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- 21. The only comparative transisthmian study (34) recorded 331 mollusk species from 58 eastern Pacific stations versus 277 species from 105 Caribbean stations, collected with the same equipment. These counts suggest that diversity in the eastern Pacific is twice as large as that in the southwestern Caribbean (11), on the assumption of similar abundance of specimens (no data are given). However, mollusks are generally much less abundant in the Caribbean than in the eastern Pacific (9, 34), so even this difference in diversity may be due to sampling bias.
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- 23. The species in common were Strombina (Strombina) colinensis, Strombina (Strombina) lessepsiana, Sincola (Sincola) chiriquiensis, and Sincola (Dorsina) pigea in the late Miocene; Strombina) (Strombina) lessepsiana and Sincola (Sincola) chiriquiensis in the early Pliocene; and none dur-

ing any other interval. The subgenera characteristic of different oceans were: *Sincola (Sinaxila)*, exclusively Caribbean; *Sincola (Sincola)*, restricted to Caribbean since the late Miocene; *Strombina (Recurvina)* and *Cotonopsis (Cotonopsis)*, restricted to the eastern Pacific except for a single Pliocene species in each.

- 24. Seven specimens of Strombina (Strombina ?) aff. pumilio from a single locality in the eastern Pacific Rio Corredor section of the Armuelles Formation are most similar to Strombina (Strombina) pumilio from the late Pliocene to modern Venezuela and a single modern specimen from the Gulf of Darien. Five specimens of Cotonopsis (Cotonopsis) cf. mendozana from four localities scattered through the late Pliocene Caribbean Escudo de Veraguas Formation are most similar to Cotonopsis (Cotonopsis) mendozana, known only from the modern eastern Pacific.
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Diversity of Atlantic Coastal Plain Mollusks Since the Pliocene

Warren D. Allmon,* Gary Rosenberg, Roger W. Portell, Kevin S. Schindler

About 70 percent of tropical western Atlantic mollusk species have become extinct since the Pliocene, which has led to perceptions of a corresponding decline in diversity. However, a compilation of gastropod species from Plio-Pleistocene faunas of the United States Atlantic coastal plain and from Recent western Atlantic faunas indicates that regional diversity has not changed since the Pliocene. Gastropod diversity in the Pliocene Pinecrest Beds in Florida approximates that seen today on either coast of Florida. Gastropod diversity is not demonstrably different in the Recent tropical western Atlantic than in the Recent tropical eastern Pacific. High extinction rates must have been balanced by high origination rates.

The late Cenozoic history of mollusk diversity in the tropical and subtropical western Atlantic has traditionally been viewed as one of a decrease in species richness by a series of extinction episodes, from a Miocene-Pliocene peak to the depauperate fauna of today (1-4). In what has been termed a regional mass extinction, 70% of

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western Atlantic mollusk species are thought to have become extinct since the late Pliocene, whereas only about 30% of eastern Pacific species became extinct over the same interval (2, 3). We have compiled lists, in part based on our own new collections, of gastropod species from the Recent and Plio-Pleistocene of the western Atlantic. We have used these data to test whether extinction has led to a substantial decrease in diversity in the region, both in absolute terms and relative to the Recent eastern Pacific fauna.

The fossil database comprises lists of species from three Plio-Pleistocene faunas (Pinecrest, Caloosahatchee, and Bermont)

W. D. Allmon, Paleontological Research Institution, Ithaca, NY 14850.

G. Rosenberg, Malacology Department, Academy of Natural Sciences of Philadelphia, Philadelphia, PA 19103.

R. Portell and K. Schindler, Florida Museum of Natural History, University of Florida, Gainesville, FL 32611. *To whom correspondence should be addressed.

at two localities in southern Florida (5-7). We concentrated on the Pliocene Pinecrest fauna because it is the oldest and is therefore critical to understanding patterns of diversity in the region (8). Data for molluscan diversity in the Pliocene of the Atlantic coastal plain north of Florida were taken from previously published sources (9-11) (Table 1). The Recent database was compiled from the published literature (12); it includes all described shelled marine species of gastropods known to occur between Cape Hatteras, North Carolina, and Rio de Janeiro $(35^{\circ}N \text{ to } 23^{\circ}S, including Bermuda).$

Our results give a tally of 460 species of gastropods >5 mm in maximum adult dimension in the Pliocene Pinecrest Beds (Table 1). Of these species, 173 (38%) are known to occur in the overlying Caloosahatchee Formation (Table 2). Of the 500 total gastropod species reported from the Pinecrest, only 150 survive today, a number that reflects an extinction rate of 70%. About 95% of the extant Pinecrest species are known to occur in depths < 30 m in the Recent fauna (12). In addition, independent paleoenvironmental evidence suggests that the Pinecrest Beds were deposited in depths \leq 30 m (13). Therefore, we used 30 m as the cutoff point in making comparisons between Recent and Pinecrest diversity.

To obtain an estimate of diversity representing all size ranges, we added to the 460 species >5 mm a proportion of species with maximum observed size \leq 5 mm that corresponds to the proportion present in the Recent Florida shallow-water fauna (22%) (Table 3). We thus obtained a total of about 590 species of gastropods. About 211 bivalve species have been reported in the Pinecrest after intensive recent study (3), giving a total for bivalves and gastropods of 801 (14).

Pinecrest bivalve diversity can be used to estimate expected gastropod diversity. There are 320 species of bivalves known to occur in water depths <30 m in Florida [compiled from Florida references in (12)]. Thus, the gastropod:bivalve ratio is 2.58:1 for the Recent Florida fauna (827:320). Applying this ratio to the 211 Pinecrest bivalves, we expect a total of 544 Pinecrest gastropods. We consider our estimate of 590 Pinecrest gastropods reliable because it is within 8% of the number expected on the basis of diversity of bivalves.

Our estimates of diversity of Pinecrest gastropods are comparable to our estimates for the Recent shallow water gastropod fauna of Florida (12) (Fig. 1). The count of 460 fossil species >5 mm is about 7% higher than the 429 species in this size range now living in western Florida; the estimate of 590 Pinecrest species of all sizes is only 8% greater than the 545 living species documented in western Florida.

Similarly, numbers for the living fauna of eastern Florida are approximately 4% greater than those for the Pinecrest fauna. Thus, although the Pinecrest fauna is about 70% extinct, it is not possible to demonstrate that Recent diversity in Florida has declined as a result of Pliocene or subsequent extinctions.

The molluscan fauna of the Pinecrest has been estimated to contain as many as 1200 species. This number exceeds that of the Recent fauna of Florida by as much as three times (2, 15, 16) and approximates Recent diversity of the entire Caribbean region (2). Estimates such as these have created the impression of a tremendous diversity for the fossil beds of Florida, while underestimating Recent diversity there, which is ≥ 1147 species. Although further work may reveal that the Pinecrest contains considerably more species than we have been able to document, species are still being discovered in the Recent fauna of Florida, through both range extensions and new species (17). Therefore, we expect that additions to the fossil fauna will be approximately balanced by additions to the Recent.

differ substantially from that reported in the overlying Caloosahatchee Formation (Table 2), which has a documented fauna of 386 species of gastropods >5 mm in maximum size. This number is about 16% fewer than the 460 Pinecrest species, a difference comparable to the 12% difference today between counts for the east and west coasts of Florida (Table 3). Documented diversity is considerably lower in the Bermont than in the underlying Caloosahatchee. The Bermont beds, however, were not recognized as containing a distinct fauna until the 1960s (8) and so have been undersampled relative to the longer known Pinecrest and Caloosahatchee faunas. Not surprisingly, recent studies indicate that the Bermont contains a considerable number of undescribed or unreported species [for example, (18)].

Comparison of diversities over time is difficult because the Recent fauna has been sampled better geographically than have the fossil faunas, whereas the latter are much better sampled temporally. The count of 333 species that we report for the Pinecrest at Sarasota (Table 1) is an overestimate of local diversity in that it includes many discrete beds, each of which may

Diversity in the Pinecrest also does not

Table 1. Diversity of gastropods in Plio-Pleistocene stratigraphic units of the Atlantic coastal plain.

Unit	Locality	Total species	Species >5 mm	Data source
Bermont	Belle Glade	261	225	7
Bermont	Leisey	95	[.] 86	30*
Bermont	Total	304	259	7, 30*
James City	Total	99	99	10
Waccamaw	Total	213	200	9
Caloosahatchee	Sarasota†	42	40	*
Caloosahatchee	Total	495	386	9
Chowan River	Total	113	113	10
Bear Bluff	Total	22	22	10
Duplin	North Carolina	211	195	9
Duplin	South Carolina	216	198	9
Jackson Bluff	Total	170	144	9
Yorktown	North Carolina	117	101	9
Yorktown	Virginia	293	275	9
Pinecrest	Sarasota‡	333	333	31*
Pinecrest	Total	500	460	9, 31*
Florida Museum of Natural History collections.		†Unit 1, APAC Quarry (13).	±Units 10 to 2.	APAC Quarry

*Florida Museum of Natural History collections. †Unit 1, APAC Quarry (13). ‡Units 10 to 2, APAC Quarry (13).

Table 2. Molluscan origination and extinction for selected Plio-Pleistocene units of the Atlantic coastal plain. See (β) for stratigraphic relations. The calculations are based only on species >5 mm (Table 1). YDJP, putative synoptic time slice of late Pliocene age comprised of the Yorktown, Duplin, Jackson Bluff, and Pinecrest units; CB, Chowan River and Bear Bluff formations; JWC, James City, Waccamaw, and Caloosahatchee formations. The gastropod data are from Table 1; bivalve data, indicated by the abbreviation b, are from (β).

	Ν	lumber of spec	Apparent	Apparent	
Unit	Older unit	Younger unit	Shared	extinction (%)	origination (%)
Pinecrest-Caloosahatchee	460	386	173	62.4	55.2
Pinecrest-Caloosahatchee (b)	211	150	110	47.9	26.7
YDJP-CB	600	100	53	91.1	47.0
CB-JWC	100	462	75	25.0	83.8

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reflect an average of diversity over time (13). The diversity of a single horizon may be considerably lower than 300 species. Two studies of restricted local shallow molluscan faunas in Florida reported 181 and 155 species of gastropods, respectively (19). Thus, local diversity may not have changed significantly in Florida since the Pliocene. Confirmation of this hypothesis would require quantitative sampling in Recent sites and fossil horizons that represent comparable environments.

There is thus no evidence that molluscan diversity has changed appreciably in Florida from the late Pliocene to the Recent: the Pinecrest, Caloosahatchee, and Recent faunas do not differ substantially in number of species. The late Neogene regional mass extinction recognized in the western Atlantic (2, 3), and best represented in strata exposed in southern Florida (3), must therefore have been a phenomenon of not only high extinction but also high origination. During the time of this extinction, origination rates exceeding 50% occurred not only in Florida but also farther north along the Atlantic coastal plain (Table 2). Further research is needed to parse these originations into components of speciation, immigration, and lineage transformation. Nevertheless, the regional mass extinction in Florida may be an example of the often observed linkage between speciation and extinction (20) and may be similar in this respect to other regional episodes of extinction in the geological record (21). Similar patterns have been documented for gastropods in the late Neogene record of the southern Caribbean, where there was also substantial origination with no decline in diversity during this time (22).

Even at the level of faunal province, we see no evidence for an overall decline in



Fig. 1. Cumulative number of species by depth for Recent Floridian gastropods >5 mm in maximum size. Each species is counted from its minimum known depth in the western Atlantic. The circle representing total Pinecrest diversity (Table 1) is placed at 30 m, on the basis of independent paleoenvironmental data (*13*). Symbols: filled squares, all Florida; open squares, East Florida; and diamonds, West Florida.

Region	Total species	Species <30 m	Species <30 m and >5 mm (%)
Florida	1171	827	645 (78.0)
W. Florida	710	545	429 (78.7)
E. Florida	778	617	490 (79.4)

diversity in the western Atlantic. The Recent fauna of the tropical western Atlantic has traditionally been viewed as less diverse than that of the tropical eastern Pacific (1-4), having declined from approximate parity before the uplift of the Central American Isthmus to a depauperate condition at present. However, our data show that the western Atlantic has almost 7% more species of shelled gastropods than in the comparable latitudinal range of the tropical eastern Pacific (31°N to 6°S) (23) (2643 versus 2475); the Gulf of Mexico and Caribbean alone (30°N to 8°N) contain 2362 species (12). The tropical fauna of the western Atlantic is therefore not demonstrably less diverse than that of the eastern Pacific. If it has suffered more extinction than has the eastern Pacific, then it has also seen more origination.

Western Atlantic faunal history farther north may have been different. In North Carolina and Virginia, diversity has declined since the Pliocene. Virginia has about 100 Recent shallow water gastropod species >5 mm in maximum size, and North Carolina has about 230 such species (12), whereas the Pliocene Yorktown Formation of Virginia has 275 species in this size range (Table 1). The reason for this difference may be different climatic histories north and south of Cape Hatteras (24). Virginia and North Carolina north of Cape Hatteras have temperate Recent faunas, and climatic cooling may have caused the loss of diversity since the Pliocene (2, 3, 25). South of Cape Hatteras, however, large degrees of Neogene extinction cannot be attributed to climatic cooling alone.

Our results suggest that any environmental change associated with Plio-Pleistocene faunal change produced not only high rates of extinction but also high rates of origination (26). Other mechanisms may have been present in this system, such as changes in sea level (25, 27) or, more probably, changes in productivity caused by variations in oceanic circulation associated with closure of the Central American Isth-

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mus, the onset of Northern Hemisphere glaciation (28), or both. These mechanisms could also be used to explain a simultaneous increase in molluscan originations and extinctions in the late Pliocene.

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Divergence in Proteins, Mitochondrial DNA, and Reproductive Compatibility Across the Isthmus of Panama

Nancy Knowlton,* Lee A. Weigt, Luis Aníbal Solórzano, DeEtta K. Mills,† Eldredge Bermingham

It is widely believed that gene flow connected many shallow water populations of the Caribbean and eastern Pacific until the Panama seaway closed 3.0 to 3.5 million years ago. Measurements of biochemical and reproductive divergence for seven closely related, transisthmian pairs of snapping shrimps (*Alpheus*) indicate, however, that isolation was staggered rather than simultaneous. The four least divergent pairs provide the best estimate for rates of molecular divergence and speciation. Ecological, genetic, and geological data suggest that gene flow was disrupted for the remaining three pairs by environmental change several million years before the land barrier was complete.

Geographic isolation is thought to permit divergence and speciation by disruption of gene flow (1). Pairs of marine sister taxa separated by the Isthmus of Panama are ideal for studying these processes (2-5)because isolation of the Caribbean and the eastern Pacific is well dated and relatively recent (6, 7). This geological framework has prompted study of transisthmian sister taxa to test the accuracy of molecular clocks and to estimate the timing of other evolutionary events (3, 4, 8). It has been difficult to interpret inconsistencies and estimate possible phylogenetic differences in divergence rates (9), however, because of the limited number of taxa and characters studied. To address these problems, we investigated divergence in allozymes, mitochondrial DNA (mtDNA), and reproductive compatibility for seven shallow water transisthmian pairs of sister taxa in the snapping shrimp genus *Alpheus*.

We used the taxonomic literature to identify transisthmian pairs that were spe-

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cifically and unambiguously described as each other's closest relatives on the basis of morphological criteria (10). Collections along the two coasts and adjacent islands of central Panama at depths less than 5 m revealed unrecognized sibling species in addition to these pairs (11). In total, we examined 17 taxa (Table 1): two unambiguous pairs (P4-C4, P5-C5), three triplets (P3-C3, P3-C3'; P6-C6, P6'-C6; P7-C7, P7'-C7), and one guartet (P1-C1, P1-C2, P2-C1, P2-C2). We used shared anatomical and color pattern character states (12) to posit relations within the triplets and quartet. The result was seven morphologically defined transisthmian sister species pairs (bold in Tables 1 and 2).

For each taxon, we characterized allozymes by using conventional starch gel electrophoresis (13) and sequenced a segment of the mtDNA cytochrome oxidase I (COI) gene (14). Aggressive behavior was used as an estimate of behavioral components of reproductive compatibility (15) because snapping shrimp attack heterospecific individuals and all conspecifics except potential mates (16). We calculated genetic divergence between transisthmian pairs using Nei's D for allozymes and Kimura's corrected percent sequence divergence for mtDNA (17). We estimated divergence in behavioral compatibility by standardizing measures of tolerance and intolerance for transisthmian pairs against values observed in intraoceanic, conspecific control matings (15).

These three measures of divergence consistently support assignments of transisthmian sister species pairs on the basis of morphology and color pattern. Within the



Fig. 1. Single most parsimonious phylogenetic tree constructed on the basis of mtDNA sequences with PAUP (*18*). Transitions were given one-quarter the weight of transversions (based on the fourfold greater abundance of transitions than transversions in our data), and trees were rooted by the P7-P7'-C7 clade. Taxon codes are as in Table 1.

Smithsonian Tropical Research Institute, Apartado 2072, Balboa, Republic of Panama.

^{*}To whom correspondence should be addressed. Alternate mailing address: Smithsonian Tropical Research Institute, Unit 0948, APO AA 34002–0948, United States.

[†]Present address: Route 2, Box 538, Joshua, TX 76058.