Preliminary cladistic analysis of the bivalve family Cardiidae

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Abstract. Phylogenetic relationships within the bivalve family Cardiidae have been examined by cladistic analysis. Thirty-six of the approximately 180 cardioid supraspecific taxa are analyzed, including members of each of the generally recognized cardiid subfamilies, plus the cardioid families Lahilliidae, Lymnocardiidae, and Tridacnidae. Data for each taxon have been taken from a single species. For the outgroup, a hypothetical ancestor has been constructed from data for the carditids *Cyclocardia ventricosa* (Gould) and *Cardita variegata* Bruguière. The data consist of 54 characters and 170 character states. Results indicate that the families Lahilliidae, Lymnocardiidae, and Tridacnidae should be given subfamilial status within the Cardiidae. *Septocardia* is removed from the Cardiinae and placed in its own subfamily, and the subfamily Protocardiinae is found to be paraphyletic. The Laevicardiinae, as proposed by Keen (1936, 1951, 1969, 1980), is shown to be polyphyletic: *Cerastoderma* is a lymnocardiine; *Dinocardium* is a cardiine; *Clinocardium* is the type genus of the Clinocardiinae. In addition to *Laevicardium*, only *Habecardium* and *Fulvia* remain in Laevicardiinae. The Trachycardiinae is found to be a monophyletic taxon within the Cardiinae. The subfamilies Clinocardiinae, Tridacninae, Lymnocardiinae, and Fraginae form a monophyletic clade. *Sawkinsia*, long considered a tridacnid, belongs with the Cardiinae.

Bivalves of the family Cardiidae, or cockles, display a wide spectrum of shell shapes, ribbing and ornamentation patterns, hinge morphologies, and numerous other conchological features. Their complex morphology, accompanied by their good fossil record, allows the cardiids to be evaluated evolutionarily, ecologically, functionally, and phylogenetically in considerable detail.

The higher-level taxonomy of the Cardiidae has been more thoroughly studied than that of most other groups of bivalves (Dall, 1901; Stewart, 1930; Keen, 1936, 1937, 1951, 1969a, 1980; Fischer-Piette, 1977; Kafanov and Popov, 1977; Popov, 1977; Wilson and Stevenson, 1977; Voskuil and Onverwagt, 1989). Kafanov and Popov (1977) made the only detailed attempt to reconstruct the phylogenetic history of the group.

Traditionally, the superfamily Cardioidea comprises: (1) the Cardiidae; (2) the extinct, southern hemisphere Lahilliidae; (3) the brackish-water Lymnocardiidae, confined to eastern Europe and southwestern Asia; (4) the Tridacnidae, or giant clams [Keen (1969b), Kafanov and Popov (1977), and Scarlato and Starobogatov (1979) have placed the giant clams in a separate superfamily]. Kafanov and Popov (1977) contended that the Lahilliidae belonged to the Arcticoidea. However, as noted by Finlay and Marwick (1937) and Marwick (1944), the hinge of *Lahillia* is of the cardiid, not arcticoid type; the lack of external ornament is apparently a case of convergence. This classification has not been taken for granted, and representatives of the Tridacnidae, Lahilliidae, and Lymnocardiidae are included in this analysis.

Additionally, *Cardium acuticostatum* d'Orbigny, 1842, is included. Wilckens (1904) placed the Cretaceous C.

acuticostatum in Cardium (Bucardium), which Keen (1980) indicates is known from only the Miocene to Recent.

Boss (1971), Kafanov and Popov (1977), Keen (1980) and Ponder *et al.* (1981) placed the enigmatic *Hemidonax* in the Cardioidea. However, Scarlato and Starobogatov (1979) argued that *Hemidonax* is aligned with the Donacidae. After examination and comparison of the shell and of the anatomy (both external and internal) of *Hemidonax* to both cardiids and donacids, I cannot justify placing *Hemidonax* as a member of the Cardioidea. However, neither can I place *Hemidonax* within the Donacidae. Instead, I favor placing *Hemidonax* as *incertae cedis* within the order Veneroida, until a phylogenetic analysis of the Veneroida is undertaken.

Virtually all cardiid taxonomy is based on hard parts, with the exception of Starobogatov's (In: Kafanov and Popov, 1977) study of stomach structure. Most of what is known about the anatomy of cardiids comes from the study of the common cockle, *Cerastoderma edule* (Linnaeus) (see Ménégaux, 1890; Johnstone, 1899; Zugmayer, 1904; Küpfer, 1915; Roche, 1925; Atkins, 1937; Graham, 1949), which is taken as a model for the entire family. Furthermore, Russian malacologists (Kafanov and Popov, 1977; Popov, 1977; Taktakishvili, 1987) place *Cerastoderma* within the subfamily Lymnocardiinae - which, less *Cerastoderma*, Keen (1969a, 1980) considered to be a separate family within the superfamily Cardioidea. The only treatment of comparative anatomy of the cardiids is that of Pelseneer (1911).

The goals of this study are to: (1) determine the status and content of subfamilies erected by previous workers; (2) propose a preliminary phylogenetic hypothesis for the family. Characters and character states are briefly described herein and will be treated fully in future publications.

MATERIALS AND METHODS

There are about 180 generally accepted cardioid genera and subgenera (see Keen, 1969a, 1980; Kafanov and Popov, 1977; Vokes, 1980; Taktakishvili, 1987). It is not feasible currently to run a computer-driven cladistic program for such a large number of taxa. The 36 taxa chosen in this study include at least one representative of each of the cardiid subfamilies accepted by Keen (1969a, 1980), Kafanov and Popov (1977), and Voskuil and Onverwagt (1989). As stated above, the tridacnids (*Tridacna*), lahilliids (*Lahillia*), and lymnocardiids (*Hypanis*) are also represented in this analysis (the suffix -ids is used in a vernacular sense until their taxonomic placement is discussed thoroughly). Kafanov and Popov's (1977) and Keen's (1980) classification scheme for the taxa analyzed in the present study is given in appendices 1 and 2.

A cladistic analysis of the 36 taxa with 54 characters comprising 170 character states (appendices 3 and 4) was made using PAUP 3.0d (Swofford, 1989). The accelerated transformation option (ACCTRAN) was used, and steps were not added to terminal taxa with polymorphisms. Synapomorphies for each node are presented in appendix 5.

Character states were encoded from a single species of each genus or subgenus. Most of the taxa are represented by their type species. Exceptions are those taxa for which

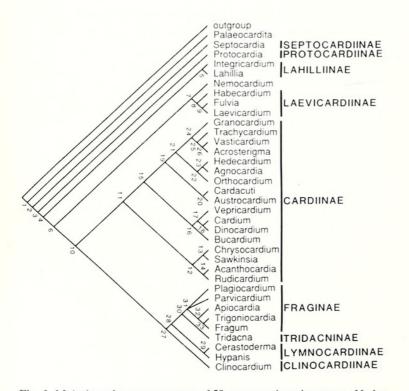


Fig. 1. Majority-rule consensus tree of 50 most parsimonious trees. Nodes 21 and 22 supported by 60% of trees; nodes 28 and 30 supported by 70% of trees; node 31 supported by 90% of trees; all other nodes supported by 100% of trees. Synapomorphies supporting each node given in appendix 5.

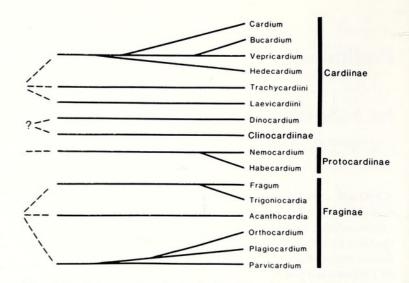


Fig. 2. Evolutionary scenario for the Cardiidae from Kafanov and Popov (1977). Only those taxa included in both the present study and Kafanov and Popov (1977) are shown.

(1) material of the type species was unavailable, or (2) anatomical material was not available for the type (e.g. *Nemocardium*, for which the type species is extinct), but was available for another species generally assigned to that taxon. Taxa represented by species other than the type are listed in appendix 6. Therefore, this analysis should be taken as a phylogeny for these species only. Character states presented may not be constant throughout all species of a given genus. Because there is considerable disagreement over what constitutes a genus or a subgenus in the Cardiidae, all terminal taxa will be considered to have equal rank as genera; no distinction will be made between genera and subgenera, except as noted in the text.

Citations in the character list (appendix 3) refer to previous discussions of that character. Except for the information on *Cyclocardia* (see below), character 12, and character 8 for *Parvicardium* (from Pelseneer, 1911), all character states were encoded from examination of specimens.

The Cardiidae are generally accepted as having been derived from a member of the Carditoidea [Cox, 1949; Keen, 1969a, 1980; Newton, 1986; but see Morris (1978) and Morris et al. (1991)]. These authors have postulated an evolutionary scenario of Palaeocardita originating from some primitive carditid or permophorid, with Septocardia then originating from Palaeocardita. Palaeocardita is usually placed with the Carditidae (Chavan, 1969). However, examination of the one species of Palaeocardita available to me, Palaeocardita silberlingi Newton et al., has led me to place this species within the ingroup Cardiidae, on the basis of its cardinal teeth, which are arranged as in Septocardia and Protocardia [see Newton et al. (1987) for a discussion of this species]. Therefore, to represent the outgroup, a hypothetical ancestor was constructed with information from the Recent carditids Cardita variegata Bruguière and Cyclocardia ventricosa

(Gould). Character states for *C. ventricosa* were taken from information in Yonge (1969). For characters 6 and 30, the two carditids provided conflicting information, and hence the states are scored as missing ("?").

Most characters are unordered. It was possible to construct character state trees based on ontogeny for characters 23 (shell shape), 24 (ribbing pattern), 29 (mosaicostracum), and 40 (rib flares).

RESULTS AND DISCUSSION

Fifty most parsimonious trees of 208 steps (consistency index = 0.566) were found. The 50% majority-rule consensus tree is presented in figure 1, which can be compared with two previously presented evolutionary scenarios. Kafanov and Popov (1977) produced a phylogram based on two key characters : stomach structure (analyzed by Ya. I. Starobogatov) and Popov's (1977) work on shell microstructure. Kafanov and Popov (1977) considered 38 taxa [only those taxa represented in both my analysis and that of Kafanov and Popov (1977) are shown in figure 2; they considered neither the origin of the Cardiidae nor the group's Mesozoic history]. Only nine of the 28 extant taxa were examined for stomach structure. Starobogatov's study of stomach structure rests heavily on the presence/absence and position of sorting areas, as described by Purchon (1960a). Purchon's (1960a) description of the cardiid stomach came from the study of Cerastoderma edule by Graham (1949). Starobogatov (In: Kafanov and Popov, 1977) stated that the SA-3, or posterior sorting area, is absent in Cerastoderma based on examination of C. glaucum (Bruguière) (Ya. I. Starobogatov, pers. comm.) and Hypanis. However, the posterior sorting area (labeled SAP) is the most prominent structure in Graham's (1949) figure of the stomach of C. edule. While promoting the utility of using stomach structure to elucidate the higherlevel phylogeny of the Bivalvia (Purchon, 1959, 1960a, b), Purchon (1960a:481) warns that "...it is not easy to make an objective analysis of the occurrence and identities of the various sorting areas. The presence or absence, and the degree of development of the various sorting areas has a profound effect on the appearance of the interior of the stomach, and could obscure more fundamental issues such as the course taken by the major typhlosole and the intestinal groove...the occurrence of sorting areas can only be used with the greatest caution for phylogenetic purposes." Although seven of the 20 anatomical characters in the present analysis concern the stomach, none relate to the sorting areas.

Popov's (1977, 1986) classification of bivalve shell microstructure conflicts with those of Carter (1980, 1989), Carter and Clark (1985), Carter and Lutz (1989) and Watabe (1984). The only systemically useful microstructural characters that I have found so far concern the relationship of the ornament to the rest of the shell (characters 28, 29 and 40).

The only cardiid phylogeny suggested by Keen is found in her description of cardiid evolution (Keen, 1980). I have constructed a phylogram (Fig. 3) based on that description.

In the present analysis, *Palaeocardita silberlingi* is located at the base of the cladogram, followed by *Septocardia*, *Protocardia*, and then the rest of the Cardiidae. This topology is in agreement with the ideas of early cardiid evolution suggested by Cox (1949), Keen (1969a, 1980) and Newton (1986). However, the monophyly of *Palaeocardita* is questionable (C. R. Newton, pers. comm.), and the more common species, including the type *P. austriaca* (Hauer) and *P. crenata* (Goldfuss) could be carditoids, whilst *P. silberlingi* is a cardiid. Due to the uncertainty of the taxonomy of *Palaeocardita*, I refrain from placing *P. silberlingi* in a higher taxon within the Cardiidae.

Septocardia was placed in its own family, the Septocardiidae, in the superfamily Tridacnoidea by Kafanov and Starobogatov (In: Kafanov and Popov, 1977). Septocardia is clearly a primitive cardiid and it does not share any of the derived features of *Tridacna*. Likewise, Septocardia does not belong in the derived subfamily Cardiinae as indicated by Keen (1969a, 1980). Septocardia is here placed in the cardiid subfamily Septocardiinae.

The subfamily Protocardiinae has been understood to include the genera *Protocardia*, *Integricardium*, *Jurassicardium*, and *Nemocardium* (Kees, 1969a, 1980). My results indicate that this is a paraphyletic group. *Integricardium* is more closely related to *Lahillia*. The paraphyly of the Protocardiinae has been acknowledged implicitly for some time. McLearn (1933) erected *Onestia* as a subgenus of *Integricardium*; the former was considered a genus by McLearn

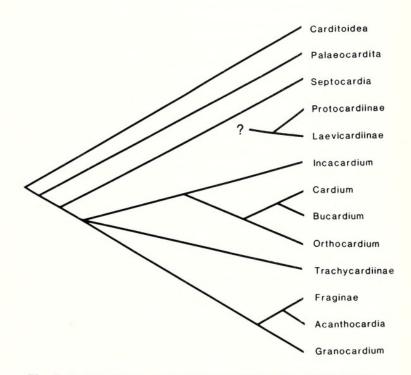


Fig. 3. Evolutionary scenario reconstructed from Keen (1980:24-30).

(1945) and Day (1978), but not by Keen (1969a, 1980). Day (1978) postulated that *Integricardium* is ancestral to *Onestia*, which is in turn ancestral to *Lahillia*. He also placed *Onestia* in the Lahilliidae. Present results indicate that the family Lahilliidae should be relegated to a subfamily (Lahilliinae) within the family Cardiidae (as originally proposed by Finlay and Marwick, 1937), and should include *Integricardium*.

Nemocardium is the sister taxon to the Laevicardiinae. This group is in turn the sister taxon to the rest of the cardiids. Because of its change in ribbing pattern from that similar to Nemocardium as a juvenile, to that of Fulvia as an adult, Habecardium has been recognized as transitional from the Protocardiinae to the Laevicardiinae (Glibert and van de Poel, 1970; Keen, 1980). Glibert and van de Poel (1970) erected Habecardium as a subgenus of Laevicardium, into which some of the species of Habecardium had been placed previously. Keen (1980) placed *Habecardium* as a subgenus of Nemocardium. Popov (1977) and Kafanov and Popov (1977) also placed Habecardium as a subgenus of Nemocardium, but did not recognize it as transitional to Laevicardium and Fulvia, placing the latter two taxa in the Cardiinae. Besides ribbing pattern (24:2), the Laevicardiinae are united by the number of ctenidial plicae (5:2), tentacles that extend only to the bottom of the posterior adductors (10:0), presence of complex eyes (12:1), a centrally located right caecum (18:1), and shape of the cardinal teeth (43:3 and 45:3).

Keen (1969a, 1980) placed all the Cenozoic protocardiines in the genus *Nemocardium*. Other authors (Fischer-Piette, 1977; Popov, 1977; Wilson and Stevenson, 1977; Noda, 1988; Voskuil and Onverwagt, 1989) have raised some of the subgenera to the generic level. It is suspected strongly that the subtraction of *Habecardium* from *Nemocardium* would leave the latter as a monophyletic group. I decline here to place *Nemocardium* within a subfamily. A systematic analysis which includes all of the subgenera of *Nemocardium* as in Keen (1969a, 1980) represented, plus the Laevicardiinae, is in progress.

Jurassicardium is a monotypic taxon known from only a few specimens. Only the type material is sufficiently well preserved to be of systematic use, and I have not examined it.

The remainder of the Cardiidae comprise those forms typically accepted to constitute the taxa Cardiinae, Trachycardiinae, Fraginae, Clinocardiinae, Lymnocardiidae, and Tridacnidae. Two monophyletic clades can be distinguished within this unnamed taxon. One clade, here considered the subfamily Cardiinae, contains the taxa placed in the Trachycardiinae and most of the taxa placed in the Cardiinae by Keen (1969a, 1980), and all of taxa placed in the Cardiinae (except for *Laevicardium* and *Fulvia*) by Kafanov and Popov (1977). In agreement with Kafanov and Popov (1977), the present results indicate that *Trachycardium* and the related taxa *Acrosterigma* and *Vasaticardium* do not constitute a separate subfamily but are members of the Cardiinae. The least derived monophyletic group within the Cardiinae contains the taxa *Acanthocardia*, *Rudicardium*, *Sawkinsia*, and *Chrysocardium*. Synapomorphies of this clade are cardiiform shell shape (23:5), tuberculate spines (28:5) and irregular cross-striae (30:1). *Rudicardium* is considered either a subgenus of *Acanthocardia* (Keen, 1969, 1980; Popov, 1977) or a synonym of it (Voskuil and Onverwagt, 1989). These two taxa are united by a suite of hinge characters: incomplete anterior cardinal socket (42:1); shape of the cardinal teeth (43:8 and 45:10); hinge plate overlapping the right posterior lateral socket (48:1).

Cox (1941) erected Sawkinsia as a genus of cardiid. Vokes (1953) placed Sawkinsia in the Tridacnidae, and was followed by Rosewater (1965), Keen (1969b), and Jung (1976). Stasek (1962) considered the resemblance between Sawkinsia and the tridacnid Hippopus to be a case of convergence. Sawkinsia does not share any of the derived characters of Tridacna, nor any of Tridacna's notable autapomorphies: (1) there is no loss of the anterior lateral teeth; (2) the spines are tubercles, not wide and gently curved; (3) nor is there any evidence of the rotation of the shell about the animal. According to the present phylogenetic hypothesis, Sawkinsia is a member of the subfamily Cardiinae.

Woodring (1982) erected the genus *Chrysocardium* in the subfamily Fraginae based on a single left valve. *Chrysocardium* shares not only a lunule flap touching the beak (25:3) with *Sawkinsia*, but three characters found nowhere else in the Cardiidae: hinge inversion (36:1) (described for *Sawkinsia* by Cox, 1941); weak myophorous buttress (38:1); double keel (41:1). *Chrysocardium* should be considered a synonym of *Sawkinsia*, however *C. aurum* Woodring, appears to be valid. Except for missing data, the characters for two taxa are scored identically (see appendix 4).

The next monophyletic group includes *Bucardium*, *Cardium*, *Vepricardium*, and *Dinocardium*, and is united by seven synapomorphies. The close relationship of the first three taxa to each other has been recognized by numerous authors (see Keen, 1969a; Kafanov and Popov, 1977). The position of *Dinocardium*, however, remains uncertain. Keen (1951, 1969a, 1980) placed *Dinocardium* in the Laevicardiinae. Kafanov and Popov (1977), in dismantling the Laevicardiinae, tentatively placed *Dinocardium* in the Cardiinae; Kafanov (1980:298) called the taxonomic position of *Dinocardium* "most mysterious."

The next node within the Cardiinae contains Austrocardium and Cardium acuticostatum, and is united by lack of lunule flap (25:0) and shape of the anterior cardinal (45:5). Freneix and Grant-Mackie (1978) erected the Cretaceous Austrocardium as a monotypic taxon. Wilckens (1904) placed the Cretaceous form C. acuticostatum in Cardium (Ringicarcium) [=Cardium (Bucardium)], which is otherwise a Miocene to Recent taxon (Keen, 1969a, 1980). The results of my analysis indicate that C. acuticostatum belongs in Austrocardium. There are three other Cretaceous species that differ little from C. acuticostatum and Austrocardium. These are: (1) C. denticulatum Baily, which was placed by Dartevelle and Freneix (1957) in Acanthocardia (Acanthocardia); (2) C. (Bucardium) lillei Freneix and Grant-Mackie (specimens of which had originally been described as C. acuticostatum); (3) Schedocardia ? waiparana Freneix and Grant-Mackie. These species should be placed provisionally in Austrocardium, as they share the apomorphies of Austrocardium but not those of either Bucardium or Acanthocardia. Except for cases of missing data, the posterior gape of C. acuticostatum (33:1) is the only character not scored identically to that of Austrocardium (appendix 4). The posterior gape is convergent with that of Cardium.

Hedecardium, Orthocardium, and Agnocardia form a monophyletic clade. Hedecardium has been considered variously as a subgenus of Vepricardium (Keen, 1969a, 1980), a genus closely related to Vepricardium (Popov, 1977), and a genus in the Protocardiinae (Marwick, 1960; Maxwell, 1978). The latter authors derived Hedecardium from Nemocardium on the basis of its discrepancy in thickness in the ribs across the shell. However, the rib discordance in Hedecardium is not comparable to that in Nemocardium. In Hedecardium, four to six ribs on the posterior slope are split with a furrow running down the middle, and the remaining posterior ribs are reduced in strength (as happens numerous times in the Cardiidae; it was the basis of Keen's [1936] subfamily Laevicardiinae, shown to be polyphyletic). This change from all ribs of equal width to the rib discrepancy seen in Hedecardium can be seen in the growth stages of a single shell. Likewise, the early growth stages of Hedecardium are circular, and circular shells are unknown in any form of Nemocardium. As a juvenile, Hedecardium would strongly resemble Orthocardium. It is recommended that Hedecardium and Orthocardium be considered as distinct genera.

Orthocardium has been considered a subgenus of Vepricardium (Keen, 1969a) or of Cardium (Keen, 1980) or a genus of fragine (Popov, 1977; Kafanov and Popov, 1977). Here, Orthocardium is united with Agnocardia and Hedecardium by concave ribs (32:1), a condition otherwise unknown in the Cardiidae.

The last clade within the Cardiinae comprises *Granocardium*, *Trachycardium*, *Acrosterigma*, and *Vasticardium*, and is united by one character, ovate shell shape (23:3). The latter three taxa, united by three synapomorphies, are usually placed in the Trachycardiinae (Keen, 1969a, 1980) or the tribe Trachycardiini (Kafanov and Popov, 1977) within the Cardiinae. Current results support the latter.

The other major clade of cardiids is united by five synapomorphies and generally contains forms that have been assigned to the Clinocardiinae, Lymnocardiidae, Tridacnidae, and Fraginae. Of these the least derived is *Clinocardium*. This taxon was placed in the Laevicardiinae by Keen (1951, 1969a, 1980), but has come to be accepted as the type of the subfamily Clinocardiinae (Kafanov and Popov, 1977; Kafanov, 1980; Voskuil and Onverwagt, 1989). The next node encountered is *Cerastoderma* and *Hypanis*, which is the sister taxon to the *Tridacna* and Fraginae. Synapomorphies of lymnocardiids + *Tridacna* + Fraginae are medium labial palps (1:1), functional byssus in adult (6:2), posterior cardinal socket angle (44:1) and shape of the anterior cardinal (45:9). Yonge's (1936) and Stasek's (1962) suggestion that the ancestry of *Tridacna's* was close to that of *Cerastoderma* is upheld by the results. Giant clams should be considered as the subfamily Tridacninae within the Cardiidae.

Cerastoderma and *Hypanis* form a monphyletic group. Therefore, as has been argued by eastern European malacologists for some time (Kafanov and Popov, 1977; Popov, 1977; Taktakishvili, 1987; Basch, 1990), the brackish-water forms should be subfamily Lymnocardiinae, and contain *Cerastoderma*. Furthermore, the results support Kafanov and Popov's (1977) contention that the Lymnocardiinae are related closely to the Fraginae.

Five of the seven characters that unite Tridacna and the Fraginae are anatomical: tentacle pattern (9:2); large valvule (11:2); centrally located style sac (15:1); presence of a raised bar on the stomach floor (17:1); posteriorly located left caecum (19:0). The Fraginae is united by the presence of ventral appendages on the foot (7:1), absence of a periphonal suture (8:1), and presence of a mosaicostracum (29:1). The least derived taxa, Plagiocardium and Parvicardium, were placed in the Fraginae by Kafanov and Popov (1977) but in the Cardiinae by Keen (1969a, 1980). Apiocardia, Trigoniocardia, and Fragum are united by eight synapomorphies, all based on hinge characters. Finally, Trigoniocardia and Fragum are united by six synapomorphies, five of which are anatomical characters: short labial palps (1:0); fewer than ten ridges on the palps (2:1); the inner palp connected to the bottom of the inner demibranch (3:1); fewer than 20 ctenidial plicae (5:0); type 4 gut (13:3); quadrate shell shape (23:2).

From the above discussion it can be concluded that the family Cardiidae includes nine subfamilies: Septocardiinae; Protocardiinae; Lahilliinae; Laevicardiinae; Cardiinae; Clinocardiinae; Tridacninae; Lymnocardiinae; Fraginae. Taxa usually assigned to the Trachycardiinae form a monophyletic group within the Cardiinae. *Dinocardium* is a cardiine closely related to *Cardium* and *Vepricardium*. *Sawkinsia* (=*Chrysocardium*) is transferred from the Tridacninae to the Cardiinae, as it is related closely to *Acanthocardia* and *Rudicardium*.

The Protocardiinae (as presented in the Treatise on Invertebrate Paleontology) is paraphyletic. *Integricardium* is a member of the Lahilliinae. *Nemocardium* is the sister taxon to the Laevicardiinae. The subfamilies Clinocardiinae, Tridacninae, Lymnocardiinae, and Fraginae form a monophyletic group.

ACKNOWLEDGMENTS

Numerous people and institutions have allowed me to examine their collections and have lent specimens. Foremost, I would like to thank the Academy of Natural Sciences of Philadelphia, my supervisor at the Academy, G. Rosenberg, and G. Davis and the rest of the staff of the depatments of Malacology and Paleontology of that institution. I would like to thank the following people and institutions for allowing me to examine their collections: T. Waller, W. Blow, R. Hershler, R. Houbrick, J. Harasewych (United States National Museum); M. Hinckley, N. Eldredge (American Museum of Natural History); C. Coney (Los Angeles County Museum of Natural History); M. Baker, M. Nitecki, J. Voight, R. Bieler (Field Museum of Natural History). I would like to thank the following people for lending specimens: W. Ponder, Australian National Museum; A. Beu, New Zealand Geological Survey; J. A. Grant-Mackie, University of Auckland; S. Freneix, J. -C. Fisher, and Y. Gayrard, Museum National d'Histoire Naturelle, Paris. I examined specimens of Palaeocardita and Septocardia at C. R. Newton's laboratory at Syracuse University. I learned techniques for the study of shell microstructure at J. G. Carter's laboratory at the University of North Carolina-Chapel Hill.

The following agencies have generously given me financial assistance for my research: The Academy of Natural Sciences of Philadelphia (Jessup Fund), the National Capitol Shell Club, the Western Society of Malacologists, the Paleontological Society, Conchologists of America, the Santa Barbara Shell Club, the Hinds Fund of the University of Chicago, Sigma Xi, the Gurley Fund of the Department of Geophysical Sciences of the University of Chicago, and the Amoco Corporation. During part of my graduate study I was supported by a grant from the National Science Foundation (EAR-90-05744) to D. Jablonski. I would also like to thank the following people at the University of Chicago who have assisted me in my research: D. Jablonski, M. LaBarbera, O. Draughn, K. Roy, P. Wagner, and D. Miller.

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Date of manuscript acceptance: 25 September 1991

Classification of taxa used in this study following Kafanov and Popov (1977). Taxa not listed are not discussed therein.

Superfamily Cardioidea Family Cardiidae Subfamily Protocardiinae Protocardium Integricardium Nemocardium Habecardium Subfamily Cardiinae Tribe Cardiini Cardium Tribe Vepricardiini Vepricardium Agnocardia Bucardium Hedecardium Tribe Trachycardiini Trachycardium Acrosterigma Vasticardium Tribe Laevicardiini Fulvia Tribe Dinocardiini Dinocardium Subfamily Clinocardinae Tribe Clinocardiini Clinocardium Subfamily Fraginae Tribe Fragini Trigoniocardia Apiocardia Fragum Tribe Acanthocardiini Rudicardium Acanthocardia Tribe Parvicardiini Plagiocardium Parvicardium Orthocardium Subfamily Lymnocardiinae Tribe Cerastodermatini Cerastoderma Tribe Hypanini Hypanis Superfamily Tridacnoidea Family Septocardiinae Septocardia Family Tridacnidae Tridacna Superfamily Arcticoidea Family Lahilliidae Lahillia

APPENDIX 2.

Classification of taxa used in this study according to Keen (1969a, b; 1980). Taxa not listed are not considered in any of Keen's (1969a, 1969b, 1980) papers.

Superfamily Cardiacea Family Cardiidae Subfamily Cardiinae Septocardia Granocardium Vepricardium (Vepricardium) Vepricardium (Orthocardium) Vepricardium (Hedecardium) Cardium (Cardium) Cardium (Bucardium) Acanthocardia (Acanthocardia) Acanthocardia (Rudicardium) Acanthocardia (Agnocardia) Plagiocardium Parvicardium Subfamily Trachycardiinae Trachycardium Acrosterigma (Acrosterigma) Acrosterigma (Vasticardium) Subfamily Protocardiinae Protocardia Integricardium Nemocardium (Nemocardium) Nemocardium (Habecardium) Subfamily Fraginae Fragum Trigoniocardia (Trigoniocardia) Trigoniocardia (Apiocardia) Subfamily Laevicardiinae Laevicardium (Fulvia) Laevicardium (Dinocardium) Cerastoderma Clinocardium Family Lahillidae Lahillia Family Lymnocardiidae Hypanis Superfamily Tridacnacea Family Tridacnidae Tridacna Sawkinsia

APPENDIX 3.

List of characters and character states.

I. Anatomy

- A. Labial palps
 - 1. Length: 0) short, 1) medium, 2) long
 - Number of ridges on palps: 0) ridges absent, 1) <10, 2) 10 19,
 3) 20 29, 4) >29
 - 3. Connection of inner palp with ctenidia: 0) connects behind inner demibranch, 1) connects with bottom of inner demibranch
- B. Ctenidia
 - 4. Inner demibranch/outer demibranch relation: 0) Outer demibranch does not overlay inner demibranch, 1) Outer demibranch partially underlain by inner demibranch
 - 5. Number of plicae: 0) <20, 1) 20 39, 2) 40 59, 3) 60 79, 4) 80 - 99, 5) >99
- C. Foot
 - Byssal apparatus: 0) absent in adult, 1) present in adult, but nonfunctional, 2) functional in adult (see Pelseneer, 1911)
 - 7. Ventral appendages: 0) absent, 1) present
- D. Siphons and tentacles
 - 8. Periphonal suture: 0) present, 1) absent (see Pelseneer, 1911)
 - 9. Tentacle pattern: 0) absent, 1) numerous, in both mantle fold and on siphonal area, 2) numerous, in mantle fold only, 3) few, in mantle fold and siphonal area, 4) few on siphonal area only
 - Dorsalmost extension of tentacles: 0) bottom of adductors, 1) to middle of adductors, 2) to top of adductors, 3) beyond top of adductors
 - 11. Valvule (see Pelseneer, 1911) 0) absent, 1) small, 2) large
 - Eyes (see Kishinouye, 1894; Nagel, 1897; Zugmayer, 1904; Weber, 1908; Pelseneer, 1911; Braun, 1954) 0) simple, 1) complex
- E. Gut
 - 13. 0) Type 1, 1) Type 2, 2) Type 3, 3) Type 4, 4) Type 5
- F. Stomach (see Graham, 1949 and Purchon, 1960a, for general descriptions of bivalve stomachs)
 - 14. T3 (tertiary typhlosole): 0) absent, 1) present
 - 15. Position of style sac: 0) posterior, 1) central
 - 16. T1 (major typhlosole) curved (see Nakazima, 1964): 0) yes, 1) no
 - 17. raised bar: 0) absent, 1) present
 - 18. Position of right caecum: 0) right side of stomach, 1) central
 - Position of left caecum: 0) posterior to right caecum 1) caeca parallel, 2) anterior to right caecum
 - 20. Accessory left caeca: 0) absent, 1) present
- II. Shell
 - A. General
 - 21. Posterior margin: 0) digitate, 1) crenulate, 2) smooth
 - 22. Rib number: 0) absent, 1) less than 70, 2) greater than 70
 - 23. shell shape: 0) carditaform, 1) quadrate long, 2) quadrate short,
 - 3) ovate, 4) circular, 5) cardiiform, 6) oval, 7) cerastiform,8) trigonal, 9) elliptical, 10) oblique

Character state tree: ((((5,(7,8)6)4,10,3)2,9)1)0

24. Anterior/central rib pattern: 0) concentric, 1) radial, equal in width to posterior ribs, 2) concentric, changing to radial, 3) rib discordance, 4) radial, thinner than posterior ribs, 5) none

Character state tree: (((2)4,5)0,3)1

- Lunule flap (see Kafanov, 1980, pp. 298-299): 0) absent, 1) raised, does not block beak, 2) blocks beak but does not touch it, 3) touches beak, 4) strongly folded over beak
- 26. Ridges on lunule flap: 0) absent, 1) present
- 27. Growth line strength: 0) strong, 1) weak
- Spines: 0) lumpy nodes, 1) none, 2) round knobs, 3) A-shaped, separate shell layer from ribs, 4) hollow posterodorsally, 5) tubercles, 6) gently curved, 7) hollow keel

Spines are defined as emanating from the top of the ribs. The "spines" of trachycardiines emanate from the side of the ribs, and are considered separately (character 40).

29. Mosaicostracum ("spines" on fragines): 0) none, 1) beads, 2) scales

Mosaicostracum was first described by Hamilton (1969). Keen (1980) and Vokes (1977, 1989) referred to this layer as the intritacalx (D'Attilio and Radwin, 1971). Carter (1989) considers intritacalx synonymous with mosaicostracum. This character is linearly ordered.

- 30. Cross-striae: 0) simple, 1) irregular, 2) absent
- 31. Internal rib expression: 0) weak, 1) strong
- 32. Ribs concave: 0) no, 1) yes
- 33. Posterior gape: 0) absent, 1) present
- 34. Dorsal nymph extension, 0) absent, 1) present
- 35. Posterior umbonal buttress: 0) absent, 1) present
- 36. Hinge inversion: 0) absent, 1) present
- 37. Nymph groove: 0) absent, 1) present
- 38. Myophorous buttress: 0) strong, 1) weak, 2) absent
- 39. Adductor scar relief: 0) strong, 1) weak
- 40. Rib flares ("spines" on trachycardiines): 0) absent, 1) strong,2) weak
- This character is linearly ordered.
- 41. Double keel: 0) absent, 1) present
- B. Hinge teeth
 - 1) Right cardinal teeth
 - 42. Anterior socket: 0) complete, 1) incomplete
 - 43. Posterior cardinal shape: 1) 1, 2) 2, 3) 3, 4) 4, 5) 5, 6) 6, 7) 7, 8) 8, 9) 9
 - 44. Angle of posterior cardinal socket to horizontal: 0) greater than 40 degrees, 1) equal or less than 40 degrees
 - 45. Shape of anterior cardinal: 0) 0, 1) 1, 2) 2, 3) 3, 4) 4, 5) 5, 6) 6, 7) 7, 8) 8, 9) 9
 - Rest of hinge
 - 46. Right anterior lateral teeth ventral tooth continues up into umbo:0) no, 1) yes
 - 47. Right anterior lateral teeth ventral tooth inserts into socket: 0) no, 1) yes
 - 48. Right posterior laterals: hinge plate overlaps socket: 0) no, 1) yes
 - 49. Anterior lateral teeth: 0) absent, 1) present
 - 50. Left posterior lateral teeth: 0) weak or absent, 1) strong
 - 51. Nymph overlies posterior cardinals: 0) no, 1) yes
 - 52. Left anterior lateral horizontal: 0) no, 1) yes
 - 53. Left anterior lateral socket: 0) absent or weak, 1) strong
 - 54. Left posterior lateral socket: 0) absent or weak, 1) strong

Data matrix for cladistic analysis. "?" signifies missing data. X indicates character state 10. A-D indicate polymorphisms. A: states 1 and 2; B: states 0 and 1; C: states 2 and 3; D: states 1, 2, and 3.

	1	2	3	4 5
Taxa	12345678901234			0 1 2 3 4 5 6 7 8 9 0 1 2 3 4
outgroup	10010?000?0001	00000111010?0	000?00000000?	0000?????00??0?
Palaeocardita	???????????????????????????????????????	??????11010?	000000000000000?	000101???00??00
Septocardia	???????????????????????????????????????	??????1111100	000000000000000?	000101???000?00
Protocardia	???????????????????????????????????????	??????1110300	0102000000020	000101000100000
Integricardium	???????????????????????????????????????	?????209010	11020?000002?	00010100010000?
Lahillia	???????????????????????????????????????	?????20950?	11020?0000020	000101??0000?00
Nemocardium	12011000111010	000000022430	1202000000020	0000202001100000
Habecardium	???????????????????????????????????????	??????11220?	1102000000020	000303000100000
Fulvia	1201200010111?	??????112230	110200B000021	001303001100000
Laevicardium	12015000121110	01010021X230	1102000000021	001303001100000
Granocardium	???????????????????????????????????????	??????013120	120200000020	0000202000110000
Cardium acusticostatum	???????????????????????????????????????	??????01410?	1102001000020	0000205000110000
Austrocardium	???????????????????????????????????????	???????1410?	110200000020	000?2??000110000
Chrysocardium	???????????????????????????????????????	??????115130	1???00000101?	??102020001?0???
Sawkinsia	???????????????????????????????????????	???????15130	150100000101?	010?020001?0???
Hedecardium	???????????????????????????????????????	??????014320	120000000000000000000000000000000000000	0001202001110000
Orthocardium	???????????????????????????????????????	??????014120	120001000020	0000202000110000
Vepricardium	22014100101021	010100014141	1400100100121	001202101100000
Cardium	???????????????????????????????????????	??????015141	1702101100121	0 0 2 8 1 X 1 0 0 1 0 0 0 0 0
Agnocardia	???????????????????????????????????????	??????0141C0	1300010010020	0001202000110000
Bucardium	???????????????????????????????????????	??????014141	1102100100021	0022?0001100000
Acanthocardia	???????????????????????????????????????	??????115120	140010000021	00180X101100000
Rudicardium	???????????????????????????????????????	??????115120	150100000020	000180X001110000
Plagiocardium	???????????????????????????????????????	??????1161D0	121000000020	0001517000110000
Clinocardium	23105110321020	010010116110	110200000020	0001508000100000
Dinocardium	23015100131020	010110115141	1102100100121	000202101100000
Trachycardium	24004100131020	0100200131D0	110000000000000000000000000000000000000	0100202001110000
Vasticardium	22004100121020	010020013110	110000000000000000000000000000000000000	0200406000110000
Acrosterigma	???????????????????????????????????????	??????013110	110000000000000000000000000000000000000	0200406000110000
Parvicardium	???????1???0??	??????118110	1110100010020	0001519000100000
Apiocardia	1201121122202?	??????118120	1110000010020	0000619010121111
Trigoniocardia	0111021121203?	??????112110	0110000000020	0000619010121111
Fragum	01112201232030	111000012130	112000000020	0001609000120011
Cerastoderma	12012A00311040	01001011710?	110200000020	0001519001100000
Hypanis	140112004?1040	010010117110	1102001000020	0001919??0000?00
Tridacna	?????200???0??	??????1181??	160000000000000000000000000000000000000	0001704??0000??0

SCHNEIDER: CLADISTIC ANALYSIS OF THE CARDIIDAE

APPENDIX 5.

Synapomorphies for interior nodes. Nodes numbered as in figure 1. Terminal taxa not diagnosed.

Node Synapomorphies (Character:State) 1 2:2, 5:1, 9:1, 11:1, 12:1, 14:0, 16:1, 20:0, 43:1 2 23:1, 25:1 3 24:0, 28:1, 30:2, 38:2, 49:1 4 27:1 5 21:2, 22:0, 23:9 6 23:2, 43:2, 45:2 7 24:4, 26:3, 48:1 8 5:2, 10:0, 12:1, 18:1, 24:2, 43:3, 45:3 9 39:1, 42:1 10 1:2, 6:1, 13:2, 19:1, 23:4, 24:1 11 5:4, 10:3, 25:2 12 23:5, 28:5, 30:1 13 25:3, 36:1, 38:1, 41:1 14 42:1, 43:8, 45:10, 48:1 15 21:0 16 18:1, 25:4, 26:1, 31:1, 34:1, 39:1, 42:2, 48:1 17 37:1, 46:1 18 2:3, 5:5, 23:5 19 4:0, 19:2, 50:1 20 25:0, 45:5 21 28:2, 30:0 22 32:1 23 42:1 24 23:3 25 25:1, 28:1, 40:1 26 10:2, 40:2, 43:4, 45:6 27 9:3, 23:6, 42:1, 43:5, 45:4 28 1:1, 6:2, 44:1, 45:9 29 13:4, 23:7 30 9:2, 11:2, 15:1, 17:1, 19:0, 23:8, 30:0 31 7:1, 8:1, 29:1 32 42:0, 43:6, 47:1, 50:2, 51:1, 52:1, 53:1, 54:1 33 1:0, 2:1, 3:1, 5:0, 13:3, 23:2

APPENDIX 6.

Taxa represented by species other than type.

- Palaeocardita, type species Palaeocardita austriaca (Hauer). Species examined: Palaeocardita silberlingi Newton et al.
- Integricardium, type species Integricardium dupinianum (d'Orbigny). Species examined: Integricardium globulum (Whitfield).
- Lahillia, type species: Lahillia angulata (Philippi). Species examined: Lahillia larseni (Sharman and Newton).

Nemocardium, type species Nemocardium semiasperum (Deshayes). Species examined: Nemocardium bechei (Reeve).

Laevicardium, type species Laevicardium oblongum (Gmelin). Species examined: Laevicardium laevigatum (Linne).

Granocardium, type species Granocardium carolinum (d'Orbigny). Species examined: Granocardium dumosum (Conrad).

Agnocardia, type species Agnocardia claibornense (Aldrich). Species examined: Agnocardia dissidepictum (Woodring).

Trigoniocardia, type species Trigoniocardia granifera (Broderip and Sowerby). Species examined: Trigoniocardia antillarum (d'Orbigny).

Tridacna, type species Tridacna gigas (Linneaus). Species examined: Tridacna maxima (Röding).

Hypanis, type species Hypanis plicatum (Eichwald). Species examined: Hypanis colorata (Eichwald).



Schneider, J A. 1992. "Preliminary Cladistic analysis of the Bivalve Family Cardiidae." *American malacological bulletin* 9, 145–155.

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