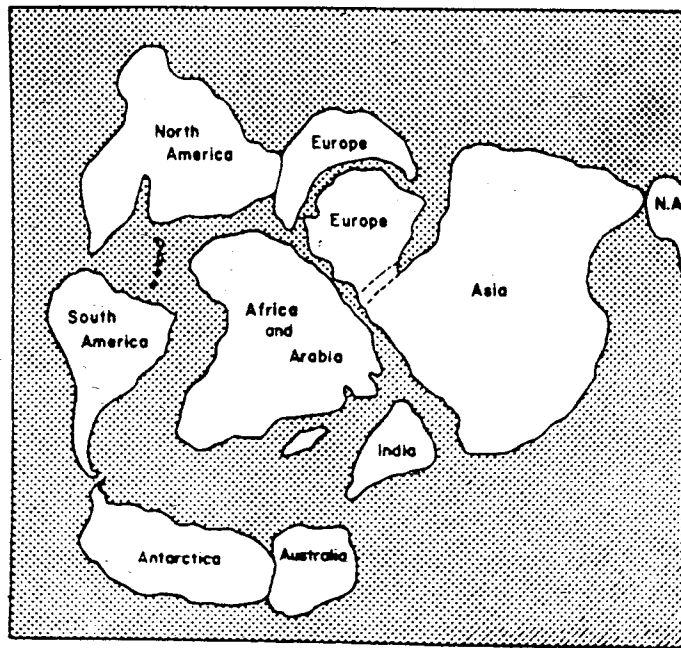


FIGURA 1

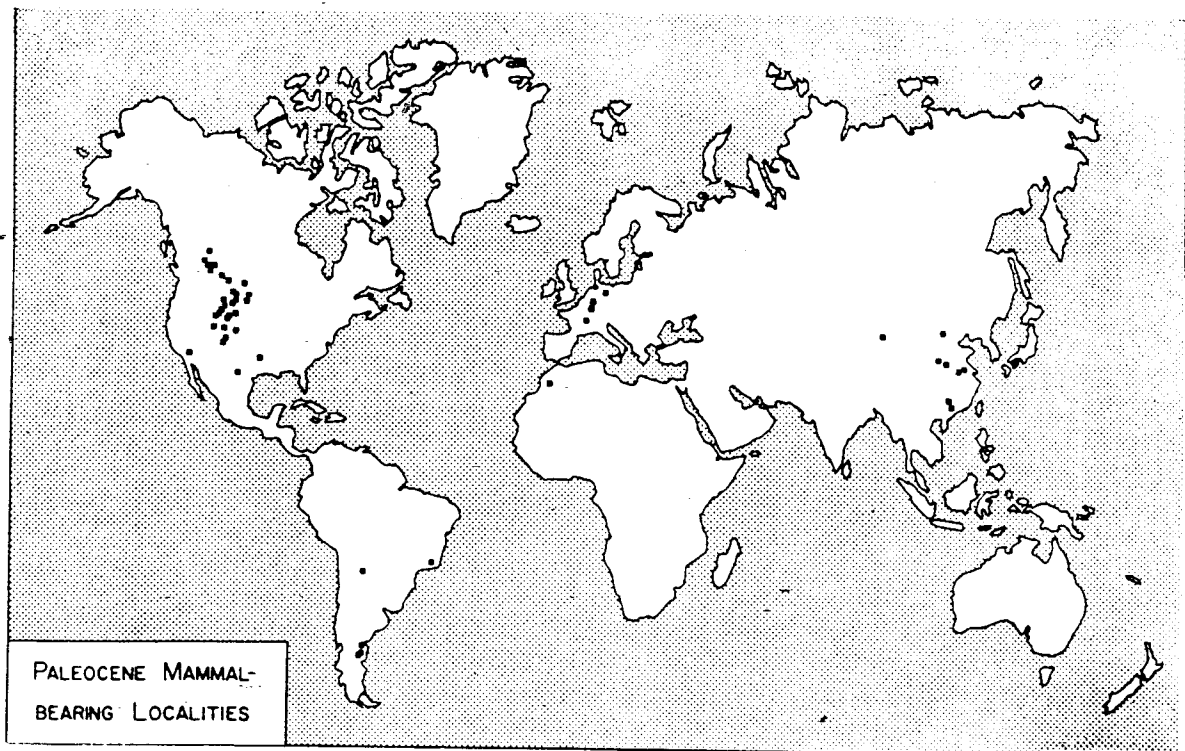


FIGURA 2

EPOCH		N. AMERICA	EUROPE	ASIA	S. AMERICA
PALEOCENE EOCENE	Early	Wasatchian	Ypresian	Lingchan	Casamayoran
		Clarkforkian		Bayanulnian	
	Late	Tiffanian	Thanetian	Nungshanian	Riochican
	Ear. Middle	Torrejonian	Dano-Montian	Shanghuan	Salamancan
	Ear.	Puercan			

FIGURE 3

TABLE 1.

TAXON	NORTH AMERICA				EUROPE		ASIA		SOUTH AMERICA	
	Pu	To	Ti	Cf	D-M	T	E-M	L	R	
Order MULTITUBERCULATA										
Family Ptilodontidae	•	•	•	•						
Family Neoplagiaulacidae	•	•	•	•		•				
Family Cimolodontidae		•	•			•				
Family Boffiidae					•					
Family Eucosmodontidae	•	•	•	•						
Family Taeniolabididae	•	•	•					•		
Order MARSUPIALIA										
Family Didelphidae	•		•	•						•
Family Borhyaenidae										•
Family Polydolopidae										•
Order PROTEUTHERIA										
Family Palaeoryctidae	•	•	•	•		•				
Family Micropternodontidae								•		
Family Leptictidae	•	•	•	•		•				
Family Pantolestidae		•	•	•		•				
Family Pentacodontidae		•	•	•		•				
Family Apatemyidae	•	•	•	•		•				
Family Mixodectidae		•	•			•				
Order LIPOTYPHLA										
Soricidae	•	•	•	•		•				
Erinaceoidea		•	•	•		•				
Order DERMOPTERA										
Family Plagiomenidae	?	•	•	•						
Order PRIMATES										
Family Microsyopidae	•	•	•	•		•				
Family Plesiadapidae		•	•	•		•				
Family Carpolestidae		•	•	•						
Family Paromomyidae		•	•	•						
Family Picrodontidae		•								
Family Saxonellidae			•			•				
Order CONDYLARTHRA										
Family Arctocyoniidae	•	•	•	•		•				
Family Periptychidae	•	•	•	•				•		
Family Mioclaenidae	•	•	•					•		
Family Phenacodontidae		•	•	•						
Family Hyopsodontidae	•	•	•	•		•		•		
Family Mesonychidae		•	•	•		•		•		
Family Didymoconidae								•		
Family Menisocotheriidae				•		•				
Family Tricuspidodontidae						•				
Family Didolodontidae										•
Order TAENIODONTA										
Family Stylinodontidae	•	•	•	•						
Order TILLODONTIA										
Family Esthonychidae			•	•				•		

TABLE 2

NORTH AMERICA	EUROPE	ASIA	SOUTH AMERICA
<u>Puercan</u> Condylarthra - 69% Multituberculata - 13% Proteutheria - 8% Total species - 84 <u>Torrejonian</u> Condylarthra - 40% Proteutheria - 20% Multituberculata - 16% Total species - 159 <u>Tiffanian</u> Condylarthra - 29% Primates - 19% Multituberculata - 15% Total species - 167 <u>Clarkforkian</u> Condylarthra - 22% Primates - 16% Creodonta - 9% Total species - 88	<u>Thanetian</u> Condylarthra - 50% Proteutheria - 17% Multituberculata - 14% Primates - 14% Total species - 36	<u>E.-M. Paleocene</u> Anagalida - 35% Pantodonta - 27% Condylarthra - 23% Total species - 48 <u>L. Paleocene</u> Pantodonta - 30% Anagalida - 27% Condylarthra - 15% Total species - 71	<u>Riochican</u> Marsupialia - 39% Notoungulata - 32% Condylarthra - 11% Total species - 57

FIGURA 4

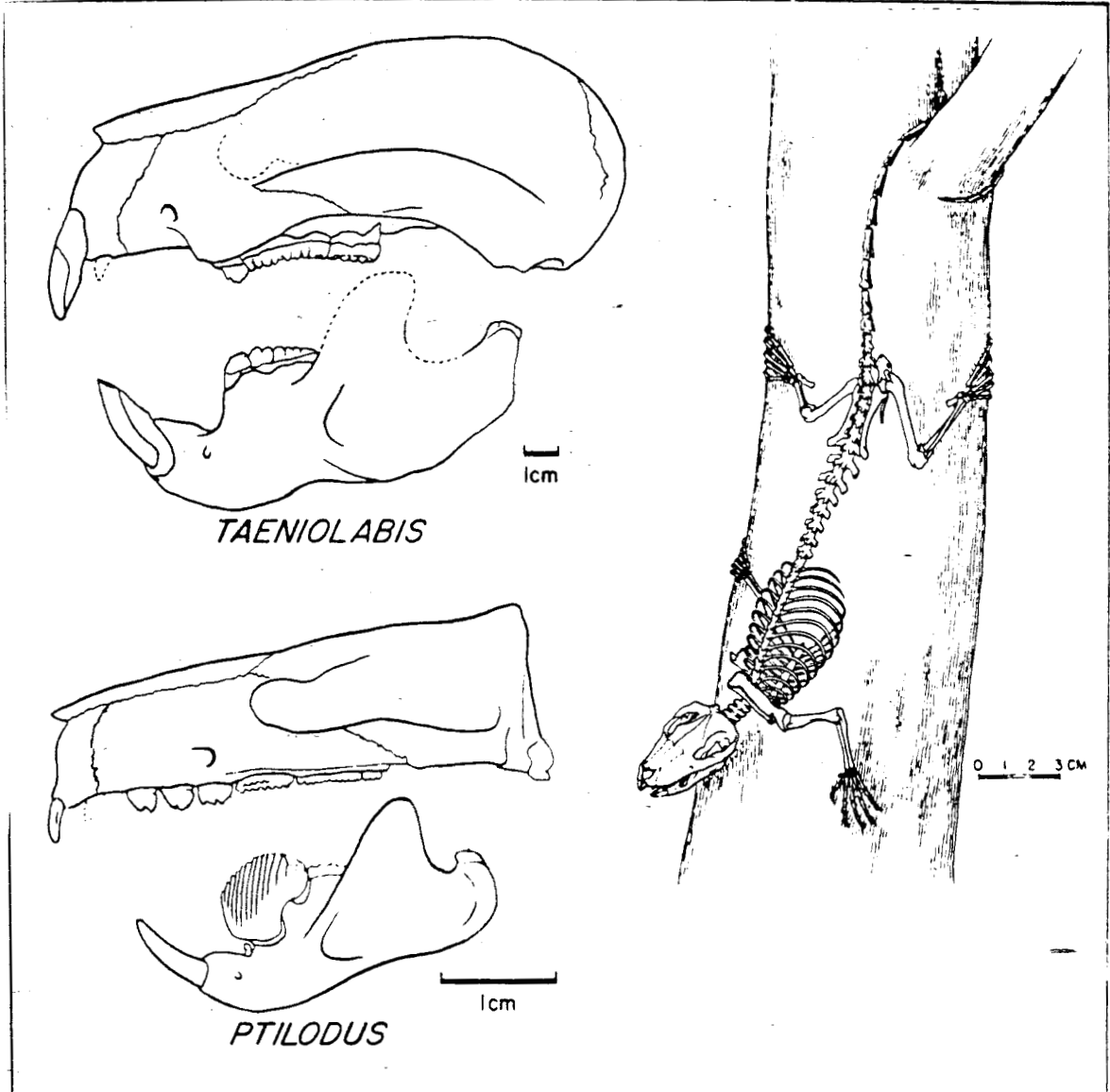


FIGURA 5

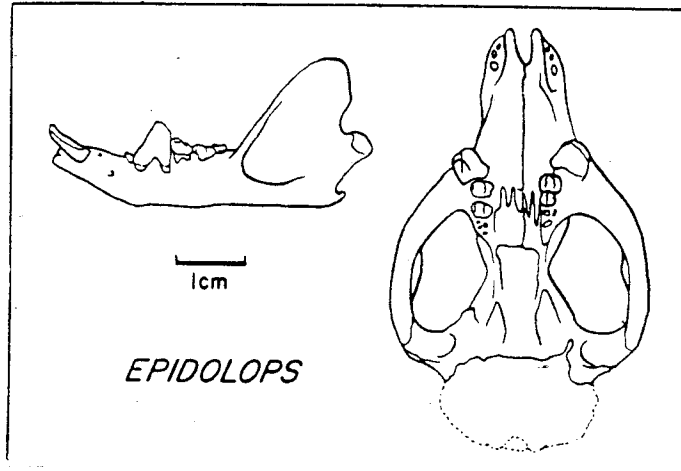


FIGURA 6

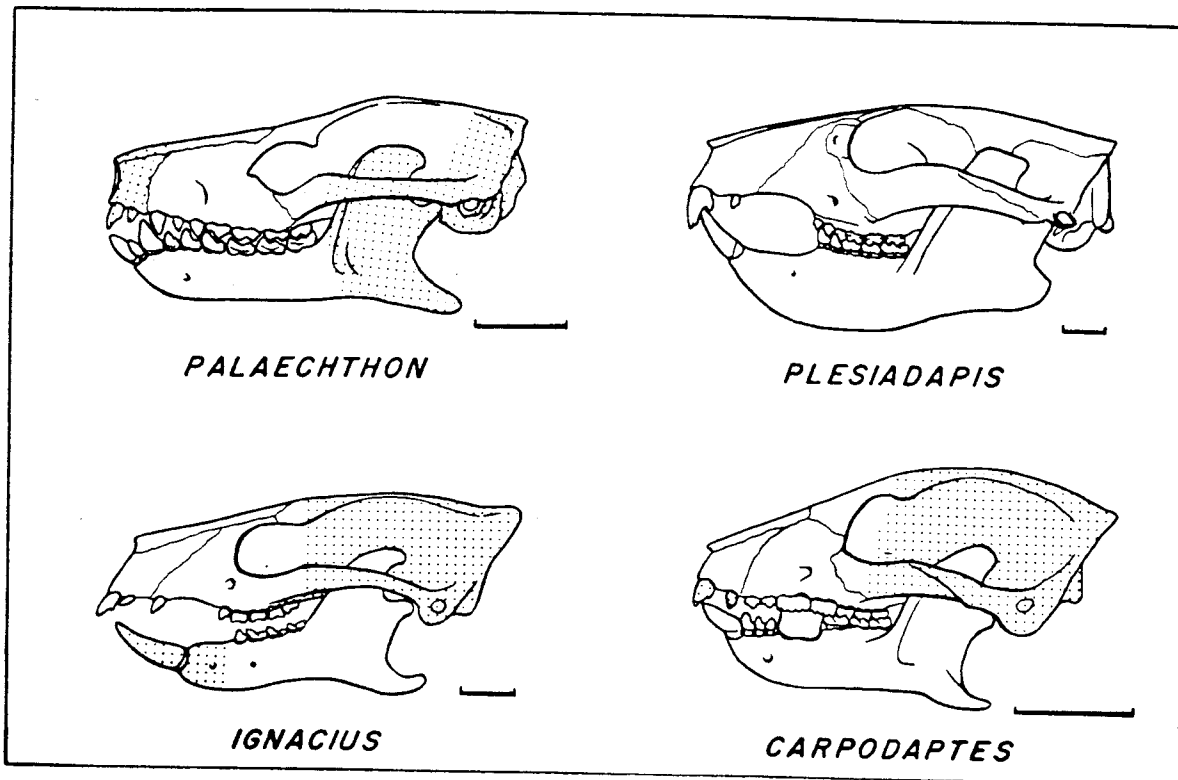


FIGURA 7

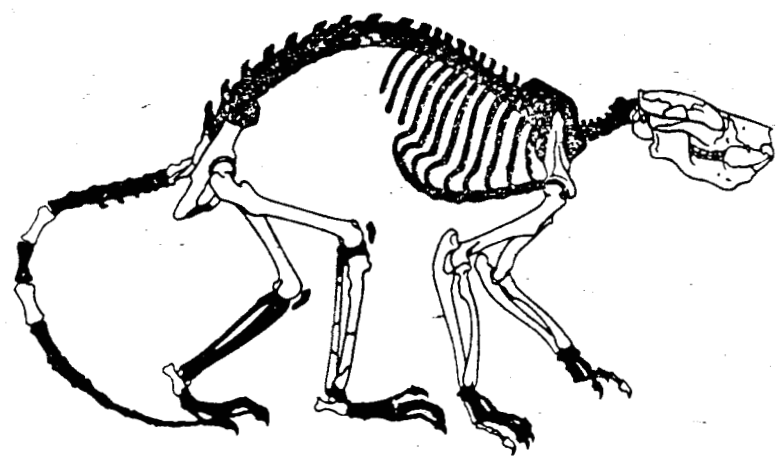


FIGURA 8

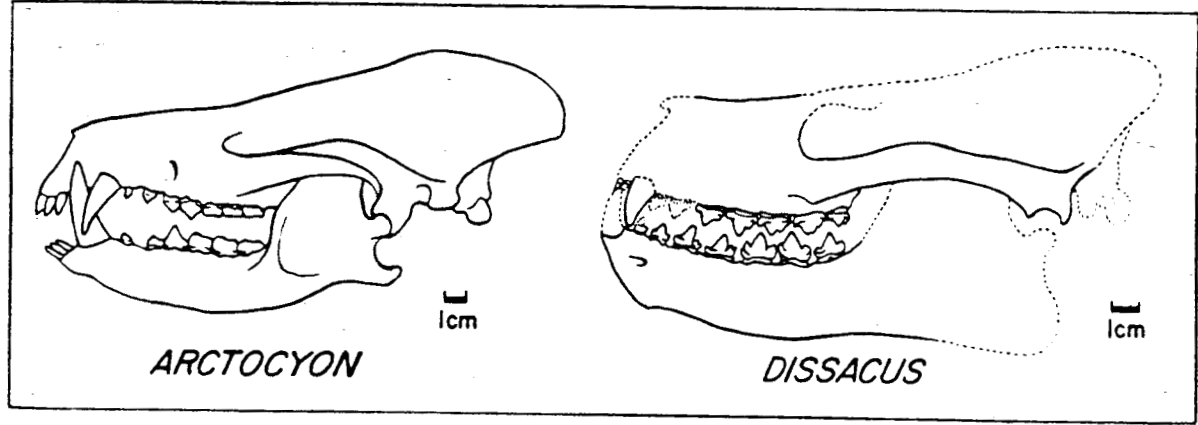


FIGURA 9

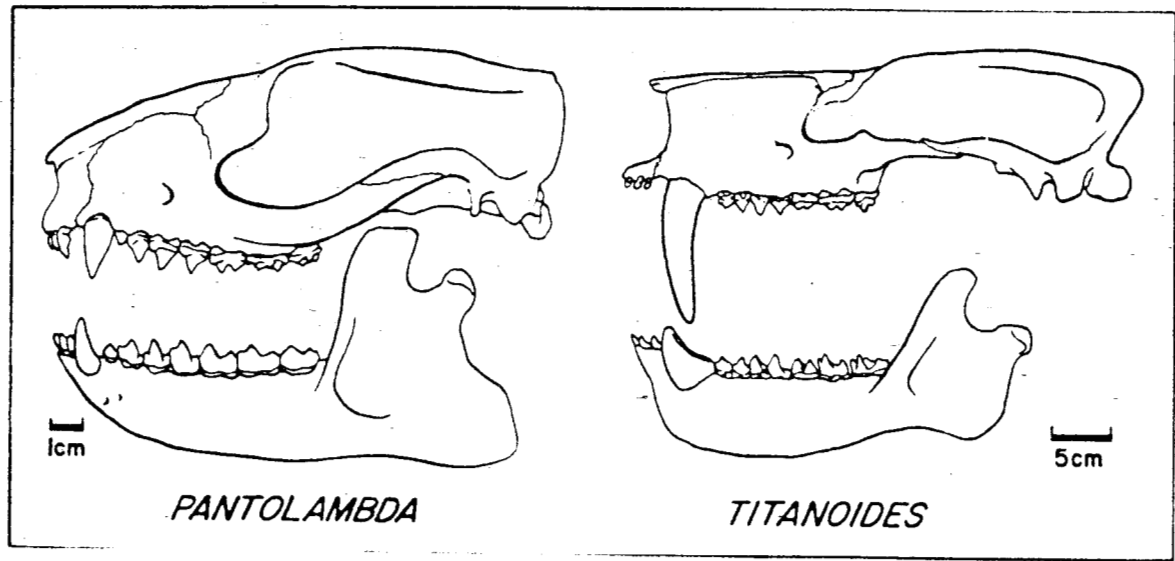
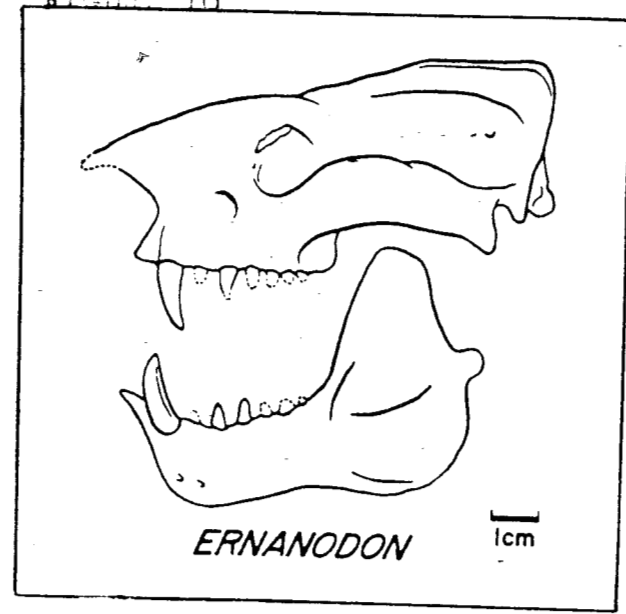


FIGURE 10



In: GINGERICH, P. D. BADGLEY, C., ORGANIZERS, 1984. MAMMALS. NOTES FOR A SHORT COURSE: KNOXVILLE, TENN., UNIV. TENNESSEE DEPT. GEOL. SCI. STUDIES IN GEOLOGY 8, AND THE PALEONTOLOGICAL SOCIETY, p. 128-138.

9.5(a)

NEOGENE FAUNAS OF NORTH AMERICA

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INTRODUCTION.

The second half of the Age of Mammals yields many stories worth telling. The one that I find grandest and most central, however is the "Savanna Story." While featuring it as the principle theme, I hope to weave in the following secondary subjects:

- 1) the adaptive radiation of four representative groups, namely hypsodont horses, camels, elephants and cricetid rodents;
- 2) intercontinental dispersal patterns cross Beringia and between the American continents;
- 3) and faunal turnover patterns.

A good general reference on these and related topics is Kurtz (1971).

SAVANNA STORY

The major climatic trend during the last fifty-million years appears to have been toward cooler and drier conditions, or less equability. And in the centers of continents, behind cordilleras in the zone of prevailing westerlies, or on coasts of cold-water upwelling, this trend has been accelerated. During the mid-Cenozoic on every continent except Antarctica, predominantly forest settings gave way to mosaics of forest and nonforest biomes; savanna, scrub and steppe vegetation expanded, and the importance of grasses increased.

The opening of savannas had a profound effect on mammal faunas (Webb, 1977). Some of the most distinctive groups of the Neogene epoch originated in such settings and some of the most rapid evolutionary outbursts were fomented there. The governing mechanism seems to have been the prevailing climate which controlled the dominant vegetation which in turn provided the principal selective force on land vertebrate faunas. These notions are represented in Figure 1.

HYPSODONT HORSES

The rise of hypsodont horses is a familiar history which illustrates several features of savanna adaptation. In their dentitions, Oligocene horses of the genera *Meshippus* and *Miohippus* developed a complete cheek series of six molariform teeth with lopate (rather than cuspidate) patterns. Within the early Miocene genus *Parahippus* the crown height of each molariform tooth neared completion by addition of a crochet and elaboration of the hypostyle. From early species such as *P. pristinus* to late species such as *P. conatus* body size doubled. Presumably these animals shifted from browsing to mixed feeding. The next step in this horse history is the genus *Merychippus* which appears in the medial Miocene. The late Heningfordian transition to primitive species such as *M. primus* from advanced *Parahippus* such as *P. leonesis* is distinguished by the final connection of the crochet from the proto-loph to the metaloph, and increasing crown height. From mixed feeding *Merychippus* had moved to grazing. The genus *Merychippus* rapidly gave rise to a major radiation of hypsodont horses in the middle and late Miocene (Figure 2). It should be noted that this radiation took place entirely within North America. Only or two hipparionine genera did reach the Old World in the late Miocene; and of course *Equus* so dispersed in the late Pliocene. In the early Miocene, however, the only equid genus that extended its range to Eurasia was *Anchitherium*, a browsing form derived with little change except size increase from *Miohippus*. Likewise in North America, such primitive browsing genera as *Hypohippus*

and *Megahippus* persisted as contemporaries of the progressive grazing genera through most of the Miocene.

HYPSODONT CAMELS

The camels also experienced a Miocene radiation as the savanna biome expanded across the midcontinent of North America. They also developed hypsodont cheek teeth with copious cement and longer, deeper-faced crania to accommodate such dentitions. The most peculiar characteristics of progressive camelids, however are found in their postcranial skeletons. The limbs which even in the Oligocene had quite elongate proportions, became larger and in many groups still longer. The most remarkable were the giraffe camels *Aepycamelus* of the mid to late Miocene, which were closely convergent in size and shape with modern Giraffa. In foot structure the hypsodont camelids experienced a peculiar reversal: instead of remaining unguligrade (on tip toes), as do all other progressive cursorial ungulates, they "fell" to a digitigrade stance, reduced their hooves, and developed a large pad around each digit. (The subordinal name for camels, Tylopoda, refers to this distinctive "cushion foot".) This and a number of related adaptation evidently provided camels with an unusually efficient mode of locomotion in open country. Figure 1. The dominant vegetation in midcontinental North America shifted from forest to savanna to steppe as rain fall declined. Savanna settings were optimum for diverse terrestrial herbivores including grazers and browsers. See text for fuller discussion. Ungulates, especially horses and camels, reached their acme in the late Miocene. Figure 2. Successive genera of horses in the Miocene of North America. Grazers, derived from *Parahippus* and *Merychippus* diversified greatly in the later half of the epoch as savanna settings expanded. With the expansion of steppe in latest Miocene and Pliocene the browsers were lost and grazers severely declined. As with hypsodont Equidae the radiation of Camelidae was confined to North America in the medial to late Miocene (Fig. 3). The more primitive browsing genera survived alongside

the progressive mixed-feeding and grazing genera until about the end of the Miocene. Camelids dispersed outside of North America only in the late Pliocene when large camelines reached the Old World and medium-sized lamines reached South America. Figure 3. Successive genera of camels in the Miocene of North America. Grazers and mixed-feeders diversified in the later half of the epoch as savanna settings expanded. With the expansion of steppe environments in the latest Miocene and Pliocene the browsers became extinct and the grazers and mixed-feeders declined.

ELEPHANTIDAE

When one speaks of large mammals images of elephants almost inevitably come to mind. And no better example of savanna adaptations can be found than among the Proboscidea. The gomphotheres had evolved from the pig-sized Oligocene genus *Phiomia* in North Africa to the rhino-sized genus *Gomphotherium* in the mid Miocene of Europe. And this group was paralleled in its history by the mammutids. Indeed both groups reached North America in the late medial Miocene together. But these early groups were mainly large browsers, not yet true savanna beasts.

The elephantids appear rather suddenly in the late Miocene of Africa; intermediate forms (such as *Stegotetrabelodon*) document the rapid transition between long-jawed gomphotheres and short-jawed elephants. The most evident changes are in the dentition and involve markedly increased hypsodonty in the molariform teeth plus the addition of copious cement. As the gomphotheres lophids are modified to elephant plates, the plate height, plate compression (from front to back) and late number per tooth increase very rapidly. Meanwhile the cranium becomes markedly deeper and, unlike horses and camels, markedly shorter. Elephantids develop "rotary eruption", a unique dental adaptation, in which only one (or parts of two) molariform teeth in each jaw half come into wear at a time, thus permitting a short jaw to accommodate a succession of massive grinders.

The limbs also became proportionally longer in elephantids. They were well-adapted to grazing or mixed-feeding in open-country. The elephants (mammoth) dispersed throughout the Old World during the late Pliocene and early Pleistocene, and reached North America in the early Pleistocene. Gomphotheres reached South America, radiated there and produced some elephant-like adaptations, but elephantids never entered that continent.

CRICETID RODENTS

No order of mammals has proved more successful in distributing widely and diversifying explosively than the Rodentia, and within this order the most successful families have been the Cricetidae and their derivatives, the Microtidae (Voles), and the Muridae (rats and mice). From their Oligocene origins the cricetids diversified in every northern continent; and by the end of the Tertiary, one group had reached South America (with its own subfamilial radiation) and another group had reached Africa and then Madagascar (with another subfamilial radiation). Meanwhile by the mid-Miocene in southern Asia, the Muridae appeared and spread to Australia by the late Pliocene.

The remarkable success of these "mice" has been attributed partly to their efficient chewing mechanism, with lophate teeth and masseter muscles extended forward onto the rostrum, and partly to their very efficient reproductive system with large litters on a monthly basis. The cheek teeth consisted of only three molars in each jaw half. Each molar progresses from a simple four-cuspid pattern (as in *Eumys*) to a lophate pattern with wide reentrant valleys (as in *Peromyscus*), and finally to a tall (sometimes rooted, often ever-growing) multiprismatic pattern with heavy cement (as in Microtidae) of the Pliocene in north temperate steppe and tundra environments. Both in their appearance and in their function, microtid teeth can be compared to elephantid teeth. Both represent grazing adaptations.

INTERCONTINENTAL DISPERSALS

Each group of savanna-adapted organisms originated and radiated in the heartland of a single continent. Only later did some members of the group disperse to other continents. One can envision the savanna fauna of each continent as a deep reservoir and the land bridges between them as spillways well up on their sides. As the reservoirs fill up or as the spillways are lowered somewhat, the contents of adjacent reservoirs mix. In such a manner intercontinental dispersals between savanna faunas took place. The secular trend toward climatic deterioration through the late Cenozoic increased the possibilities of such dispersals.

In Figure 4, I have attempted to summarize the Neogene intercontinental dispersals between North and South America, and also between North America and the Old World. In this simplified scheme only two North America and Asia and the Isthmian Link between North and South America. We may focus first on Beringian interchanges and secondly on Isthmian interchanges.

BERINGIAN INTERCHANGES

The Bering Land Bridge permitted several sets of dispersals between the New World and the Old World. Although the exact tabulation of land mammals genera will be subject to further improvements, the following general points seem secure: 1) a major set of dispersals reached North America from Asia in the Hemingfordian (about 20 million years ago);

2) another major set of dispersals moved reciprocally in both directions during the Hemphillian especially in the late Hemphillian (about 5 million years ago);

3) the rest of the Pliocene and Pleistocene saw frequent, nearly continuous dispersal events;

4) the direction of movement changed markedly from a nearly balanced reciprocal flow in the Mio-Pliocene to a predominance of taxa moving from Asia to North America in the Pleistocene; and,

5) the habitat preference of the mammals moving through Beringia shifted from forest and savanna in the Miocene and Pliocene to steppe and in the Pleistocene.

THE GREAT AMERICAN INTERCHANGE

The history of South America's mammals fauna took place almost entirely in isolation, and because of that, the few limited episodes of immigration take on great importance. By the beginning of the Cenozoic three major groups of mammals are recorded, in South America, namely marsupials, edentates, and ungulates. A second introduction of new mammals, evidently by island-hopping or "Noah's Ark" across the Caribbean, brought ceboid primates and hystricognath rodents in the late Oligocene. A third invasion consisted solely of the procyonid (raccoon) *Cyonasua* in the late Miocene. To account for the entire Tertiary mammal fauna of South America, one need postulate only six immigrant taxa, there at the beginning two in the late Oligocene and one in medial Miocene.

Figure 4. Neogene immigrants to North America. From the left, land mammal genera from Eurasia; from the right, land mammal genera from South America. An early peak of Asiatic immigrants marks the Hemingfordian (20 million years ago); nearly continuous peaks also mark the Pliocene and Pleistocene record.

The Great American Interchange produced late Pliocene and Pleistocene immigrations from South America. The great American Interchange introduced new species of mammals into South America on a dramatically augmented scale. In the late Pliocene a continuous land bridge through Panama was produced by convergence of the Cocos Plate and central America, and this rapidly led to a dozen species of land mammals moving in each direction between the American continents. The results in South America included extensive replacements of native fauna and intense radiation of immigrant groups, including canids, llamas, and most notably, cricetid rodents. Half of the present

land mammal genera of South America are descendants of Plio-Pleistocene immigrants (Marshall et al., 1982). The reciprocal side of the interchange was less impressive, although it carried South American mammals into North America for the first time since the beginning of the Cenozoic. Most emigrants from South America diversified little, all but three became extinct, and none extended beyond North America into the Old World. They did, however, overwhelmingly colonize Central America, which is now assigned by zoogeographers to the Neotropical Realm.

FAUNAL EQUILIBRIUM PATTERNS.

The species-equilibrium theory was introduced by MacArthur and Wilson for island biotas over ecological time, but it has since found wide application in tracking the biota of continents, ocean basin, or the whole globe. The faunal equilibrium hypothesis can be put to an especially striking test by the South American fauna. The turnover rates and numbers of genera may be compared before, during and after the interchange. Generic diversity, which had been near 72 for at least 6 million years, rose to 84 in the Uquian (at the peak of the interchange) and to 107 in the Ensenadan (also during the interchange); turnover rates also increased markedly as predicted by equilibrium theory.

Equilibrium theory also predicts that the number of immigrants moving from a source fauna to an "island" will depend on the diversity of that source fauna. The same principle applies to a reciprocal intercontinental interchange. For example North America contributed 60 percent more genera to the great American interchange than South America, and that is the approximate difference in their respective faunal diversities in the Pliocene. A similar imbalance occurs across the Bering Strait in the latest Miocene and Pliocene, with Asia contributing a larger number of immigrants to North America than the reverse.

Equilibrium theory recognizes a dynamic interrelationship between the immigration rate (supplemented by new cladogenetic taxa) and the extinction rate. It thus predicts that waves of

755 765

new immigrant taxa will surely produce waves of extinct taxa. In unequal interchanges between continents of different sizes and carrying capacities, the smaller continents will suffer more. Thus the theory predicts many extinctions throughout the Pliocene and Pleistocene, and proportionally heavier impacts in South America and in North America than in the Old World.

LATE CENOZOIC EXTINCTIONS.

Increasing severe and frequent extinctions swept through the land mammal faunas of the late Cenozoic. The largest episode in North America came in the late Hemphillian when nearly 40 genera vanished. The final late Pleistocene extinction wave was more intense than the others of the late Cenozoic, and thus seems to demand a more direct explanation than the others. The two regularly cited causal hypotheses are human predation and climatic change, and the new book by Martin and Klein (1984) still does not wholly resolve this debate between them. I will conclude by chiefly advocating the climatic hypothesis of the Pleistocene extinctions.

Several difficulties beset the human hunting theory. Most of the Pleistocene extinctions in Europe, Asia, and Beringian parts of north America came long after they were inhabited by bands of hunting peoples well equipped with lithic tool kits. In temperate North America, however, there is close coincidence between the first extensive Pale-Indian records and the last records of most species of mega fauna. Even so major faunal and floral reorganization indicate severe climatic shifts at the time, and several groups disappeared at high latitudes but survive at low latitudes. In a longer view, it is quite clear that several major extinction events resulted from climatic deteriorations of the last ten million years. The increasingly severe climates of the late Cenozoic led to the replacement of the old native savanna faunas by an increasingly cosmopolitan steppe fauna, especially rich in grazing microtine rodents and remarkably impoverished in large mammals.

FIGURE 1

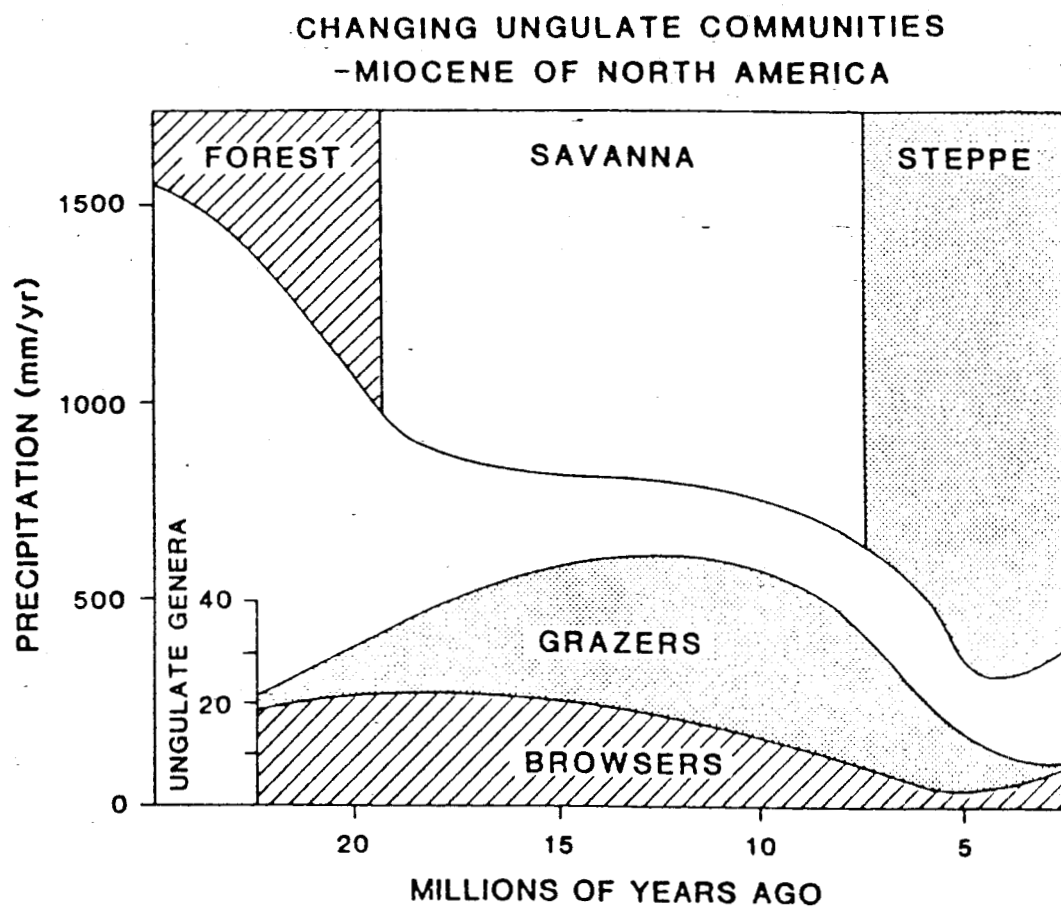


FIGURE 2

Abb

MIOCENE EQUIDAE OF NORTH AMERICA

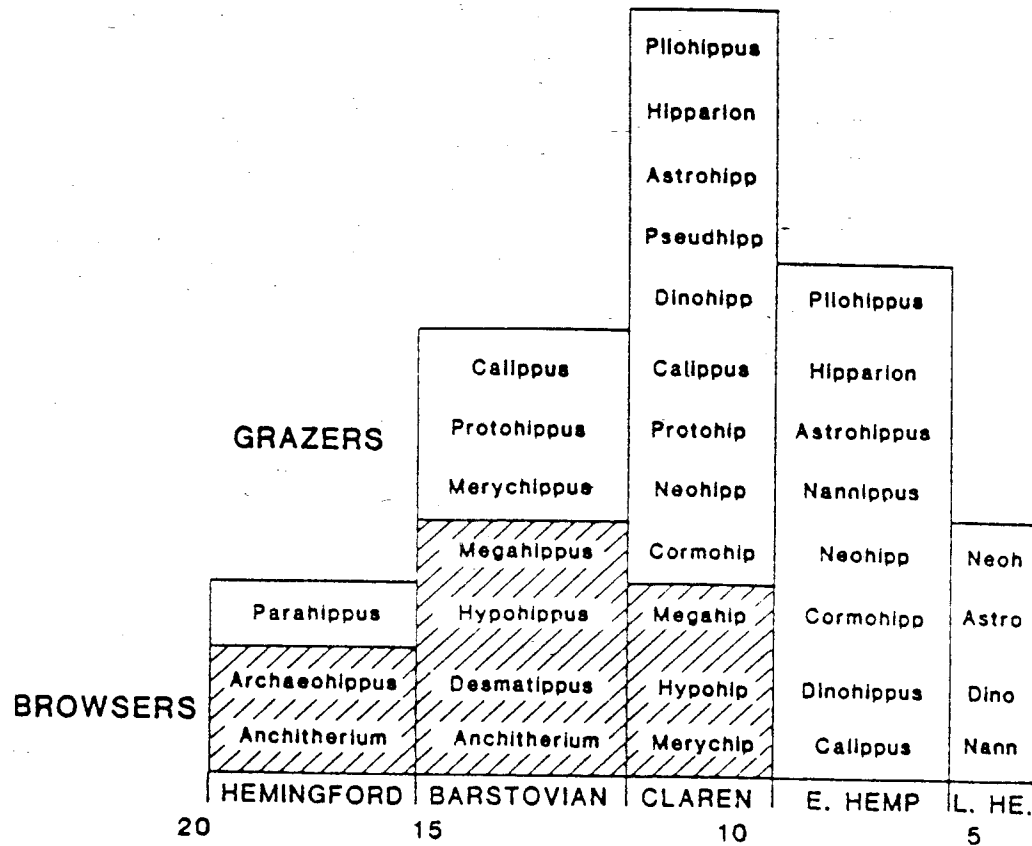


FIGURE 3

MIOCENE CAMELIDAE OF NORTH AMERICA

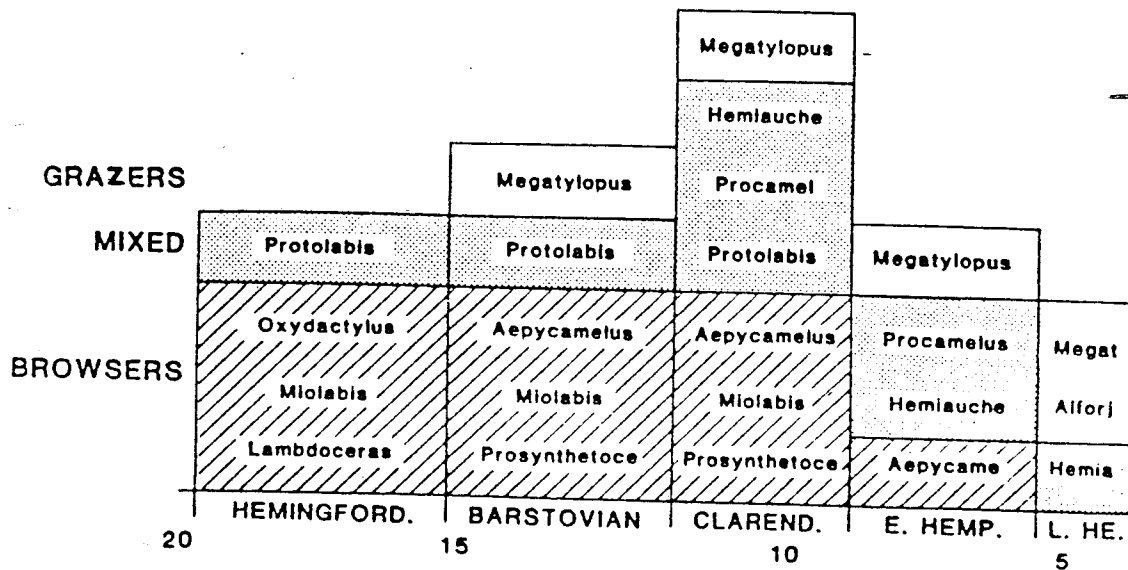
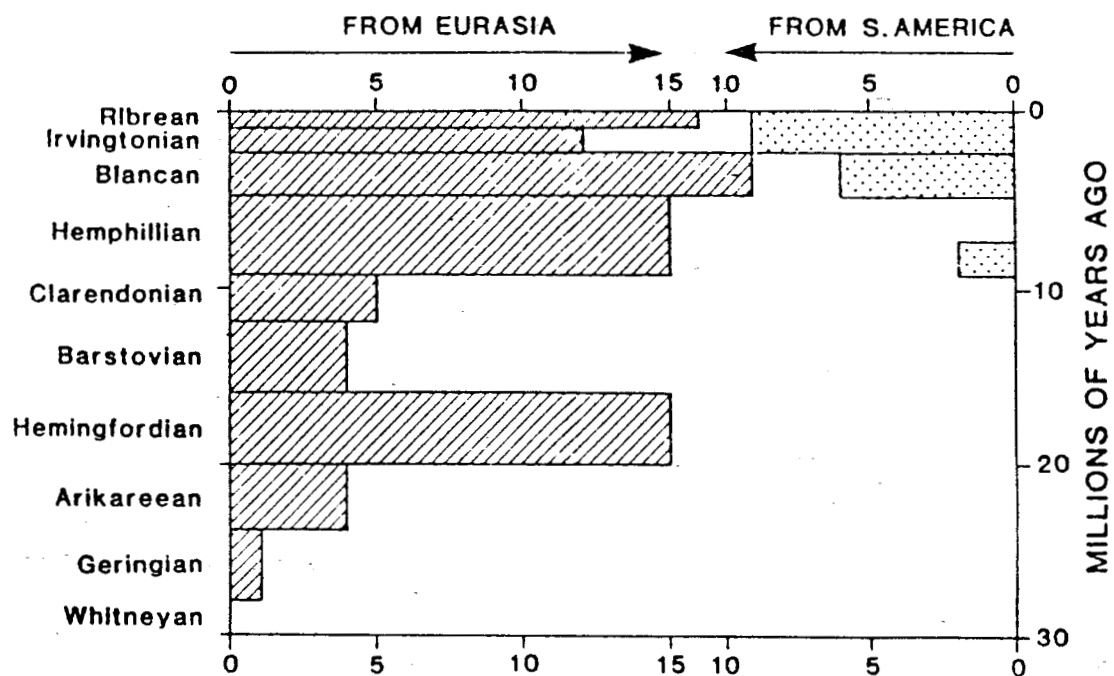


FIGURE 4

IMMIGRANT LAND-MAMMAL GENERA IN NORTH AMERICA



LA ASCENDENCIA DEL HOMBRE UNA INTRODUCCION.

In: GINGERICH, P.D. BADGLEY, C., ORGANIZER, 1984. MAMMALS. NOTES FOR A SHORT COURSE: KNOXVILLE, TENN., UNIV. TENNESSEE DEPT. GEOL. SCI. STUDIES IN GEOLOGY 8, AND THE PALEONTOLOGICAL SOCIETY, P.182-198.

"Human evolution. Catherine Badgley Museum of Paleontology university of michigan Ann Arbor, Michigan 48109.

INTRODUCTION.

The evolutionary history of humans is well understood in outline, compared to that of many other groups of mammals. But human evolution remain enigmatic in its details, and these are compelling both scientifically and personally because they relate to the biological uniqueness of humans. Humans are placed in the primate family hominidae, which, in traditional classifications, contains a single living species, *Homo sapiens*. The closest living relatives of humans are great apes: the chimpanzees *Pan paniscus* and *Pan troglodytes*, the gorilla *Gorilla gorilla*, and the orangutan *Pongo pygmaeus*. These apes have traditionally been placed in the family pongidae as the sister group of hominidae. Living Hominidae and Pongidae, together with Hylobatidae (gobbons) comprise the modern representatives of the primate suborder Hominoidea.

In the last two decades, interpretations of the fossil record pertaining to human evolution have changed significantly (Pilbeam, 1980); and there is no consensus viewpoint even now (Ciochon, 1983). A dramatic increase in the recovery of vertebrate fossils from continental, Old World sites has revealed the presence of moderate taxonomic diversity of hominid since the early Miocene. Nonetheless, the hominoid fossil record is fragmentary- in terms of the completeness of individual specimens, the number of body part

represented, and the distribution of fossil through geologic time. Studies in molecular evolution of proteins and nucleic acids have provided a consistent "molecular" view of the relative divergence times of living hominoids. Current general agreement about broad cladistic patterns among the hominoidea reflects a synthesis of the branching pattern indicated by comparative anatomy and molecular systematic, with absolute divergence times loosely based on the temporal distribution of fossil.

It is difficult to place humans in the proper perspective of mammalian biology, for both cultural and biological reasons. Religious views advocating a separate origin and purpose of humans versus the rest of "nature" are deeply embedded in western belief systems. Even though these views no longer prevail intellectually, they are still widespread implicitly in culture and politics. For over a century, scientific view of hominid origins and of man's place in nature have reflected preconception and prejudice as much as natural history and fossils (Pilbeam, 1980; Landau et al, 1982). From the standpoint of anatomy and behaviour, humans possess the shared derived features of mammals and many characteristic primate features. But, humans are unquestionably striking and typical mammals. Both the evolutionary enhancement of typical primate attributes and the evolution of features unique to hominids have resulted in a primate lineage of extraordinary behavioral and ecological character.

Below, I review the comparative biology of living hominids to examine similarities and differences between humans and their close relatives, then outline the hominoid fossil record to document where and when characteristically human features were acquired. Phylogeny follows that combines information from hominoid molecular systematic and fossil. This review concludes with a brief consideration of the environmental context of human evolution. Useful general references on human evolution include Lee and Devore (1968), Butzer and Izacc (1975), Jolly and Plog (1976), Legros Clark (1978), Wolpoff (1980), Zohman (1982), Ciochon and Corruccini (1983), and Pilbeam (1984).

HUMANS AND PRIMATES.

Humans exhibit many anatomical and behavioral features characteristic of primates in general. Manual dexterity, high visual acuity, diurnal activity patterns, relatively long maturation times and life spans, relatively large brain volumens, and elaborate social behavior are common among living primates. Many features viewed as characteristically human--e.g., intelligence, language, culture, tool-making, and long-lasting social bonds--are but extensions of trends already widespread within the primate order. It is important here to distinguish attributes that are original in human evolution from those that are not original but are nonetheless fundamental.

Skeletal features that humans share with most primates include the retention of five digits on both the hand and the foot, an opposable thumb (many primates also possess an opposable big toe), flat nails instead of claws on the terminal phalanges, short faces, and a high ratio of brain size to body size relative to other mammals. All catarrhines (Old World monkeys, apes, and humans) share the same dental formula: $2/2, 1/1, 2/2, 3/3$. The premolars retained are considered to be the third and fourth premolars. Hominoids lack tails and possess relatively larger brains than other primates. Their cheek teeth are low-crowned; hypoconulids are present on the lower molars, although these are occasionally, with attendant anatomical modifications in the habitually bipedal, with attendant anatomical modifications in the pelvis, hip, knee, ankle, foot, vertebral column, and skull. Dentally hominids differ from pongids (figure 1) in having a less heterodont dentition, with the anterior teeth (incisor and canines) smaller in relation to cheek teeth than in pongids. Pongids, hylobatids, and Old World monkeys have sexually dimorphic canines: males have larger canines in relation to cheek-tooth size than females. The canines of modern humans exhibit little dimorphism in this trait. Finally, hominids are distinguished by large brain sizes relative to body sizes. Living humans have a ratio of brain size to body size over twice that of a chimpanzee (Jerison, 1973; Zihlman, 1982). Figure 1. The skull

of male gorilla (a) and of modern *Homo sapiens* (B) 1 (not to scale), to illustrate differences in dentition and relative brain size (After LeGros Clark, 1978).

Living primates vary in many aspects of natural history, and in some instances, humans represent a departure from the prevailing trend. Most living and extinct primates are associated with forested habitats in tropical to sub-tropical climatic regimes. Hominids, including the earliest australopithecines, occur in more open, seasonal habitats. By the end of the Pleistocene, humans had come to inhabit all the climatic extremes of the terrestrial environment.

Most primate diets are opportunistic and omnivorous in terms of the range of foods eaten, but are heavily vegetarian in terms of the volume of food consumed. Primate foods include fruits, flowers, seeds, leaves, stems, gum tubers, bird eggs, and insects. Among the hominoids, gibbons, orangutans, and chimps are predominantly frugivorous with leaves and other plant parts as secondary components. Gorillas rely mainly on leaves, shoots, and stems, with fruit as a smaller dietary component. Chimps are known to eat meat on occasion (Teleki, 1975). Humans depart from all of these dietary patterns in being omnivorous: their diets are still predominantly vegetarian, but a significant portion is also meat. Living hunter-gatherers have diverse vegetarian diets that include nuts, fruits, berries, leafy greens, tubers, and resins; hunted meat rarely exceeds 40% of the diet (Lee, 1968). Many sedentary societies rely heavily on grains (the seed product of the Graminae, the Grasses), such as wheat, rice, and corn. This dietary dependence is distinctive among the hominoid and would not be possible among humans without food preparation such as grinding and cooking. Primates are very selective foragers and consume food on the spot. Humans differ in transporting, storing, and sharing a significant portion of their food.

Most primates live year-round social groups, comprising one or more adults of both sexes and the offspring of the female(s). Most species are polygamous, although styles of polygamy and the amount of interaction among males and females

varies widely. For many species, females remain in their natal groups and breeding males are immigrants from other groups. Wrangham (1980) proposes an ecological model for these female-bonded groups in which the primary determinants of group size is feeding competition. Species that are polygamous are also sexually dimorphic in body size, with males always the larger sex. Size dimorphism reaches its greatest extreme in the orangutan and gorilla; males may weigh twice as much as females (Zihlman, 1982). Size dimorphism may have been even more extreme among some Miocene hominoids, similar cheek teeth of *Ramapithecus* and *Sivapithecus* at several late Miocene fossil sites (de Bonis, 1983).

Among living hominoids, the great apes are polygamous and gibbons are monogamous. Orangs are the most solitary of the apes. Groups consist of adult females and their offspring in adjacent feeding areas. The territory of a dominant male typically overlaps the smaller territories of several females. Gorillas live in small groups comprising a dominant male, several females, and their offspring. Chimps live in multi-male, multi-female groups, in which males tend to be more closely related than females (Nishida, 1979). Chimp intermale aggression is highest among the apes; aggression is greatest in inter group conflicts, and next over intra-group dominance hierarchies. Gibbons live in small social units of adult male and female plus offspring. Humans are usually monogamous, with a similar social unit. But these families typically aggregate in loose or tight social, economic, and political units. These units may shift composition and size, depending on resource distribution including seasonal variation.

Relative to other primates, human infants are altricial (underdeveloped) at birth. Both the central nervous system and the immune system are immature, and there is long period of postnatal brain growth. Chimp and human have approximately equal ratios of brain weight to body weight in newborns. But, brain advancement (newborn) world. The African apes inhabit forest of central Africa; no fossils are known for *Pan* and *Gorilla*. Gibbons live today only in southeast Asia; they have a sparse and am-

biguous fossil record in Africa and Eurasia. The Orangutan lives today only in Indonesia; the only fossils attributable to *Pongo* are from the Pleistocene of southeast Asia. With the exception of *Homo sapiens*, hominoids are less "successful" -- in terms of taxonomic diversity, abundance, and distribution -- now than during most of the Neogene. Figure 2 The distribution of Neogene Hominoid localities in the Old world.

The distribution of Neogene hominoid fossil localities (figure 2) exhibits a distinctive geographic and temporal pattern. Early Miocene hominoids of the *Proconsul* group are known exclusively from East Africa, primarily from sites located in the western Rift Valley (Andrews and Van Couvering, 1975). During the middle Miocene, the East African record is very poor, but hominoids (the *Dryopithecus* group and *Ramapithecus* group) occur widely through Eurasia, from Spain to China. Hominoids persist until about 7 mybp in eastern and southern Asia, then are unknown in Eurasia until the middle Pleistocene. In Africa, a few hominoid fossils of uncertain affinity are known at the end of the Miocene. Then, the fossil record is fairly dense from about 3.7 mybp on in East Africa, mainly from the eastern Rift Valley, and from about 3.0 mybp on in southern Africa (figure 3). The temporal distribution of hominoid fossils from East Africa matches that of other vertebrate fossils and reflects mainly the disparate times of rifting and rapid sedimentation in the two arms of the Rift Valley, with the eastern Rift Valley experiencing more recent tectonic activity (Behrensmeyer, 1982). The virtual absence of hominoid and other vertebrate remains in East Africa from 14-4 mybp cannot be taken as evidence for the absence of living hominoid. In Eurasia, however, vertebrate faunas are known through the Neogene, hence, the absence of hominoids from the late Miocene until the Pleistocene apparently reflects the disappearance of the Miocene forms. Figure 3 The temporal and geographic distribution of Neogene hominoid localities.

Much of the hominoid fossil record consists of jaws and isolated teeth, although cranial and post-cranial elements are finally becoming known for a few taxa. With the rare exception,

hominoid fossils are uncommon at the sites where they are known. The predominance of dental material, especially cheek teeth, has not permitted unambiguous evaluation of taxonomic diversity at the genus and species levels (Badgley et al., 1984; Pilbeam, 1984). For the present discussion, I employ the taxonomically neutral terms "dryomorph" and "ramamorph," after Ward and Pilbeam (1983), with reference to the miocene Homonoids. The relationships of these terms to a conventional taxonomy is given below. It should be stressed, however, that no taxonomic scheme is in general agreement, and this one can be considered a representative frequency distribution rather than a definitive scheme. Information is drawn from Freagle and Kay (1983) and Kelly and Pilbeam (in press).

SYSTEMATICS OF FOSSIL AND LIVING HOMINOIDEA

Propliopithecidae: *Aegyptopithecus*,

Propliopithecus, *Pliopithecus*.

Proconculidae (African dryomorphs), *proconsul*, *Rangwapithecus*, *Dendropithecus*, *Limnopithecus*, *Micropithecus*.

Dryopithecidae: (European dryomorphs), *Dryopithecus*, *Rudapithecus*, *Bodvapihthecus*.

Ramapithecidae: (ramamorphs), *Ramapithecus*, *Sivapithecus*, *Renyapithecus*, *Ouranopithecus*, *Gigantopithecus*

Hylobatidae: *Hylobates*, *Symphalangus*

Pongidae, *Pongo*, *Pan*, *Gorilla*

Hominidae: *Australopithecus*, *Homo*

A morphologically distinctive suite of early to middle Miocene hominoids, the dryomorphs, is known from East Africa and Europa. The East African record begins at about 23 mybp with the earliest specimens of *Proconsul* 1 from Meswa Bridge, Kenya, and continues until about 14 mybp at ft. Ternan (Andrews, 1981). The African material includes at least eight species: three small-bodied ones the size gibbons (*Limnopithecus*, *Dendropithecus*, *Micropithecus*) and five medium sized forms (2kg) three species

of *Proconsul* 1 and two species of *Rangwapithecus* 1. In dental features, the African dryomorphs exhibit variable enamel thickness (most forms have thin enamel); canines are dimorphic; incisors are large relative to cheek teeth; molars bear cingula. These hominoids had moderately long faces; they moved quadrupedally in an arboreal setting (Rose, 1983). The European dryomorphs, including *Dryopithecus*, *Rudapithecus* 1 and *Bodvapihthecus* 1 (Kelley and Pilbeam, in press), have thin enamel, molar cingula, dimorphic canines, small to moderate incisors, and a sectorial P. The earliest Old World monkeys are known from about 19 mybp. probably, the earliest dryomorph material postdates the split of hominoids and cercopithecoids (Pilbeam, 1984).

A second distinctive suite of miocene hominoids, the ramamorphs, is known primarily from the middle to late Miocene of Eurasia. Common taxa in this group are *Ramapithecus* 1 and *Sivapithecus* 1, although there is currently dispute and ambiguity about the taxonomic diversity represented by these taxa (Kay, 1982; Badgley et al., in press). The earliest ramamorph material consists of recently discovered specimens attributed to *Sivapithecus* 1 from Bylux in Northern Kenya (Walter and Leakey, 1984); this site is dated at 17 mybp and is thus contemporaneous with the period of maximum diversity of the African Dryomorphs. The body size range of ramamorphs is 15-17 kg, with "*Ramapithecus*" 1 referring to the smaller form (15-20 kg) and "*Sivapithecus*" 1 referring to material mainly in the range of 40-50 kg. (Pilbeam, 1980). These hominoids exhibit very thick enamel on the cheek teeth, moderate canine dimorphism, megadonty (cheek teeth very large in relation to body size) and a somewhat molarized. Molar cingula are absent. Incisors and canines are relatively small in *Ramapithecus* 1 and larger in *Sivapithecus* 1 represent female and males, respectively small in *Ramapithecus* 1 and larger in *Sivapithecus* 1. It is plausible, at some sites, that *Ramapithecus* 1 and *Sivapithecus* represent females and males, respectively, of one species; if so, then the sexual dimorphism in body size is greater than that of any living primates (DeBonis, 1983; Wu et al.;

1983). Postcranially, ramamorphs are not well known, except for unassociated skeletal elements of the forelimb, hindlimb, and foot of *Sivapithecus*, from Siwalik sites of Pakistan (Pilbeam et al., 1980). The functional interpretation of these elements indicates that *Sivapithecus* moved quadrupedally, with the main emphasis on climbing, and lacked the locomotory specialization for knuckle-walking, suspension, or bipedalism of living large hominoids (Rose, 1983, in press). The only cranial material known from ramamorphs is from Lufeng, a late Miocene site in southwest China; nearly complete skulls of both the large and small hominoid have been recovered (Lu et al. 1981; Wu et al. 1982). They are crushed flat and it is not possible to estimate brain volumes.

The hominoid fossil record is almost blank between 7 and 4 mybp. The earliest undisputed hominoid fossil comes from Laetoli in northern Tanzania, at 3.7 mybp and Hadar in Ethiopia at around 3.0 mybp, and are considered represent one taxon, *Australopithecus afarensis* (Johanson and White, 1980). This hominid is fully bipedal, as evidenced not only by the morphology of hindlimb elements, but also by a remarkable set of three hominid trackways at Laetoli (Leakey and Hay, 1979; Hay and Leakey, 1982). Recently, Stern and Susman (1983) have argued that this hominid was also capable of considerable arboreal maneuverability as well. *Australopithecines* are known from East Africa and South African localities, from 3.7 - 1.0 mybp. At least four species are known (*A. afarensis*, *A. africanus*, *A. robustus*, and *A. boisei*), with precise ancestor-descendant relationships unresolved at present. One of the major controversies is whether *A. afarensis* or *A. africanus* lies at the base of the split between later *australopithecines* and *Homo* (White et al., 1983). The youngest *australopithecine*, *A. boisei*, robust form from East Africa, disappeared around 1 mybp. Brain volumes of the early *australopithecines* are 450 - 550 cc, 20 - 30 % larger than that of ape of similar body size. Cheek teeth are megadont with thick enamel, and the P is molarized. Canines are small, incisiform, and exhibit little sexual dimorphism. Size dimor-

phism is moderate, greater than in modern humans.

The first stone tools are known from Koobi Fora and Olduvai Gorge at about 2.1 mybp. While tools have been considered to be associated only with *Homo*, their appearance antedates the earliest record of *Homo* by about 0.3 my. The distribution of vertebrate remains and stone tools at these early sites suggests that hominids used tools for food-processing, possibly for hunting, and may have established long-term occupation sites (Isaac, 1978).

The first appearance of *homo* (*H. habilis*) is from Koobi Fora at about 1.8 mybp. The earliest *Homo* is distinguished by a considerably larger brain (700 cc) and by different tooth proportions; cheek teeth are smaller relative to the anterior dentition, compared to *Australopithecus*. By about 1.75 mybp, *H. habilis* had disappeared and was replaced by *H. erectus* in East Africa. *H. erectus* is characterized by larger body size and greater robustness than *H. habilis*, a further expansion of brain size (850-1000 cc), and a further diminution of cheek tooth size. This species was the first to leave Africa and become widely distributed throughout the Old World. Fossils attributed to *H. erectus* are known from Java about 1 mybp and from China at about 0.8 mybp. Around 0.3 mybp, *H. erectus* was succeeded by "Archaic" *H. sapiens*, including the Neanderthals. "Modern" *H. sapiens* appeared about 40,000 years bp, succeeding archaic *H. sapiens* abruptly. Morphological trends in the lineage leading from *H. erectus* to modern *H. sapiens* include an increase in brain size, a decrease in tooth size, and in the robustness of skeletons and skulls. Changes in distinctive stone-tool industries from Africa and Europe match.

Figure 4 represents an attempt to synthesize the hominoid fossil record and molecular systematics into a provisional phylogeny. In this scheme, hominoid evolutionary history contains multiple radiations from the early Miocene onward. The radiation of large hominoids begins around 16 mybp, with a split into a predominantly Eurasian clade (ramamorphs + *Pongo*) and predominantly African clade

(leading to *homo*, *Pan*1, and *Gorilla*1). The common ancestor of the clades leading to *Pongo* 1 and *Homo* 1 is undocumented by fossils, as is the common ancestor of *Homo* 1 and the African apes. Thus, much remains to be learned, even of this well studied case of mammalian evolutionary history. Figure 4. Phylogeny of Neogene Hominoids, incorporating molecular and anatomical data with the temporal and geographic distribution of fossils. Based on information in Pilbeam (1984) and (Kelley and Pilbeam, in press).

THE ENVIRONMENT AND HUMAN EVOLUTION.

The ecological significance of the changes in geographic distribution and morphology of hominoids can be understood, in part, in terms of global paleoenvironmental history. In the last two decades, there has been considerable progress in reconstructing the positions of land masses and oceans, ocean temperatures and circulation patterns. These physical factors are important determinants of climate. Geological, floral, and faunal studies of the continental Neogene of the Old World indicate changes in geographic barriers and in the distribution and composition of terrestrial plant and animal communities.

Hominoids probably arose in Africa, since they are known only in Africa from the time of *Aegyptopithecus* 1 through the radiation of the African *Dryomorphs*. The first hominoids outside Africa are known from Arabia and Turkey in the middle Miocene (Figure 3). Until the early Miocene, Afro-Arabia was an isolated land mass, lying South of its present position and separated from Eurasia by the Tethyan seaway. During the period 19 - 16 mybp, contact occurred between Arabia and Eurasia in a pattern of intermittent land bridges, large island, and narrow seaways through what is now the eastern Mediterranean and the Middle East. Somewhat later, contact occurred in the western Mediterranean. These land connections facilitated the dispersal of land animals from Eurasia into Europe and vice versa. Two main dispersal phases are recognized for mammals, one around 18 mybp and

another around 15 mybp. In the first dispersal, more taxa moved from Asia to Africa; in the second, more taxa moved from Africa to Asia (H. Thomas, paper presented April 6, 1984, at the "Ancestors" conference, American Museum of Natural History, New York). Hominoids may have expanded into Eurasia in multiple dispersal events. The existence of morphologically heterogeneous hominoids--i.e. the European *dryomorphs* and the African *ramamorphs*--both with morphological counterparts in Africa--suggests that large hominoids dispersed from Africa more than once.

Within East Africa, tectonic activity centered in the western Rift Valley and orogenic uplift of the East African Highlands led to the establishment of complex biogeographic provincialism in the early Miocene (Andrews and Van Couvering, 1975). Mammalian faunal endemism occurs as early as 18 mybp between sites in the western Rift Valley (e.g., Rusinga) and the eastern Rift Valley (e.g., Buluk), with *dryomorphs* of the *Proconsul* 1 group to the west and the earliest fossils with *ramamorph* affinities to the east. Climatic, floral, and faunal differences occur as well (A. Walker, paper presented April 6, 1984, at the "Ancestors" conference, American Museum of Natural History, New York), with the western sites being closer to the lowland tropical forest block (Andrews and Van Couvering, 1975).

Once out of Africa, the hominoids of the middle Miocene dispersed throughout Eurasia. The earliest record of *ramamorphs* in Siwalik localities of the Indian subcontinent is 12 mybp, from the type Chinji Formation of northern Pakistan (s.m. Raza and N.M. Johnson, pers. comm.). *Ramapithecus* and *Sivapithecus* 1 persist in Siwalik localities until about 7 mybp. From this time on, fossils of the *ramamorph* group disappear from Eurasia, although it now appears likely that an eastern population of *ramamorphs* gave rise to *Pongo* 1 (Andrews and Cronin, 1982; Ward and Pilbeam, 1983). What these thick-enamelled hominoids ate is an unresolved issue, although their diet is probably in the general domain of large hominoid herbivory. The biostratigraphic association of Siwalik *ramamorphs* with presumed forest-dwelling

mammals such as tragulids and deinotheres (Barry et al, 1982) suggests that the almost simultaneous disappearance of all these taxa is related to a dwindling vegetation association such as the South Asian tropical forest that today occupies the Malay region (Badgley et al., in press).

Substantial changes in global climates occurred during the Miocene. The northward movement of the Africa plate brought about both the gradual closing of the Tethys Sea and the opening of Drake's passage, the latter leading to the formation of the circum-Antarctic ocean current. The establishment of this current led to a pronounced intensification of latitudinal temperature differentiation. From the early Miocene until the present, ocean surface temperatures have declined by 10°C (Kennett, 1977). The consequence of this climatic cooling on the forest and eventually by woodland and grassland (Bonnefille, 1979). By the end of the Miocene, the vegetation type known as savanna—a grassland with less than 20% forest canopy—prevailed in much of east Africa. Modern savanna environments are characterized by highly seasonal rainfall; as a result of the seasonal availability of primary productivity, many species of savanna mammals migrate.

At Laetoli, where the first undisputed hominids occur, is a series of ash layers containing thousands of vertebrate footprints. This sequence, known as the Footprint Tuff, was deposited over a period of a few weeks spanning the end of the dry season and beginning of the rainy season (Hay and Leakey, 1982). In the footprint record, three hominid trackways occur with mammals whose modern counterparts are the earliest wet-season immigrants in a savanna environment. The "cause" of bipedalism is an ancient and unresolved puzzle in human evolution. The Footprint Tuff offers a plausible ecological hypothesis—that bipedalism enabled these large hominoids to migrate among food resources that displayed an increasingly heterogeneous spatial and temporal distribution.

In the Pliocene-Pleistocene, intense climatic fluctuations—related to the growth of the Antarctic ice cap and the onset of continental glaciation in the northern hemisphere—were added to the global

trend in climatic cooling. For primates with a dentition inherited from frugivorous and folivorous ancestors, the savanna must have posed a serious challenge in terms of diet. One of the most curious aspects of human evolution is how, through the hominoid-hominid transition, diets changed from the inferred more typical anthropoid diet of the Miocene hominoids to a condition of omnivory in which meat and diverse plant parts other than leaves and fruit are significant components. This dietary shift is puzzling because it is not reflected in modification of the dentition to the extent that is present in other mammals. The hominid dentition possesses neither high-crowned nor sectorial teeth. The development of tools—probably first in wood, later in stone and bone—may have served the roles of food-gathering and processing that are served by the dentition in other mammals.

Tools may have been developed initially for procurement of plant foods or for defense or for scavenging carcasses or for some combination of these activities. The taphonomy of early artifact sites indicates that hominids carried the raw materials considerable distances (up to 20 km) from their geological sources to the location where tools were manufactured and employed (Isaac, 1976; 1978). The repeated associations of stone-tool concentrations with concentrations of vertebrate remains suggest that many tools were created for the processing, although not necessarily the hunting, of animal food.

The expansion of the range of *Homo* 1 through the Old World occurred during the intense climatic fluctuations of the Pleistocene and is perhaps related to the diversity of potential food resources facilitated by the development of tools. Human populations, while spreading rapidly through the Old World and into the New World around 13,000 bp, remained relatively small until quite recently. The rapid increase in population size achieved by modern *Homo sapiens* is coincident with the control of food production, manifest ultimately in the cultivation of plants and domestication of animals. With these measures, human populations were no longer dependent on following the ever-changing food resources on a seasonal or longer basis.

Agriculture permitted sedentism and sedentism preceded cities.

CONCLUSION

The ecological perspective offered in the preceding section is based primarily on the relation of hominoids and hominid to food resources. Although this synthesis is plausible, it is speculative and will probably remain so. Two aspects of the hominoid fossil record hinder the resolution of some of the major issues in human evolution. The first is a sampling problem as pertains mainly to the Miocene portion of the fossil record: the fossil record, although much improved compared to twenty years ago, is still scant. The worldwide gap from 7-4 mybp, the larger gap in the African record, and the absence of fossils of the African apes obscure the transition from hominoid to hominid. The Plio-Pleistocene record, by comparison, is rich. But, the development of most characteristically hominid features are not documented in skeletal remains. Most of these features are behavioral - e.g., the emergence of language, art, and ranked societies. Even the changing relationship to food resources is not reflected in dental features as much as in population sizes and social organization.

Since most aspects of behavior are unobservable in fossils, many essential transitions in becoming human must be reconstructed--if at all--on the basis of the information from other living systems. It is for this reason that the study of human origin has forested so much research on the ecology and behavior of diverse human populations, non-human primates, and other mammals. These efforts to examine human ecology and behavior by analogy have made substantial contributions to the biology of mammals in general, as well as to our own evolutionary history.

ACKNOWLEDGMENTS

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J. Dudgeley

FIGURA 1

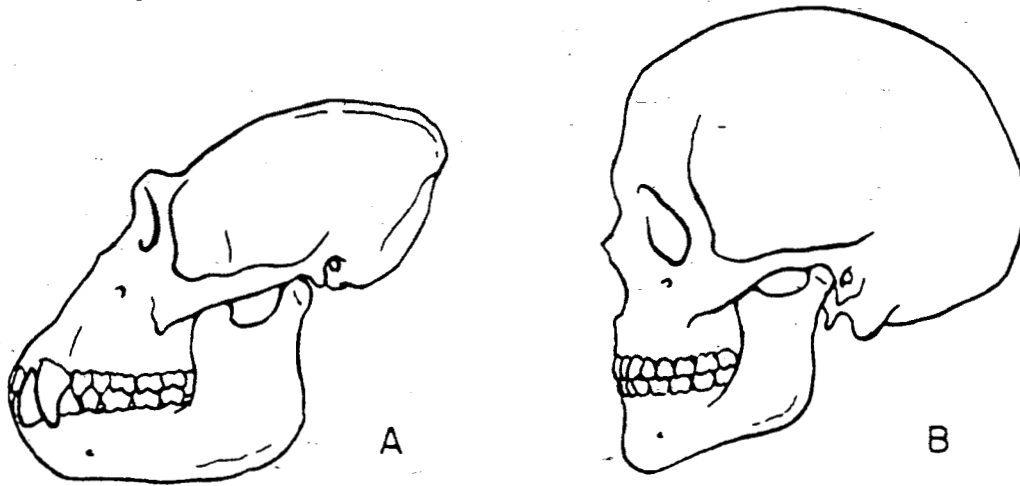
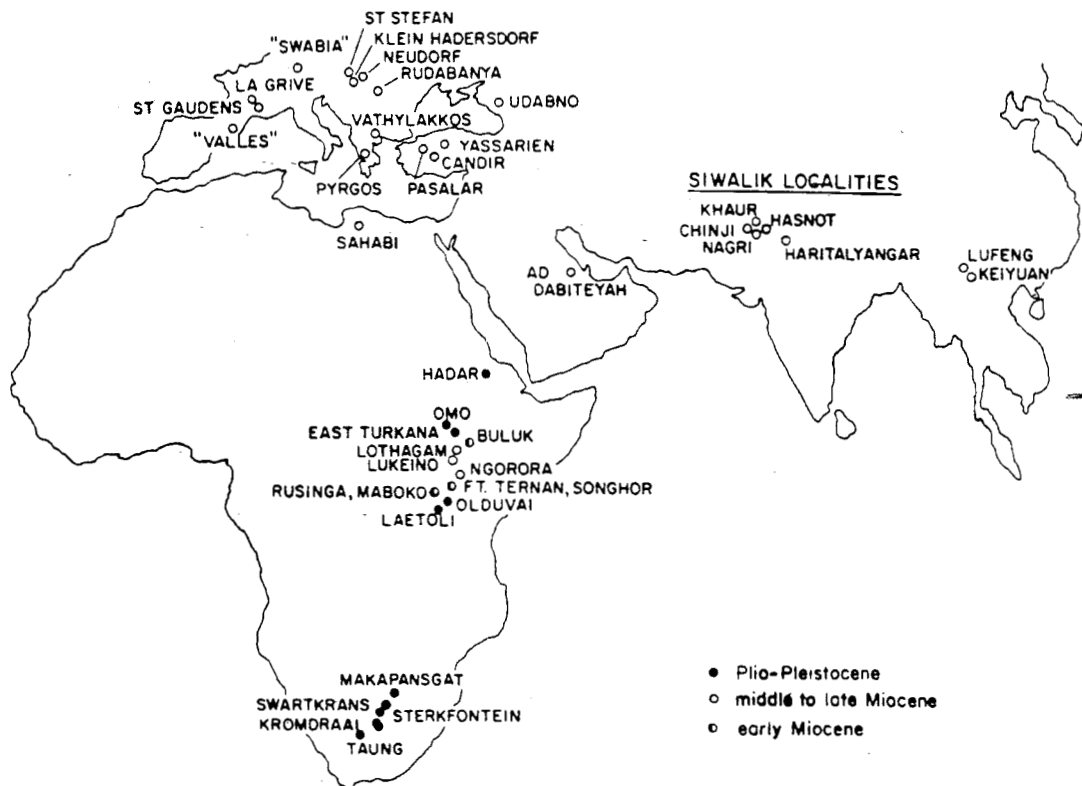
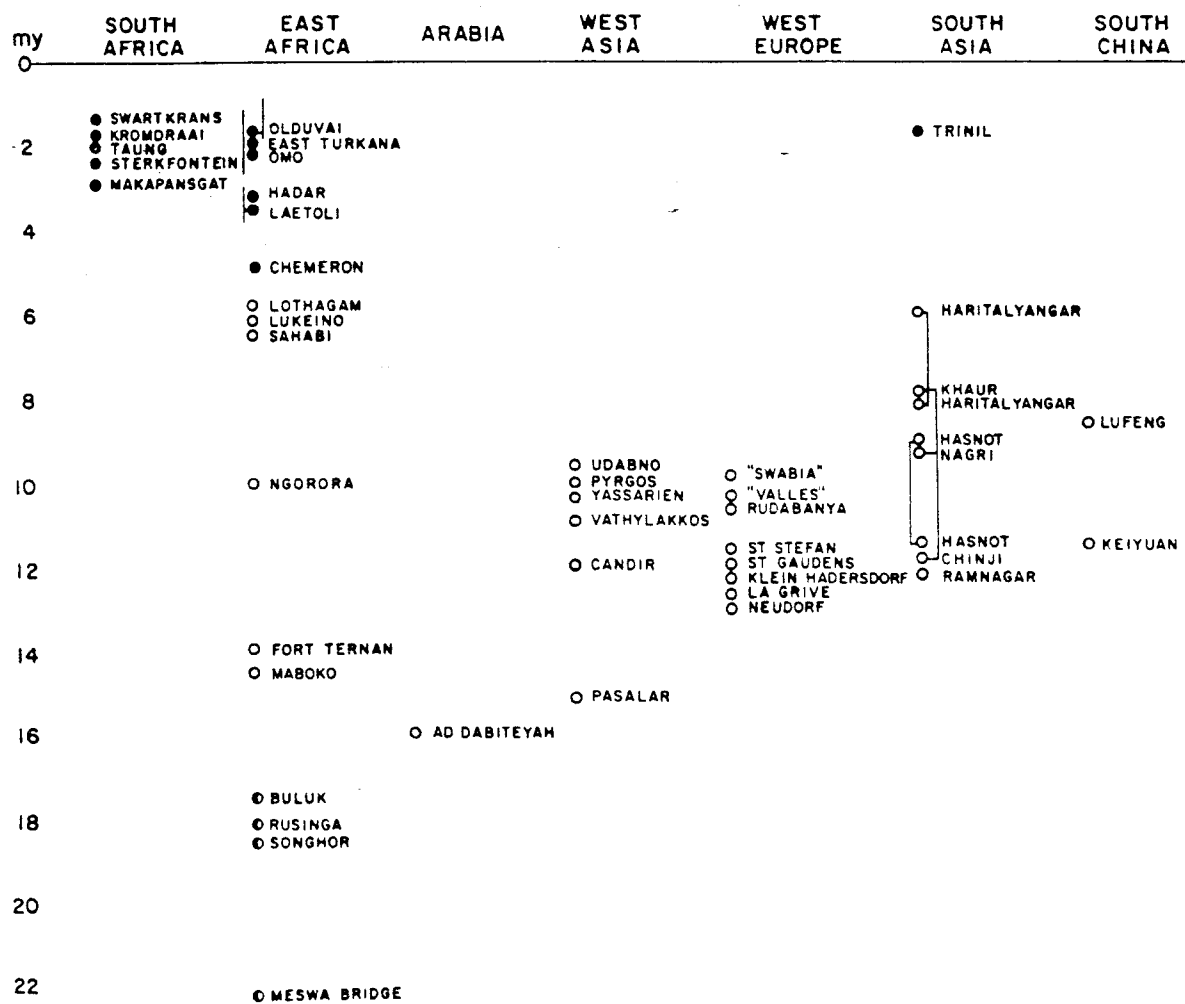


FIGURA 2



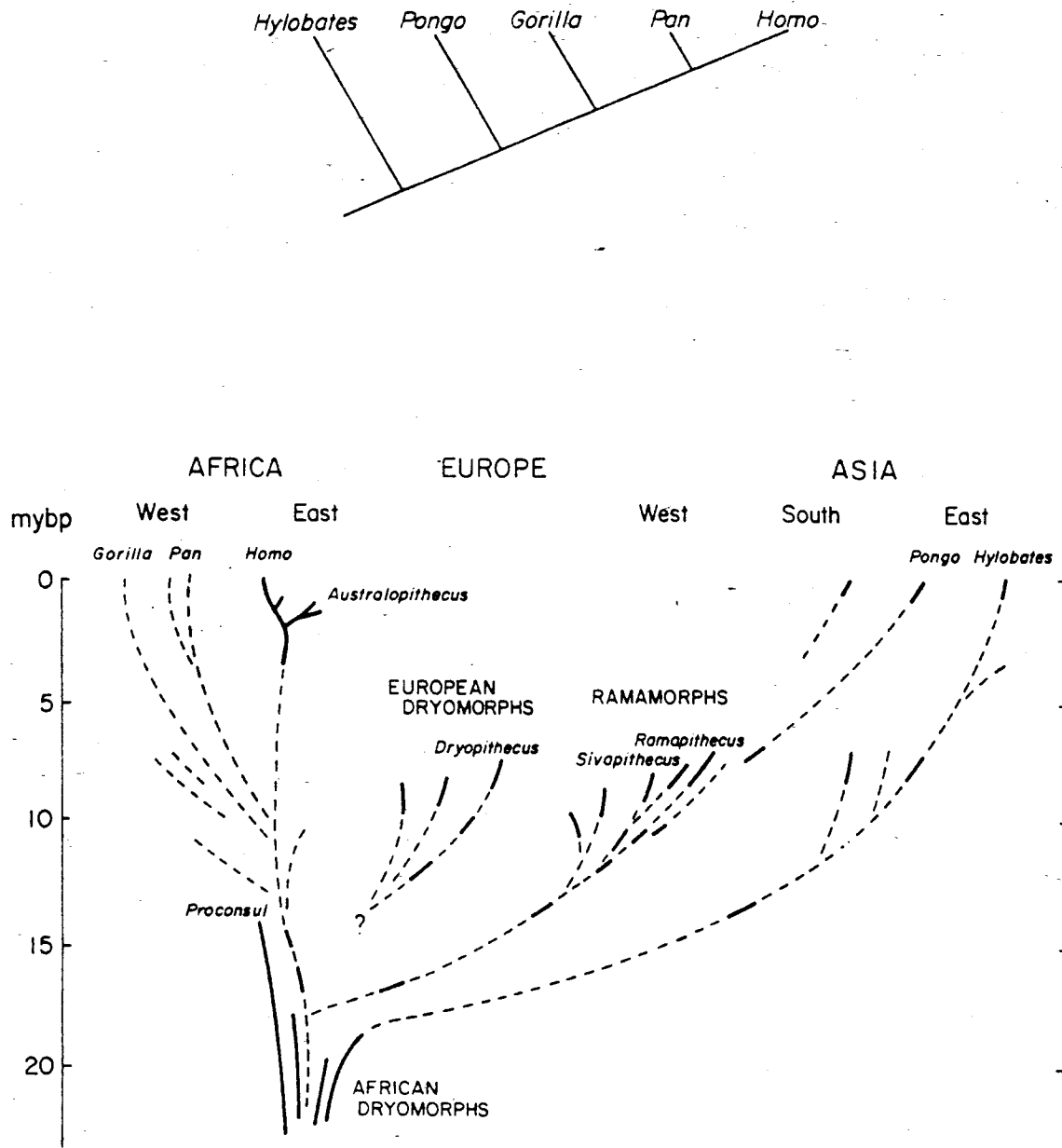
J. Badgley

FIGURE 3



J. Badgley

FIGURA 4



SECCION IV: RELEVANCIA DE LA INFORMACION PALEONTOLOGICA EN LA TEORIA BIOLOGICA.

Capítulo 10. Contribuciones teóricas diversas.

10.1 PREAMBULO

El propósito de esta sección, es destacar la relevancia que ha tenido la información paleontológica en la formulación de nuevas ideas, conceptos o teorías biológicas, así como en la reformulación de otras. Las contribuciones han sido muy numerosas, en campos variados que se extienden desde la Biología Evolutiva, hasta la Ecología; Todas tienen en común que constituyen una reflexión o interpretación del registro fósil- frecuentemente en relación estrecha con el registro estratigráfico- vehículo que permite asomarnos al pasado y que muestra la secuencia de cambios que han tenido el intrincado camino de la evolución orgánica. Esta posibilidad es privativa de las ciencias Histórico-geológicas, y no puede tenerse en las neontológicas, por lo que debe aprovecharse al máximo.

En el capítulo que nos ocupa, se han seleccionado sólo algunos tópicos en donde las contribuciones han sido importantes, sin pretender mostrar completo, el amplio espectro conceptual que han cubierto. La primera parte del mismo está dedicada a cuestiones de Biología Evolutiva, e incluye estos artículos: En el primero, David Raup diserta sobre el origen geológico temprano de los grandes taxos, señalando mediante un análisis probabilístico de la distribución de los tiempos de divergencia (que es el tiempo transcurrido desde que existió el ancestro común) teóricos en comparación con los observados, que el surgimiento temprano de los grandes grupos es un rasgo derivado de la geometría del árbol evolutivo, es decir una propiedad topológica de los conjuntos sucesivamente divergentes. En cambio, la interpretación de la individualización morfológica de los distintos phyla, todavía no ha

sido ecológicamente explicada de manera satisfactoria.

En el segundo trabajo, Douglas Erwin discute el tema de relojes moleculares, filogenias moleculares y el origen de los phyla, señala que aunque las secuencias de proteínas, RNA y DNA se han usado ampliamente para elaborar filogenias o para determinar el tiempo de divergencia, en especial cuando la información paleontológica es insuficiente o falta del todo, los resultados con frecuencia son ambiguos, equívocos o peor aun, conducen a errores, debido a que la redundancia inherente del código genético, frecuentemente enmascara la homoplasia, imposibilitando así reconstruir inequívocamente las secuencias de divergencia. Concluye indicando que un uso más afortunado de esta técnica sería su aplicación en el discernimiento de los patrones de la estructuración genómica de caracteres.

En ello para explicar como la evolución direccional de caracteres a nivel específico, puede surgir a darse por heterocronia (que es el cambio en la regulación del desarrollo morfológico, p. ej. pedomorfosis.). Argumenta que las discontinuidades morfológicas- a lo largo de un gradiente- han surgido por ser heterocronia. Así mismo señala que el establecimiento de gradientes morfológicos heterocronizados, sólo es posible en un gradiente ecológico adecuado.

En el cuarto y último trabajo de este apartado, Jack Sepkoski presenta el concepto de faunas evolucionarias; las describe como conjuntos de taxa de alta jerarquía- particularmente clases- que tienen historias similares de diversificación y que conjuntamente han dominado la biota por un lapso geológico extenso. A su vez este concepto (la expansión y declinación de las faunas evolucionarias en particular) puede aplicarse

para describir megavariaciones en la dominancia faunística y en la interpretación de los cambios cronológicos en la diversidad taxonómica global. Se han identificado tres de estas faunas en el dominio marino y la aplicación de este concepto se han extendido al dominio terrestre con éxito moderado.

Cabe destacar que Sepkoski le da una formalización y cierta caracterización matemática, a una idea que ha estado en el campo de la Paleontología desde el siglo pasado: La singularidad de "faunas cronológicas", es decir de faunas caracterizadas por ciertos grupos dominantes, claro están restringidas geocronológicamente, y que por tanto a su vez permite definir o caracterizar bioestratigráficamente a tales lapsos geocronológicos; por ej. es del dominio público que la fauna marina del Paleozoico Temprano, es significativamente diferente de la del Paleozoico Tardío, y que cada una de ellas tiene una composición taxonómica particular. Para los fines que nos ocupan, cada una de ellas sería una fauna evolucionaria.

En el resto de la sección, se abordan dos temas más, biodiversificación y extinción. El primero es tratado por Jack Sepkoski y Mike Hulver, quienes han compilado un extenso conjunto de diagramas sobre diversidad paleontológicamente registrada de clases o linajes filéticos (familias a clases), tanto de organismos marinos como terrestres, para el Fanerozoico; ellos presentan esta rica información en un atlas de indudable valor.

En el último artículo Antoni Hoffman diserta sobre las diversas concepciones o puntos de vista que han tenido sobre el fenómeno de las extinciones masivas. Señala, que aunque este tema en la última década (sin duda como resultado del interés que sobre este fenómeno despertó la hipótesis del impacto meteorítico como causa de la extinción masiva del Cretácico terminal), y la historia de la investigación sobre el mismo, es compleja y se remonta al nacimiento mismo de las ciencias geológicas. La concepción catastrofista ha sufrido un ciclo en la responsabilidad y aceptación que han tenido en el mundo científico. Originalmente, Cuvier y otros,

invocaron catástrofes diversas, para explicar la desaparición de especies y aún de linajes complejos, observables en el registro fósil. Este punto de vista resultó respetable, tanto por la autoridad académica de Cuvier y de los otros proponentes de esta tesis, como por su aparente concordancia bíblica, que hacía del llamado diluvio universal, una de tales catástrofes.

Poco antes en Inglaterra, James Huntton propuso una tesis esencialmente gradualista, ordinaria y común para describir a los procesos geológicos y para explicar los resultados de los mismos, observables en el registro rocoso de la historia de la Tierra. Esta tesis fue ampliamente desarrollada por Charles Lyell y otros, quienes mostraron que aún los grandes cambios observables en este registro, podían explicarse gradualísticamente, concibiéndolos como el resultado de la acumulación de muchos "microcambios" al través de un largo tiempo geológico-sin que se le vea principio o fin, según escribía el propio Huntton.

Puntos de vista tan opuestos ocasionaron la controversia principal del primer tercio del Siglo XIX. La concepción gradualista se fue imponiendo y constituyó uno de los apoyos fundamentales de la teoría.

103 BIOLOGIA EVOLUTIVA

103 (a)

On the early origins of major biologic groups

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NOTA: Información temporalmente
extraviada.

Paleobiology, 9(2), 1983, pp. 107–115

10.3 BIOLOGIA EVOLUTIVA

10.3(a) On the early origins of major biologic groups

David M. Raup

Abstract.—The commonly observed fact that the principal biologic groups appear early in an adaptive radiation is shown to be an artifact of the geometry of the evolutionary tree. For marine invertebrates of the Phanerozoic, total diversity and species turnover rates combine to make early origins of major groups inevitable. This is demonstrated through an analysis of the theoretical probability distribution of divergence time (time to most recent common ancestor) in comparison with observed distributions. Whereas the fact of early origins does not require ecological explanations, the interpretation of the morphological distinctness of major groups when they first appear remains an ecological problem.

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Introduction

Students of evolutionary history have observed repeatedly that in an adaptive radiation, the major subgroups appear early and at about the same time. In a review of this phenomenon, Nicol et al. (1959, p. 351) said, "The major subdivisions of a group (e.g., classes of a phylum or orders of a class) generally originate early in its history, whereas new subdivisions of lesser rank (e.g., genera and species) may arise at any time throughout the group's geologic history."

Documentation of the generality of early origins of major groups is straightforward. The phenomenon is most striking in marine invertebrates. Nearly all living phyla of marine invertebrates that have reasonably good fossil records have first occurrences either in the late Precambrian or early to middle Cambrian. At the class level there are 27 paleontologically important living groups and all have documented occurrences which are Silurian or older (data from Sepkoski 1982). The mean geologic age of first occurrence for these 27 classes is 533 Myr with a standard deviation of only 51 Myr. Fully 20 of the 27 have first occurrences earlier than 500 Myr (Cambrian or older). These data are shown graphically in Fig. 1. For orders of marine invertebrates, the mean time of first occurrence is 328 Myr BP and there is a strong frequency peak at about 500 Myr (Fig. 2).

The same relative pattern can be seen in the geologic records of vertebrates and land plants

although origins are generally displaced in time toward the Recent. For example, nearly all orders of living mammals have origins during a fairly narrow interval between 50 and 70 Myr BP.

An important consequence of the foregoing is that standing diversity at the higher taxonomic levels (phyla, classes, and orders) tends to be stable or to decline through geologic time following the initial radiation. Curves for standing diversity of marine shelf invertebrates prepared by Valentine (1969) show that the number of phyla is essentially constant, with virtually no origination or extinction. The standing diversities of classes and orders decline steadily so that present-day levels are 25%–35% lower than in the late Ordovician. This is due to a substantial excess of extinctions over originations and it further supports the generalization that originations of the higher-level groups are concentrated early in geologic history.

These generalizations have been the subject of considerable theoretical analysis. Interpretations have varied greatly. At one extreme, Willis (1940, p. 191) said "evolution goes on in what may be called the downward direction from family to variety, not in the upward, required by the theory of natural selection." In other words, the general phenomenon of early origins has been considered by some workers to be counterintuitive or against general Darwinian theories of evolution. Many of the interpreta-

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tions have been largely ecological, and much excellent analysis of this aspect has been carried out by Valentine (1973, 1980). Valentine and others have argued that the phenomenon results from the exploration of the empty adaptive zones that exist early in a radiation. Valentine postulates that a few distinctive bauplans invade, or originate in, empty niche space, thus establishing the major classes and orders and only later is diversity filled out at the lower levels. The general observation that major groups increase in genus/family ratios through geologic time has been used to support ecological explanations.

In this paper I will explore an alternate view: that the early origin of major groups is an inevitable product of the geometry of the branching evolutionary tree. I will argue that the simple fact of early origins is an artifact and requires no biological explanation. The analysis will be confined to purely cladogenetic aspects of the problem. Consideration of the morphologic distinctness of groups when they first appear, important to Valentine's analysis, will not be investigated.

Divergence Time as an Appropriate Metric

The analysis of early origins could be developed from purely empirical data on the ranges in time of higher taxa. This has been done by others, especially Nicol et al. (1959), but is somewhat less than satisfactory. There is no rigorous mathematical basis for differentiation of

the several levels in the taxonomic hierarchy. Also, the geologic range data for higher taxa are often subject to considerable uncertainty, due in part to the ever-present problems of definition of particular taxa. In the present analysis, I will use divergence time as a metric, and this will avoid many of the problems of definition of higher taxa.

Divergence time is defined as follows. Given two living species, how far must one go back in geologic time to find the single species which is the most recent common ancestor of both living species? This time, in millions of years, is the divergence time for that pair of species.

Hypothetical evolutionary trees are shown in Figs. 3-4. Let us assume that they are correct and complete representations of the evolution of imaginary biologic groups through time. In these trees, each vertical line is a species lineage. We need not be concerned with questions of punctuated equilibrium or gradualism. In Figs. 3-4, the vertical lines may be unchanging species enjoying complete stasis or they may be lineages undergoing phyletic transformation through a succession of chronospecies. In either case, the end of the vertical line indicates complete extinction of a genome, and the start of a vertical line indicates a speciation event. What happens in between is not relevant to the present analysis. In the diagrams, the arrangement of lineages left to right has no significance.

In a real-world situation, nested sets of taxonomic groups could be superimposed on each of

the trees. Clusters or clades could be recognized, and their number would depend on the methods used and on the morphologic differences between the species. Probably no two classifications developed by different workers would be exactly the same. Fortunately, divergence time can be used as a convenient proxy, thereby avoiding many problems.

Divergence times for a few cases are indicated in Fig. 3. Figures 5-6 show frequency distributions of the divergence times for all possible pairs of surviving species in the trees. The phenomenon of early origins under consideration in this paper is manifested by a concentration of divergence times early in the time scale. The link between early origins and divergence time will be developed below.

Behavior of Divergence Time

The divergence time distributions in Figs. 5-6 are very different, but they have an important characteristic in common. The distributions are not smooth. Rather, they are dominated by spikes representing the points of common ancestry for major clusters. This spikiness has been noted by Sibley and Ahlquist (1982) and by others working with genetic distance data. It is log-

ically inevitable that if we have two distinct branching clusters with a single common ancestor, any species pair that draws one species from each cluster will have the same divergence time. This is illustrated in Fig. 3 by the fact that any pair of species involving cluster A and cluster C shows a divergence time of precisely 47 time units before present. This produces a major spike in the frequency distribution of divergence times at 47. The spikes are present throughout the hierarchy of taxonomic levels. Sibley and Ahlquist have suggested that concentrations of genetic distances reflecting these spikes can be used to delimit taxonomic rank.

Although both divergence time distributions in Figs. 5-6 show the spikiness just discussed, the distributions differ. Figure 5 has a concentration of young divergence times and Fig. 6 has a concentration of older ones. Both distributions are highly skewed and would appear even more so were the vertical scale not logarithmic. The obvious difference between the trees in Figs. 3-4 is in species turnover rate. Turnover rate is a function of two quantities: speciation rate and extinction rate. Mean divergence time is also influenced by the total elapsed time, but this can be subsumed in the turnover rate. In other words, the important element is the geometric fabric of the tree, the relationship between turn-

over rate and elapsed time. The contrast between the trees in Figs. 3-4 suggests that a multitude of relatively short branches leads to a low mean divergence time (Fig. 5). As will be shown later, standing diversity is also an important element determining the divergence time distribution.

Figures 3-6 present some of the characteristics of the divergence time distribution in relation to branching trees, but they can tell us nothing about the real world because they are not scaled to approximate natural rates of turnover or real world diversity. As Stanley et al. (1981) have emphasized, scaling for diversity is often important in the analysis of the dynamics of evolutionary trees.

An Equation for Divergence Time Distribution

Assume that at some time, t , before present there are S species lineages. In Fig. 3, eight such lineages exist at the bottom of the diagram. During the 90 time units leading to the top of the diagram ("Recent"), all of the starting lineages die out but their progeny persist for varying lengths of time as clusters or clades. In the case of Fig. 3, only the starting lineage on the extreme left has descendants surviving to the Recent. Thus, all living species must have divergence times less than 90 time units. In Fig. 4, on the other hand, several of the starting lineages have living descendants, so that diver-

gence times for many living species pairs will exceed the 90 time units.

The problem is to develop an equation which will predict the expected divergence time distribution for randomly chosen pairs of living species. The relevant constants are S_0 , the standing diversity at time = 0 ("Recent"); p , the average rate of branching, expressed as speciation events per lineage per time unit; and q , the average rate of extinction, expressed as terminations per lineage per time unit. In deriving an equation, I will use the mathematics of random branching processes and assume time homogeneity. That is, it will be assumed that p and q do not change through time. This does not prejudice the question of whether speciation and extinction rates are constant; it is only a convenient way to describe differences in branching patterns. The resulting equation could be elaborated to include changing p and q values.

Where branching rate is greater than extinction rate ($p > q$), standing diversity increases through time. This is the general condition in evolution over long periods of geologic time, and for evolutionary radiations in particular. If we look backward in time from the Recent, standing diversity will appear to decay. Given p and q and an assumption of time homogeneity (above), we can compute the expected value of

diversity (S_t) at some time = t before present (BP) as follows:

$$S_t = S_0 e^{-(p-q)t} \quad (1)$$

It should be noted that S_t is an expected value and has a high variance where samples are small. Therefore, the estimate of past diversity, S_t , in the derivation that follows is a reasonable approximation only for situations in which one is dealing with thousands or tens of thousands of species, and the reader should be cautioned that application to cases with smaller samples should be avoided.

Now let us specify two Recent species at random and imagine the evolutionary tree that produced these two (and all other surviving species). Each of the two specified species can be traced back along a unique path from lineage to lineage until the base of the tree is reached. At some point the ancestral paths of the two species will meet, and this point represents the divergence time for the species pair. The two paths will meet at a speciation event, that is, a cross-link in the tree. In each small time interval, the total number of possible cross-links is

$$C_2 = \frac{S_t!}{(S_t - 2)!(2)} = \frac{S_t(S_t - 1)}{2} \quad (2)$$

The expected number of cross-links (speciation events) that actually occur in a given time interval is the expected diversity times the speciation rate. If we consider a time interval of 1 Myr, the expected number of speciation events, N , is

$$N = pS_t = pS_0 e^{-(p-q)t} \quad (3)$$

Any one of these speciation events has a probability of successfully connecting the two ancestral paths of interest, and this probability is the reciprocal of the number of possible different cross-links (2). And the probability of not making a successful link is one minus the reciprocal of (2). Consequently,

$$\text{probability of no cross-links in } n \text{ tries} = \left[1 - \frac{2}{S_t(S_t - 1)} \right]^n \quad (4)$$

The probability of at least one successful cross-link in t successive time units is one minus the product of a series developed from (4). This may be written as follows:

$$W_t = 1 - \prod_{j=1}^t \left[1 - \frac{2}{S_j(S_j - 1)} \right]$$

In this equation, S is defined as in (1) and as in (3). W_t is the probability that a randomly chosen pair of Recent species has a divergence time less than some time = t before present. numerical value of W_t approaches 1.0 as base of the evolutionary tree is approached (single ancestor for all Recent species).

Given values of the three constants (p , q , S_0), the equation can be solved for a series of values to develop the distribution of W_t , from which a predicted frequency distribution of divergence times can be developed.

Divergence Time Distributions

The W_t distribution depends on the particular combination of p , q , and S_0 , but some generalizations are possible. All other things being equal, increasing diversity increases mean divergence time. Thus, although the tree in Fig. 3 yields a fairly low mean divergence time (Fig. 5), a tree with the same overall fabric (relatively low p and q) but higher diversity would yield a higher mean divergence time. This makes especially important that divergence time calculations be scaled properly for number of species. It also means that computer simulations are totally impractical for most real-world situations.

Increasing turnover rate (numerical value of p and q) decreases mean divergence time, as can be seen from a comparison of the trees in Figs. 3-4. Increasing the rate of growth of divergence ($p - q$) decreases mean divergence time by concentrating more of the divergence points to be concentrated higher in the tree.

The divergence time distributions shown in Figs. 5-6 are monotonic and highly skewed; this is not typical. It is logically true that frequency of divergence times must drop to zero as the Recent is approached (time = 0) and go to zero as the ancestor of the entire group is approached (high t). All complete, ideal distributions are unimodal. This can be seen if calculations such as those in Figs. 3-4 are run repeatedly with the same input constants and results summed. Whether the frequency distributions are skewed toward or away from the Recent depends on p , q , and S_0 .

It was noted above that empirical divergence time distributions are collections of spikes (Figs. 5-6). This will be true of any distribution based on a single tree. Equation (5) does not produce spikes because it describes a generalized probability density distribution which approximates the combined result of many separate trees built from one set of constants.

Predicted Divergence Times for Marine Invertebrates

To use W_t to predict the divergence time distribution for a real-world situation, we must have good estimates of final diversity and of branching and extinction rates.

The literature contains a variety of estimates of present-day diversity (see Valentine 1970; Valentine et al. 1978). We will be concerned here with numbers of readily fossilizable marine invertebrates on the continental shelf. Estimates of present-day diversity range from roughly 100,000 to 300,000 living species, and I will use 150,000 as the standard for this paper although, as will be shown, the effects of changes in this number are relatively minor in the context of divergence time distributions.

Extinction rate estimates are on reasonably firm ground, but the uncertainties are such that we must also consider a range of values. Raup (1978) used data on generic survivorship to develop an average species extinction rate of $q = .09$. Since the average duration of a species in geologic time is the approximate reciprocal of

q , the mean duration for shelf invertebrate species should be about 11 Myr. For a variety of reasons this estimate may be high, but it may be used as a starting point for analysis.

An estimate of mean speciation rate can be obtained from the extinction rate and modern diversity estimates. If diversity has increased over the Phanerozoic (Sepkoski et al. 1981), then speciation rate, p , must exceed the extinction rate, q . The fundamental quantity is $(p - q)$ in (1). Equation (1) can be rearranged as follows:

$$S_0/S_t = e^{-(p-q)t} \quad (6)$$

The ratio S_0/S_t is the factor by which diversity decreases as one goes backward in time. Thus

$$(p - q) = -\frac{\ln(S_0/S_t)}{t} \quad (7)$$

and

$$p = (p - q) + q. \quad (8)$$

Solving equations (7) and (8) yields an estimate for mean p .

Table 1 shows a set of calculated values for W_t for q ranging from .05 to 5.0 (equivalent to species durations from 20 to 0.2 Myr) and for $(p - q)$ from .00019 to .00461 (equivalent to factors of diversity increase from Cambrian to Recent of from 1.1 to 10). All computations in Table 1 use an estimate of 150,000 species for present-day standing diversity and $t = 500$. The important result of the computations is that the numbers in Table 1 are all extremely small. This means that for any randomly chosen pair of living species, the probability is very low that they belong to the same clade. Or, to put it another way, the probability is high that the common ancestor of the two species is to be found in rocks older than 500 Myr. Only in the lower right-hand corner of the table are the numbers significantly different from zero. One would have to postulate a mean species duration of 200,000 yr or less and also postulate a Paleozoic to Recent diversity increase approaching an order of magnitude to find more than a few percent of the divergence times less than 500 Myr.

Parallel computations based on modern diversities of 100,000 and 300,000 species did not give appreciably different results. The compu-

tations were also done for $t = 400$. The resulting W values were slightly higher, but the basic pattern was unchanged.

The computations resulting in Table 1 show that an evolutionary tree that has the basic geometry of the real tree for marine invertebrates will be one in which the overwhelming percentage of randomly chosen species pairs will have pre-Ordovician common ancestors. This says, in turn, that randomly chosen pairs of living species are most likely to involve intergroup comparisons, where the constituent groups have Cambrian or Precambrian origins. Although this is in qualitative agreement with the general observation that most major groups have early origins, it must be checked with actual distributions of divergence times of living marine invertebrates.

It should be noted in passing that equation (5) could be used to predict the time of the start of the whole radiation or the origin of the tree itself ($W_t = 1.0$). If this is done for the range of p and q values used for Table 1, the predicted start of the invertebrate radiation is deep in the Precambrian. But to carry the computations this far would be invalid because there is ample evidence that the major invertebrate radiation was a late Precambrian phenomenon (Valentine 1973, 1980; Sepkoski 1979). The early stages of the radiation were characterized by much higher branching rates than those used here, and they represent a significant departure from time homogeneity of the branching process.

Divergence Times in the Real World

Surveying divergence times with real species is extremely difficult, even though in theory it should be simple. One could select pairs of living species at random and determine from the paleontological literature an estimate of the time to most recent common ancestry for each pair. But problems abound because of the many uncertainties in reconstruction of phylogenetic patterns, of missing stratigraphic records, and other problems. But we can get a general picture of actual divergence time distributions and set some general limits.

Let us consider the 27 paleontologically important classes of marine invertebrates that are living today. Rough estimates of the numbers

of living species in each are available, and they can be grouped into four categories: 20,000–30,000 species (Rhizopodea and Gastropoda), 10,000–20,000 species (Bivalvia and Malacostraca), 1000–10,000 species (9 classes), and fewer than 1000 species (14 classes) (data from Valentine [1970] and Van Valen [1973]). These numbers are generalized in Table 2. If we assume for simplicity that the average number of species in each of the size groups is the midpoint of its range, then the total number of species is 136,500 (see Table 2).

As we have seen (Fig. 1), all of the 27 classes have first occurrences in the geologic record in rocks of 400 Myr BP (top of the Silurian) or older, with a mean time of first occurrence of 533 Myr BP. The common ancestors of many of the younger classes (Ordovician and Silurian) are in the Cambrian or Precambrian. Let us generalize that 500 Myr BP is the minimum divergence time for a species pair drawn from any two of the classes. It may be that this is actually a conservative estimate because experience has shown that as more fossils are discovered, the first occurrences of major groups tend to be pushed back in time.

Consider a class that has 15,000 living species. Any one of these species can join with another living species to form a species pair. Some pairs will be intraclass and some will be interclass, depending on whether they have species from the same class or different classes. The number of possible intraclass pairs for a class having 15,000 species is $\{(15,000)(14,999)\}/2 = 1.125 \times 10^8$. Extending this procedure to all 27 classes,

the total number of intraclass pairs is 9.86×10^6 (Table 2). The total possible pairs (intraclass and interclass) is $[(136,500)(135,499)]/2 = 9.32 \times 10^9$.

The ratio of the two totals just given is the proportion of intraclass pairs: 0.106. Thus, only about 10% of all possible species pairs are intraclass. If the minimum age of origin of the 27 classes is taken as 500 Myr, then it follows that at least about 90% of the possible pairs of living species have divergence times greater than 500 Myr and only 10% are less than 500 Myr. But because many of the subgroups within classes have divergence times as old, or almost as old, as the classes themselves, simple counts of numbers of intraclass pairs almost certainly underestimate the number of pairs with high divergence times. This means that the actual proportion of divergence times greater than 500 Myr is almost certainly greater than 90%.

Let us now turn to a somewhat higher resolution analysis and consider divergence time distributions within a single class, the class Echinoidea. Using the data set of Raup (1975), there are approximately 1000 living echinoid species classified into about 15 orders and 50 families. Using the reasoning that was applied above, there are about 500,000 possible species pairs of living echinoids. Some of these are pairs within families or within orders, some are between families or between orders. Obviously, species related at the lower taxonomic ranks are more likely to have lower divergence times. Consider a family such as the Arbaciidae. There are 31 living species and thus 465 intrafamily comparisons are possible. But there are about 30,000 possible comparisons between arbaciid and nonarbaciid echinoids. Thus, of all the possible pairs of living echinoid species involving arbaciids, only about 1.6% are within-family pairs. Now, the Arbaciidae is the only living family in the order Arbacioida, and thus 98.4% of echinoid pairs involving at least one arbaciid are also interordinal and imply high divergence time. Mean divergence time for echinoid orders ranges between 190 and 345 Myr. Therefore, if we were to pick an arbaciid species at random and pair it with any other echinoid species, the overwhelming probability (0.984) is that this pair is an interordinal pair and has a divergence time measured in hundreds of millions of years.

We can go somewhat further and consider randomly selected pairs which include an arbaciid echinoid and any other marine invertebrate. As we have seen, the total number of possible pairs of invertebrate species is about 9.3×10^9 . If one member of a species pair is an arbaciid echinoid, there are only 465 possible pairs which would also include a second arbaciid. It is clear that if we were to choose an arbaciid echinoid at random and link it with any other marine invertebrate, the probability is negligible that the second species is even an echinoid. More likely, it will be in another invertebrate class, which again gives us a high divergence time.

Discussion

The theoretical predictions of divergence time distribution for marine invertebrates (Table 1) indicate that nearly 100% of divergence times should be 500 Myr or older. Even allowing for considerable error in our estimates of turnover rates and past diversities, it is difficult to lower this estimate to less than 98% or 99%. The analysis of empirical divergence times gives minimum estimates of these old divergence times of about 90% (Table 2). While both the theoretical and empirical analyses support the basic contention that major groups appeared early, there remain several interesting questions. One of these is the significance of the disparity between 99% and 90% for theoretical and empirical analyses, respectively. There are two possible explanations for this. The first is purely artificial. The estimated numbers of high divergence times in the real world are probably too low because the numbers of intraclass pairs with high divergence times have been underestimated. The other possible interpretation is that the real-world estimate of 90% is actually valid and represents a real departure from the theoretical predictions of the stochastic model used as a basis for the equations.

The derivation of the equations assumed that speciation and extinction rates are constant through time. We know that this is not the case. In particular, we know that the evolutionary system is occasionally perturbed by mass extinction. That is, there are short intervals of time wherein the extinction rate, q , rises markedly. This results in a drop in standing diver-

sity. Following the mass extinction there is a fairly rapid recovery of diversity, which implies that speciation rate is much higher relative to extinction rate for the time necessary to bring diversity back up to pre-mass extinction levels (Sepkoski 1979). The recovery from a mass extinction often is associated with adaptive radiations; that is, bursts of cladogenetic activity which either produce new major groups (families and orders) or which see the rapid diversification of groups that already existed. One of many examples is the rapid and extensive evolutionary radiation of the scleractinian corals following extinctions at the end of the Cretaceous. Mass extinctions have the effect of forcing the biota through a bottleneck and, to some degree, "restarting the clock" in the branching system of evolution. One effect of such a perturbation should be that modern taxa originate at a time later in evolutionary history than would be predicted by the purely stochastic model. And this may well explain a mean divergence time which is less than predicted. Any sequence of mass extinction and recovery will produce a new spike in the divergence time distribution and lower the overall mean divergence time for the group as a whole. This explanation of the observations is speculative, of course, because we do not know for sure how much the real-world mean divergence times are underestimated. This is a direction for research which has considerable potential. One can even imagine that major adaptive radiations are expressed as spikes in the divergence time distribution and that this could be a useful way to detect major bursts of cladogenetic activity.

Conclusions

The combination of theoretical and empirical analyses presented in this paper demonstrates that the early origin of major groups is the expected condition. This would not be true for any evolutionary tree, for there are many combinations of p , q , and S which produce a divergence time distribution quite different from what we observe in the real world. Therefore, we can conclude that the fact of early origins of major groups does not require an ecological or other biological explanation beyond accommodating the simple inputs of starting diversity, speciation rate, and extinction rate.

This paper has been devoted entirely to the cladogenetic or branching aspects of the problem. The fact that new groups are usually distinct morphologically as soon as they appear in the fossil record has been completely ignored. The distinct morphology suggests an unusually rapid diversification early in the history of a biologic group, and this may well result from the ecological causes suggested by Valentine. Thus, whereas the present analysis indicates that early origins are a nonbiological artifact, the analysis cannot speak to the most intriguing problem of the morphologic divergence that takes place so quickly in an adaptive radiation.

Acknowledgments

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FIGURE 1

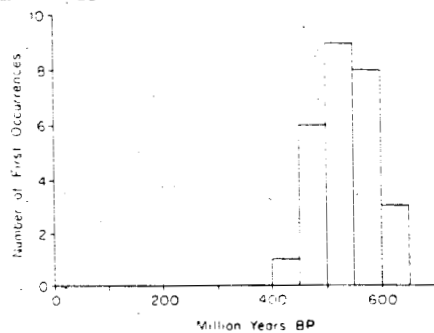


FIGURE 1. First occurrences of 27 modern classes of paleontologically important marine invertebrates. Stratigraphic range data from Sepkoski (1982).

FIGURE 2

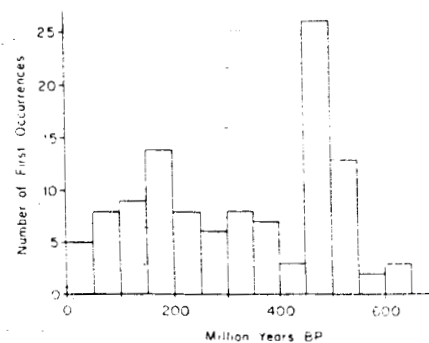


FIGURE 2. First occurrences of the modern orders in the 27 classes represented in Fig. 1. Stratigraphic range data from Sepkoski (1982).

FIGURE 3

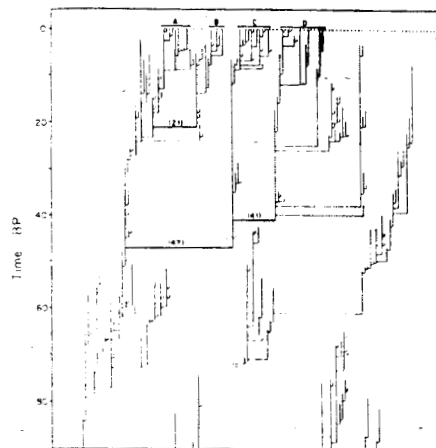


FIGURE 3. Hypothetical evolutionary tree based on relatively high rates of species turnover. A few divergence times are indicated, such as 47 for any species pair involving clusters A and C.

FIGURE 4

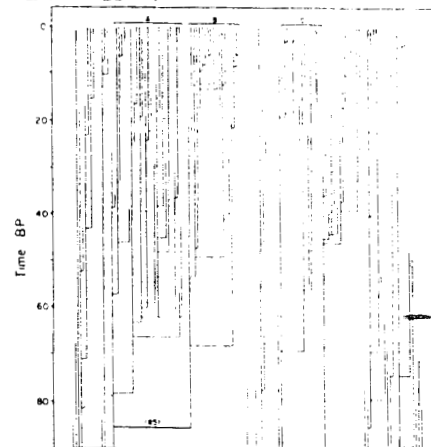


FIGURE 4. Hypothetical evolutionary tree based on relatively low rates of species turnover.

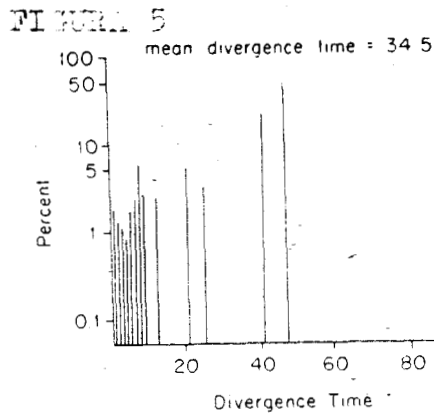


FIGURE 5. Frequency distribution of divergence times for all possible pairs of surviving species in Fig. 3. Spikes at 41 and 47 reflect concentrations of those divergence times in Fig. 3.

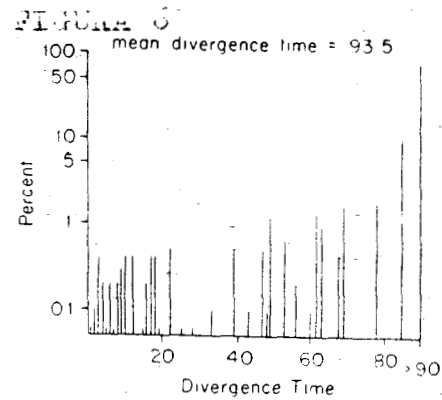


FIGURE 6. Frequency distribution of divergence times for all possible pairs of surviving species in Fig. 4. High frequency of divergence times greater than 90 reflects the fact that many of the common ancestors of surviving species existed at the base of Fig. 4.

TABLE 1

Duration of species (Myr)	Diversity increase factor (Cambrian-Recent)					
	1.1	2	3	4	5	10
20	.000	.001	.001	.001	.001	.002
10	.001	.001	.001	.002	.002	.003
5	.002	.002	.003	.003	.003	.005
2	.004	.005	.006	.007	.008	.013
1	.007	.010	.012	.014	.017	.026
0.5	.014	.019	.024	.029	.033	.051
0.2	.034	.047	.059	.070	.080	.122

TABLE 1. Predicted proportion of divergence times less than 500 Myr. Equation (5) for W is solved with an assumed modern diversity of 150,000 and various combinations of p and q . Extinction rate (q) is the reciprocal of species duration, and speciation rate (p) is determined by the rate of diversity increase, extinction rate, and modern diversity (eq. [7] and [8]).

TABLE 2

Classes	Median species per class	Total species	Total intraclass pairs
2	25,000	50,000	6.24×10^8
2	15,000	30,000	2.24×10^8
9	5,500	49,500	1.36×10^8
14	500	7,000	1.75×10^6
27		136,500	9.86×10^8
Total species pairs (intra- and interclass)			9.32×10^8
Proportion of intraclass pairs:			0.106
Proportion of interclass pairs:			0.894
			1.000

TABLE 2. Generalized numbers of intraclass species pairs for modern classes of marine shelf invertebrates (limited to groups generally recognized as fossilizable).

10.3 (b)

MOLECULAR CLOCKS, MOLECULAR PHYLOGENIES AND THE ORIGIN OF PHYLA.

Douglas H. Erwin

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Protein, RNA sequences have been widely used to construct phylogenies and to calculate divergence times using a molecular clock. Reliance on molecular information is particularly attractive when fossil evidence is missing or equivocal, as in the Cambrian metazoan radiation. I consider the applicability of molecular clocks and phylogenetic analysis of molecular data to the origin of metazoan phyla, and conclude that molecular information is often ambiguous or misleading. Amino acid sequences are limited use because the redundancy of the genetic code masks patterns of descent, while of a nucleotide sequences only four potential states exist each site (the four nucleotide bases). In each case, homoplasy may often go undetected. The application of a molecular clock to resolve the timing of the metazoan radiation is unwarranted, while molecular phylogenetic reconstruction should be approached with care. A potential more useful technique for phylogenetic reconstruction would be the use of patterns of genome structure and organization as characters. Molecular clock, phylogenetics, metazoan radiation, origin of phyla.

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The explosive growth of molecular biology has led to the estimation of the time of divergence events using a molecular clock and the determination of phylogenetic patterns by comparison of

amino acid sequences (from proteins) or nucleotide sequences (from RNA and DNA). Such techniques have been applied to paleontological problems where the fossil evidence is either missing or equivocal in an attempt to resolve the timing and pattern of divergence. Runnegar (1982a, b, 1986) has suggested that molecular information may resolve the question of the timing and duration of the Cambrian Metazoa radiation. I address here the reliability of the molecular clock as well as other applications of molecular data (phylogenetic analysis using protein, DNA or RNA sequences) to resolve the rate and timing of the Cambrian radiation. It is not the goal of this paper to discuss more restricted (either temporally or phylogenetically) uses of molecular data, or to argue that such data is without interest, but rather to discuss the application of molecular data to a particular paleontological problem.

THE MOLECULAR CLOCK.

Zuckerkandl & Pauling (1962, 1965) noted that the substitution of amino acids within a protein follows a Poisson process. They suggested the characteristic substitution rate for individual sequences is related to absolute time rather than generation length, which led to an absolute time scale by calibrating substitution rates for individual amino acid sequences against known divergence times determined from the fossil record. The molecular clock model has been applied to both amino acid and nucleotide sequences to estimate phylogenies and divergence times (for reviews see Wilson et al 1977; Thorpe 1982; Gillespie 1986; Li et al 1987; Andrews 1987, and papers in Jukes 1987, particularly Zuckerkandl 1987).

Applications of the clock to globins, cytochrome C and 5S RNA sequences produced a date of 1000 to 900 millions years ago (Ma) for the initial radiation of the animal phyla (Fig. 1 Runnegar 1982 a, b, 1986) this work supports claims of a lengthy period of metazoan diversification prior to the appearance of late Precambrian metazoan body fossils (Runnegar 1982 a, b, 1986) and casts doubt upon arguments for a more rapid origination of phyla during the latest Precambrian (Valentine 1986; Valentine & Erwin 1987 and references therein).

The molecular clock model can only be applied when certain assumptions are met. These are: constancy of substitutions rate for the genes under investigation within lineages, and dependence of substitutions rate on absolute rather than generation times. Procedures which test the validity of these assumptions for individual studies are in wide use. However, the reliability of the molecular clock model has been questioned by the discovery of highly variable substitution rates while broad phylogenetic analyses indicate that substitution rates may be more closely effected by generation length than previously appreciated.

The questions of rate constancy.- A constant substitution rate requires incorporation of mutations into a population at a rate equal to their rate of mutations (Kimura 1983; constant is in terms of a Poisson process). The number of variable sites (NVS) controls the number of sites at which substitutions is permissible, and produces a characteristic substitutions rate for each sequence (Kunisawa et al 1987). Since each protein differs in NVS, the rate of substitutions varies between sequences. Furthermore, NVS may vary through time. Consequently, deviations from rate constancy may reflect changes in the rate of incorporations or production of mutations, in the selective value of substitutions or in NVS.

Statistical studies have revealed deviations from rate constancy, but until recently these deviations have been considered insufficient to reject the clock (Langley & Fitch 1974; Fitch & Langley 1976; Hudson 1983). More recent studies suggest the deviations are significant. Moo-Lee et al (1985) compared amino acid substitution rates of superoxide dismutase in *Drosophila* and mammals, indicating a highly erratic clock. However, Moo-Lee et al. may have misinterpreted the saturation effect of substitutions and the data may be equally consistent with a generation time effect. A comparison of echinoid and vertebrate mitochondrial and nuclear DNAs nuclear DNA substitution rates while mitochondrial rates have been stable. Other studies have reached similar conclusions (Gingerich 1986; Britten 1986).

Variations in the rate of substitution in SS RNA were detected in a comparison of arthropod se-

quences with variety of metazoa species (Hendricks et al 1986). The authors concluded that the random deviations in the evolutionary rate SS RNA are too great to be used in analyzing metazoa phylogeny. Field et al (1988) have successfully used 18S RNA to analyze metazoan phylogeny (see below). Gillespie's statistical analysis of substitution rates (1984 1986) demonstrates a highly episodic pattern of substitution, with short, with short periods of rapid change interspersed with longer periods of slower replacement. He suggests that the clock cannot be modeled using a Poisson process in which the rate of the process is itself randomly changing.

Support for rate-constancy has been provided by a determination of the number of variable sites (NVS) in cytochrome C and hemoglobin by Kunisawa et al. 1987. They calculated the square root of the variance and the average number of amino acid substitution differences for about 100 different amino acid sequences of each protein demonstrated little variation in NVS. They concluded that the substitution rate per variable site is close to uniform from bacteria through mammals, although a combination of rapid and slow sites is required to explain the substitution rates of cytochrome C. A constant NVS does not preclude a highly episodic substitution pattern producing a long-term appearance of rate-constancy.

Goodman (1981) has argued that sequence comparison reveal increased rates of protein evolution immediately following establishment of new clades, followed by a decrease in substitution rates as adaptations to a new role increase. In particular, Goodman claims evidence for increased selections during the initial appearance of jawed fish and tetrapods and later during the early Cenozoic mammalian radiation. Increased selective pressure would cause a change in the substitutions immediately following gene duplication (as in hemoglobins and cytochrome C), are also discussed by Ohta (1988) and Li (1985). Jukes (1987) noted an increased rate of transitions (purine-purine or pyrimidine changes) relative to transversions (purine-pyrimidine changes) during the early stages is clearly greater than anticipated by early proponents of the clock, yet whether this variation invalidates the clock entirely,

as suggested by some authors (Hudson 1983; Gillespie 1984), is unclear.

The problems generations times.- If the rates of substitutions is dependent on generation time rather than absolute time, calculated substitutions rates may differ simply because of differences of generation times between lineages, or within a single lineages over time. Wu & Li 1985), Li et al. 1987 and Li et al 1985) compared nucleotide substitution rates for 11 genes (including globins) using a relative-rates test and compared the result several out groups (different outgroups were required because of insufficient sequence information). They concluded that the substitution rate in rodents has been higher than in man since the split between lineages leading to rodents and primates. Synonymous substitutions rates (those in which there is no change in the amino acid) occurred two times more frequently in mice than in humans, while non-synonymous changes occurred 1.3 times more often. Comparison of homologous genes formed by gene duplication provided a directed test of the role of generation time, with only a weak effect if large change in generation time occur. However, the generation time effect is significant for synonymous changes (see also Easteal 1985; Li & Wu 1987; Wu & Li 1985; Easteal 1987; Fisch 1987).

THE MOLECULAR CLOCK AND THE ORIGIN OF ANIMAL PHyla.

Runnegar's molecular clock date for the origin of animals phyla is based largely on amino acid sequences from vertebrate and invertebrate globins and to a lesser extent cytochrome C and 5S RNA nucleotide sequences (Runnegar 1982 a 1986) The gene duplication event following the separation of the agnatha (jawless fish) from jawed fish (and all other vertebrates; Fig. 2) established two related sequences, the alpha globins and beta globins, with independent evolutionary histories. This duplication provides a calibration point for the clock at 450 Ma from fossil evidence. Since the alpha and beta sequences diverged at the same point in all vertebrates. Runnegar has made pairwise comparisons of the amount of substitution in alpha and beta amino acid sequences and demonstrated an approximately normal

distribution of percentage differences (Runnegar 1982a, 1986) Rate-constancy of substitutions in vertebrate globins was examined through pairwise comparisons of alpha and beta chains within species to the appropriate human globin chains. this suggests that while the rate of substitutions has varied between lineages, the rates has been approximately constant within a lineage, as required by the clock. Between-phylum comparisons were then used to calculate... an approximate minimum date for the initial radiation of the animal phyla (Runnegar 1986; 13; Fig. 1).

By assuming that the rate of globin substitutions has been uniform for the past 450 Ma and than this uniform rate may be extrapolated an additional 450 Ma, Runnegar (1982a, 1986) calculated a date for the origin of phyla. Three-way comparisons of annelid, mollusc and vertebrate sequences demonstrate that between phyla differences are greater than those within vertebrates (as expected). Corrected values (which account for superimposed substitutions) are used to calculate the initial radiation of animal phyla and the divergence between alpha globin and myoglobin and beta globin and myoglobin.

There are several reasons for questions the 1000-9000 Ma date. First. the annelid, mollusc, and vertebrate divergence dates the actual divergence of the phyla only in molluscs, annelids and vertebrates are each others closest relatives, or if all three diverged simultaneously (perhaps along with other taxa) from a common ancestor. Other wise, the date, if correct, may simply reflect the divergence of pre annelid, pre-molluscs and pre vertebrate lineages during the evolution of metazoa. In this case the date is for an earlier event in metazoa phylogeny and may not reflect the radiations of existing animal phyla. More importantly, this application of the molecular clock extrapolation of evolutionary rates beyond calibration points rather than interpolation between known points (W.M. Fitch, pers. Comm., 1988) Since there are not methods to test the accuracy of such extrapolations, the result of such studies are unreliable. third, as discussed above, substitutions rates appear to be highly episodic, a feature which will be masked by the sort of long-term analysis used by Runnegar. As noted, increased substitutions rates are particularly

common during gene duplication rates is likely to have greater than calculate rate, and the divergence times less.

A more telling problem lies in the cellular role and early evolutionary history of the globins are likely to have been under intense selective pressure during the metazoa radiation and again during the early evolutions of the vertebrates. These pressure stemmed from the conversion from diffusion to active respiratory transport as body size increased (Runnegar 1983), from increases in metabolic activity and from possible increases in atmospheric O₂ (Runnegar 1982 a, 1982c; see also Towe 1981; Berkener & Marshall 1964).

MOLECULAR PHYLOGENIES

The use of molecular sequence data in phylogenetic analysis is in principle no different from using any other characters. homologous sequences of amino acid, DNA, or RNA are compared and the differences noted, a transformation series (describing the sequences of changes) constructed and the resulting character matrix analyzed using must have a substitutions rate appropriate for the question being addressed, slow for comparisons of phyla and classes with long divergence times, but fast for species within a genus.

A major source of difficulty is constructing transformation series and resolving potential homoplasies. Table 1 contains a portion of the flanking (non-coding) and coding region of the H4 histone gene from an echinoid and an asteroid. Noted that the amino acid sequence of the coding region is constant, despite seven changes in the nucleotide sequence. Each of the nucleotide changes occurred at the third codon position, and due to the redundancy of the genetic code no changes in amino acid sequence occurred. more realistically the sequence is highly conserved and changes in amino acid sequence were selected against.

The impossibility of determining the actual nucleotide sequence from amino acid data is demonstrated by the number of possible nucleotides which could produce most amino

acid. When changes in amino acid sequence do occur, it is frequently difficult to determine the transformation series. In the lower portion of Table 2 the most parsimonious nucleotide changes are listed which would produce the change from the alpha chain to the beta chain. Note that there is no reason to believe that the shortest possible transformation route was actually followed, not that the transformation route was actually followed, nor that the transformation occurred in the direction illustrated. the final residue, Tryptophan, is the only one also found in myoglobin and thus presumably conserved from the ancestral sequence of myoglobin and hemoglobin. [These and other problems are discussed at length by Patterson (1987)].

Field et al. (1988) recently proposed a set of phylogenetic relationships for the metazoa based on 18S ribosomal RNA. Their result suggest three radiations of metazoa (the first actually a radiations of protist lineage, since the multicellular animals appear to be polyphyletic), and rapid phylogenetic splitting of major groups soon after their appearance. While Field et al's, result may well correct, no phylogeny can be based on a single sequence, any more than a biological systematist would use external patterns or internal morphology but ignore developmental patterns or internal morphology. Their work needs to be substantiated using sequences other than 18S RNA and which have substitution rates appropriate for phylum-level comparisons. Finally, the branching order within major groups probably lies within the levels of resolution of the technique and may well be modified by future studies.

DISCUSSION.

The inherent ambiguity in most molecular phylogenetic techniques limits their ability to resolve outstanding evolutionary problems, particularly those involving broad phylogenetic analyses. DNA-DNA hybridization has been used in the phylogenetic studies of birds (Sibley & Ahlquist 1983) and hominoids (Sibley & Ahlquist 1984). Hybridization studies are less informative than sequence studies since the former only produces a measure of overall similarity from the temperature of dissociation. They produce no in-

formation on the patterns of evolutionary change within sequence, nor do they provide any means of separating convergence from shared descent.

Despite the comments above, molecular data may prove enormously important in future broad phylogenetic studies. Phylogenetic characters are presents within the structure and organization of the genome itself, and like morphometric characters have a large number of possible states, are easily distinguished and transformation series constructed. The divergence history of human globin in Fig. 2, was derived from comparative molecular studies and can easily be used as a series of phylogenetic characters. This applications of molecular data has been relatively infrequent, but has greath potential. the major obstacle preventing wider application is tha lack sufficiently diverse information on genome structure and organization.

Determining tha rate and duration of major radiation events is an important paleontological problem, butg the difficulties inherent in variable substitutions rates requiere substantial input from the fossil recor. It is not clear taht tha two fundamental assumption of the molecular clock; rate-constancy and generation-time independence, are vaild in general. nor is it clear that Runnegar's data are applicable to the origin of phyla, or data their extrapolations actually dates the appearance of the extant phyla. The clock may be valuable in analyses of restricted phylogenetic groups with a similar generation lengh, but ist extention to the broad sweep of metazoan history is clearly unwarranted. jan Bergstrom pointed out that if there is a strong differential in substitutions rates during a divergence event, as suggested by Goodwin (1981), use of the molecular clock within a single clade may be flawed as well. Runnegar's conclusion that the initial divergence of metazoa phyla occurred 900-1000Ma (Runnegar 1982 a, b, 1986) appars insuportable. Molecular evidence does not presently allow us to discriminate between a lengthy. precambrian divergence of metazoa and a rapid burst of the diversification close to the actuall appearance of fossil (Valentine & Erwin 1987; Valentine 1986).

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Fig. 1. Molecular clock dates produced by Runnegar for the divergence of metazoa phyla as determined from amino acid sequences of vertebrate globins, cytochrome C and nucleotide sequences of 5S RNA. Calibration points are taken from the divergence between the alpha and beta vertebrate globins, a pairwise comparisons of divergences in amino acid sequences of invertebrate globins between mollusc classes (M), annelid classes (A), and echinoderms classes (E). The point F-A is derived from the average difference between fungal and animal sequence of 5S RNA. The solid region denotes the range of points derived from between phylum comparisons cytochrome C. Since several substitutions may occur at the same site, percentage divergence may be greater than 100%. Fig 6 from Runnegar (1986).

Fig. 2. The evolutionary relationships between members of the human globin gene cluster based on comparisons with other vertebrate groups. The present human globins genes were derived by gene duplication events. The fossil record provides an estimate of the time of the split between the Agnatha and all other fish (and thus all other vertebrates) thus calibrating this duplications even. The remaining gene duplication events have been calibrated in a similar fashions.

Tabla 2. Comparison of initial portion of N-Terminal ends of a lpha and beta chains of humans hemoglobins. The upper portion of the figure shows the DNA triplet codons which could produce the amino acid in the alpha chain, illustrating the redundancy of the genetic code. In the lower portion the shortest possible transformations series are indicated, assuming the alpha chain is primitive. The variety of possible transformations indicates that transformations series cannot be constructed from amino acid data. Capitalized amino acid are identical in each chain. the final residue, tryptophan also occurs in myoglobins. A- adenine, T-thyamine, C- Cytosine, G- guanine, X- any nucleotide.

FIGURE 1

D. H. Erwin

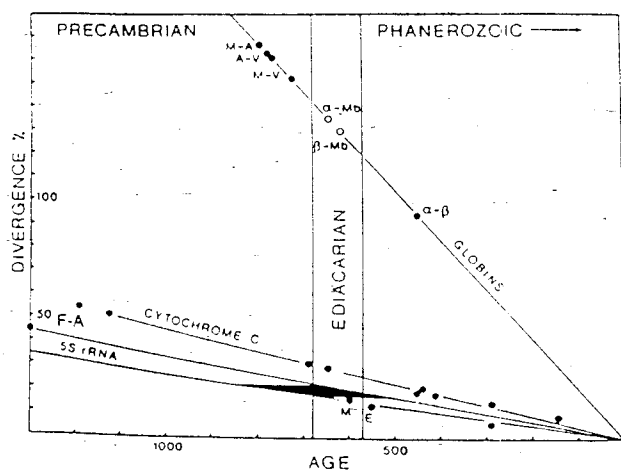


FIGURE 2

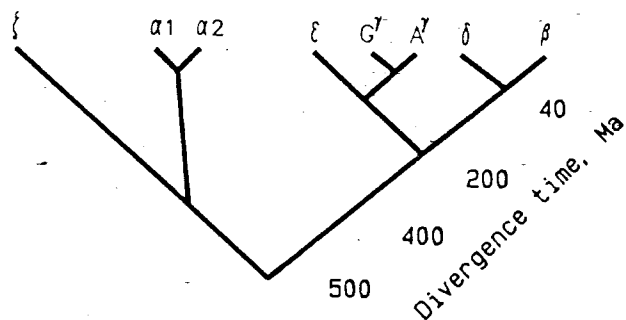


Table 1

	5'	-10	0	5	10	3'									
Sp	CATCATCATG	TCA	GGT	CGA	GGA	AAA	GGA	AAG	GGA	CTC	GGA	AAA	GGT	GGT	
Po	CACAACTATG	TCT	GGT	CGC	GGT	AAA	GGT	GGA	AAG	GGG	CTC	GGC	AAA	GGG	GGT
		ser	gly	arg	gly	lys	gly	gly	lys	gly	leu	gly	lys	gly	gly

Table 2

N-terminal end		
α	β	
VAL	VAL	Possible nucleotides for α-chain:
	His	GTT GTC GTA GTG
LEU	LEU	CAC
Ser	Thr	TTA TTG CTT CTA CTG CTC
PRO	PRO	AGU AGC TCT TCA TCG TCC
Ala	Glu	CCT CCC CCA CCG
Asp	Glu	GCT GCC GCA GCG
LYS	LYS	GAT GAC
		AAA AAG
Thr	Ser	Possible transformation series:
Asn	Ala	ACX → AGT, AGC or ACX → ICX
VAL	VAL	ΔAT, ΔAC → GCX
Lys	Thr	AAA, AAG → ACA, ACG
ALA	ALA	
Ala	Leu	GCX → CTX or GCX → TTG, TTC
TRP	TRP	

HETEROCHRONY AND PHYLOGENETIC TRENDS. *

Kenneth J. McNamara

INTRODUCTION.

Morphological evolution has been considered by some authors (Schindewolf 1936; Goldschmidt 1940; Waddington 1962; Zuckerkandl 1968, 1976; Van Valen 1974; Stanley 1975, 1979; Gould 1977; Alberch 1980) to occur as the result of changes in the regulation of morphological development (heterochrony). Such developmental changes may from part of normal phenotypic variation (Cock 1966; Larson 1980; Travis, 1981), but if the resultant morphological novelties are adaptive significance, selection may favor the establishment of the variant as a genetic isolate, and speciation may result.

Aphylogenetic trend may be defined as directional morphological change between more than two species. Such trends have been interpreted in a number of ways; as little more than elements in the fossil record, chosen by investigators as they apparently conform to a morphologically graded series (Eldredge and Gould 1974); as the result of directional speciation (Grant 1963; Stanley 1979); as due to the action of species selection (Stanley 1973, 1979); or as the result of evolution toward increased speciation in species-specific characters-the effect hypothesis (Vrba 1980).

Ede (1978), Gould (1980), Alberch (1980) and Levinton and Simon (1980) have suggested that heterochrony may direct morphological variation of a species along particular pathways. It is the aim of this paper to pursue this line of argument by examining a series of recently described examples, mainly from the fossil record, which have been interpreted as phylogenetic trends, in which heterochrony has been the principal factor in

directing the course of the morphological evolution (Ruckards 1977; McNamara 1978; in press a, b; McNamara and Philip 1980a; Wright and Kennedy 1980).

A model is proposed in which directional morphological change at the species level arises by repeated selection of phenotypic variants which develop by heterochronic processes. In the model, selection for adaptively significant heterochronic morphological novelties along an environmental gradient resulting in the establishment of a phylogenetic trend.

HETEROCHRONY

Changes in ontogenetic sequences through time occur by contraction, extension, or a shift in timing of rates of morphological development. Reduction in morphological developments through successive ontogenies (paedomorphosis) can occur by neoteny, that is, a reduction in rate of development; progenesis, in which precocious sexual maturation reduces the period of juvenile allometric growth; or by post-displacement, a retardation in onset of growth of particular organs. Increased morphological development during ontogeny (termed peramorphosis by Alberch et al 1979) can occur by; acceleration, an increase in growth rate; hypermorphosis, in which delay in onset of maturity lengthens the period of juvenile allometric growth; or by pre-displacement, that is the earlier onset of development of particular structures.

As Alberch et al (1979) have noted, an ontogenetic trajectory is regulated by a number of parameters; onset age of growth; cessation of growth (generally at onset of maturity); growth rate during the growing period; and size at the commencement of the growth period. Growth through the juvenile phase of development often occurs by allometry, and growth rates are generally much higher than in the adult. Large scale morphological changes may therefore occur during an organism's ontogeny. Consequently any alteration in growth rate or timing of onset of growth, particularly early in the ontogeny, will profoundly change the adult morphology. This will result in the rapid development of the descendant morphological novelties without the development

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of intermediate morphotypes between the ancestor and descendant.

The morphological characteristic of an individual are generally appropriate for the mode of life of its particular stage in development. The environment occupied by the juvenile may be very different from that occupied by the adult. For example, many marine invertebrates have free-living planktic larval stages, but immobile denthic adult. As the ontogeny of an individual in purely directional, the evolution of descendant paedomorphs or peramorphs provides an inbuilt, pre-existing directional component. If selection of such morphotypes occurs periodically along an environmental gradient this will result in directional speciation and the development of a phylogenetic trend.

PAEDOMORPHOCLINES AND PERAMORPHOCLINES.

Let us first consider a series of ontogenies which change with time by paedomorphosis. The ancestral apaedomorph (i.e. non-paedomorphic form) may be said to pass through morphological stages A to M during its development to an adult (fig 1). If a descendant species which evolves from the ancestral form is neotenic or progenetic, the descendant will pass through fewer morphological stages during growth, say only A to K. By the postulates of the model there will subsequently be selection of a paedomorph which passes through fewer stages still during its ontogeny, say A to I. Similarly, this form may then itself give rise to a descendant species which passes only through stages A to G, and so on to the last species to evolve, the most paedomorphic with respect to the ancestral apaedomorph, which passes only through morphological stages A to C.

It can be seen by considering just the adult morphological stages of the six hypothetical species that they constitute a morphological gradient through time: M-K-I-G-E-C. This temporal morphological change between adults thus follows the opposite pathway to the ontogenetic development of the earliest species, the apaedomorph, which passes through all morphological stages, A to M. The sequence of adult

morphologist displaying such a temporal morphological gradient will be a phylogenetic trend; it is herein termed a paedomorphocline.

The evolution of a descendant paedomorphic species need not result in the extinction of its direct ancestor. Indeed, in the situation described above all six species could coexist temporally, though not spatially (see below); the paedomorphocline could then be considered on a single time plane as a purely ecological phenomenon.

As Gould (1977) has pointed out, there is no reason why paedomorphosis should occur any more often than peramorphosis. Thus one can envisage the converse to the paedomorphocline in the peramorphocline. This, by definition, will possess the opposite pattern of morphological change to the paedomorphocline. An ancestral species passing through morphological stages A to C during ontogeny may give rise by one of the peramorphic processes to a descendant species which passes through morphological stages A to E during ontogeny (fig 2); this in turn may give rise to a species passing through morphological stages A to G, and so on if the resultant morphologies are adaptatively successful along the environmental gradient which the peramorphocline is invading. For a similar reason to those for the paedomorphocline, if the ancestral species persists in its particular niche and if suitable niches are available along the environmental gradient, speciation will be unidirectional (see discussion).

THE TEGULORHYNCHIA-NOTOSORIA PAEDOMORPHOCLINE.

The earliest known species of the rhynchonellide brachiopod Tegulorhynchia, an undescribed species from the Early Paleocene to Early Eocene of Western Australia (Mcnamara in Press a), may be considered to be the ancestral apaedomorph and the earliest member of the Tegulorhynchia-Notosoria paedomorphocline (fig 3). This ancestral species of Tegulorhynchia undergoes more morphological change during its ontogeny than any other species on the paedomorphocline. Small juveniles possess a narrow shell with shallow valves of similar depth; a rectimarginate commissure; 25 costellae at a shell length of 2 mm; and long, pointed beak with a large

foramen, and widely disjunct, small deltidial plates. Through the ontogeny, during which the shell grows to a maximum length of about 18 mm, the principal morphological changes involve a broadening of the shell; a large increase in valve depth; development of a strongly uniplicate commissure; an increase in the number of costellae by frequent branching early in ontogeny, producing up to 80 fine costellae in the adult; reduction in height of beak and increase in umbonal angle; and diminution in size of foramen by the deltidial plates becoming conjunct, then almost closing the foramen.

The second species in the paedomorphocline is the New Zealand Late Eocene to Early Miocene *T. squamosa* (Hutton 1873). Lee (1980) considers the living *T. Doederleini* (Davidson 1886) to be a separate species from *T. squamosa* on account of stratigraphical (ranging from the Pliocene to the present day) and geographic discontinuities, even though morphologically there seems little to warrant their separation. These species may be considered to be neotenic with respect to the apaeomorph, as they have fewer ribs in the adult (up to 60); slightly narrower shell; less strongly developed uniplication; larger foramen, though the deltidial plates remain conjunct; and larger shell size (length up to nearly 25 mm).

The Australian Tertiary species *T. coelata* (Tenison Woods 1878) and *T. thomsoni* Chapman and Crespin 1923 continue this paedomorphic trend. They have fewer costellae than *T. squamosa*; narrower shell; disjunct deltidial plates; larger foramen; more pointed beak; and weaker uniplication of the commissure. Stratigraphically *T. coelata* and *T. thomsoni* appeared later than *T. squamosa*, *T. coelata* being first recorded from the Early Miocene. The younger of the two, *T. thomsoni*, has fewer costellae than *T. coelata* (28 as opposed to 36-40); more prominent beak; valves of similar depth and larger foramen, consistent with its position further along the paedomorphocline.

The end members of the paedomorphocline, *Notosaria antipoda* (Thomson 1918), which first appeared in the Late Oligocene, and *N. nigricans* (Sowerby 1846), which ranges through to the present day, possess the most juvenile characters

in the adult; narrowest shell; weakest uniplication; fewest costellae (25); most disjunct deltidial plates; and largest foramen.

These end members seem to have appeared before the preceding species on the paedomorphocline, *T. thomsoni*, illustrating that, on occasions, some paedomorphs may not necessarily have evolved from the immediately preceding species on the paedomorphocline. In other words, the position of *T. thomsoni* on the paedomorphocline was bypassed by species of *Notosaria*, which probably evolved from *T. squamosa*. *T. thomsoni* is morphologically close to *Notosaria*, and taxonomic revision of the species might even result in its emplacement within that genus. Geographic separation of *Notosaria* and *T. thomsoni* indicates evolution of similar morphotypes from different ancestors, which have different positions on the paedomorphocline. The southern Australian *T. thomsoni* evolved from *T. coelata*; the New Zealand species of *Notosaria* evolved from *T. squamosa*. There is greater morphological separation between *Notosaria* and *T. squamosa* than between the Australian species, *T. thomsoni* and *T. coelata*. Be that as it may, the overall phylogenetic trend has favored selection of paedomorphs.

Ecological significance of the paedomorphocline. The living species of Tegulorhynchia, *T. doederleini*, occurs only in water deeper than 100m (Lee 1980) and to as deep as 635m (Dall 1920). Lee (1980) considers that fossil Tegulorhynchia inhabited deep or relatively quiet water. The paedomorph *Notosaria*, however, commonly occurs in the intertidal zone (Allen 1960; Percival 1960; Rudwick 1962; Richardson 1981) and is most commonly found in water less than 200m deep (Lee 1978), although it has been recorded from deeper water. Lee and Wilson (1979) consider the extinct *N. antipoda* also to have inhabited a shallow water, high hydrodynamic environment.

I believe that the neotenic retention by species of *Notosaria* of a large foramen and, consequently, a large, muscular pedicle (Richardson 1981), which permits strong attachment of the animal to its substrate (Richardson 1979), enabled this morphotype to inhabit a much shallower water, thus higher hydrodynamic environment than Tegulor-

hynchia. The latter, in its quiet water environment, can obtain adequate attachment with a smaller teth ering pedicle. The juvenile Tegulorhynchia, with its relatively large foramen, would possess a relatively thicker pedicle to provide secure attachment to the substrate, as the smaller juveniles are more susceptible to dislodgement by current activity and require firm attachment (Richardson 1981).

Other morphological character possessed by the paedomorphic Notosaria, such as fewer, much coarser costellae, are also suited to occupation of a shallow water environment. The relatively shallower valves of Notosaria indicate possession of a shallower spirolophous lophophore than Tegulorhynchia. Such a lophophore would be adequate in higher hydrodynamic environment. Occupation of quieter water by Tegulorhynchia necessitates possession of a deeper spirolophous lophophore. This correlation of habitat water depth with lophophore size, deduced from shell depth, has been noted by Hurst and Watkins (1978).

The intermediate species of Tegulorhynchia along the paedomorphocline may be predicted to have been adapted to intermediate conditions along the environmental gradient between the predominantly nearshore, high hydrodynamic habitat of Notosaria, and the deep water, low hydrodynamic environment occupied by the apaedomorph.

The apaedomorph shows no indication of any morphological drift during its 10 Myr range from the Early Paleocene to the Early Eocene (McNamara in press a). Lee (1980) has similarly noted that the morphotype *T. squamosa-doederleini*, although exhibiting a certain degree of phenotypic variation, underwent no directional morphological change for over 40 Myr from the Late Eocene to the present day. The same is true for Notosaria, Lee and Wilson (1979) having observed that there is, indeed, less variation between population means from the Miocene to present day than between extremes of living populations of *N. nigricans*.

Olenellus Paedomorphocline.

A paedomorphocline may be observed in the fossil record on what is, essentially, a single time plane. In the Early Cambrian Fucoid Beds of northwest Scotland, five species of the trilobite *Olenellus* have been identified, and interpreted as having arisen by paedomorphosis (McNamara 1978). In contrast to the neotenic Tegulorhynchia-notosaria paedomorphocline which is observable both spatially and temporally, the *Olenellus* paedomorphocline is an example of a spatial progenetic paedomorphocline. A slight increase in size, the progenetic nature of the *Olenellus* example, discussed in detail by McNamara (1978), is demonstrated by the progressive reduction in maximum adult size along the paedomorphocline.

Although all five species of *Olenellus* occur in the same rock unit, *O. lapworthi* Peach and Horne 1892, is considered to be the ancestral apaedomorph, because its morphology is typical of that of most other areas. Even within the Fucoid Beds it is a widespread species, whereas *O. armatus* Peach 1974, which lies at the other end of the paedomorphocline (fig 4), occurs at only one locality. This restricted distribution of *O. armatus* is characteristic of progenetic species (McNamara 1981). *O. lapworthi* followed along the paedomorphocline by *O. reticulatus* Peach 1894, *O. hamolucus*, Cowie and McNamara 1978, *O. intermedius* Peach 1894 and finally, *O. armatus*.

Although the detailed ontogeny of the apaedomorph, *O. lapworthi*, which would have undergone most morphological development during its ontogeny, is incompletely known, its adult form is very similar to that of other species of *Olenellus* in which ontogenies have been well documented (Palmer 1957; Hu 1971). Essentially, the apaedomorph would have undergone large-scale morphological change involving an appreciable broadening of the cephalon, largely due to widening of the extraocular area; posterior migration of the frontal lobe, resulting in increased glabella furrow sinuosity; posterior migration of eye lobe and genal angle, resulting in a rounding of the anterior cephalic margin; migration of the intergenal spine toward the genal angle, and its degeneration; and an increase in thoracic pleural width as new segments were added (McNamara 1978).

As trilobites, like other arthropods, grew by periodic moulting, their ontogeny consisted of periods of morphological stasis, interspersed by rapid saltations to new morphologies. This is quite unlike the morphological development of the branchiopods which proceeded continuously through to the adult phase. Even so, species of both groups of the animals appear to have periodically evolved along paedomorphocline by morphological saltations. The potential existed for precocious maturation to have occurred in species of *Olenellus* at any moulting event. However, only five paedomorphs appear to have become establishing as separate species, each occupying and adaptive peak on the paedomorphocline. Palmer (1957) estimates that *Olenellus* underwent ecdysis at least twenty-nine times during ontogeny—twenty-nine potential opportunities for progenesis. If the fossil record in the Fucoid Beds is a reasonably accurate indication of the number of the species, can occur only when sufficient morphological distance is established between the ancestral and descendant morphotypes such that an adaptive threshold is crossed. Competition for resources is thus avoided.

ECOLOGICAL SIGNIFICANCE OF THE PAEDOMORPHOCLINE.

The environmental gradient along which this paedomorphocline is considered to have developed is again a deep to shallow marine situation. In this case, however, it is believed that higher water temperatures in shallow water might have triggered premature maturation in the pelagic larvae of the benthic adult apaeomorph, which is thought to have inhabited deeper water (McNamara 1978). Control by temperature in inducing and inhibiting maturation in vertebrates is well documented (Singh-Pruthi 1924; Newell 1948; Wigglesworth 1954; Boner 1968). If Jell 1978 is correct in interpreting the function of genital caeca, which cover the extraocular area of the apaeomorph, as a secondary respiration base, the progressive reduction in extraocular area along the paedomorphocline may have restricted the successive paedomorphocline may have restricted the successive paedomorphs to progressively

more oxygenated waters, this interpretation is supported by the character of the last paedomorph, *Olenellus armatus*. It retains many overt larval characteristics, suggesting it led an entirely pelagic existence in shallower, warmer, more oxygenated waters than its progenitor.

OTHER PAEDOMORPHOCLINES.

The living spatangoid *Breynia*. Spatial paedomorphocline can be observed not only species. A recent study (McNamara in press b) of living Australian species of the spatangoid echinoid *Breynia* indicates the presence of a paedomorphocline involving three living species. Despite its widespread occurrence in Miocene and Pliocene rocks of the Indian Subcontinent, Japan, Taiwan and Indonesia, *Breynia* is recorded in Australia only from Miocene rocks in the northwest of the continent and is absent from the Miocene deposit of the southern part. The three living species which occurs along the western coast, are *B. desorii* Gray 1851, which occurs along the western coast and the western part of the northern coast; *B. australasica* (Leach 1815), which occurs along the eastern coast and in the Arafura Sea.

The three species form a paedomorphocline. The western species, *B. desorii*, which is known to have existed at least as far back as the late Pleistocene and was derived from the Indian Ocean genetic pool, is considered to be the apaeomorph. It shares more features in common with Miocene species of *Breynia* than either of the other two living Australian species, principally in its greater aboral tuberculation, longer petals and smaller peritome. Some variants even possess a lambrum which is separated from the plastron by ambulacral plates, a feature otherwise found only in Miocene species of *Breynia*.

B. australasica is interpreted as having evolved by paedomorphosis from *B. desorii* and spread to the east coast; absence of both fossil and living *Breynia* from southern Australia suggests that *Breynia* spread around the northern part of the continent, *B. australasica* attains maturity later than *B. desorii* and retains juvenile characters. *B. desorii* into the adult phase; it is therefore neotenic. The juvenile characters retained in-

clude fewer primary tubercles; shorter petals; broader, shorter internal fasciole; relatively larger peristome and periproct; and fewer subanal pore pairs, *B. n.sp.* is the end member of the paedomorphocline. It is an uncommon species and retains earlier juvenile characters into the adult than *B. australasie*, from which it is believed to have evolved. It has a low test: fewer primary tubercles; shorter petals; smaller plastron; broader periplastral area; and fewer subanal pore pairs within a narrower fasciole.

Phenotypic variation between adults of species of *Breynia* involves slight changes in rates of development of particular structure. For instance, in a population of the apaedomorph *B. desorii* from off Dirk Hartog Island, Western Australia, many adult specimens possess a wider internal fasciole and fewer primary tubercles than is normal for the species. In retention of these more juvenile characters, the individuals are reminiscent of the paedomorphic *B. australasie*. However, in all other characters, such as peristome and periproct shape, the Dirk Hartog population is like *B. desorii* from elsewhere. It seems likely that such phenotypic variation led to the evolution of *B. australasie* from *B. desorii*.

The environmental gradient along which this paedomorphocline has developed, is not known. There are insufficient data on the life habits of the adults, and nothing is known of the mode of life of the juveniles. The differences between species could be related to differences in depth of burial in the sediment; preference for different sediment type; or perhaps occupation of benthic habitats at different water depths. Development of a larger plastron and more subanal tubercles in *B. desorii* suggest that it may burrow deeper than either of the other two species.

The Cenomanian ammonite *Protacanthoceras*. Wright and Kennedy's (1980) study of the acanthoceroid ammonite genus *Protacanthoceras* has revealed the evolution of a number of paedomorphic species through the Middle to Upper Cenomanian *Protacanthoceras tuberculatum* Thöni 1972-P.T. mite Wright and Kennedy 1980-*P. arkelli* Wright and Kennedy 1980-P. *tegulicium* Wright and Kennedy 1980 form the early Middle

Cenomanian *costatus* Zone to the mid Upper Cenomanian *naviculare* zone. Wright and Kennedy (1980) record that while speciation appears to have occurred by series of saltations, there is also evidence in the *P. tuberculatum* tootstock of some apparently gradual phylogenetic changes between the successive paedomorphs involved a reduction in strength of ornament and development of a compressed shell. The final species in the paedomorphocline, *P. tegulicium*, lacks siphonial and outer ventrolateral tubercles. These are present from the nuclei of *P. arkelli* which precedes *P. tegulicium*. Tuberculation is most strongly developed in the ancestral apaedomorph of the paedomorphocline, *P. tuberculatum tuberculatum*.

Evolution of the small (15-30 mm diameter) *Protacanthoceras* from the large (up to 500 mm) *Acanthoceras* was by progenies. The small species *P. asgerri* Wright and Kennedy 1980, which does not lie on the paedomorphocline, probably arose from *P. tuberculatum* by progenies. The environmental gradients along which the paedomorphocline developed is not known, although Dr. Kennedy (litt. comm., 10 Jun 1981) suggest that the adaptive significance of the paedomorphosis may have lain either in the shorter juvenile phase, reducing the time between hatching and spawning, or in development of a smaller size, allowing exploitation of new niches.

The Silurian graptolites. *Petalograptus* and *Cephalograptus*. Rickards (1977) has described a series of paedomorphs in the monograptis *Petalograptus* and *Cephalograptus*, which evolved from *P. onatolengatus* through three intermediate distinct paedomorphs with overlapping time ranges, along a paedomorph is a species with much enlarged early thecae. The progressive increase thecal size along the paedomorphocline implies neoteny was the paedomorphic process. Again, the environmental gradient associated with the paedomorphocline is not known, although the development of progressively longer thecae along the paedomorphocline may indicate adaptation to different feeding strategies.

THE PARASTER-SCHIZASTER PERAMORPHOCLINE.

The spatangoid echinoid genus *Schizaster* is generally considered to consist of two subgenera: *Schizaster* and *Paraster*. Which lie at opposite ends of a morphological gradient (McNamara and Philip 1980a). The *Paraster* morphotype appeared before the *Schizaster* morphotype and has, like *Schizaster*, persisted to the present day. The basic *Paraster* morphological characteristics of relatively small test; centrally positioned apical system and apex shallow ambulacrum III, which aborally bears few, widely spaced, pore pairs; shallow anterior notch; widely divergent anterior petals; short labrum, which projects little anteriorly; and relatively small plastron, are suitable for living in coarse sediment. Chesher (1966) has noted that living *P. florinfiensis* Kier and Grant 1965 burrows in coarse calcareous sand to a depth of 25 cm. Fossil species of *Paraster* from the Tertiary of Australia occur in calcarenites (McNamara and Phillip 1980a), suggesting they had similar substrate preferences.

A series of the species occurs through the Australian Tertiary which I interpret as a peramorphocline from the *Paraster* morphotype to the *Schizaster* morphotype (Fig 5). The Middle Miocene *S. sphenoides* Hall 1907 developed at the end of a peramorphocline which comprises the ancestral aperamorph (i.e. non-peramorphic form) *P. carinatus* McNamara and Phillip 1980a (Middle to Late Paleocene), followed by *P. tatei* McNamara and Philip 1980a (Late Oligocene to Early Miocene) and *S. abductus* Tate 1981 (early Miocene). The principal morphological changes which occur along the peramorphocline involve; the attainment of a larger, longer test; a more posteriorly situated apex and apical system; less anteriorly divergent anterior petals; deeper petals; deeper anterior notch and ambulacrum III, which bears an increasing number of more closely spaced funnel-building tube feet (McNamara and Philip 1980a, Fig 8); more anteriorly positioned and sunken peristome; more anteriorly projecting labrum; and relatively larger plastron. The extreme *Schizaster* morphology developed in the Middle Miocene by *S. sphenoides* has persisted to the present day.

Ontogenetic study of a living species of *Schizaster*, *S. myorensis*, from Moreton Bay, Queensland (McNamara and Philip 1980b) revealed that small juveniles of the *Schizaster* morphotype possess the same morphological characteristics as juvenile *Paraster* and are very similar to adult *Paraster*. Ontogenetic development of *Schizaster* therefore parallels phylogenetic development of the *Schizaster* morphotype from the *Paraster* morphotype through the Tertiary. Species possessing the extreme *Schizaster* morphotype are well adapted to living in very fine-grained sediments. Indeed, *S. myorensis* burrows to at least 10 cm depth in the muddiest sediment in Moreton Bay. Furthermore, extinct forms such as *S. sphenoides* occur only in fine-grained marly beds.

By what is believed to be a combination of hypermorphosis (the *Schizaster* species becoming mature later than *Paraster*) and acceleration in development of some features, particularly the rate of production of funnel-building tube feet, the *Schizaster* morphotype is capable of occupying a very fine-grained sediment. Living buried in such a sediment would necessitate the possession of morphological characteristics which can most efficiently utilize a single source of water drawn down the funnel. *Paraster*, although building a funnel (Chesher 1966) which also assists in surface feeding, would have its test bathed by water percolating through the much coarser sediment which it inhabits.

The peramorphic development in *Schizaster* of deep petals, posteriorly positioned apex and apical system (which enabled lengthening of the anterior petals and ambulacrum III), deeper peristome and elongation of the labrum, all reflect development of adaptations which contributed toward efficient utilization of a localized water source, and consequently permitted colonisation of a much finer-grained sediment than that inhabited by ancestral morphotypes. The increase in area of palastron, which was accompanied by an increase in number of burrowing plastral spines, facilitates burrowing in a fine-grained sediment. The ability of the evolving peramorphs to inhabit progressively finer sediments reflects the adaptations of the species on the morphological gradient to the environmental gradient of

coarse to fine-grained sediment. Exploitation of new sedimentary environments by morpho types evolving along the peramorphocline resulted in spatial separation of the peramorphs and resultant geographic and genetic isolation.

DISCUSSION.

The model of paedomorphoclines and peramorphoclines developed from the example discussed above, proposes that directional morphological evolution may arise by heterochrony. How can such a model be reconciled with evolutionary theory which predicts that speciation is an essentially random event? What are the principal factors controlling the speciation? Why should the speciation be directional? The following discussion will deal with paedomorphoclines, but equivalent arguments apply to peramorphoclines.

The establishment of an extreme paedomorphic phenotype as a separate species is ultimately dependent upon its possession of morphological novelties adaptive in an environment which the ancestral form is unable to occupy because of its unsuitable morphological characteristics. The morphological novelties arising from paedomorphosis may be regarded as preadaptations (sensu Simpson 1944) which enable colonization of a new environment.

Larson et al. (1981) have discussed how presadaptations arising in plethodontid salamanders arose by developmental alterations. Once initiated, complex morphological changes were able to be archived relatively rapidly. The morphology of a descendant paedomorph must therefore be sufficiently different from its ancestor to avoid competitive exclusion and allow ecological separation of the ancestor and descendant. The extreme paedomorphic phenotype may be regarded as having the ability to overcome an adaptive threshold in order to attain the ecological separation. This is likely to ensure reproductive and genetic isolation of morphotypes, so facilitating the evolution of a new species. As a consequence, the ancestral species would be expected to persist temporally, through not spatially, with the paedomorph.

If selection act only on morphotypes derived by changes in regulation of development, the persisting ancestral species effectively inhabit reverse speciation to ancestral morphologies, by competitive exclusion, and so direct speciation along the paedomorphocline (fig. 6). Selection on successive paedomorphs along a paedomorphocline can only occur if the adaptations that are developed are suited to the environmental gradient, such as a relatively larger pedicle and coarser ribs in the Tegulorhynchia-Notosaria paedomorphocline developed in response to an environmental gradient of deep to shallow water.

The importance of an environmental gradient in controlling directional speciation has been recognized by Grant (1963), who observed the presence of five species of the herbaceous plant *Polemonium* along an environmental gradient toward lower temperatures at higher elevation. The position occupied by each species on a paedomorphocline may be equated to an adaptive peak (Wright 1932). Variation in gene regulation will provide the changeable genetic composition of adaptive peaks. Only some of the paedomorphic phenotypes will reach the adaptive peaks. These will be the few extreme paedomorphs which are capable of overcoming the adaptive threshold. The position of the adaptive threshold will depend upon the adaptive significance of the morphological novelty and the nature of the environmental gradient; and the nature of the environmental gradient; but it need not remain static through time. Consequently, if the same range of paedomorphic phenotypes were to exist over a period of time, it is only when there is a shift in the position of the adaptive threshold that extreme paedomorphic phenotypes would be able to cross it (Fig. 6). In all probability, however, the range of paedomorphic phenotypes will also be a variable factor through time. The coincidence of the two, perhaps both by a slight shift in the adaptive threshold and by an extension of the range of paedomorphic phenotypes, will be necessary for the successful occupation of the next adaptive peak along the paedomorphocline.

A paedomorphocline may therefore be considered as a series of adaptive peaks lying along

a unidirectional morphological axis. The model predicts that these peaks generally become established sequentially along an environmental gradient. The position of each peak on the environmental gradient will be dictated by the position of the ancestral paedomorph, and the degree of morphological separation between each descendant and ancestor pair necessary for them to avoid significant competition for resources.

Variation in rates of morphological development in the paedomorph, or changes in the timing of onset of maturity, might be expected to be sufficient to produce enough morphotypes to occupy all the adaptive peaks on the heterochronic gradient. In such a situation the last species in the paedomorphocline could therefore potentially develop directly from the paedomorph by a large change in the timing of onset of maturity. Other, intermediate, forms on the heterochronic gradient would subsequently develop at random. It is clear, however, from the examples of temporal paedomorphoclines and paedomorphoclines described above, that the adaptive peaks are not occupied in this random fashion.

The clearly sequential evolution of species indicates that there are constraints imposed on the development or the heterochronic phenotypes. Either each species can undergo only a limited degree of developmental retardation or acceleration, sufficient only to produce occasional phenotypes capable of overcoming the adaptive threshold and occupying the next adaptive peak along the paedomorphocline, or ecological constraints may restrict more than a limited distance along the environmental gradient. The most probable situation is that both of these factors operate.

It might be expected that the model of heterochronic directional speciation would only fit a gradualistic interpretation, as both ontogeny and the environmental gradient along which the species is developing may be gradually changing, through space and time respectively. However, as has been outlined above, this does not appear to be the case. Of the almost infinite number of morphotypes which could potentially develop along the paedomorphocline, only a

small number, those able to cross the adaptive threshold and occupy new adaptive peaks, become established as viable species. All are separated by significant morphological unit, with little or no morphological shift through time once established as a separated species.

The species which lie along a paedomorphocline are therefore characterized by their temporal morphological stability, their temporal morphological stability, Their distinct morphological separation from one another and by their sudden, generally sequential, appearance in the fossil record. The pattern of evolution of species along heterochronic morphological gradients is therefore consistent with the "punctuated equilibria" model of macroevolution (Eldredge and Gould 1972; Gould and Eldredge 1977), rather than the gradualistic model.

The importance of heterochrony in rapid speciation has been discussed by Valentine and Cambell (1975); Gould (1977); McNamara (1978); and Stidd (1980). However, the role of heterochrony in the development of phylogenetic trends and assimilation into the "punctuated equilibria" framework has received little attention. Stanley (1975, 1979) has stressed the importance of "species selection" in the punctuational model and its role in the development of a phylogenetic trends. He considers that the direction takes by speciation has a strong random element, a view taken by Rensch (1959), Wright (1956, 1967) and Mayr (1963). Speciation, if essentially random, would have equal probabilities of moving in either direction along an environmental or morphological axis. STANLEY (1979) considers that apparently directional speciation occurs as a consequence of environmental factors acting on initial random speciation. Recently, however, Gould (1980) and Alberch (1980) have questioned the validity of the concept of random speciation in all situations and stressed the non-random appearance of some morphological novelties. Maynard Smith (1981) has gone so far as to state that "there is no evidence to show that the direction of punctuational changes is random with respect to major trends".

When selection acts only on descendant paedomorphs it is immediately restricted to a single

morphological axis of the organism's own ontogeny. As outlined above, persistence of the ancestral morpho type on the paedomorphocline blocks evolution back to the ancestral form by competitive exclusion. Selection in that situation is channelled in one direction: away from the ancestral morphology. Directional speciation will thus ensue, as long as the environmental gradient persists. Termination of the gradient will result in the cessation of the phylogenetic trend.

CONCLUSIONS.

A number of recently described examples of heterochrony in diverse groups of the organisms indicate that directional speciation, resulting in phylogenetic trends, may develop because the ontogeny of an individual can provide an built, directional component. Any heterochronic variation, by extension, contraction, or change in timing of onset, of morphological effects on the descendant adult. Paedomorphs or peramorphs may thus be morphologically quite distinct from their progenitors and can potentially evolve rapidly. Selection for these morphotypes can lead to rapid speciation and the evolution of morphologically distinct descendants, without the appearance of intermediate morphotypes.

Providing that the descendant morphotypes are suitably adapted to conditions along an environmental gradient, a phylogenetic trend, in the form of a paedomorphocline or peramorphocline, can become established. Persistence of the ancestral form with little or no directional morphological drift through time limits selection along the heterochronic morphological gradient to one direction; that is, along the environmental gradient, away from the ancestral species. Over a period of time directional speciation may thus occur and result in the development of a phylogenetic trend.

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Fig. 1. The paedomorphocline; a discontinuous morphological gradient of progressively more paedomorphic species through time. Time 1 represents a situation whereby all paedomorphs and the apaedomorphs could coexist temporally; ecological separation of morphotypes will ensure they do not coexist spatially, A to M represent arbitrary ontogenetic stages. Species time range is for the adult morphology.

Figure 2: The peramorphocline a discontinuous morphological gradient of progressively more peramorphic species through time; successive species evolving along the peramorphocline pass through more morphological stages during ontogeny. A to M represent arbitrary ontogenetic stages. Species time range is for the adult morphology.

Fig 3. Tegulrhyndia, Notosaria, paedomorphocline by neoteny; illustrating temporal narrowing of shell, reduction in number of costellae, decrease in beak angle and increase in relatively foramen size by progressive reduction in rate of growth of deltidial plates. These adaptations enabled colonisation of progressively shallower water environment. All adult reconstruction x 2 (after McNamara in press a, tex-fig 3).

Fig 4 A spatial paedomorphocline by progenesis in species of olenellus from the lower Cambrian of north west Scotland. The paedomorphocline is considered to have followed an environmental gradient of increasing temperature and oxygen content into shallower water. Bars by reconstruction represent Imm, illustrating the progressive reduction in maximum adult size as a consequence of progenesis.

Fig 5. Peramorphocline from *Scrizater* (paraster) to *Schizaster* (schizaster) through the Australian Tertiary to recent. Peramorphic establishment of the *Schizaster* ss. morphotype is thought to reflect adaptation to occupation of finer grained sediments. S. (paraster) adult

reconstruction X 1; S (schizaster) adult reconstruction x 0.5; juvenile reconstruction x 1.

Figure 6. Suggested mechanism for the development of a paedomorphocline. It is not until a population of species 1. Such as population B, has paedomorphic phenotypes capable of crossing the adaptive threshold and occupying a new adaptive peak, that a paedomorphic species may arise. Shifting adaptive thresholds and variation in range of paedomorphic phenotypes affects timing of evolution of species 2. For instance, even though the range of paedomorphic phenotypes is greater in population A than B, the position of the adaptive threshold is such that it is not crossed by phenotypes of population A. Extreme paedomorphs of population C of species 1 are limited by comparative exclusion. There is no adaptive threshold. The persistence of such population, however, effectively blocks reverse speciation from species 2 to species 1. Selection of extreme paedomorphic phenotypes of species 2 which can cross and adaptive threshold results in the establishment of a further paedomorphic species, species 3. The paedomorphocline is therefore established and unidirectional speciation ensues. Broken lines represent adaptive thresholds.

FIGURE 1

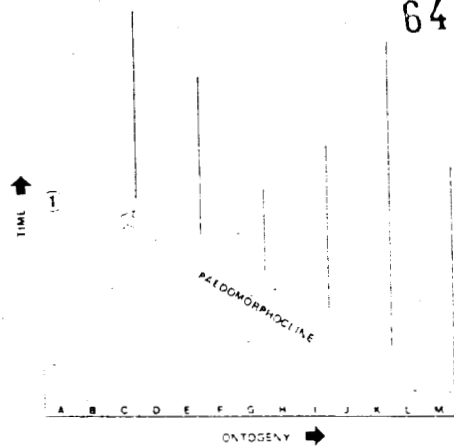


FIGURE 2

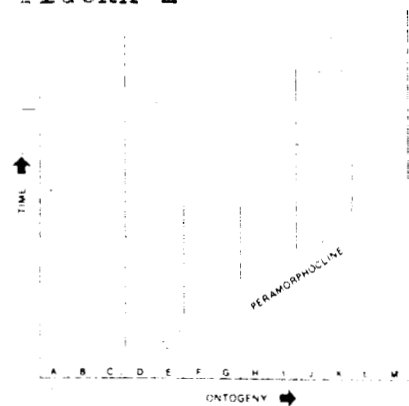


FIGURE 3

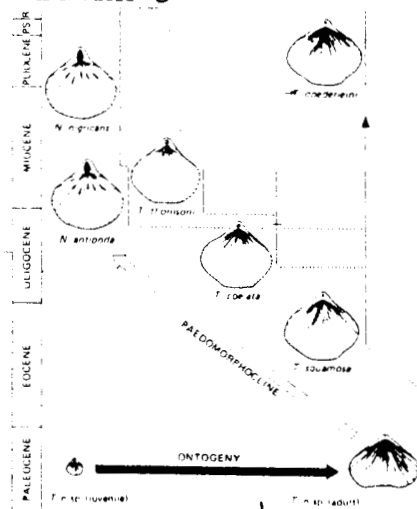


FIGURE 4

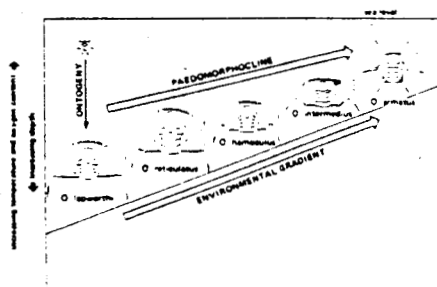


FIGURE 5

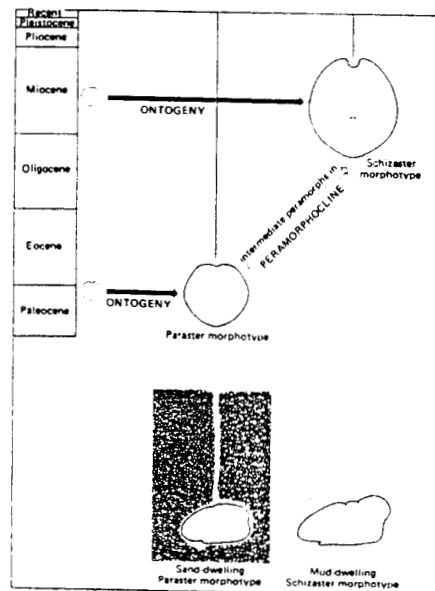
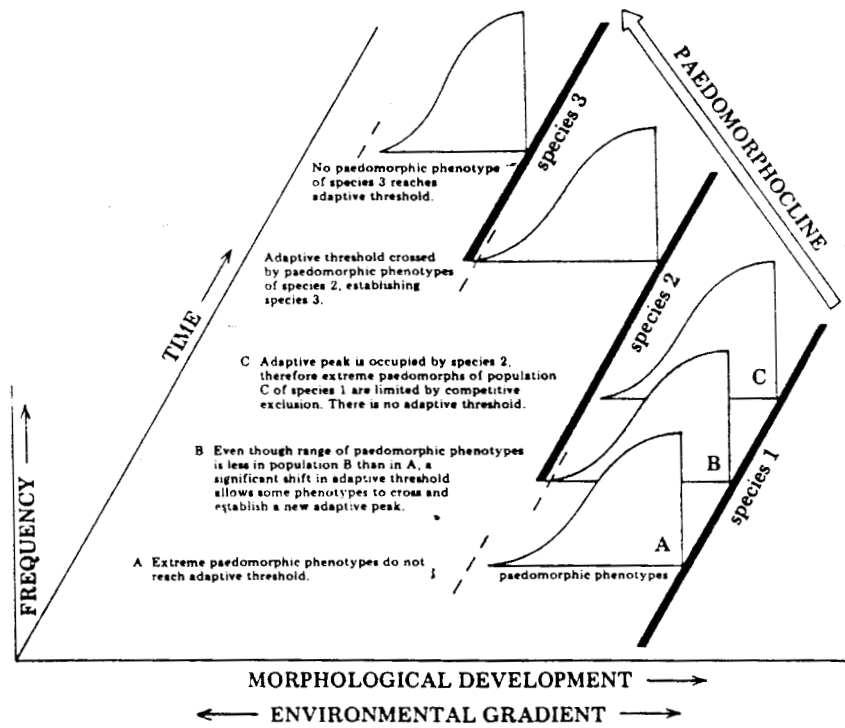


FIGURA 5



10.3 (d)

EVOLUTIONARY FAUNAS.

J.J. Sepkoski, Jr.

In: D.E.G. BRIGGS & P.R. CROWTHER, EDITS., 1990. PALEOBIOLOGY, A SYNTHESIS, OXFORD, INGL., BLACKWELL SCIENT. PUBLIC., P 37-41.

Evolutionary faunas are sets of higher taxa (especially classes) that have similar histories of diversification and together dominate the biota for an extended interval of geological time. The expansion and decline of evolutionary faunas can be used to describe large-scale variations in faunal dominance and to interpret temporal changes in global taxonomic diversity in the fossil record. The concept was introduced by Sepkoski (1981), who identified three 'great evolutionary faunas' in the Phanerozoic marine record. These faunas were defined statically in a factor analysis of familial diversity within taxonomic classes, which grouped together classes that attained their maximum diversities around the same time. The analysis permitted the histories of the aggregate faunas to be traced from initial diversification through dominance and into decline. This treatment of the faunas as units throughout their histories distinguishes the concept of evolutionary faunas from that of "dynasties", used by some authors for assemblages of dominant taxa during specified intervals of geological time.

MARINE EVOLUTIONARY FAUNAS.

Characteristics. The three evolutionary faunas identified in the marine fossil record are the Cambrian fauna, important during the Cambrian Period, the Paleozoic Fauna, dominant from Ordovician to Permian, and the Modern, or Mesozoic-Cenozoic Fauna, dominant in the post-Paleozoic (fig. 1A). The classes in each faunas share a number of characteristics, or central tendencies, suggesting that they are not randomly assembled groups of taxa. The most striking characteristic is that the classes tend to diversify

together, each successive fauna displaying a slower rate of diversification but higher level of maximum diversity than those preceding it. These properties lead to a sequential expansion of evolutionary faunas and resultant step-like pattern of increase in global marine diversity (with the step between the Paleozoic and Modern faunas disrupted by the Paleozoic and Modern faunas disrupted by the massive Late Permian extinction event-Section 2.13.4). This pattern is present even though most marine classes originated early in the Phanerozoic and the total number of classes has remained virtually constant since.

The expansion of each evolutionary fauna is associated with the decline of the previously dominant fauna. The declines are much slower than the initial diversifications, giving the faunas very asymmetrical histories. Such a pattern is difficult to simulate in 'random' models of diversification (Sepkoski 1981) but can be described with coupled logistic equation of the form:

$$dD_i/dt = r_i D_i (1 - D_j/D_i)$$

where D_i is the diversity of the i th evolutionary fauna at time t , r_i is its initial diversification rate, D_j is its maximum or 'equilibrium' diversity, and D_j is the summed diversity of all faunas at time t (Sepkoski 1984; Kitchell & Carr in Valentine 1985). This equation states that an evolutionary fauna only if its initial diversification rates is lower and equilibrium diversity is higher. If r_i is higher, the evolutionary fauna will expand so rapidly that the preceding fauna will never appear to diversify; if D_i is lower, the evolutionary fauna will never be able to expand and replace the preceding one. Thus, the coupled logistic equation suggests a certain inevitability in the sequential diversification of evolutionary faunas, although it does not specify their timing or relative differences in maximum diversity.

Classes within evolutionary faunas tend to have similar mean rates of taxonomic turnover. Classes in the Cambrian Fauna tended to have high turnover rates, those in the Paleozoic Fauna compar

atively low rates (with some exceptions in all cases). These differences are reflected in the responses of the faunas to mass extinctions (Sepkoski 1984): The Cambrian Fauna suffered large proportional reductions in diversity relative to the Palaeozoic fauna during mass extinctions in the Ashgillian and Frasnian, and the Paleozoic Fauna suffered more than the Modern at all major mass extinctions of the Phanerozoic. This differential reaction seems to have led to the great change in faunal dominance associated with the Late Permian mass extinction (section 2.13.4.).

Evolutionary faunas also seem to have differing ecological characteristics. The Cambrian Fauna tended to be assembled into broadly intergrading communities that were dominated by generalized deposit feeders and grazer and had low epifaunal and infaunal tiering (Bottjer & Ausich 1986; see also Section 1.7.1). Communities of the Paleozoic Fauna were dominated by epifaunal suspension feeders with complex tiering; many other ecological guilds were also represented so that the faunal as a whole seems to have occupied more 'space' than the Cambrian Fauna (Bambach in Tevesz & McCall 1983). Finally, the Modern Fauna is represented by yet more guilds and is characterized by large numbers of durophagous predators (Vermeij 1987) and mobile deep infauna (Trayer in Tevesz & McCall 1983); epifaunal tiering is reduced in most communities.

Sepkoski and Miller in Valentine (1985) demonstrated that evolutionary faunas tend to form coherent assemblages within shelf environments throughout the Paleozoic Era. Members of the Cambrian Fauna were spread across the entire shelf early in the Paleozoic Era but became progressively restricted to deeper-water environments during the Ordovician as members of the Paleozoic Fauna expanded across the middle and finally outer shelf. At the same time, early members of the modern Fauna came to dominate inner shelf environments and later, deeper, low-oxygen environments. The Late Permian mass extinction eliminated dominance of the Paleozoic Fauna from middle and outer shelf environments and led to expansion of the Modern Fauna across the entire shelf.

It must be emphasized that none of these evolutionary and ecological differences between the faunas is absolute. In a sense, the faunas are 'fuzzy bounded sets' with their characteristics overlapping and some members of each fauna mimicking members of other. The characteristics thus represent nodes on a continuum. Major unsolved problems are why such nodes should exist and why they seem to change so little through the Phanerozoic.

COMPOSITION AND HISTORY. The individual histories of the marine evolutionary faunas are illustrated in Fig 2. The Cambrian Fauna was dominated by trilobites along with inarticulate brachiopods, on the placophorans, hyoliths, and eocrinoids; most of the problematic taxa of the so-called 'small shelly faunas' of the Tommotian are also included. Various of these classes are paraphyletic, with descendent monophyletic taxa belonging to other evolutionary faunas; however, in most cases the paraphyletic classes either declined long before their descendent taxa diversified (e.g. Monoplacophora) or contained monophyletic subtaxa that diversified in parallel with the rest of the evolutionary fauna (e.g. the inarticulata). The Cambrian Fauna diversified very rapidly from the latest Vendian into the early Cambrian and was the principal constituent of the evolutionary explosion across the Precambrian-Cambrian boundary (see also section 1.5). Its maximum diversity was attained in the late Middle and early late Cambrian. Beginning in the latest Cambrian, the fauna began a long, gradual decline, accentuated by the Ashgillian and Frasnian mass extinctions (section 2.13.2, 2.13.3).

The Paleozoic Fauna initiated its expansion as the Cambrian fauna began to decline; this combination resulted in nearly stable global diversity throughout the late Cambrian. The Paleozoic fauna was dominated by articulate brachiopods with important contributions from crinoids, corals, ostracodes, cephalopods, and stenolaemate bryozoans. These groups were major components of the Ordovician radiations, which tripled global taxonomic diversity over a 50 million year interval. The Paleozoic fauna attained its maximum diversity from the late Ordovician to Devonian and then began decline. During the Carboniferous and Permian, this decline was matched

by a slow expansion of the modern fauna so that again global diversity remained nearly constant. The Paleozoic Fauna was severely reduced by the late Permian mass extinction (section 2.13.4) but in the Mesozoic underwent two radiations: one in the Triassic, terminated by the Norian mass extinction (2.13.5), and a second, slower expansion in the Jurassic. The Jurassic expansion was reversed in the Cretaceous when global diversity exceeded Paleozoic levels, and the remnants of the Paleozoic fauna again went into decline.

The modern fauna is dominated by gastropod and bivalve molluscs, osteichthyan and chondrichthyan fishes, gymnolaemate bryozoans, malacostracans and echinoids. Most of these classes appeared but diversified only slowly through the Paleozoic era. They suffered minor extinction relative to the Paleozoic fauna during the late Permian and became the dominant fauna in the Triassic. Through the Mesozoic and Cenozoic, the modern fauna continued the rather slow and steady diversification initiated earlier, producing the long post Paleozoic increase in global taxonomic diversity.

Throughout their histories, the three great evolutionary faunas experienced considerable internal turnover, with continuous change in ordinal and lower-level taxonomic composition. This was particularly true of the Cambrian fauna, which underwent very rapid changes during its initial radiation. It may prove useful to subdivide this fauna and define two more evolutionary faunas: an Ediacaran fauna, encompassing the distinctive soft-bodied animals fossil of the Vendian (section 1.3.1.5), and a Tommotian fauna, comprising the mostly problematical skeletal taxa of the earliest Cambrian (sections 1.4, 1.5, 5.2.5). These possible faunas seem to fit into the general progression of evolutionary rates and diversity levels observed for the three great evolutionary faunas. The Ediacaran and especially Tommotian taxa appear to have had higher diversification rates and more rapid evolutionary turnover than the remainder of the Cambrian Fauna, and seem to show successive increases in diversity leading into the Cambrian period. Further analysis of diversity pattern and faunal change in the Vendian and early Cambrian are needed to assess whether such additional

evolutionary faunas are useful for describing the early metazoan radiation.

TERRESTRIAL BIOTAS.

The concept of evolutionary faunas has proved useful for organising faunal turnover and diversity change in the marine record and has been extended with varying success to other evolutionary systems, specifically terrestrial vascular plants and tetrapod vertebrates. Niklas et al (1983) identified four major plant groups, which can be termed evolutionary floras, in species-level data on tracheophyte diversity (fig 1B) there are: (1) an initial Silurian-Devonian flora of early vascular plants that radiated and then disappeared during the Devonian; (2) a Pteridophyte dominated flora, including ferns, lycopods, sphenopsids, and progymnosperms, that diversified in the Late Devonian and early Carboniferous and dominated plant communities to the end of the Paleozoic era; (3) a gymnosperm-dominated flora of seed plants that appeared in the late Devonian and rose to dominance in the Mesozoic; and (4) an angiosperm flora that originated in the early Cretaceous and rapidly radiated to dominance thereafter, replacing the preceding gymnosperm flora. As in the marine system, each of these floras (excepting the angiosperms) originated early in the history of vascular plants and radiated sequentially to produce step-like increases in global tracheophyte diversity.

Three assemblages of terrestrial tetrapod families have been identified by Benton (1985) in the vertebrate fossil record (fig 1C). These comprise: (1) the labyrinthodonts, anapsids, and synapsids, which appeared during the middle Paleozoic and completely dominated the terrestrial vertebrate record to the end of the Paleozoic; (2) the early diapsid, dinosaurs, and pterosaurs, which arose in the Triassic, attained maximum diversity in the late Jurassic and Cretaceous, and disappeared at the terminal Cretaceous mass extinction (section 2.13.7); and (3) the lissamphibians, turtles, crocodiles, lizards, birds and mammals, which originated in the Triassic and Jurassic, expanded through the Cretaceous, and then diversified to very high levels in the Cenozoic. Although these assemblages have

some similarities to marine evolutionary faunas, there are some important differences: the assemblages do not all appear early in the history of tetrapods and their sequential diversifications are not all associated with step-like increases in global diversity. It remains to be seen whether such patterns could be identified if more terrestrial taxa (e.g. the arthropods) were included and analyses performed at lower taxonomic levels. If so, evolutionary faunas and floras would appear to be a general proprietary of the development of Phanerozoic biotas.

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Fig 1. Diversity curves, A, marine animal families. B terrestrial vascular plant species; C terrestrial tetrapod families. Each curve is divided into fields that illustrate the diversities of the constituent evolutionary faunas and floras, A, After Sepkoski (1984); Cm = cambrian evolutionary fauna, Pz = Paleozoic fauna, Md = modern fauna; stippled field represents known diversity of families with rarely preserved members that lack heavily mineralized skeletons, B, After Niklas et al (1983); numbered fields as in text. C. After Benton (1985) numbered fields as in text.

Fig 2. histories of the three great evolutionary faunas of the marine fossil records as represented by their familial diversities through the Phanerozoic Representatives of the important classes in each fauna are illustrated above the diversity curves (After sepkoski 1984).

Figure 1

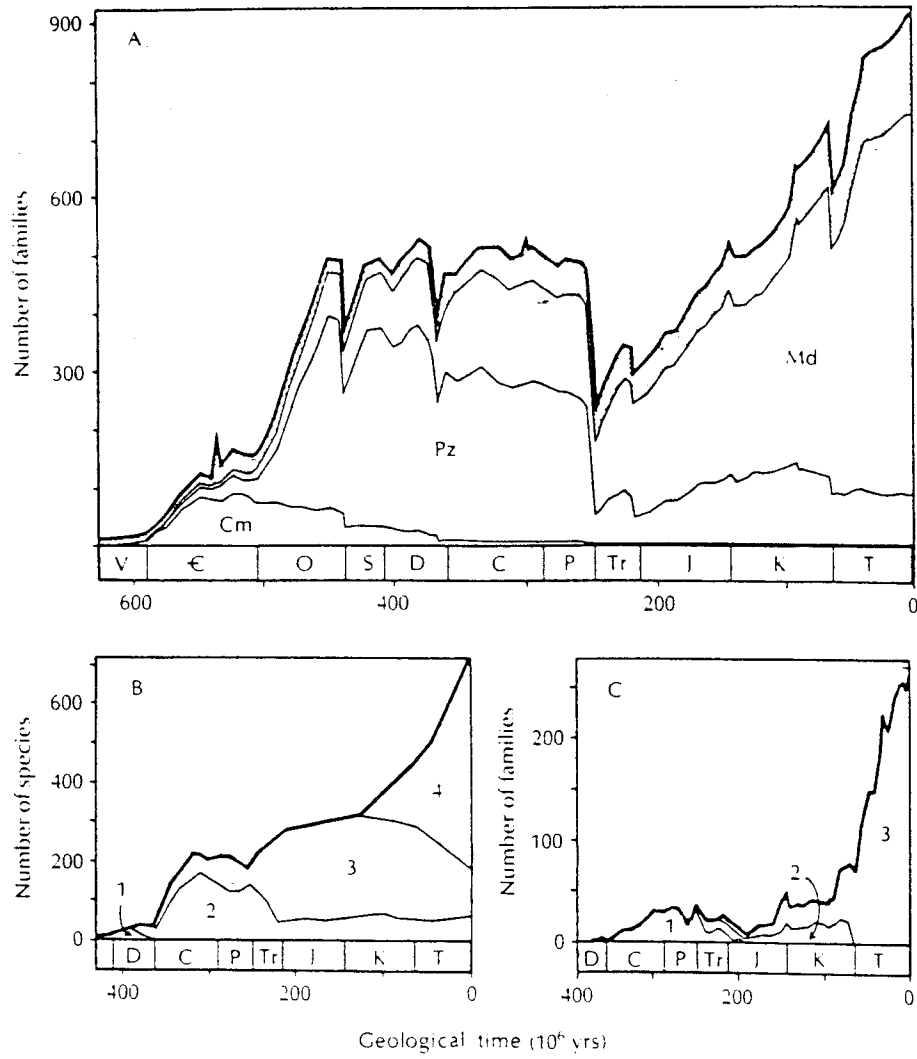
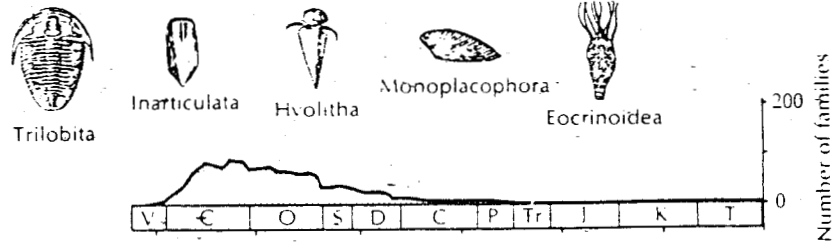
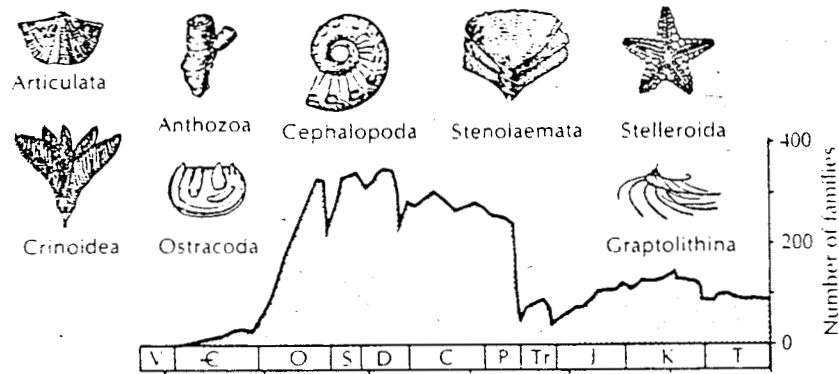


Figure 2

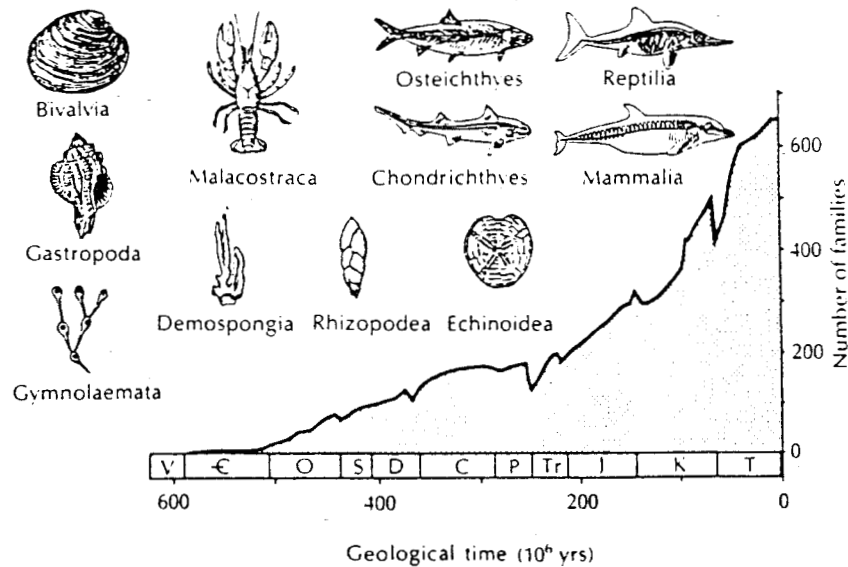
CAMBRIAN FAUNA



PALAEOZOIC FAUNA



MODERN FAUNA



In: Valentine, J.W. Edit., 1985 Phanerozoic diversity patterns. profiles in macroevolution: Princeton, N.J.: & San Feo., Cal Princeton univ. Press, Chapt 1, p 11-39.

10.4 BIODIVERSIFICACION.

10.4 (A)

An Atlas of Phanerozoic Clade Diversity Diagrams.

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Clade diversity diagrams are spindle-shaped graphs that summarize patterns of taxonomic evolution within higher taxa through geologic time. Most clade diversity diagrams are constructed about a central axis that represents time (scaled either metrically or ordinally, by stratigraphic interval). Some measure or estimate of taxonomic diversity (or richness) is then plotted symmetrically about the axis to give the diagram an overall spindle shape (e.g., Fig 1).

Diversity diagrams for individual clades convey information about their size, shape and variability in the fossil record (cf. Gould et al, 1977). Such morphologic information is valuable for assessing how evolutionary rates (that is, rates of origination and extinction) vary within the taxa through geologic time. Clade diversity diagrams for groups of higher taxa hypothesized to be related by phylogeny or by function are useful for comparisons of the histories of the taxa. Common pattern of expansion or contraction may relate to general factors governing all taxa, whereas reciprocal patterns may be interpretable as negative interactions between pairs of ecologically similar taxa (e.g. Simpson, 1953; Banbach, this volume). Sets of clade diversity diagrams also are useful for summarizing variation among large numbers of clades for the purpose of testing general macroevolutionary models (e.g., Raup et al, 1973; Gould et al., 1977).

This chapter presents a collection of clade diversity diagrams which we hope will be useful

for examining the general histories of a wide variety of animal taxa. The main body of the chapter is a series of 12 figures displaying spindle diagrams for orders, classes, and phyla of both marine and nonmarine (or continental) animals for the whole of the phanerozoic (including the Vendian). Nearly all of the diagrams are plotted at a uniform taxonomic and temporal resolution, specifically that of familial diversity per stratigraphic stage. The taxonomic rank of family is used simply because comprehensive data with good stratigraphic resolution can be obtained for all animal groups at this level. Although families do not display all of the detail of the fossil record they should be sufficiently sensitive to show major evolutionary trends and patterns with characteristic timescales of five to tens of million years (see also Sepkoski, 1979, 1982a; Raup and Sepkoski 1982).

The clade diversity diagrams in most of the figures are formatted in strips that have time in the vertical dimension. Most of the strips are scaled from 625 myr at their bottoms to approximately 1 myr BP at their tops. (No data on Recent diversity are directly included in the diagrams). Geologic eras and systems are indicated at the left-hand ends of the strips, with eras denoted by Cz = Cenozoic, Mz Mesozoic, Pz Paleozoic and pe = late Precambrian; systems are denoted by standard symbols, with V = Vendian. The widths of the clade diversity diagrams in each strip indicate the numbers of families known from direct fossil evidence or from interpolation between known occurrences to be present in the clades in each of 80 stratigraphic stages or comparable intervals (see Table 1 in Sepkoski, 1982b for a listing of the stages used). A scale for the familial diversities appears in the lower right-hand part of most of the figures. All of the diagrams were produced with an IBM personal computer and Epson dot-matrix printer.

The first two figures in this chapter contain class-level summaries of the entire Phanerozoic fossil record. Figure 1 displays clade diversity diagrams for the 87 classes and 15 unique, problematic genera that have representatives in the marine fossil record. This illustration is an updated version of figure 1 in Sepkoski (1981) with corrections based on new data in Sepkoski

(1982b). The second figure in this chapter summarizes the continental fossil record. The diversity diagrams displays numbers of freshwater and terrestrial families within the 39 animal classes known from the nonmarine fossil record; data on the classes were compiled from the literature sources listed in table 1. Also shown at the bottom of figures 2 are clade diversity diagrams for numbers of species within the 13 taxonomic divisions of the tracheophytes and bryophytes; the data for these diagrams were taken from Niklas, Tiffney and Knoll (this volume), same relative scale as in fig 1 and 2 in order to facilitate comparison. Figures 3 and 4 display family-level clade diagrams for orders within the moderately diverse marine phyla: the protozoa, porifera, coelenterata, bryozoa, brachiopoda, annelida and hemichordata. (the set of clade diversity diagrams for the Annelida includes several taxa of questionable affinities which might best be considered incertae sedis; these are in the group of diagramss beginning with Cribriocyathida and ending with Courculitidae) The more diverse marine phyla are represented in Figures 5 to 8. Figure 5 displays orders of marine molluscs. Figure 6 orders marine arthropods; Figure 7 orders of echinoderms; and Figure 8 orders of marine vertebrates.

The two large phyla of continental animals, the nonmarine Arthropoda and chordata, are featured in fig 9 and 10. Nonmarine taxa have been segregated from their marine relatives because we believe that the land and sea are best treated as separate major arenas of evolution (see also Boucot, 1983). Despite the fact that some continental clades contain secondary species which alternate between marine and nonmarine habitats and that all clades ultimately had their origins in the ocean, the great majoritariness of continental animals evolved in situ, isolated from evolutionary activity in the seas. Thus, the segregations of marine and continental taxa enhances assessment of evolutionary patterns within axes for the continental clade diversity diagrams in figures 9 and 10 have been truncated below the Silurian; this is because there is virtually no nonmarine fossil record prior to the mid Paleozoic (see Boucot and Janis, 1983).

The final pair of figures in this chapter (fig 11 and 12) contains 14 diversity diagrams for families within entire phyla, split between marine and con-

tinental. These diagrams are formatted somewhat differently than in the preceding figures. The splindles have been cut in half and rotated so that the time axis runs horizontally. This arrangement permits easier assessment of the times and magnitudes of diversity change but impedes comparisons of changes between groups.

The use of a single level of taxonomic and stratigraphic resolution in all clade diversity diagrams is intended to aid interpretation and comparison of patterns among the various taxa. However, the consistency of resolution does not imply a uniformity of quality throughout the data. The accuracy of the taxonomic and stratigraphic information varies considerably among the taxonomic and stratigraphic information varies considerably among the taxonomic groups. In general, the quality is much better for marine taxa than for nonmarine taxa. Also, as should be expected, the fossil data are much better (and much more complete) for heavily skeletonized animals than for soft-bodied and lightly sclerotized animals. In fact, many of the diagrams for the latter groups reflected little more than the geologic distribution of Lagerstätten that preserve unusual fossils. This is particularly evident in the long, thin clade diagrams for such extant groups as the Nemertinea and Priapulida (figure 1); these diagrams show only the extension of stratigraphic ranges from the recent to the one or more Lagerstätten that happen to contain the groups' early members.

Much of the characters of the diversity diagrams for some large clades, such as insect (figure 2 and 9), also represents the effects of Lagerstätten. For the insects, the more important Lagerstätten include the upper Carboniferous siderite concretion deposits of North America and Europe, the mid-permian lake deposits of Kansas and Kazakhstan, the Eocene Green River deposits of Wyoming, and especially the Oligocene Baltic Amber of northern Europe. The Baltic Amber alone contributes most of the Cenozoic bulge in the clade diversity diagrams for both insects and other lightly sclerotized terrestrial arthropods (figures 2, 9, and 12). The effects of Lagerstätten, or of their non-occurrence, are even seen in some well-skeletonized groups with fairly extensive fossil records. The drop in the diversity of continen-

tain vertebrates in the Jurassic (fig 12), for example; probably reflects largely a paucity of fossiliferous continental deposits between the Raethian and Tithonian (see also Carroll, 1977; Padian and Clemens, 1985, this volume).

These shortcomings of the fossil record, along with the problems associated with family-level data and 5 to 10 myr-long stages, do limit the value of the clade diversity diagrams presented here. However, we believe that a great deal still can be learned from them about the apparent order, or disorder, in their radiations and extinctions. Thus we hope that this atlas will aid in the assessment and interpretation of evolutionary history as well as serve as a baseline for the compilation of more accurate and detailed diversity data.

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Fig. 1. THE MARINE FOSSIL RECORD
Families within classes of fossil marine mammals.

Fig. 12. Familial diversity of selected marine and
continental animal phyla.

TABLE 1. principal literature sources of information on the taxonomy and stratigraphy of continental animal families.

Fig. 2. THE CONTINENTAL FOSSIL RECORD. Families within classes of continental (i.e. terrestrial and freshwater) fossil animal and species within divisions of continental plants.

Fig. 3. Families within orders of fossil Protozoa, Porifera and Coelenterata.

Fig. 4. Families within orders of fossil Bryozoa, Brachiopoda, Annelida, and Hemichordata.

Fig. 5. Families within orders fossil marine Mollusca.

Fig. 6. Families within orders of fossil marine Arthropoda.

Fig. 7. Families within orders of fossil Echinodermata.

Fig. 8. Families within orders of marine Vertebrata.

Fig. 9. Families within orders of fossil nonmarine Arthropoda.

Fig. 10. Families within orders of fossil nonmarine vertebrata.

Fig. 11. Familial diversity of selected marine animals phyla.

FIGURA 1

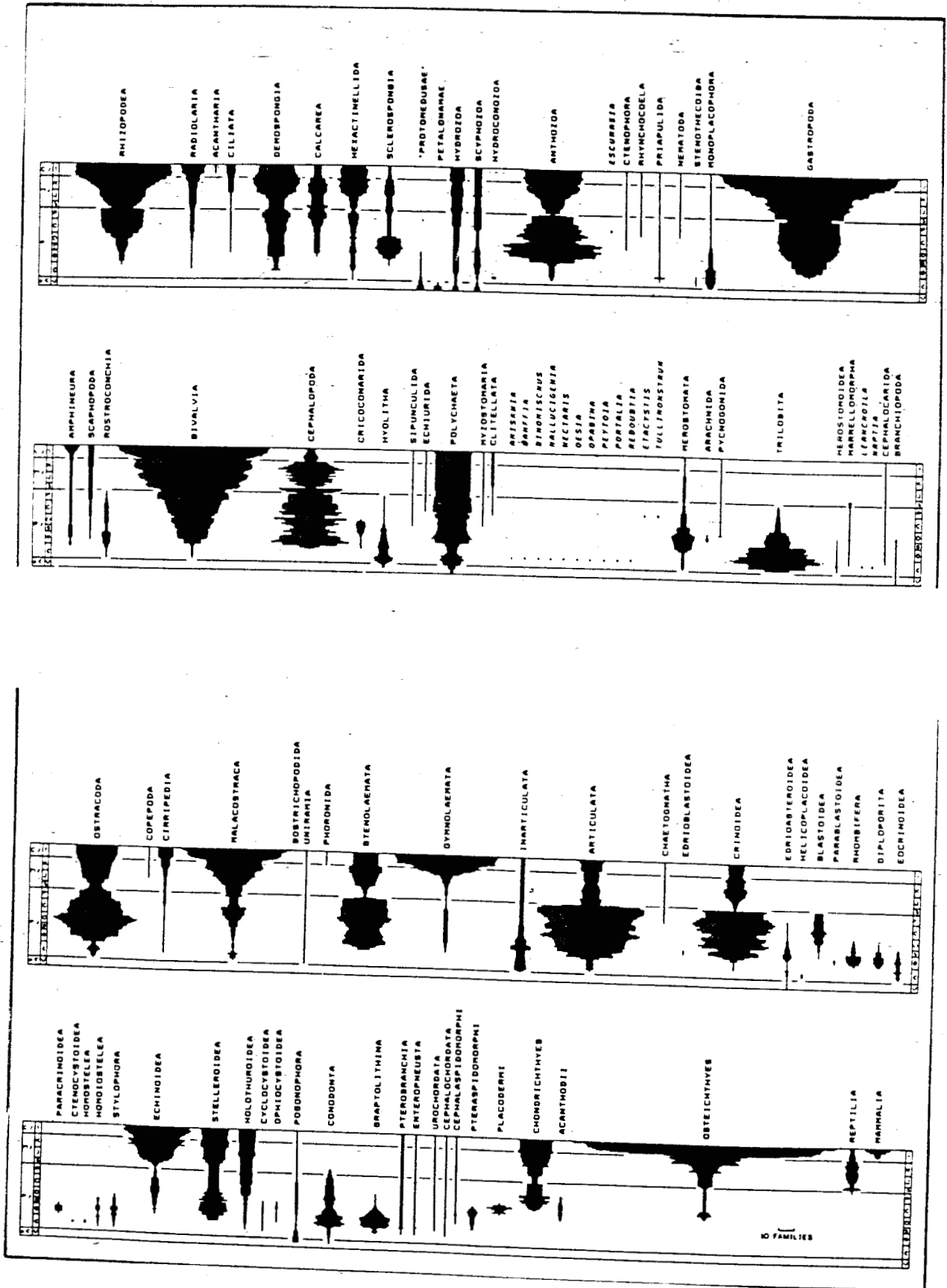
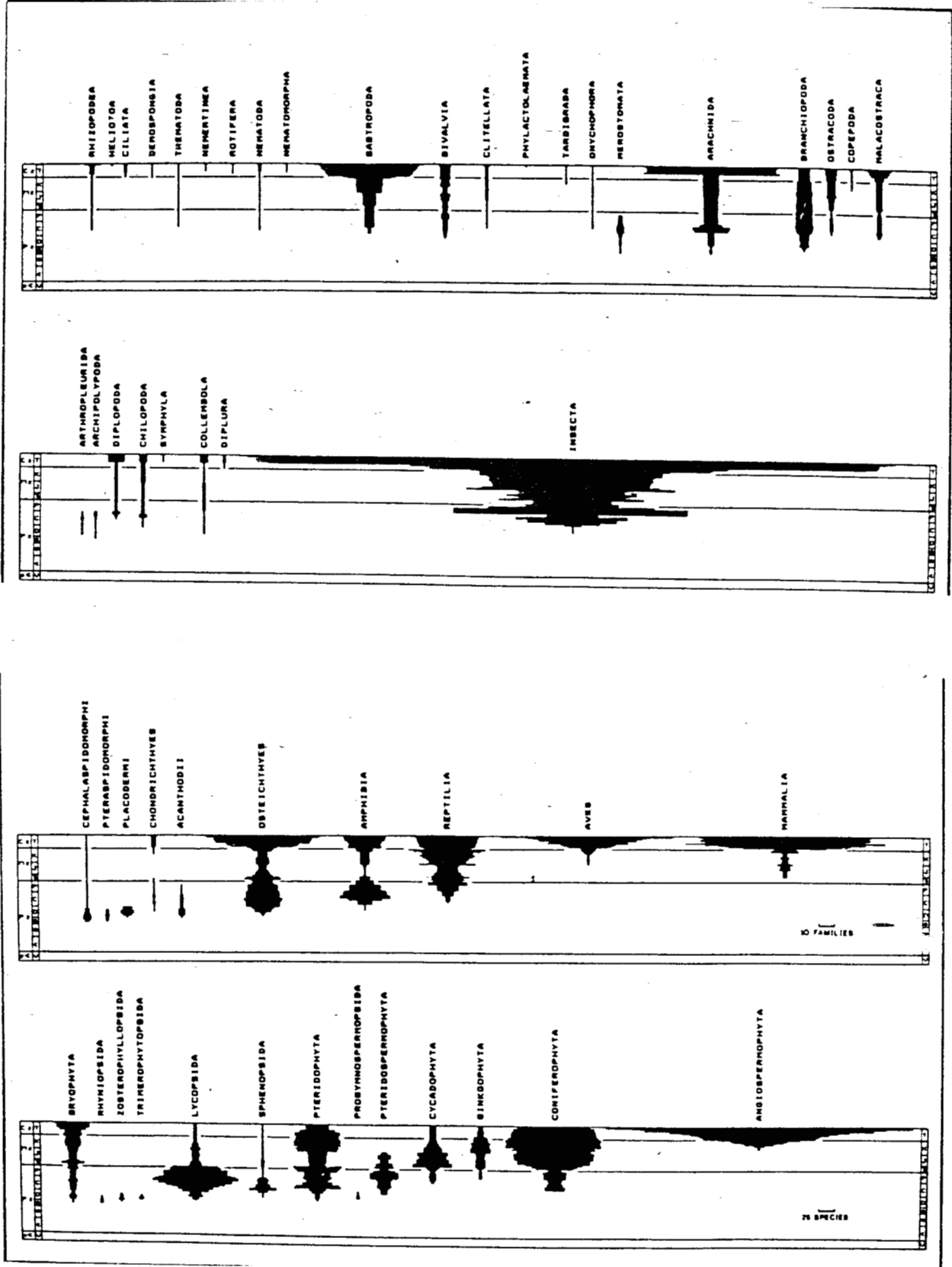


FIGURA 2



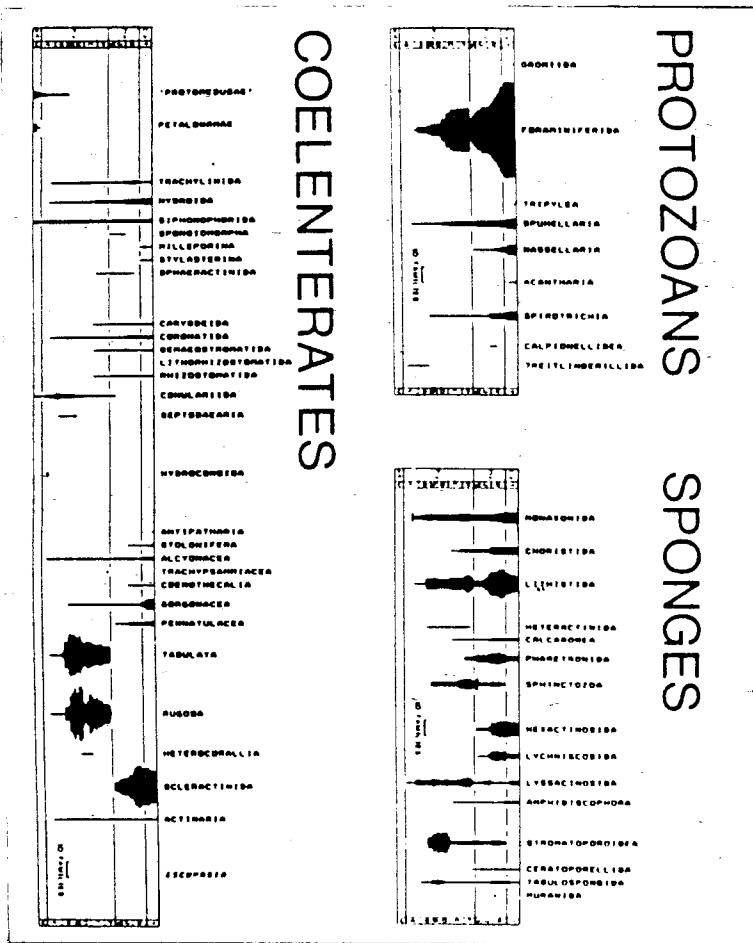


FIGURA 4

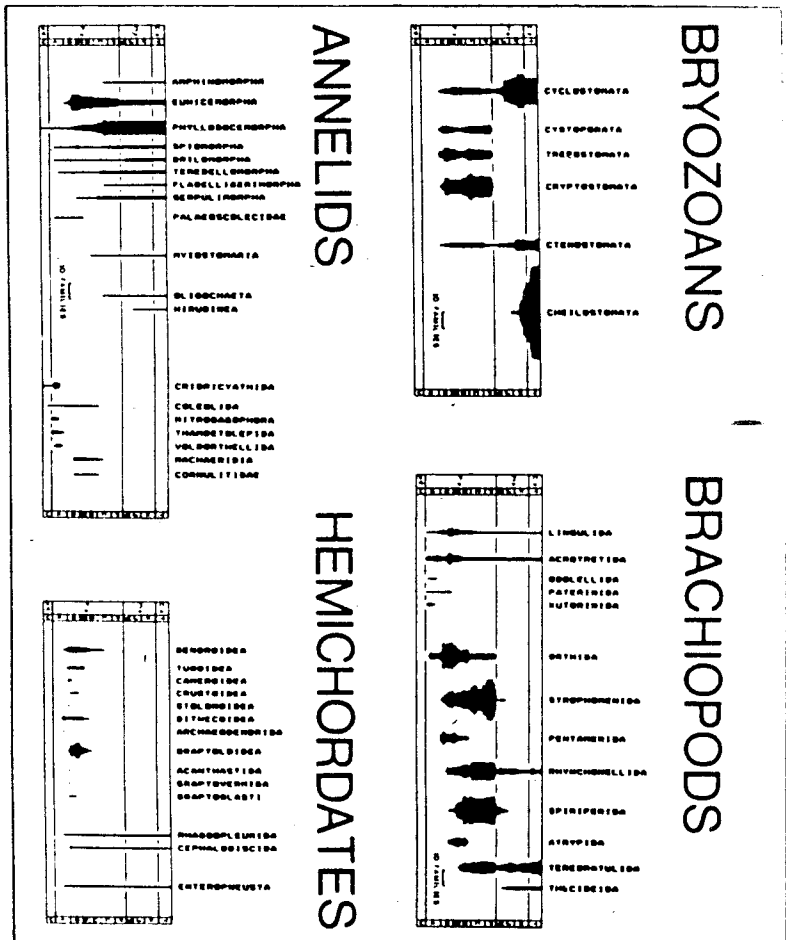


FIGURE 5

MARINE MOLLUSCS

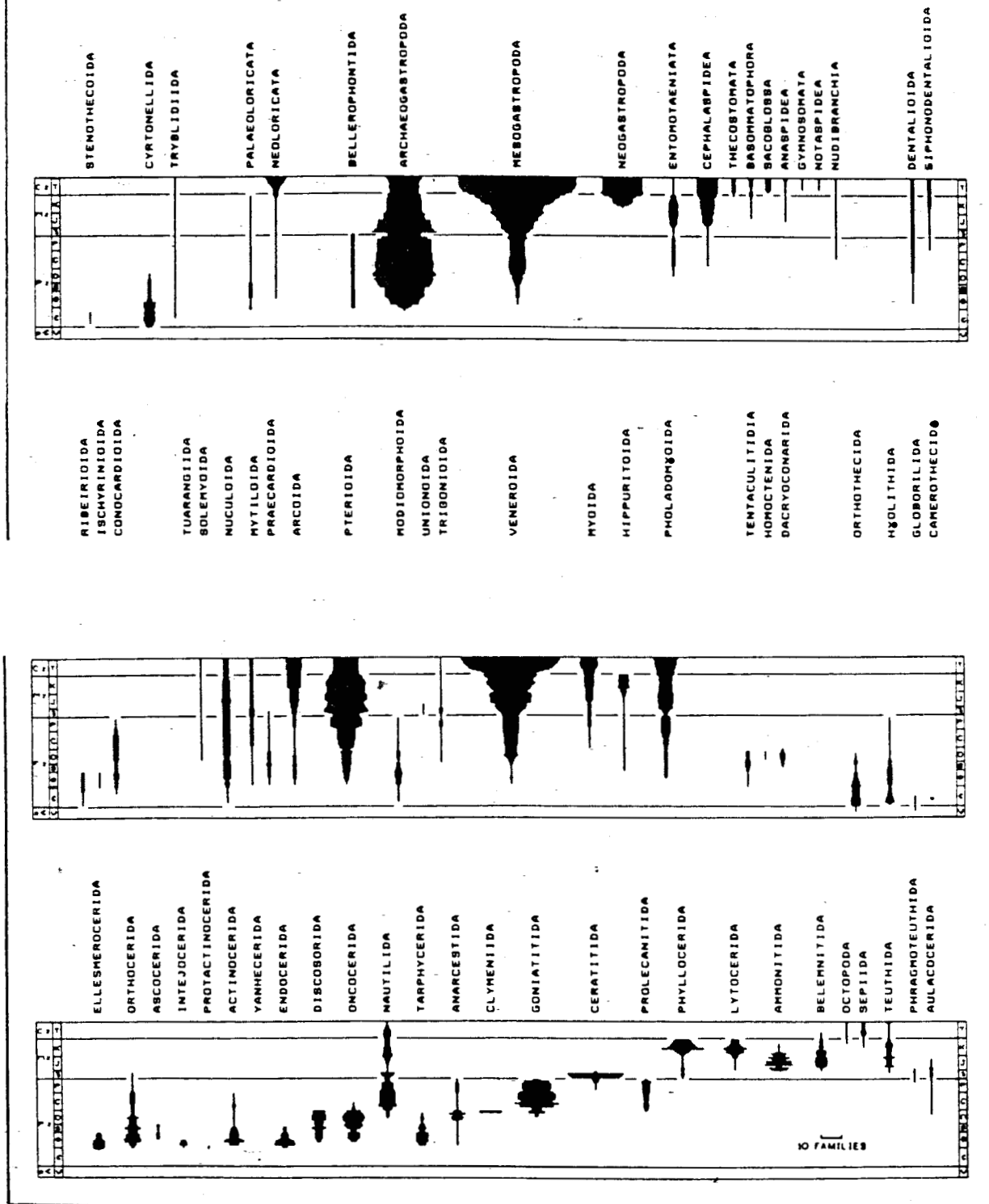


FIGURE 6

MARINE ARTHROPODS

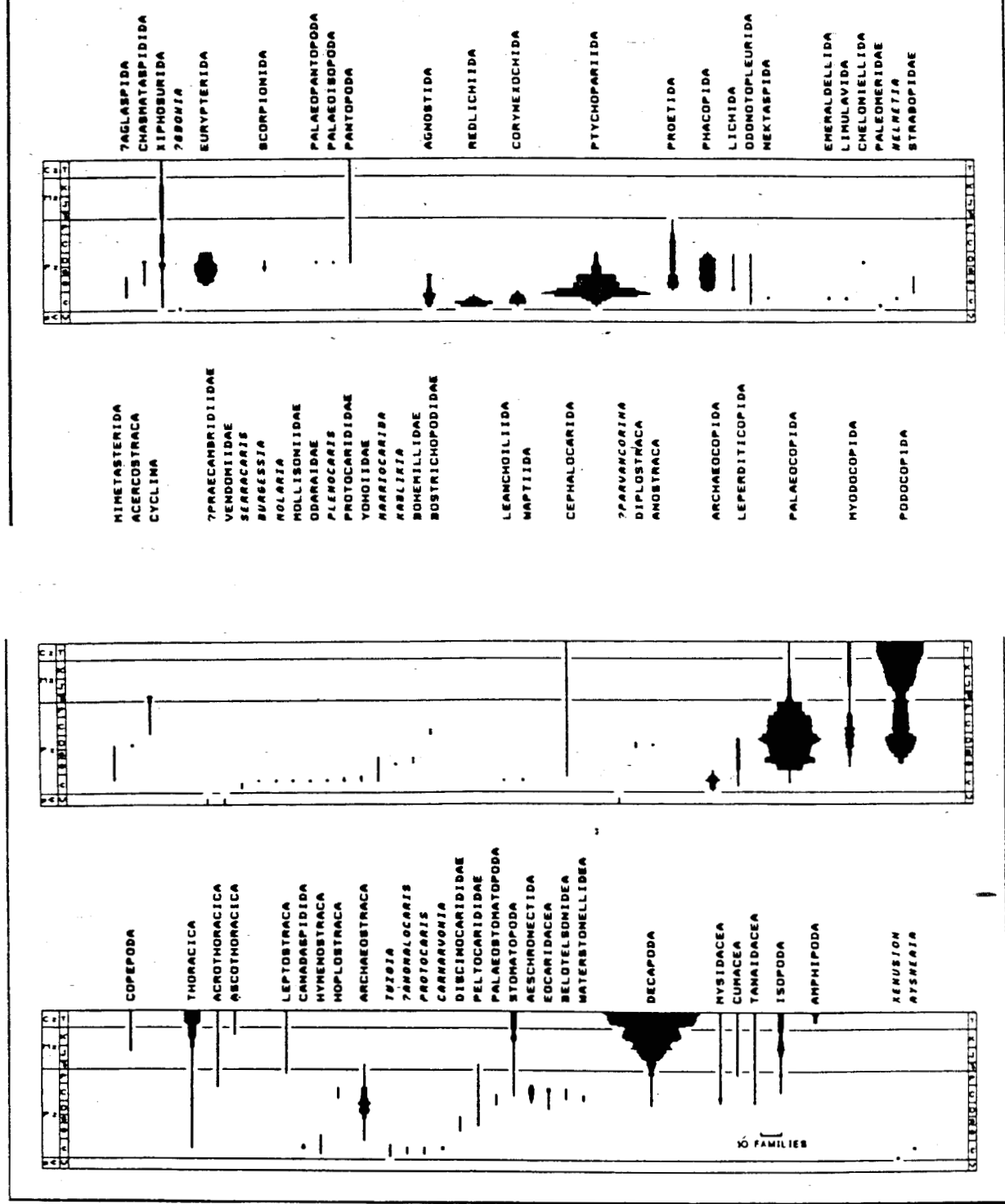


FIGURA 7

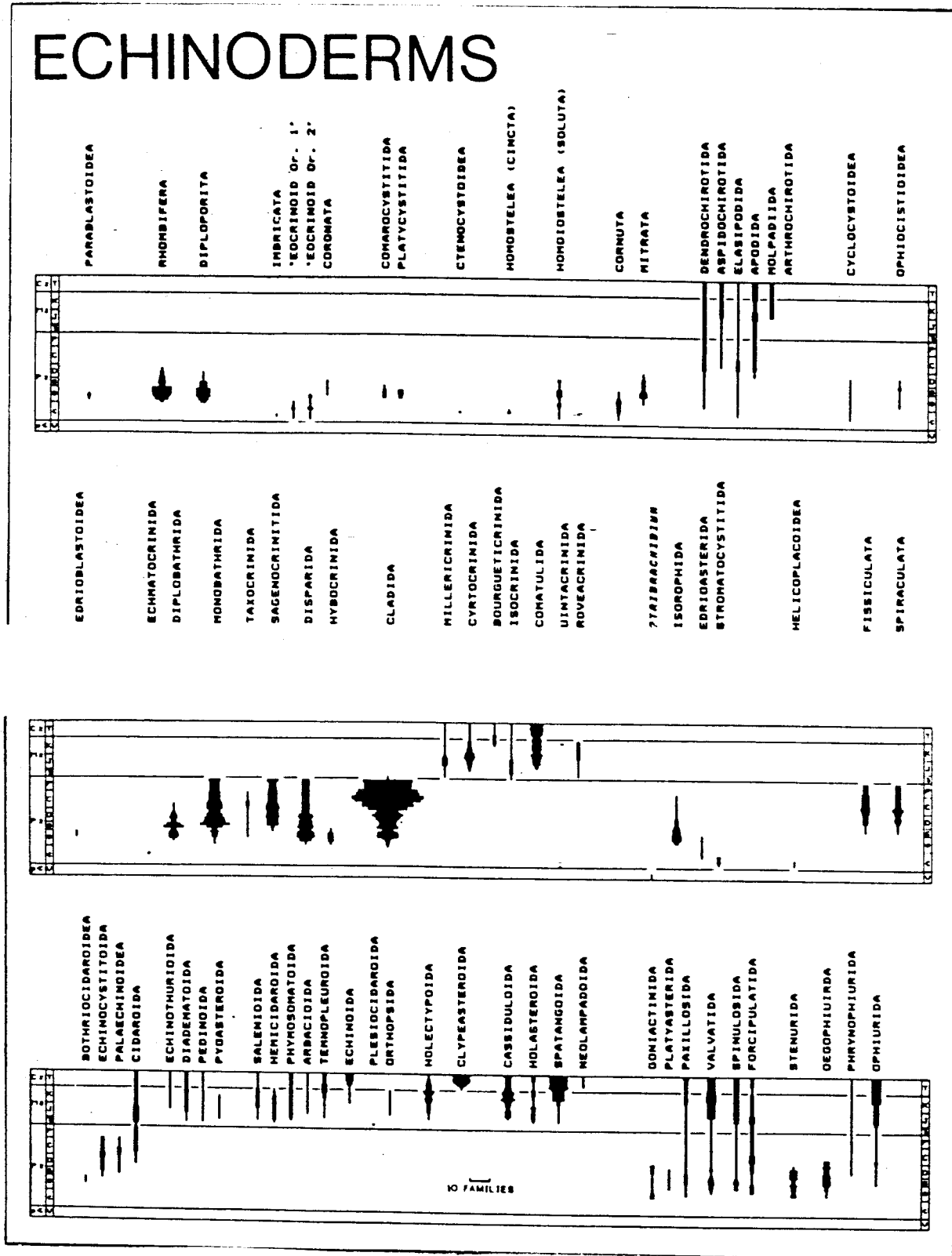
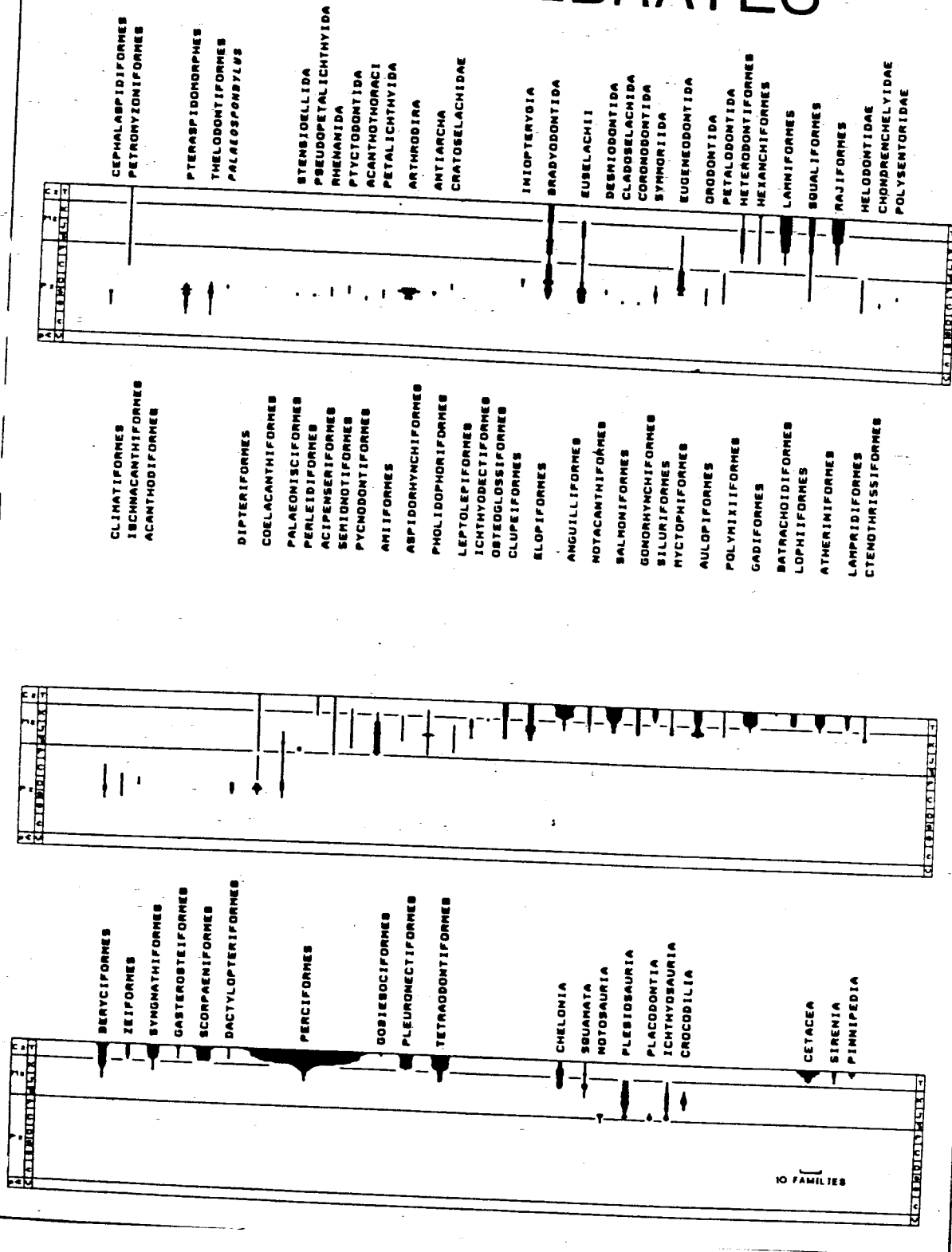


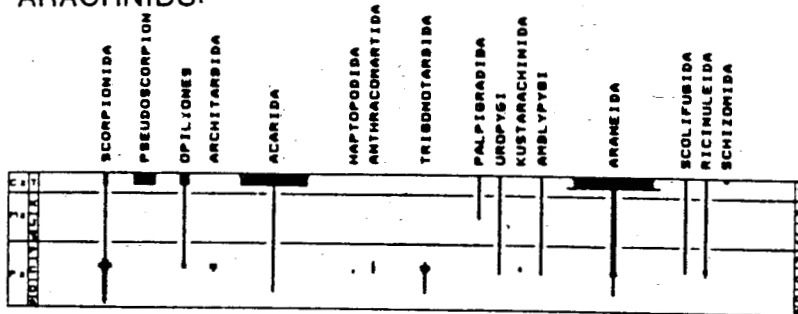
FIGURA 6

MARINE VERTEBRATES

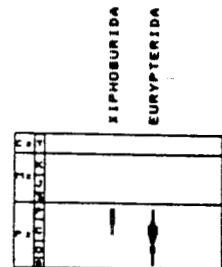


CONTINENTAL ARTHROPODS

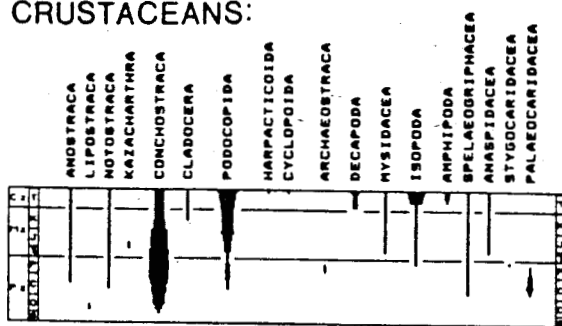
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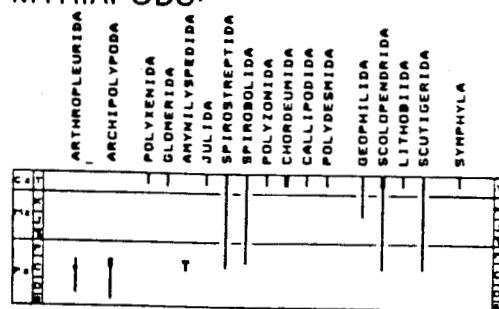
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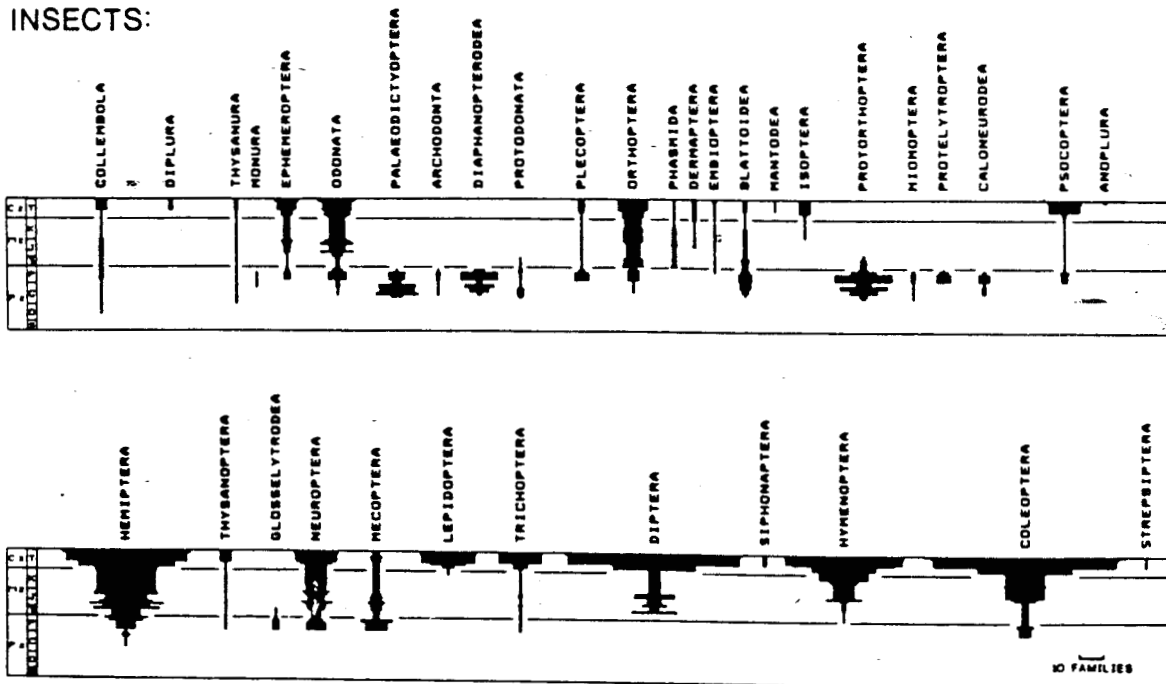
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MYRIAPODS:



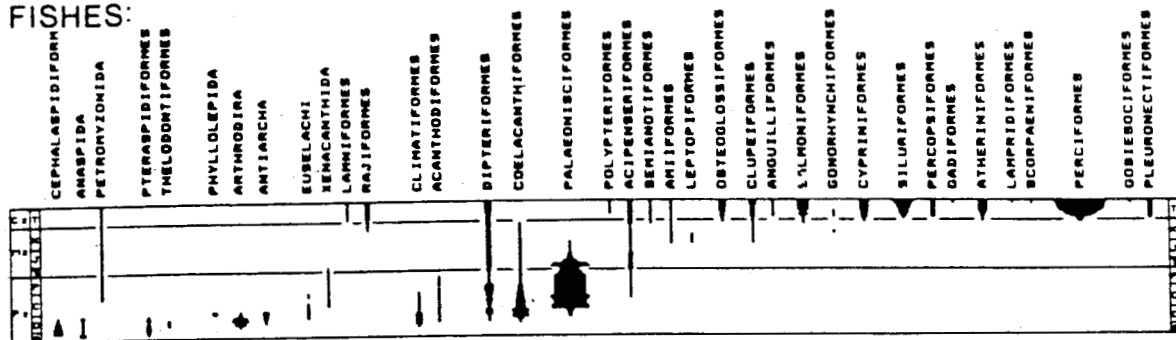
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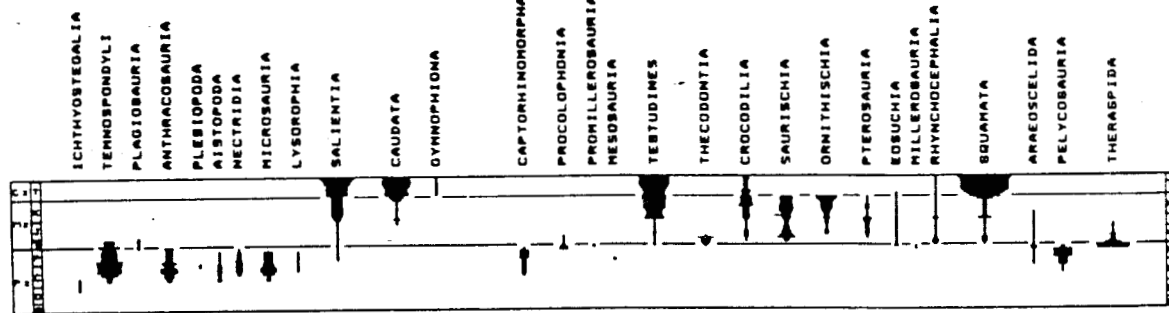
PIWUL 10

CONTINENTAL VERTEBRATES

FISHES:

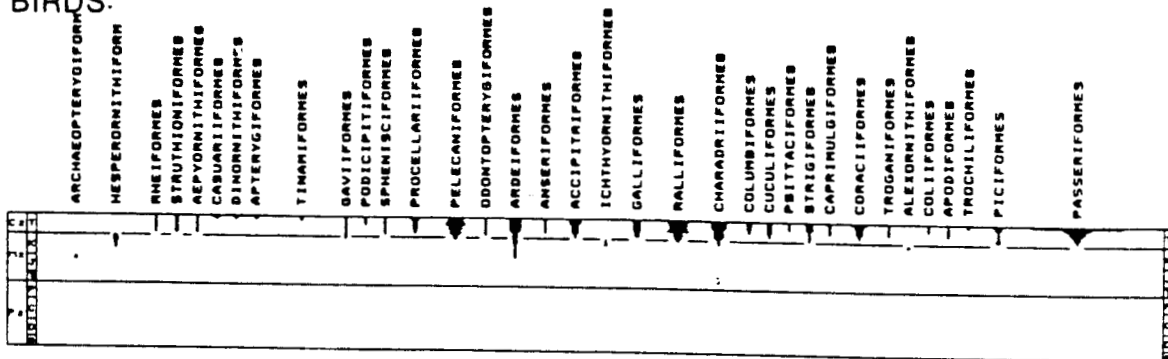


AMPHIBIANS:



REPTILES:

BIRDS:



MAMMALS:

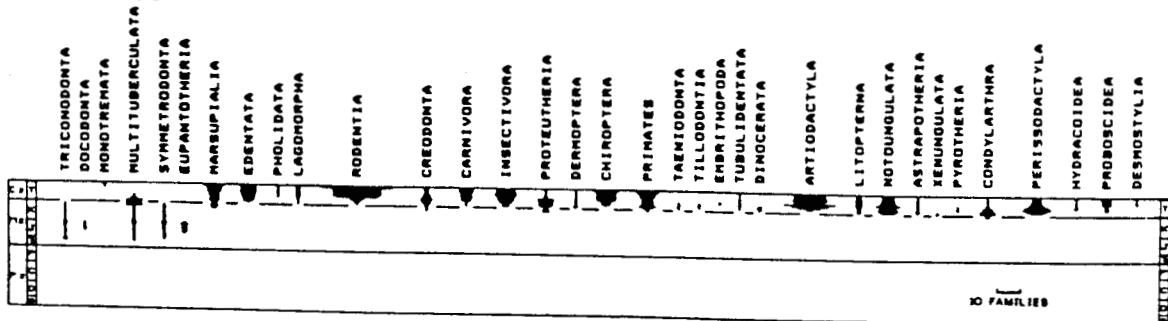


FIGURE 11

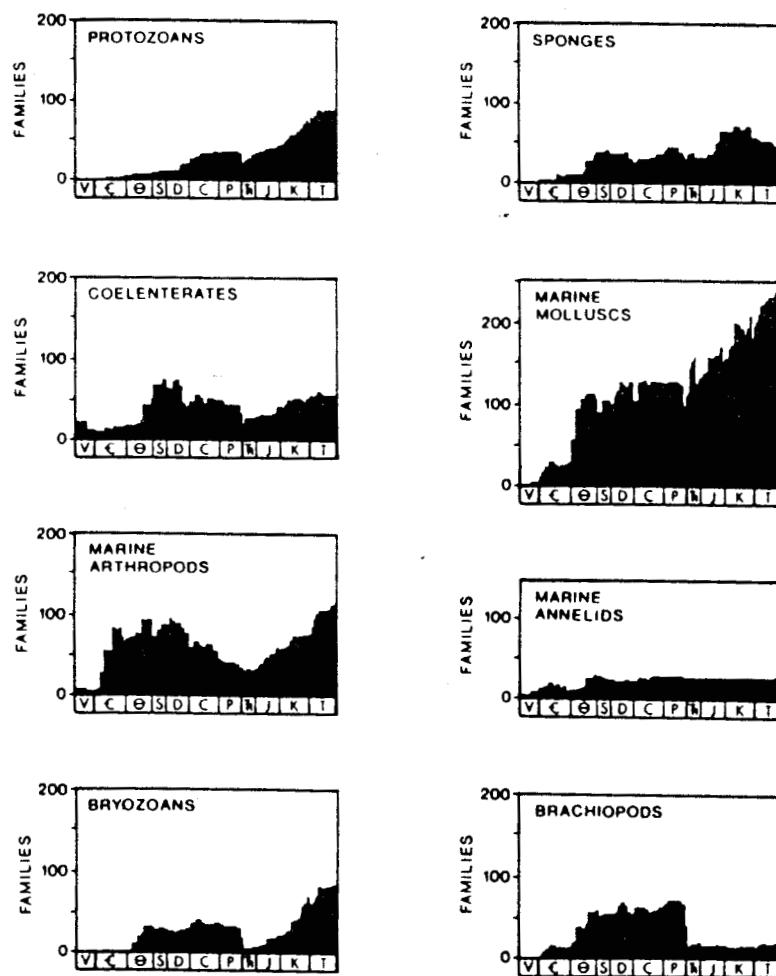
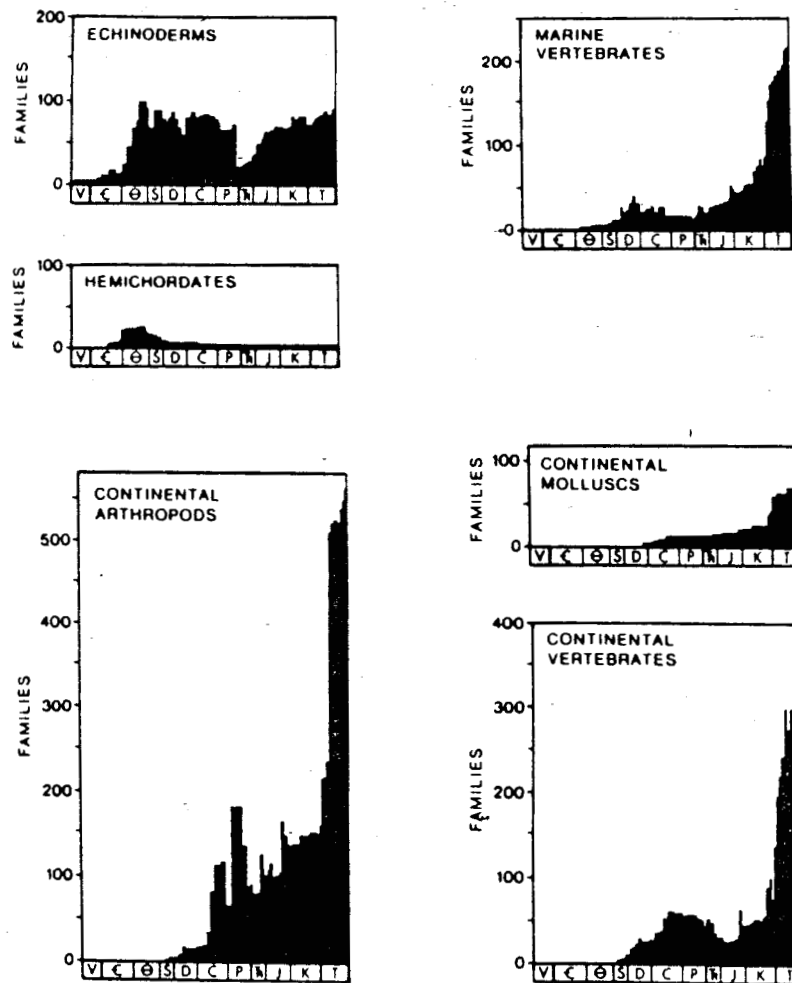


FIGURE 12



10.5 EXTINCTION.

10.5(a)

CHANGING PALEONTOLOGICAL VIEWS ON MASS EXTINCTION PHENOMENA.

Antoni Hoffman

In: DONOVA, S.K., EDIT., 1989. MASS EXTINCTIONS: PROCESSES AND EVIDENCE: NEW YORK, COLUMBIA UNIV. PRESS, CHAPT. 1, P 1-18.

A HISTORICAL SKETCH.

Even a quick perusal of the Nature and science magazine over the last decade or two early indicates that problem of mass extinctions- that is to say, the search for an adequate causal explanation of the disappearance of a large number of fossil groups within reatively short periods periods of geological time at such stratigraphic intervals as the Permian-Triassic, Cretaceous-Tertiary, Ordovician-Silurian, and Frasnian-Famennian transition-has only recently become an important topic in the earth and life sciences. The history of research in this area is nenetheless quite long and rather complex.

The ancient Greek concepts of plenitude, or fullness of the natural world, implies that no organisms that ever existed on the Earth could ultimately disappear fom its surface, because their final extinction would leave an unbridgeable gap in the Great Chain of Being. Species extinction, therefore, was for long considered impossible. Eighteenth-century naturalist kenw, of course, that a wide variety of fossil had no counterparts among living organisms, but this apparent anomaly was commoly explained by an as yet inadequate knowledge of life on the Earth. It ap peared perfectly reasonable to asume at the time that am-

monites or trilobites could still be discovered alive somewhere in tha world ocean, as Neopilina molluscs and Latimeria fishes indeed have been in the twentieth century. It is only after Cuvier (1799) had firsrt described fossil proboscideans that species extinction was eastablished as fact, for it was hard to believe that such large and prominent mammals could roam the Earth with out being ever observed by travellers. Yet tha concept of plkeni tude could still be reconciled with the fact of species extinc tion by assuming-as in fact Lamarck (1809) did man is the sole agent responsible for species extinction. According to Lamarck, species are essentially inmortal because they are alway able to adapt perfectly to any environmental configuration and man alone is capable of violationg the order of the natural world. Under this assumption is was fully understable that the species that underwent extinction were such large animals as the woolly mam moth, which could be easily considered as a valuable prey for human hunters.

With rapidly grwing knowledge of the inventory of living animal species, however, and also with rapidly increasing awareness that fossils such as ammonites or trilobites, without known living counterparts, were by no means oddities but rather very common phenomena, it become undeniable in the early nineteenth century that species extinction did indeed often take place inthe geolog ical past. Within a few decades three main causal explanations for this phenomenon were put forth. Brocchi (1814) proposed that each species was created with a specified, predetermined life- span, in close analogy to individual longevity. He believed that the extinction of species simply marks the end of their life time and is therefore essentially independent of extrinsic, environ mental factors-just as death will inevitably and the life of each person, regardless of fatal accidents or disease. Lyell (1832), in turn, regarded species extinction as a fully natural process caused primarily by the unescapable effects of continuosly chang ing environmental conditions. In his view, each species must sooner or latter encounter such a hostile configuration of environmental factors that it will become extinct, simply because its individuals will be unable to

cope successfully with this new environment context.

Both these concepts implied that species extinction should generally occur independent of one another, without a clear-cut, orderly pattern in time. Cuvier (1825) observed, however, that fossil species in the Paris Basin disappear from the record in large clusters at certain horizons-all or at least a great many of them, at once, as if wiped out by a single catastrophic agent at each horizon. He therefore interpreted species extinctions as due to local catastrophes which destroyed life entirely in an area, thus vacating the ecological space necessary for the area's repopulation by immigrants that survived the cataclysm elsewhere.

Cuvier's perspective on species extinction was developed into a more extreme position by Buckland (1823), who regarded the faunal breaks such as those described by Cuvier as a clear sign of world-wide catastrophes, and particularly by D'Orbigny (1852), who established a whole times series of such global holocaust as providing the main reference points for stratigraphy and the history of life. This emphasis on environmental catastrophes as the main causes of species extinction could, of course, be contested or even rejected outright, but even the most devoted followers of the Lyellian gradualism had to agree that some major events did indeed occur in life's physical environment on the Earth and must have had some impact on the fate of organic species. Darwin (1846), for example, noted that many large terrestrial mammals had lived until recently in South America and he attributed their extinction to the effects of rapid climatic changes during the ice age.

In the *Origin of species* Darwin (1859) wrote very little of species extinction which he regarded, very much in the same vein as Lyell, as due to various environmental factors which drive species first to rarity and then to extinction. He was inclined to emphasize the role of biotic factors, foremost among them interspecific competition, much more strongly than Lyell did, but he was also explicit in his scepticism about our insight into the nature of extinction of any particular species. In spite of his gradualistic prejudice. Darwin was ready to admit that some extinctions of large

taxonomic groups-for instance, the trilobites at the end of the Paleozoic and the ammonites at the close of the Mesozoic-were wonderfully sudden. Contrary to the Cuvierian tradition, however, which viewed these phenomena as an incontrovertible indication of mass extinction events when many organic groups had been exterminated simultaneously, Darwin suggested that their apparent rapidity might really be only artefactual, due to world-wide gaps in the fossil record which clumped together events that had actually been quite widely stretched over geological time.

This view was generally shared by the most orthodox Darwinians among paleontologists-for example, Neumayr (1889), Andrusov (1891) and Davitashvili (1969)-who conceived of species extinction as caused primarily by interspecific competition, various physical factors, and regional environmental catastrophes, whereas they regarded the so-called mass extinctions as nothing but artefacts of the fossil record. In turn, the follower of Brocchi-for example, Beurlen (1933), Zunini (1933) and especially Schindewolf (1950)-tended to interpret the extinction of individual species in terms of their specific life cycle. Though they certainly did not deny the occurrence of regional environmental catastrophes and their implications for species extinction, they largely agreed with the Darwinians that mass extinction phenomena are wildly exaggerated by the notorious imperfection of the fossil record.

The idea of the specific life cycle was later rejected by modern evolutionary biology. The interpretation of mass extinctions-in particular the Paleozoic-Mesozoic and Mesozoic-Cenozoic ones-as possible artefacts of gaps in the fossil record, however, has become one of the two rival perspectives on these phenomena. The other has, of course, envisaged mass extinctions as caused by real environmental catastrophes and it has been based on a more literal reading of the fossil record. The contrast between these two interpretations has largely shaped the history of debate about mass extinctions, but it has always hinged on divergent views on the precision and reliability of stratigraphic correlation among very distant areas.

For so long as the fossil record was only poorly known. Darwin's gradualist perspective appealed to the majority of geologists and palaeontologist. The subsequent progress in stratigraphy, however, has led in the twentieth century to the idea of mass extinction processes operating over the entire Earth instead of being confined to single geographical regions, Marshall (1928) and Henning (1932), for example, regarded the Cretaceous-Tertiary boundary extinctions as a pronounced and abrupt catastrophe of global dimensions and explained them by involving a sudden wave of cosmic radiation as the appropriately catastrophic causal factor. The stratigraphic record, however, still could be interpreted both ways and hence several workers—for example, Povlova (1924) and Sobolev (1928)—considered the same extinctions as extended over millions of years; therefore, they referred to more mundane causal processes, such as the various geographic, climatic and biotic effects of diastrophic cycle.

This divergence in palaeontological interpretations of the completeness of the record at various stratigraphic boundaries, and consequently of the rapidity and nature of mass extinction phenomena, has persisted until today. On the one hand, several prominent geologists and palaeontologists accepted the catastrophic nature of mass extinctions; in fact, even Schindewolf (1954) and Beurlen (1956) became at some point convinced that these phenomena could not be reasonably accounted for by an accidental clustering of the ends of the life cycle in a great number of species. As a result, a ever more imaginative narrative scenarios were presented which invoked a very wide variety of terrestrial and extraterrestrial causal factors as the ultimate culprits of extinction. Thus, Schindewolf (1954), Krasovsky and Shklovskiy (1957, Liniger (1961), and Russell and Tucker (1971) proposed that mass extinctions are caused by waves of cosmic radiation produced by supernova explosion in relative proximity to our planetary system. Dysssa et al. (1960) claimed that intense volcanism could cause earthly radioactivity to exceed lethal levels and thus lead to extinctions. De Laubenfels (1956) invoked a bolide impact on the earth at the Cretaceous-Tertiary boundary to explain the demise of the dinosaur.

McLaren (1970) put forth the hypothesis that a whole suite of environmental consequences of a bolide impact in the ocean could best account for the pattern of Frasnian-Famennian extinctions. Hays (1971) suggested that mass extinctions could be causally related to geomagnetic field reversals. Beurlen (1956) thought that a dramatic and virtually instantaneous change in seawater chemistry had poisoned and thus exterminated the majority of marine organisms at the end of the Paleozoic. Gartner and Keany (1978) claimed that a billover of cold brackish water from the previously isolated Arctic Basin on the world ocean had happened at the Cretaceous-Tertiary transition and caused mass mortality among the pelagic plankton; according to McClean (1978), this event should have led to a greenhouse effect due to carbon dioxide buildup in the atmosphere, and hence of further extinctions on a dramatic climatic change. There were no limits to imaginations, simply because there was no hard empirical evidence either firmly to corroborate seriously to contradict any of these stories. Schindewolf (1954), indeed, explicitly wrote that his hypothesis was merely a 'desperate move' to explain the mystery he could not resolve in a more scientific way.

On the other hand, the advocates of more gradualistic interpretations of mass extinction phenomena could at least base their explanatory scenarios on the apparent coincidence between these large-scale biotic changes and various major geological processes. Following from such evidence, global marine regressions were often regarded as the prime causes of mass extinctions (Lichkov, 1945; Newell, 1967), even though a clear mechanism of species extinction by regression was not identified. It is only after the advent of the theory of island biogeography that Schopf (1974) could causally interpret the striking correlation he observed between the pace of Permian-Triassic extinctions in the sea and the decline in the total area of continental shelves inhabitable by shallow-water marine organisms of the time. The island biogeographic theory predicts that the smaller the available area, the smaller the number of species it can harbour; hence, the origin of the single supercontinent at the end of the Paleozoic should lead to numerous extinctions. Given this

mechanism of extinction, however, mass extinctions should be very protacted in time and thay should also nonselectyve affect all the organic groups present in the ocean; and Schopf was very emphatic on this point. Other authors suggested other extinctions machanisms triggered by marine regressions. For inatance, Johnson (1974) suggested that the Frasnian-Famennian extinction of reef biotas had been caused by their particyulary vulnerable palaeogeographic position on broad shalves where even a minor regresion must have led to major enviromental consequences of global sea-levels fluctuations-from climatic to oceanographic to purely biogeografic ones.

Global marine regressions, however, are of course not the only sort of large-scale environmental phenomenon that could be corre lated with mass extinctions. Many geologist and paleoecologists have always regarded climatic change, and particularly cooling or even glaciation, as the main cause of mass extinction. The empir ical evidence in support of this view is best presented by Stanley (1987; 1988). The trouble is that, given the protacted dura tion of mass extinctions phenomena-as requiered by this kind of explanatory scenario- and given also the apparently great poten tial of various marine organisms to adapt rapidly to changes in seawater temperature, a climatic cooling hardly seems to be a sufficient cause for mass extinction; except, that is, for such unusual palaeographic situations as the West Atlantic or the Mediterranean in the Neogene.

As clearly shown by Jablonski (1986), however, in his comparative analysis of the phenomena traditionally recognized for the main mass extinctions in the history of life on the Earth-that is, the Ordovician-Silurian, Frasnian-Famennian, Permian-Triassic, Triassic-Jurassic, and Cretaceous-Tertiary transitions-neither eustatic sea-level falls, nor global climatic changes, nor any phases of the diastrophic cycle are consistently associated with mass extinction; and none of these explanations is adequte to explain all these five mass extinctions, let alone all the other geological time intervals that are sometimes interpreted to include events belonging to this category (the Eocene-Oligocene, Cenomanian-Turonian, earliest Toarcian, latest Precambrian, and so on). As theories aimed to explain all mass extinctions,

these traditional gradualist scenarios appear, then, also unsatisfactory. This is perhaps why new empirical developments were necessary substantially to rejuvenate this field of research in the 1980s.

THE CURRENT CONTROVERSY.

The finding by Alvarez et al. (1980) and Smit and Hertogen (1980) that geochemical anomalies, including a pronounced iridium spike, are associated with the mass extinction horizon of pelagic plankton at the Cretaceous-Tertiary boundary has for the first time provided hard evidence that could be interpreted as indicative of a causal link between biotic events and extraterrestrial influences. Iridium is a very rare element in the Earth's crust and its considerable concentrations at widely distant locations in the same stratigraphic horizons could be best explained as fallout from the dust cloud raised by impact of iridium-enriched bolide on the Earth. Such an impact, in turn, would also have a wide variety of environmental consequences, beginning with an initial shock and heat wave and ending with nuclear winter-type events (pollack et al 1983) and seawater intoxication by trace elements (Erickson and Dickson, 1987), wich could easily explain mass extinction of marine as well as terrestrial organism. The hypothesis of Cretaceous-Tertiary mass extinction by a huge bolide impact is, therefore, very appealing. Its proposal hasdratically changed the pace of research, and the tone of debate, on mass extinctions (for critical reviews of ideas and evidence see Sepkoski and Raup, 1986; Jablonski, 1986; Hoffman, 1989).

The hypothesis of an impact at the Cretaceous-Tertiary boundary is strongly supported by literally dozens of reports on geochemical (iridium and other siderophile element enrichment) and mineralogical (shocked quartz, fluffy carbon, microspherules comparable to altered impact droplets) fingerprints associated with either the marine or the palynological Cretaceous-Tertiary boundary. An iridium anomaly, associated in addition with microtektites, has also been discovered at the Eocene-Oligocene transition (Ganapathy, 1982). Therefore, when Raup and Sepkoski (1984) analysed the record of family

extinctions among marine animals and found that in may indicate a 26-million-year periodicity in extinction intensity of these taxa, it was logical to assume that the uniformity of extinction mechanism, suggested by the apparent periodicity of extinction events, points to extraterrestrial impacts as the culprits of mass extinction.

A search was immediately undertaken for astronomical mechanism that could cause such an impact periodicity, and astronomical mechanisms that could cause such an impact periodicity, and astronomer have rapidly come up with quite an impressive array of hypotheses; the Nemesis, a twin star of our Sun, could periodically disturb the Oort cloud of comets and throw some of them toward the inner planets; the same effect could be produced by the undetected tenth planet of our solar system; oscillations of the solar system about the galactic plane could also lead to increased frequency of our planet's encounters with comets, and so on. None of these hypotheses created more or less serious troubles, but none of them could be ultimately refuted (Sepkoski and Raup, 1986). All of them hinge, however, on the assumption of a uniformity-of-mass-extinction mechanism by extraterrestrial impacts. This assumption is in turn substantiated by the argument that if the peaks of extinction known from the fossil record are periodic, and if one or even two of these peaks are caused by impacts, then all the other peaks are most likely to be also caused by impacts.

The argument has largely shaped the controversy on mass extinctions which has continued relentlessly in the 1980's. For, on the one hand, it critically depends on the presumed validity of the hypotheses proposing the extinction periodicity and the impact causing of the Cretaceous-Tertiary (and to lesser degree also the Eocene-Oligocene) extinctions. On the other hand, it implies that all mass extinctions in the history of life on the Earth should be associated with impacts fingerprints and have a pattern compatible with a sudden and dramatic catastrophe. The latter implications has simulated several research projects that focused on the individual biotic events which could be construed as mass extinctions, but I shall not discuss this topic here, since the evidence will be presented and evaluated in detail in the other chapters of this book. Suffice

it to note here that no evidence has thus far been presented that would impacts; and the impact also is highly contentious. The question of extinction periodicity, moreover, is no less controversial.

Raup and Sepkoski (1984, 1986) conducted a series of statistical analyses of rate and intensity of extinctions of marine animals families and genera during the Phanerozoic. They concluded that the observed pattern of temporal distribution of extinction peaks is best explained by assuming a significant contribution from an approximately 26 million-year periodic signal in the later Phanerozoic, and perhaps a longer-period signal in the Paleozoic. This conclusion and my subsequent suggestion that this empirical pattern may in fact reflect nothing but a random variation in extinction intensity through time (Hoffman, 1985) have triggered a heated debate. It is important to realize that the empirical pattern of extinctions peaks evidently is not strictly periodic, so that the question of the likelihood that a random process leads to a strictly periodic pattern is irrelevant. What is actually at issue in this debate is whether the empirical pattern deviates so little from periodicity that it could only be obtained with the contribution of a strong periodic signal, or whether the observed amount of regularity is so small that the pattern could also result from a random process.

The issue is far from being ultimately resolved. On the one hand, Sepkoski and Raup (1986), Sepkoski (1986) and Forx (1987) present new analyses to reinforce the hypothesis of extinction periodicity. On the other

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On the other hand, a recent cathodoluminescence study of sheared quartz grains at the Cretaceous-Tertiary boundary demonstrated their non-volcanic origin (Owen and Anders, 1988); and the rhodium distribution in the boundary clay is very different from that found in terrestrial rocks, but closely similar to the one recorded in meteorites (Bekov et al., 1988). Moreover, tsunami deposits have been recently discovered at the

cretaceous-tertiary boundary (Bourgeois et al., 1988). These data, then strongly suggest that an impact did indeed take place at the very end of the Cretaceous, presumably during, or very soon after, a period of extremely intense volcanic activity.

The relative contribution of the environmental consequences of impact and volcanism to the Cretaceous-Tertiary mass extinction can hardly be disentangled. The fossil record, however, documents a more complex pattern than just a severe across-the-board extinction of many organic groups at the Cretaceous-Tertiary boundary. Among the pelagic plankton the foraminifer and coccolithophorid extinctions occur at demonstrably different stratigraphic horizons (Perch-Nielsen et al., 1982; Gartner and Jiang, 1984; Smit and Romein, 1985; Lindinger and Keller, 1987). Marine macroinvertebrates, in turn, were certainly undergoing significant extinctions over the last few million years of the Cretaceous (Birkelund and Hakansson, 1982; Kauffman, 1986; Ward et al., 1986; Hallam, 1987; Zinsmeister et al., 1987), thus ruling out their attributions solely to the terminal Cretaceous impact. This is perhaps why Alvarez et al. (1984) accepted the view that many organic groups, dwindling for whatever unidentified reasons toward the end of the Cretaceous, fell ultimately victim to the impact event at the Cretaceous-Tertiary boundary.

It thus appears that the Cretaceous-Tertiary mass extinction cannot be plausibly interpreted as caused solely by a bolide impact and its environmental consequences, and it is more feasible to invoke a coincidence of at least two different factors which were causally unrelated to each other (since the impact occurred at the end of a period of intense volcanism and extinctions). Hence, even if there were good reasons to assume the uniformity of extinction mechanisms for all mass extinctions, there would be no reason to accept extraterrestrial impacts as the ultimate cause for these biotic events. In the absence of strong evidence for extinction periodicity, however, the hypothesis of a uniform causation of all mass extinctions—either by impact, as envisaged by the hypothetical astronomical mechanism, or by episodes of extraordinary volcanism caused by

mantle processes, as proposed recently by Loper et al. (1988)—appears absolutely unsubstantiated.

This conclusion fully pertains to the recent article by Hut et al. (1987) who note that the Cenomanian-Turonian, Cretaceous-Tertiary, and Eocene-Oligocene extinction episodes, and who then propose that these mass extinction periods—and presumably also the others, for which they also hypothesize such stepwise nature—are caused by comet showers. The concept is derived from the fact that multiple impacts occurred during the Eocene-Oligocene transition and that an impact also took place during the extinctions at the Cretaceous-Tertiary boundary. The assumption of a uniformity of mass extinction mechanism then forces Hut et al. to envisage impact causation also for the Cenomanian-Turonian episodes of extinction and for the several latest Cretaceous episodes prior to the terminal event. There is, however, no evidence to justify this extrapolation. Moreover, both the Cenomanian-Turonian and Eocene-Oligocene data employed now by Hut et al. were previously explained by the same palaeontologists by reference to entirely were associated in time with the biotic events subsumed under the heading of mass extinction (Keller, 1983; Kauffman, 1984; Elder, 1987; Hansen, 1987).

It thus appears that all one can safely say at this point about the Cretaceous-Tertiary, Eocene-Oligocene, and Cenomanian-Turonian extinctions is that they occurred in clusters of extinction episodes, some of them perhaps causally related to bolide impacts on the Earth, but others apparently not. There is, moreover, strong evidence that some other extinction peaks, including the traditionally recognized Ordovician-Silurian and Frasnian-Famennian mass extinction, are also clusters of extinction episodes rather than single catastrophes (Brenchley 1984; House, 1985; Benton, 1986; Farsan, 1986; Hallam, 1986). This has led me to the conclusion that the major peaks of extinction may in fact be clusters of separate events more or less accidentally aggregated in time (Hoffman, 1989). They may be caused by global environmental events of various sorts—palaeoceanographic, climatic, or volcanic ones, as well as bolide impacts. Undoubtedly, extraterrestrial impacts on the Earth, enormous volcanic eruptions, oceanic anoxic event, dramatic sea-

level changes and climatic fluctuations occurred repeatedly during the Phanerozoic; yet only very few of these phenomena were actually implicated in the causation of mass extinctions. Perhaps then mass extinctions are not the biotic consequences of any single phenomenon of one or another sort, but rather rare incidences of more than one major changes in the physical environments accidentally cumulated together within relatively short intervals of geological time, say, 2-4 million years in duration.

This hypothesis of mass extinctions as coincidences of lesser episodes caused by a variety of physical factors may superficially resemble Schindewolf's (1954) desperate move to explain the mystery that could not be resolved in a more scientific way. Occurrence of such coincidences, however, is not all implausible or unlikely.

Consider, for example, w different kinds of physical events that occur at random, each with the probability of $1-p$ per million years. The following formula:

$$[p^{wk} + wp^{(w-1)k} (1-p^k)]^{n/k} = 1 - C$$

gives then the probability $1-C$ that, during the period of n million years, none of the consecutive k -million-year intervals will contain more than one event. of at least two events will occur within n million years. Assuming that, say, 5 different kinds of events may cause considerable extinctions (extraordinary volcanic eruptions, rapid climatic cooling, major transgressions and regressions, oceanic overturns, and large bolide impacts) and occur each with the average frequency of one every 50 millions year interval, C equals under such assumptions 0.73 for 100 million years and 0.96 for 250 million year. If the average frequency of events of each kind is decreased to only 1 per 100 million year, C still reaches 0.40 for at least one coincidence in a 4-million-year interval occurring within 100 million years and 0.60 for one such coincidence within 250 million years.

Recall that C seriously underestimate the actual probability of coincidence of events, as it does not take into account the probability of coincidences

that would span two consecutive time intervals. Clearly, then, coincidences (on the geological time-scale) of major environmental events accidentally aggregated within a few million years are quite likely to occur, and they may well account for at least some of the mass extinctions. This is not to say, however, that all mass extinctions are nothing but such clusters of separate extinction episodes.

For example, because of a major marine regression, the fossil record at the Permian-Triassic transition is so poor that the detailed pattern of extinction cannot be recognized on the global scale. In any event, however, no distinct episodes of extinctions can be identified, and contrary to the traditional opinion that the extinctions were spread over as much as 10 million year (Schopf, 1974), more recent studies on continuous stratigraphic sections suggest a much more rapid, though by no means catastrophic, process (Streng et al., 1984). The oceanic carbon and oxygen isotope records at the Permian-Triassic transition-spanning this greatest of all mass extinctions-indicate a dramatic oceanographic event, apparently more profound than any other observed thus far in the Phanerozoic. These isotopic records have been recently reviewed by Holser and Magaritz (1987, see also Magaritz et al., 1988), but Gruszczynski et al. (1989, Malkowski et al., 1989) present crucially important new data which demonstrate that, after a rapid shift toward extremely positive values, the oceanic $\delta^{13}C$ declined by more than 10 per mille, that is, down to distinctly negative values. The oceanic oxygen isotope essentially mimics the carbon one.

The trigger for the latest Permian changes in $\delta^{13}C$ trend is unknown. The mass balance calculations made by Gruszczynski et al. (1989; Malkowski et al., 1989) demonstrate, however, that the late Permian rise in $\delta^{13}C$ must have been caused by a rapid removal from the ocean-atmosphere system of huge amounts of organic carbon-more than 30 times the total amount of carbon in the presently living biosphere. The subsequent drop in $\delta^{13}C$ indicates that the ocean received three times more of organic carbon than it had previously lost. The amounts of organic carbon, which is reflected by the carbon isotope spike, must have left much oxygen free; the reduc-

ing conditions at the se-bottom contributed to nutrient recycling in the ocean and thus to an increase in the standing crop of the marine biosphere. However, the subsequent oxidation of organic matter, which is reflected by the carbon isotope records at the Permian-Triassic transition thus indicate a considerable decline in atmospheric oxygen and oceanic nutrient levels, which could well be the prime causes of mass extinction.

The isotopic records thus identify an oceanographic event as the cause of the Permian-Triassic extinctions, whereas no other causal factors have been recognized which could significantly contribute to these extraordinary exotic phenomena. Perhaps, then, this is a rare case of mass extinctions phenomenon that really deserves this name-triggered indeed by a single process and, consequently, confined in duration to a reasonably coherent time interval. The other mass extinctions, however, may be caused by various independent processes. It seems though that, apart from the Cretaceous-Tertiary mass extinction, their palaeoceanography is too poorly known to rule out the possibility of their, at least partial, causation by processes similar to the one operating at the Permian-Triassic transition. Much more palaeoceanographic effort, particularly on the Palaeozoic, is needed to decipher the palaeoenvironmental context of those mass extinctions.

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SECCION V: LA PALEONTOLOGIA EN MEXICO.

CAPITULO 11. EL PANORAMA PALEONTOLOGICO.

11.1. Preambulo.

A pesar de que la investigación paleontológica en nuestro país se remonta al primer tercio de siglo XIX, y de que a partir de 1986 existe una institución nacional dedicada a las ciencias geológicas-incluida la Paleontología no se ha escrito al presente una historia de la misma; por lo que resultaría imposible realizar aquí esa tarea. En consecuencia, se ha preferido abordar algunos aspectos de su problemática, destacando el entorno socio académico en que se ha insertado el desarrollo de esta ciencia en México.

Las fuentes bibliográficas específicas, son claro está muy reducidas, por lo que el presente autor se vio en la necesidad de echar mano de dos de sus propios trabajos y de notas que al respecto se han preparado con otros fines.

En el primero de ello, que fue una entrevista hecha al presente autor, se expone en lo general la gran importancia científica nacional e internacional que tiene el registro fósil con que cuenta nuestro país, y de la trascendente labor que se desarrolla en el Instituto de Geología, para darlo a conocer a la comunidad científica, por ser éste, el principal centro de investigación en esta especialidad que existe en México.

En el segundo, se aborda la problemática de la investigación paleontológica, mostrando que tanto el entorno socioacadémico de la profesión paleontológica, como la naturaleza geológica del territorio nacional, han influido poderosamente para conformar la clase de investigación paleontológica que se hace en el país, con su marcado énfasis en el trabajo sistemático y bioestratigráfico.

Con el objeto de suplementar la información presentada, se ha decidido agregar lo siguiente:

Docencia.- La paleontología General y algunas de sus especialidades como la Micro paleontología, Paleontología Estratigráfica, Palinología, Paleontología de vertebrados y otras, así como ciencias afines, p. ej., la Geología Histórica, forman parte del curriculum académico de las licenciaturas de Biología, Ingeniería Geológica y Oceanografía entre otras. Cabe destacar que estas carreras se imparten en instituciones de educación superior- principalmente universidades- ubicadas en 18 (Biología), 10 (Ingeniería Geológica) y 4 (Oceanografía) entidades federativas de la República. Ello teóricamente implica una amplia distribución geográfica de las instituciones educativas donde puede obtenerse cierta formación paleontológica. Sin embargo, las condiciones en términos de número de estudiantes, infraestructura de apoyo docente-colecciones, literatura, laboratorios, etc.- y nivel académico así como especialización de los profesores, son muy heterogéneas; siendo significativamente mejores en el Distrito Federal que en los Estados, y en conjunto, su calidad es de modesta a pobre, si se le compara con las necesidades de la formación académica que se supone debe atender satisfactoriamente.

Por lo que respecta al perfil académico del Profesor de Paleontología en este país, es el de un Licenciado en Biología, que en parte esta disciplina por tiempo parcial, ya que se dedica profesionalmente a otras actividades; aunque en el Distrito Federal, un número relativamente alto de profesores, tienen un grado académico superior al de la Licenciatura y/o se dedica profesionalmente a la práctica paleontológica en la investigación y/o en la docencia.

En relación a la formación académica de los paleontólogos, es necesario destacar que la Paleontología no es una especialidad formal en ninguna de las licenciaturas, y es una especialidad "informal" en los posgrados que se ofrecen en el

país, restringidos a la Universidad Nacional Autónoma de México y en el Instituto Politécnico Nacional, en donde las maestrías y doctorados más próximos, son en Ciencias, con la especialidad de Biología o Geología. En la Práctica, la mayor parte de los paleontólogos mexicanos con posgrado especializado en esta disciplina, lo han adquirido en el extranjero, y aún persiste esta tendencia.

LA PROFESION PALEONTOLOGICA.

En el mercado profesional en esta disciplina se restringe principalmente a la docencia, a las pocas instituciones donde se realiza investigación científica formal, que son el Instituto de Geología, el Instituto Mexicano del Petróleo y el Instituto Nacional de Antropología e Historia; así como a Petróleos Mexicanos, donde de manera cotidiana se realiza investigación aplicada y trabajo académico de servicio profesional. El desarrollo socioeconómico del país, no permite que exista de manera viable, empresas consultoras privadas, por lo que ningún paleontólogo profesional, podría subsistir con la prestación privada de susser vicios técnicos o científicos. Las perspectivas en este terreno, tal vez mantengan esta misma tendencia en el futuro inmediato.

LITERATURA CIENTIFICA.- La literatura especializada que se produce en México, consiste de una serie reducida de revistas, entre las que destacan Paleontología Mexicana, y la Revista de la Sociedad Mexicana de Paleontología. En las publicaciones Geológicas, tales como el Boletín de la Sociedad Geológica Mexicana, el Boletín de la Asociación Mexicana de Geólogos Petroleros y las Revistas del Instituto de Geología y del Instituto Mexicano del Petróleo, aparecen frecuentemente trabajos paleontológicos. Mención aparte merece el esfuerzo de producir textos y obras de consulta, principalmente por la carencia universitaria; el trabajo apenas se inicia y merece una amplia oportunidad de desarrollo, para subsanar el vacío y la dependencia que ello implica. Esta Antología es un paso en esa dirección.

LA SOCIEDAD MEXICANA DE PALEONTOLOGIA.- En 1986, se organizó por vez primera en México, una sociedad científica dedicada a promover el interés en la Paleontología, dotando por fin a la comunidad respectiva de un foro permanente, para realizar el necesario intercambio de ideas y experiencias que posibiliten un desempeño académico satisfactorio.

A la fecha, en esta sociedad, que se agrupa a la mayoría de los profesionales del ramo en el país, así como a colegas extranjeros que tienen interés en México, ha desplegado una actividad inusitada, habiendo celebrado ya tres congresos nacionales, publicando varios números de su importante revista, así como otras labores académicas. Es de desear que la Sociedad se fortalezca más, para que pueda incidir mejor en la solución de la compleja problemática que al respecto enfrenta el país.

EPILOGO.- La actividad paleontológica en México, acusa una problemática similar a la de muchas disciplinas científicas básicas, es decir dispone de un presupuesto limitado - más bien reducido- y el número de sus profesionales es pequeño. Empero ellos han realizado una amplia labor, dando a conocer una buena parte de la importantísima biota fósil presente en nuestro territorio. El avance real en este campo, requiere de la participación entusiasta de toda la comunidad, tanto en el aspecto docente, como en el de investigación científica y aplicada, así como en el de la difusión; ello permitirá alcanzar y sostener el nivel de calidad que este país necesita y merece.

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11.3 ASPECTOS GENERALES.

11.3(A)

LA RIQUEZA PALEONTOLOGICA DE MEXICO, VALIOSA A NIVEL MUNDIAL.*

El registro fósil documenta la evolución orgánica, permite conocer como vivieron ciertas especies e inferir relaciones filogenéticas.

En México, la importancia de los hallazgos paleontológicos que se realizan y se desarrollarán en el futuro es enorme, tanto a nivel nacional como mundial, por la peculiar ubicación geográfica e historia geológica que tiene nuestro país: es decir, nuestro territorio ocupa una área clave para entender la historia geológica y biológica de América y de los continentes africano y europeo.

El Dr. Ismael Ferrusquía Villafranca, miembro de la academia de la Investigación Científica y de la comunidad paleontológica, habla sobre esta ciencia, así como del esfuerzo permanente de la UNAM en el campo de la difusión cultural y de la actividad científica.

La paleontología es una ciencia natural y como tal, dijo el especialista del Inst. Geol. de la UNAM, responde a interrogantes que el hombre se plantea acerca de la naturaleza, particularmente en torno a los seres vivos que existieron en el pasado geológico: su pertenencia biológica, su escenario geográfico, su relación con los seres actuales y los vestigios que dejaron de su existencia etc.

Estas son algunas de las cuestiones que el paleontólogo trata de resolver y las actuales constituyen el núcleo de su actividad profesional. En este sentido, agregó, las respuestas tienen un doble valor: por un lado intentan resolver interrogantes trascendentes que el hombre se hace como expresiones de su curiosidad innata.

Por otro lado, la información paleontológica contribuye a resolver problemas de otras ciencias, como la biología evolutiva, en donde el registro fósil documenta al fenómeno de la evolución orgánica, permite conocer cuándo y dónde vivieron ciertas especies, inferir sus relaciones filogenéticas y entender su distribución biogeográfica, entre otras.

En el terreno de la geología, por el hecho mismo de la evolución orgánica, los seres vivos han dejado un registro fósil individualizable cronológicamente. Este posibilita que, en función de los fósiles de algunas rocas, se pueda conocer su edad geológica relativa y con base en ello establecer su correlación estratigráfica y, desde luego, conocer con relativa certidumbre la historia geológica de un lugar.

Los fósiles, aseveró el doctor Ferrusquía, profesor de la Facultad de Ciencias y autor de múltiples artículos publicados tanto a nivel nacional como internacional, son testimonios materiales de seres vivos, reconocibles como tales por su configuración y estructura, que existieron en el pasado. En términos convencionales se habla de una antigüedad mínima de 10 mil a 20 mil años. Estos testimonios incluyen desde organismos completos hasta huellas de su actividad. Ello implica la existencia de una gran diversidad de fósiles y, en consecuencia, métodos muy variados para estudiarlos. En función de su tamaño existen macroscópicos y microscópicos; y pueden tener una antigüedad de 3 mil 500 millones de años. Por su naturaleza biológica incluyen representantes de los 5 reinos que se reconocen en el presente. En cuanto a su origen, los fósiles se generan con base en procesos que excluyen a ciertos individuos de la destrucción y reciclaje de material que naturalmente ocurre en los seres vivos. Este fenómeno se denomina como proceso de fosilización y en general involucra la cobertura rápida por sedimentos, evitándose así la destrucción del resto orgánico.

Es de esperarse, afirmó el doctor Ferrusquía, que estos procesos sean más frecuentes en las cuencas sedimentarias que en otros medios, pues la presencia de un esqueleto mecánico mineralizado favorece la posibilidad de la fosilización. Por estas razones el registro fósil está

*Entrevista al Dr. I. Ferrusquía V., publicada en la Gaceta de la UNAM, Num. 2,478, Junio 11, 1990.

constituido mayoritariamente por animales esqueletóforos marinos, que son los que tienen la máxima posibilidad de fosilizarse. Aún así, es sólo una mínima parte de los individuos que forman una especie dada los que se fosilizan y de ellos sólo unos cuantos son reconocidos, colectados, estudiados y eventualmente publicados.

Es por esta razón, considero el paleontólogo, que los fósiles constituyen un recurso patrimonial, cultural, natural finito y no renovable, que debe ser reconocido y apreciado por su obsolescencia, así como protegido del abuso y la depredación.

Sobre la actividad que acerca de esta ciencia se realiza en nuestro país, el doctor Ismael Ferrusquía señala que la comunidad de paleontólogos de México es relativamente pequeña - dada la magnitud y complejidad del registro fósil nacional - y labora principalmente en el sector público, en especial en instituciones de educación superior, donde realiza actividades de docencia y en menor grado de investigación; en Petróleos Mexicanos se encuentra el mayor número de paleontólogos profesionales en áreas de exploración geológicas-petrolera.

El desarrollo económico-industrial en México es tal que ningún paleontólogo podría subsistir en la práctica profesional privada, pues no existiría suficiente demanda de sus servicios. Esta situación, considero el investigador, ha ocasionado que el investigador, ha ocasionado que la investigación paleontológica en lo general se realice como una actividad de apoyo y que, en consecuencia, no se haya gozado de libertad de llevarla a cabo por sí. Ello ha influido en el conocimiento sobre el registro fósil que se genera en México; se ha hecho énfasis en aquello que resulta de interés para la geología petrolera, en detrimento de las otras áreas.

En el país solamente el Instituto de Geología de la UNAM realiza de manera sistemática investigación paleontológica básica; por ello la importancia y trascendencia de la labor que aquí se practica. El Instituto, dijo, tiene ya una historia más que centenaria, y es junto con los Institutos de Biología y Astronomía uno de los tres fundadores de la investigación científica profesional en la UNAM. En él se aloja la colección más extensa de fósiles de México y en su biblioteca se

encuentra el mayor acervo bibliográfico, no sólo del país, sino de muchos de Latinoamérica. Sin embargo, no existe un servicio geológico o un museo nacionales específicamente encargados de mantener este tipo de acervos, desarrollar y coordinar la investigación en esta disciplina en todo el país. En la práctica el Instituto ha sumido esta especialidad a nivel nacional con las limitaciones del caso y sin el apoyo legislativo correspondiente.

Sobre la legislación que rige en esta disciplina, el doctor Ismael Ferrusquía informó que en México el Estado considera a los fósiles como parte del patrimonio cultural y por decreto presidencial desde 1986 su cuidado recae en el Instituto Nacional de Antropología e Historia de la Secretaría de Educación Pública, al mismo nivel que el material histórico o el Arqueológico. Entre otros aspectos, los fósiles mexicanos no pueden ser objeto de comercio; las colecciones privadas deben estar registradas ante el INAH y la exportación del material fósil requiere del permiso respectivo. El establecimiento práctico de este decreto está lleno de dificultades y la comunidad de paleontólogos cuestiona ya su conveniencia real para lograr los fines que se propuso el Estado al expedirlo.

Acercar el panorama actual de la paleontología en México, sus problemáticas y sus perspectivas, el especialista dijo que el registro fósil de nuestro país es muy grande y variado. Incluye elementos de precámbrico a pleistocénicos, burdamente de 1,700 millones a 5,000 10,000 años de antigüedad; animales y vegetales, macroscópicos y microscópicos, marinos y continentales; pero en general aun es casi desconocido, dado que el tamaño de la comunidad científica, especializada es proporcionalmente muy pequeño. Tal vez no exceda de 300 números de paleontólogos profesionales de México.

Por lo tanto, en el presente y en el futuro inmediato gran parte del esfuerzo de esta comunidad se centrará en la necesidad de describir este registro. Debido al escaso número de programas de investigación, estrictamente paleontológica, esta tarea tomará más tiempo; por ello una de las responsabilidades principales de la comunidad será lograr su incremento. La diver-

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sidad y complejidad del registro requiere de especialistas muy diversos que debiera en n tiempo corto, si se pretende realizar la mencionada labor para paleontólogos nacionales y en un lapso razonable. Atender esto es tambien una tarea prioritaria. Al igual que en otros segmentos de la comunidad científica que la tasa de reemplazamiento es inferior a la necesaria para mantener su dinamismo y operatividad; problema que implica, una situación paradójica no se puede proporcionar un empleo satisfactoriamente remunerativo, en número suficiente, a los escasos jóvenes que atendiendo a su vocación se esta formando académicamente en esta ciencia o que, piensan hacerlo, con todo para los entusiastas y los esforzados siempre habra un lugar, finalizado.

Jaime R. Villagrana.

II.4 INVESTIGACION

11.4 (a)

SOBRE LA INVESTIGACION PALEONTOLOGICA EN MEXICO. *

Ismael Ferrusquia Villafranca.

RESUMEN:

El marco y las características de la profesión geológica, determinan la naturaleza de los estudios paleontológicos desarrollados en México. En el territorio nacional, las rocas calcáreas jurásicas, cretácicas y cenozoicas son las sedimentarias más abundantes; constituyen asimismo importantes rocas productoras o almacenadoras de petróleo. En consecuencia, la mayor parte del trabajo paleontológico se ha hecho sobre fósiles procedentes de rocas.

La estructura geológica-paleontológica es tal, que de las diez instituciones principales solamente una realiza investigación paleontológica básica. Aproximadamente el 4% (unos 100) de los geólogos que trabajan en México, se dedica profesionalmente a la paleontología; la mayoría de ellos está en los petróleos mexicanos, Instituto Mexicano del Petróleo y el Instituto de Geología.

Debido a la gran necesidad de trabajo sistemático y bioestratigráfico básico, el énfasis de las investigaciones se da en estos aspectos. Sin embargo se promueve también la investigación paleontológica avanzada sobre otras cuestiones.

ABSTRACT

Mexico's geologic framework and the make up of the geologic profession are the main factors shaping the nature of the paleontological work. The jurassic, cretaceous and cenozoic carbonates

rocks are the most abundant sedimentary rocks, as well as the chief sources of oil. Hence the bulk of the paleontological work is done on their fossil

The geological-paleontological community works for the most part in ten major institutions; eight are devoted to applied geologic studies and only two are concerned with basic research. About 4% (some 100) of the geologists working in Mexico take up paleontology as a profession. They work mainly for the Mexican Petroleum Institute, the Mexican Petroleum Institute and the Institute of Geology.

A large amount of straight systematics and biostratigraphic work remains to be done, therefore these aspects are and will be emphasized in the paleontologic research for a long time. Advanced research is promoted also, however.

Introducción:

Para entender el alcance, la naturaleza y el desarrollo de la investigación paleontológica en México es necesario examinarla bajo la perspectiva dada por el marco geológico y la estructura de la comunidad y la profesión en México. Influye desde luego otros factores. Se suman a continuación, las características de estos factores.

EL MARCO GEOLOGICO.

Una de las maneras más sencillas de resumir descriptivamente la geología del país es el enfoque de provincialización morfotectónica, esto es, dividir al país en amplias regiones que tengan rasgos e historia geológica similar. Hay varios arreglos posibles (cf. Guzman y de Cserna, 1963) el que se presenta a continuación (fig 1), es el más apropiado para el objetivo presente.

Se reconocen aquí diez provincias.

1.- PENINSULA DE BAJA CALIFORNIA.

La mitad norte está constituida por rocas intrusivas mesozoicas que intrusieron y flaquearon rocas sedimentarias marinas y continentales. La mitad sur incluye principalmente andesitas y basaltos cenozoicos, y rocas sedimentarias marinas y continentales del

jurásico al reciente, la punta de la península está formada por rocas intrusivas y sedimentarias que aparentemente tienen una historia geológica muy distinta al resto de la península.

2.- CUENCAS Y SIERRA NOROCCIDENTALES

Es la extensión meridional de la provincia Basin and Range de Estados Unidos. Está constituida por rocas paleozoicas marinas falladas en bloques, que constituyen las sierras. Las cuencas están llenas de sedimentos continentales cenozoicos. El fallamiento en bloques ha expuesto en algunas partes a las rocas cristalinas precámbricas; es aquí donde se encuentra las rocas más antiguas de México, que corresponden al proterozoico medio y tardío.

3.- SIERRA MADRE OCCIDENTAL

Esta constituida por rocas volcánicas de edad cenozoica media y tardía que incluyen principalmente tobas e igníbritas riolíticas a riolíticas y derrames andesíticos. La escasez de caminos dificulta la cartografía geológica detallada esta provincia es la más grande de todas.

4.- FAJA NEOVOLCANICA TRANS-MEXICANA

Está constituida principalmente por rocas andesíticas a basálticas del cenozoico medio y tardío. La composición cambia cronológicamente de andesítica a basáltica. Se aprecia un fallamiento en bloques y rifting moderado, dirigido de oriente a poniente que ha producido pilares y fosas tectónicas, en donde se han desarrollado lagos y sedimentos lacustres asociados. La interpretación de esta faja volcánica es incierta todavía, pero claramente constituye un rasgo geológico mayor.

5.- SIERRA MADRE DEL SUR

Es la provincia más complicada e incluye tres dominios, Guerrerense, Oaxaqueño y Chiapaneco. Estos dominios corresponden a cuen-

cas sedimentarias de compleja evolución geológica, desarrolladas sobre rocas cristalinas proterozoicas y paleozoicas de evolución geológica compleja también. Las rocas de esta provincia incluyen el espectro de litología y su edad va del proterozoico tardío al reciente.

En la región suroriental, las estructuras geológicas de las rocas mesozoicas cambian gradualmente su dirección principal de norte a sur noroeste-sureste.

6.- PLATAFORMA DE YUCATAN

Es una plataforma calcárea cenozoica, acrecionada desde el Eoceno, donde se ha desarrollado una extensa topografía carstica.

7.- PLANICIE COSTERA DEL GOLFO

La mitad norteña está constituida por rocas clásticas marinas y transicionales que representan una fase regresiva relacionada con la genesis y evolución del golfo de México. La mitad oriental incluye rocas calcáreas cenozoicas tardías en la superficie, que cubren a rocas cenozoicas y mesozoicas en el subsuelo, portadoras de hidrocarburos abundantes, de donde se desarrollan los nuevos e importantes campos petroleros.

8.- SIERRA MADRE ORIENTAL.

Esta constituida por rocas calcáreas marinas jurásicas y cretácicas plegadas, falladas y levantadas durante la Orogenesis Laromide. Las sierras que ocupan la región oriental, cambian gradualmente su orientación nore-sur hacia este-oeste, mediante una flexura mayor, que constituye uno de los grandes enigmas de la geología mexicana. Las cuencas intermontanas están rellenas de rocas clásticas cenozoicas.

9.- SIERRAS Y ALTIPLANICIES NORTEÑAS

Están formadas por sierras orientadas noroeste-sureste, constituidas por rocas calcáreas jurásicas

y cretácicas plegadas, falladas y levantadas, separadas entre sí por cuencas intermontañas confluyentes, rellenas de clásticos cenozoicos. En la región occidental están asociadas con rocas volcánicas andesíticas, silíceas del cenozoico medio y tardío, que contienen yacimientos minerales económicamente importantes.

10.- ALTIPLANICIE CENTRAL

Es una cuenca endorreica, rellena parcialmente por clásticos cenozoicos derivados de las sierras que la bordean. En su región meridional, está asociada a rocas volcánicas andesíticas y silíceas en donde se han desarrollado yacimientos minerales muy importantes. En el subsuelo de esta provincia semiárida, se aloja extensos acuíferos.

Resumiendo la geología se tiene que, aproximadamente un tercio del país está constituido por rocas volcánicas; otro tercio está compuesto por rocas calcáreas marinas, jurásicas y cretácicas en su mayoría plegadas y falladas complejamente, que forman la Sierra Madre Oriental, parte de la Sierra Madre del Sur y las sierras nortecentrales; el otro tercio incluyen los otros tipos litológicos, entre los que destacan por su abundancia, las rocas cristalinas plutónicas y metamórficas, que se representan un 60% de este tercio y, las rocas clásticas marinas y continentales que constituyen un 30% de este. Es notoria la escasez de rocas paleozoicas sedimentarias no metamorfozadas.

Con esto en mente, resulta claro que el trabajo paleontológico en México se centra volumétricamente en las rocas mesozoicas marinas, en menor escala en las rocas cenozoicas marinas y mucho menos en las meso y cenozoicas continentales.

LA COMUNIDAD Y PROFESIÓN GEOLOGICA.

Como en otros países, los estudios geológicos en México se han encaminando principalmente a la búsqueda de recursos naturales no renovables tales como, petróleo, yacimientos minerales de valor económico, acuíferos, uranio, fosfato, etc. Esto se ha logrado principalmente a través de

instituciones gubernamentales de diferente tipo. En México trabajan al presente unos 2 mil a 2500 geólogos (López Ramos, 1974), en varias instituciones mutuamente independiente, de rango equivalente, pero de tamaño, estructura y objetivos diferentes, las siguientes diez son las principales:

PETROLEOS MEXICANOS (PEMEX)

Constituye el grupo mayoritario, dedicado obviamente a la explotación geológica petrolera. Su personal incluye 1200 geólogos, (50% de la comunidad geológica nacional).

INSTITUTO MEXICANO DEL PETROLEO (IMP)

Es el grupo de investigación de PEMEX, el trabajo geológico enfatiza la investigación básica orientada a la geología petrolera. Su personal incluye unos 120 geólogos.

CONSEJO DE RECURSOS MINERALES (CRM).

Es el segundo grupo mayoritario, dedicado principalmente a estudios de geología económica. Su personal incluye 250 geólogos.

SECRETARIA DE AGRICULTURA Y RECURSOS HIDRAULICOS (SARH)

Grupo dedicado a la prospección geohidrológica. Su personal incluye unos 200 geólogos.

COMISION FEDERAL DE ELECTRICIDAD. (CFE)

Grupo dedicado a estudios geológicos aplicado a la ingeniería civil; recientemente han desarrollado estudios sobre carbón y recursos geotérmicos, en un esfuerzo por encontrar fuentes alternativas de energía, su personal incluye unos 150 geólogos.

DIRECCION DE ESTUDIOS GEOLOGICOS Y GEOGRAFICOS DEL TERRITORIO NACIONAL (DEGETENAL).

Este grupo esta haciendo el inventario de los recursos naturales del pais; presenta sus resultados en mapas dispuestos en cuadrangulos de 20'x 30', escala 1:50000. La serie topográfica es la mejor. Los mapas geológicos se apoyan ampliamente en estudios fotogeológicos. Su personal incluye unos 80 geólogos.

URANIO MEXICANO (URAMEX).

Este grupo se dedica a la prospección geológica del uranio, etc., tiene unos 60 geólogos.

SECRETARIA DE ASENTAMIENTOS HUMANOS Y OBRAS PUBLICAS. (SAHOP)

Su grupo de geólogos se dedica a estudios aplicados a la ingeniería civil, Su personal incluye unos 50 geólogos.

FOMENTO MINERO (FM).

Este grupo realiza estudios de geología económica para la industria minera pequeña y mediana. Su personal incluye unos 50 geólogos.

INSTITUTO DE GEOLOGIA, UNAM (IG)

Grupo pequeño dedicado a la investigación geológica básica. Su personal incluye 50 geólogos.

Existen otros grupos mas pequeños en las universidades estatales, en la industria privada, principalmente minera y en la consultoría. Solamente las compañías mineras grandes, tienen personal geológico de base.

RESUMIENDO:

Las diez instituciones geológicas principales, donde presta sus servicios aproximadamente el 85% de los geólogos mexicanos, son de tipo gubernamental, de tamaño, estructura y objetivos muy distintos, mutuamente independientes y de rango equivalente. Su trabajo implica una superposición considerable; que difícil de evitar, por la nula coordinación, causada por la falta de una institución nacional rectora o de un programa maestro de trabajo geológico de alcance nacional. Únicamente dos instituciones, el IMP y el IG, realiza sistemáticamente investigación básica; los otros ocho desarrollan investigación aplicada. Aproximadamente un 4% de los geólogos que trabajan en México, estos son unos 100 aproximadamente, realizan trabajos paleontológicos; estos estudios se desarrollan principalmente en PEMEX, IMP e IG.

OTROS FACTORES QUE INFLUYEN EN EL DESARROLLO Y NATURALEZA DE LA INVESTIGACION PALEONTOLOGICA EN MEXICO.

L.- Formación profesional.

No existe en México ninguna institución dedicada a la formación profesional paleontológica; debe mencionarse sin embargo, que en los tres últimos lustros se han desarrollado programas de estudio de postgrado que incluyen especialización en paleontología. En consecuencia, el profesional de la paleontología ha adquirido su formación mediante estudios formales de postgrado en instituciones extranjeras o nacionales; o bien se ha habilitado como tal, mediante un entrenamiento capacitatorio informal y un gran esfuerzo autodidacta. Por ende, el resultado del trabajo refleja en cierta medida esta formación. Aproximadamente un tercio de los paleontólogos de México tienen estudios formales.

2.-SUBORDINACION DE LA INVESTIGACION PALEONTOLOGICA A OTROS OBJETIVOS.

Dedido a que el desarrollo histórico y económico del país se ha centrado en torno a la exportación de materias primas y energéticos, y a que el desarrollo industrial que se está realizando requiere de estos insumos, la investigación geológica se ha encausado en gran medida a su búsqueda. Consecuentemente, la labor paleontológica ha quedado subordinada a estos objetivos prioritarios, con la natural falta de incentivos para desarrollar programas que tiendan a la solución de problemas paleontológicos que no tengan una relación directa con objetivos económicos. Solo recientemente se ha comprendido, que la búsqueda y hallazgo de los recursos mencionados, es en realidad el resultado de una síntesis cognoscitiva apoyada en resultados de investigación básica de naturaleza muy diversa, y de las cuales la paleontología pura es una parte importante. Aun así, hay mucho camino por recorrer, para liberar a la paleontología de esta posición subordinada.

LA INVESTIGACION PALEONTOLOGICA EN MEXICO.

Tal es el marco en el cual se desenvuelve la investigación paleontológica en México. Comprensible la mayor parte esta dedicada al estudio de las rocas mesozoicas marinas, una parte menor a las rocas cenozoicas marinas y continentales y aun una menor, a las rocas paleozoicas. Comprensiblemente también, los esfuerzos principales se desarrollan en el Instituto Mexicano del Petróleo y en el Instituto de Geología.

INSTITUTO MEXICANO DEL PETROLEO Y PEMEX.

Se señala ya que el IMP es el grupo de investigación de PEMEX. El personal incluye unos 35 paleontólogos. De ellos, los micropaleontólogos constituyen el grupo mayoritario. El énfasis está en estudio sobre foraminíferos, tendientes a establecer zonación bioestratigráfica detallada, parámetros ambientales diversos, correlación regional y otros aspectos. Existe también un importante grupo palinológico, así como algunos especialistas en otros taxa de interés inmediato en

la prospección petrolera, como son los amonitas y conodontes.

En PEMEX, la labor paleontológica es principalmente de tipo rutinario, tendiente a la ubicación estratigráfica, apoyándose principalmente en el reconocimiento de foraminíferofaunas de las distintas zonas establecidas. Aproximadamente de 35 a 50 paleontólogos desarrollan esta labor, realizándola en las Oficinas Regionales de PEMEX, localizadas en distintos estados de la república.

Los resultados de los proyectos de investigación del IMP y de PEMEX son confidenciales. Algunos de ellos se publican en el IMP boletín del IMP o en otras revistas. Sin embargo, a pesar del enorme interés que tienen las investigaciones sobre la bioestratigrafía de la mitad sur de la planicie costera del Golfo, de la sierra madre de Chiapas y de la región de Vizcaino en la península de Baja California, debido a su relación obvia con los recientemente desarrollados e importantes campos petroleros, los resultados de estas investigaciones son muy confidenciales.

INSTITUTO DE GEOLOGIA, UNIVERSIDAD NACIONAL AUTONOMA DE MEXICO.

Es la única institución dedicada a la investigación geológica básica. Su objetivo es contribuir al conocimiento sobre la evolución central geológica del país. En ausencia de una institución geológica central y de un museo nacional, el instituto ha adoptado informalmente estas dos responsabilidades, aunándolas a su función primaria como una institución universitaria de investigación científica. De suerte que su naturaleza es "sui generis". Este triple papel se refleja en los distintos departamentos que lo integran.

El departamento de paleontología tiene como objetivo central, estudios bioestratigráficos y paleontológicos de varias regiones y grupos. Se desarrollan programas de Paleontología de varias regiones y grupos. Se desarrollan programas de Paleontología de invertebrados, Vertebrados, Micropaleontología y Paleobotánica.

El Programa de Paleontología de Invertebrados incluye estudios sistemáticos de gasterópodos, pelecípodos, equinodermos, corales, amonitas y braquiópodos meso y cenozoicos principalmente, así como estudios bioestratigráficos en varias regiones tales como Baja California, La Cuenca Mixteca, el Sureste de México y la Planicie Costera.

El Programa de Paleontología de Vertebrados incluye estudios faunísticos, de tiburones y mamíferos principalmente y estudios bioestratigráficos en Baja California, La Añtliplanicie Mexicana el Sureste de México y algunas cuencas INtermontanas de la Sierra Madre Oriental. La naturaleza exploratoria y prospectiva del trabajo, obliga a la descripción de material muy diverso.

El Programa de Micropaleontología tiene sus énfasis en estudios faunísticos y bioestratigráficos sobre ostrácodos y foraminíferos cenozoicos de la Planicie Costera del Golfo y de la Península de Baja California.

El programa de Paleobotánica incluye estudios paleoflorísticos en las cuencas carboníferas cretácicas de Coahuila, Jurásicas de Oaxaca; paleofloras triásicas de Sonora e Hidalgo y recientemente, ficolflora precámbricas en Sonora. En el Departamento de Geología, se desarrollan estudios palinoestratigráficos de Coahuila y Sonora, así como foraminíferoestratigráficos en Baja California y Sonora.

Además de estos, el Departamento de Paleontología tiene a su cargo el Museo, que funge como Depositario Nacional de facto de los fósiles mexicanos, particularmente de los tipos. El Museo de Paleontología aloja la colección de fósiles más extensa del país e incluye más de 500 000 ejemplares.

EVALUACION Y CONSIDERACIONES GENERALES.

Debido a que el número de paleontólogos ha sido siempre muy reducido y se ha visto obligado a laborar en condiciones y programas de investigación, subordinados a otros objetivos, no

se ha dispuesto ni de la masa crítica ni de la libertad de acción para desarrollar programas ambiciosos de Investigación paleontológica básica; en consecuencia, gran parte de la paleobiota mexicana es desconocida. De ahí que una gran cantidad de trabajo descriptivo-sistemático permanezca por hacer, que seguramente mantendrá ocupada a la comunidad paleontológica por mucho tiempo. Por ejemplo, el desarrollo de la Paleontología de Vertebrados, está principalmente en la fase prospectiva. El problema es importante, por que no se cuentan con especialistas en varios grupos importantes y la tasa de crecimiento de la comunidad es relativamente baja.

En cambio la Paleontología de Invertebrados es la más desarrollada, principalmente debido a su relación ovía con la geología petrolera.

Se necesita desarrollar una gran cantidad de trabajo bioestratigráfico, para producir cartas de correlación más confiables, esquemas detallados de zonación, mapas de bioespecies, etc. De hecho el Cenozoico Continental no ha sido subdividido estratigráficamente.

A pesar de la gran cantidad de trabajo descriptivo que de necesidad se realiza, se está desarrollando investigación avanzada en algunas áreas tales como morfología funcional de tiburones, patrones de distribución en mamíferos y moluscos, modelos paleoecológicos, taxonomía teórica y filogenia, entre otros.

La comunidad Paleontológica está consciente de la enorme significación de México, para verificar hipótesis o buscar soluciones a problemas científicos trascendentes tales como, biogeografía regional de diversos grupos, rutas de dispersión; origen radiación adaptativa y dispersión de ciertos grupos, cambios climáticos latitudinalmente relacionados, endemidad y distribución extrema de ciertos grupos; evolución y tectónica regional, reconstrucciones paleogeográficas y paleonepásticas, etc.

Como un ejemplo de las amplias implicaciones del trabajo paleontológico que se realiza actualmente, se resumirá uno (Ferrusquía et al., 1978 (1981)), en el que el autor estuvo directamente involucrado: "Las Huellas de dinosaurios más

australes de Norteamérica y su significación geobiológica."

En Playa Azul, Michoacán aflora una secuencia volcanosedimentaria ¿Jurásica media-cretácica temprana? constituida por derrames andesíticos, piroclastitas riolíticas, biomicritas marinas someras y filovolcanitas rojas inmaduras, buzadas al SSE y están intrusionadas por mantos, "stocks" pequeños y diques que muestran plegamientos y fallamientos. Este es el primer hallazgo de rocas volcanosedimentarias principalmente continentales de edad jurásica-co-cretácica en el margen pacífico de México.

La dinosauriofauna incluye siete morfotipos asignables a terópodos y ornitópodos; constituye un primer registro de saurionautas en México y es el más meridional en Norteamérica; testimonia la comunidad de dinosauriofauna Norteamericana del Jurásico Medio-Cretácico Temprano por lo menos hasta el Sur de Michoacán y concomitantemente, a la continuidad territorial que hizo posible esta extensión faunística.

Los resultados de estas investigaciones se publican principalmente en las series del Instituto: Revista, Boletín y Paleontología Mexicana, algunos estudios se publican en revistas extranjeras.

EL FUTURO DE LA PALEONTOLOGIA EN MEXICO.

Aunque es difícil hacer afirmaciones definitivas, examinando las necesidades principales, es posible plantear algunas expectativas plausibles:

1.- Considerando que hay deficiencias importantes de personal para cubrir algunos grupos, tópicos y áreas, resulta imprescindible generarlo.

2.- La gran cantidad de trabajo sistemático-descriptivo por hacer, mantendrá a la comunidad paleontológica ocupada por muchas décadas. Por tanto el trabajo futuro continuará enfatizando este aspecto.

3.- El trabajo bioestratigráfico que falta también es enorme, consecuentemente, se enfatizará en la futura labor tendiente a desarrollar y perfeccionar la zonación bioestratigráfica y la correlación.

4.- A pesar de esto, es deseable promover el desarrollo de investigación avanzada, para realizar una síntesis conceptual significativa y no quedar a la zaga en esta disciplina científica.

5.- Todo ello será posible en la medida que la comunidad paleontológica desarrolle una estrategia que libere de su papel subordinado entre las Ciencias de la Tierra, así podrá avocarse al planteamiento de programas ambiciosos de alcance nacional y cooperación internacional, que aseguren el desarrollo armonioso y balanceado de las distintas especialidades paleontológicas, el ataque de problemas relevantes por sí mismos o por su relación con el desarrollo socioeconómico del país y la generación del personal debidamente entrenado, al ritmo idóneo.

6.- Para ello se debe cambiar inteligentemente la importación de personal extranjero, el entrenamiento formal dentro y fuera del país, de personal nacional seriamente interesado y el establecimiento de programas de cooperación internacional, con países que cuenten con una comunidad paleontológica ampliamente desarrollada. Los mejores programas incluirían entrenamiento e investigación suplementaria en áreas o tópicos en donde se han detectado problemas específicos relevantes, pero, no se cuenta con especialistas. También se podrían intercambiar especialistas, para resolver problemas inmediatos.

7.- En último término, pero no de menor importancia es el contar recursos materiales y una infraestructura adecuada, que permita a la comunidad paleontológica desarrollarse y enfrentar exitosamente el reto que representan las tareas señaladas. La comunidad debe avocarse a esto.

SUMARIO Y CONCLUSIONES.

1.- El marco geológico, la organización de comunidad geológica y el desarrollo socioeconómico del país, han determinado en gran medida la naturaleza de los estudios paleontológicos en México.

2.- Las rocas calcáreas jurásicas y cretácicas son las más abundantes fosilíferas y también las de mayor interés petrolero, consecuentemente, en

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torno a ellas se ha desarrollado la mayoría del trabajo paleontológico.

3.-De las diez instituciones geológicas principales de México, solamente en dos se desarrolla investigación básica; ésta incluye Paleontología.

4.-Únicamente el 4% (unos 100) de los geólogos que trabajan en México, se dedica profesionalmente a la Paleontología; la mayoría están en PEMEX, IMP y el IG.

5.- El Instituto de Geología desarrolla programas de investigación en Paleontología de Invertebrados, Vertebrados, Micropaleontología, Paleobotánica y Palinología; cuenta también con el Museo Nacional, poseedor de la más extensa colección paleontológica del país.

6.-Debido a la gran necesidad de trabajo sistemático y bioestratigráfico básico por hacer, las publicaciones actuales y futuras de la comunidad paleontológica mexicana enfatizarán estos aspectos.

7.-Un programa de cooperación internacional que involucre entre enseñanza e investigación suplementaria, sería el más adecuado para desarrollar completamente y brevemente, la Ciencia de la Paleontología en México.

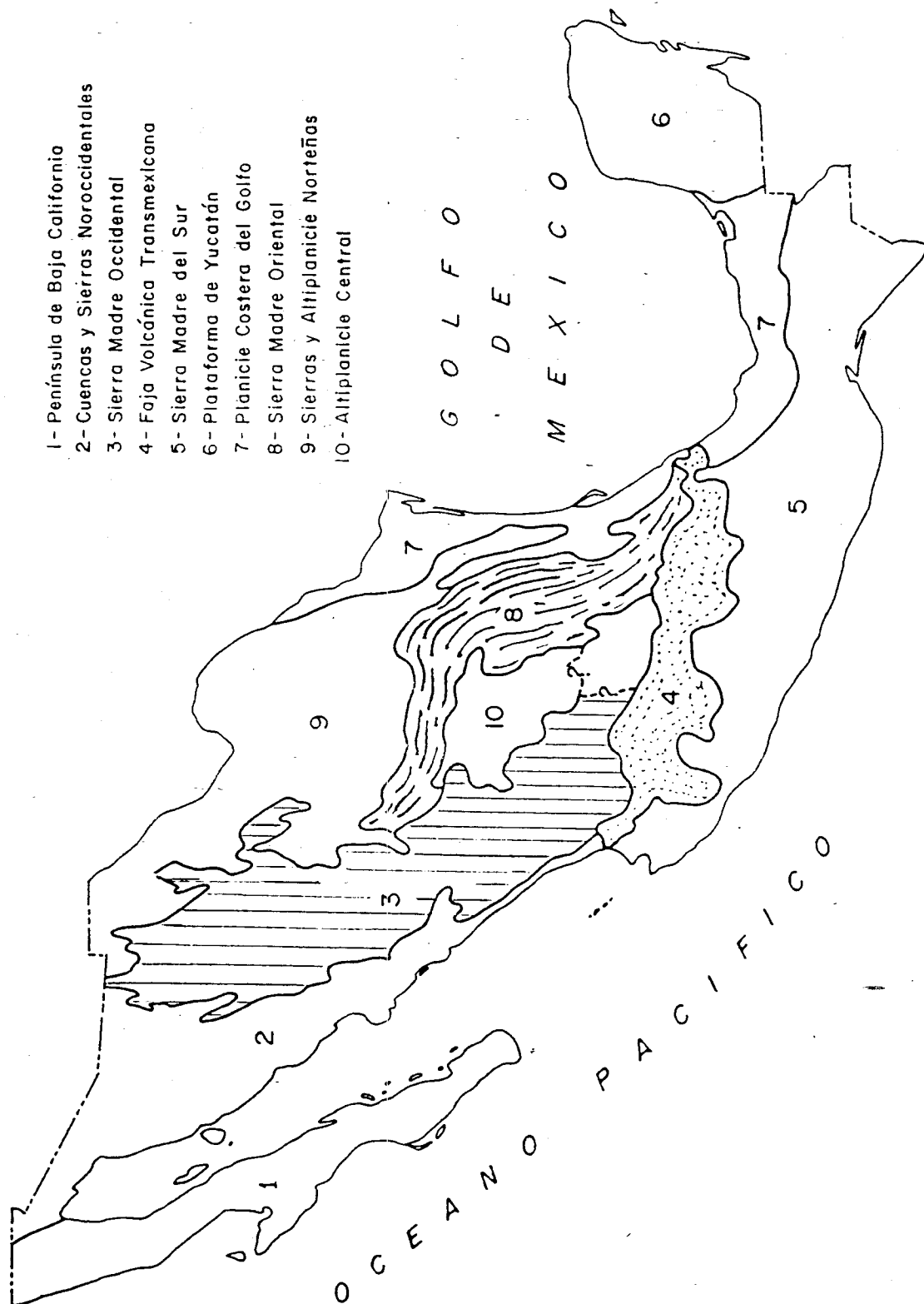
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