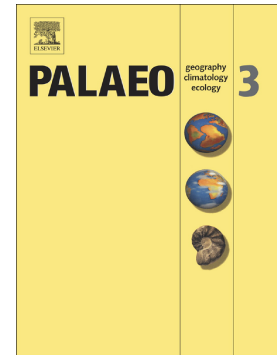


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Linking Western Tethyan Rhynchonellide morphogroups to the key post-Palaeozoic extinction and turnover events

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Abstract

Genetic control undoubtedly plays the most decisive role in the development of characters in rhynchonellides, especially in those structural features such as shell microstructure and brachidial architecture. However, considering the relationship between shell microstructure and type of crura in post-Palaeozoic rhynchonellides supported by recent growing research, the present study links this striking matching to the distribution of rhynchonellide morphogroups around the main post-Palaeozoic mass extinction and diversification events, innovatively introducing the ribbing pattern as a feature widely argued as indicator of deeper/shallower habitats and particular ecological conditions. The analyses performed on the Western Tethys rhynchonellide database and, directly on 28 species from the peri-Iberian margins around selected post-Palaeozoic extinction/diversification episodes, reveal a recurrent distribution pattern according to the habitat considered (epicontinental vs. epiocceanic). The smooth, eurinoid, septifal/arcual body plan is the distinctive inhabitant of deeper epiocceanic environments, also being the last to withdraw from the epicontinental platforms when the environmental conditions were adverse, and the pioneers or opportunistic taxa conducting repopulation of these epicontinental habitats when the conditions were re-established after crises.

The ribbed, eurinoid, falciform morphogroup seems to be the more resilient stock on epicontinental platforms in relation to ecological crises. Conversely, the ribbed, leptinoid, raducal group is typical inhabitant of epicontinental seas when normal conditions prevail and shows an uneven distribution unlinked to critical events, suggesting a more specialist pattern with a diversification related with a secondary differentiation. It is suggested that this distribution dynamic of morphogroups was controlled by the availability of requirements to generate each type of crura and microstructure in epicontinental/epioceanic habitats, such as metabolic or oxygen restrictions, or the filter-feeding efficiency of the lophophore. A research line starting-point is thus herein introduced which can lead to shed light into the adaptive strategies of rhynchonellides, relating particular morphotypes with potential triggering factors of mass extinction events.

Keywords: Brachiopoda; extinction events; adaptive strategies; environmental distribution; Western Tethys.

1. Introduction

Representatives of phylum Brachiopoda are frequently used in macroevolutionary studies (e.g. Ager 1956a; Johnson 1975; McNamara 1982; Ager 1983; Raup 1986; Valentine and Jablonski 2010; Carlson 2016; Stigall 2013) due to their wide-ranging record throughout the Phanerozoic, their considerable diversity and abundance, and relatively simple body plan, endowing this group with a reasonable morphological stability range (Kowalewski et al. 1997; Krause 2004; Harper et al. 2017; Topper et al. 2017; Vörös et al. 2019). Owing to their close connection with substrata and short mainly lecithotrophic larval stage (at least in post-Palaeozoic times), they are also exceptional proxies to shed light on the response of marine benthic ecosystems to environmental perturbations. In this sense, a number of marine biotic crises and mass extinctions recurrently affected this filter-feeding biota influencing the diversity dynamics and the evolutionary trajectories of the different

brachiopod clades (Gould and Calloway 1980; Chen et al. 2005; Curry and Brunton 2007; García Joral et al. 2011; Harper et al. 2014; Carlson 2016; Vörös et al. 2016, 2019).

Notwithstanding the well-known homeomorphy experimented by several constituents of this phylum, among brachiopod clades, the post-Palaeozoic representatives of the Order Rhynchonellida Kuhn, are relatively standardised in both the external form (shape, beak features, rhizo- to pleni-pedunculate shells, *sensu* Ratcliffe 1991 and Harper and Moran 1997 respectively) and internal architecture (crural developments as skeletal supports of spirolophous lophophore), accordingly constituting effective ecological and evolutionary tools.

Concerning the external attributes, one of the most challenged features on the ecological interpretations in rhynchonellides is the ornamentation. It is stated that smooth brachiopods are more prolific in both deeper and warmer environments (Agee 1965; Rudwick 1970; cf. Fürsich and Hurst 1974; Vörös 2005). Smooth rhynchonellides stock is supposed to play a significant role in several crucial extinction and/or diversification events in the evolutionary history of brachiopods (Vörös 2005). In fact, several smooth evolutionary lineages mainly arranged into the families Basiliolidae and Norellidae were invoked by Vörös (2005) to be crucial in the Mesozoic marine diversification, hypothesizing that they survived the End-Triassic extinction and were resilient to the major post-Palaeozoic marine crises as inhabitants of bathyal deep-sea environments, thus facilitating the pioneering re-colonization of habitats in the aftermath of these crisis events.

On the other hand, after experimental substantiation, Manceñido and Motchurova-Dekova (2010) postulated a close correspondence between types of crura and the microstructure pattern of the secondary layer of the shell in most of the rhynchonellid groups. Using combination of both internal diagnostic criteria has been growing as innovative tool on the suprageneric taxonomy, also revealing major evolutionary lineages within rhynchonellides (Motchurova-Dekova and Simon 2007; Radulovic et al. 2007; Motchurova-Dekova et al. 2008, 2009; Manceñido and Motchurova-Dekova 2010; Schreiber et al. 2014; Simon et al. 2018, among others).

In this paper, both aforementioned inter-related hypotheses are coupled and tested. Firstly, new data on the hypothesized positive crural architecture-shell microstructure relationship (Manceñido and Motchurova-Dekova 2010) are empirically evaluated in several smooth and ribbed rhynchonellides from the Mesozoic-Cainozoic peri-Iberian platforms-basins system, where brachiopod faunas are widely represented virtually in all periods and series, being specially studied in their Jurassic heydays and in crucial events (e.g. Choffat 1880; Jiménez de Cisneros 1923; Dubar 1925; 1931; Comas-Rengifo and Goy 1975; Calzada 1975, 1984, 1997; Calzada and Pocoví 1980; García Joral and Goy 1984, 2000, 2004, 2009; García Joral et al. 1990, 2011, 2018; Alméras et al. 1996; Alméras and Fauré 2000; Bitner and Martinell 2001; Reoli et al. 2005; Andrade 2006; García-Ramos 2006; Baeza-Carratalá 2011, 2013; Comas-Rengifo et al. 2006, 2013, 2015; Baeza-Carratalá et al. 2015, 2016 a, b, 2018; Colás 2015; Berrocal-Casero et al. 2017, 2020; Giannetti et al. 2018). Subsequently, the external ribbing pattern is taken into account to perform an equivalence analysis coupling this external feature with the internal architecture/microstructure compiling an updated Western Tethys Meso-Cainozoic rhynchonellide database for that purpose.

Linking both internal and external primary systematic criteria with the occurrence of the different post-Palaeozoic rhynchonellide taxa, we can extend the hypothesis that substantial turnover events in the diversity dynamics of post-Palaeozoic rhynchonellides were undergone by smooth morphotypes (cf. Vörös 2005) also involving the role of other morphogroups in these events, in turn phylogenetically related to several evolutionary lineages (Vörös 2005; cf. Manceñido and Motchurova-Dekova 2010). Thus, comparing the rhynchonellide distribution curve derived from the global dataset of the Paleobiology Database (PaleoDB) with the possible matching predicted by two hypotheses, it is possible to deduce if particular rhynchonellide morphogroups could recurrently develop in the evolutionary history of brachiopods and if this may be explained in terms of ecological strategies or behaviours in response to ecological perturbations.

2. Materials and Methods

A number of 102 Western Tethyan post-Palaeozoic rhynchonellide genera dataset was built up in a matrix (supplementary material) with an initial geographical discrimination from the revised *Treatise* (Kaesler and Selden 1997-2007) combined with raw data of the Paleobiology Database (PaleoDB, <http://paleobiodb.org/> accessed via the FossilWorks gateway, <http://fossilworks.org/>), and new recent generic attributions mainly from the peri-Iberian basins. Discrimination also took into account the availability of reliable documentation about internal features and microstructure in previous selected literature revealing crural and microstructure criteria (44 genera) either directly in descriptions or in the pictures. New data on 28 species (22 genera) from the peri-Iberian basins were analysed. The matrix dataset was supplemented by addition of 37 genera on the basis of their suprageneric attribution, when crural architecture and shell microstructure were properly guaranteed from different confamiliar taxa.

The internal structure and microstructure of the shell was studied on 34 peri-Iberian rhynchonellide specimens using the conventional method of taking acetate peels after making transverse serial sections. The distance between serial sections was 0.1 mm. Using Scanning Electron Microscope (SEM) has been promoted as better exploratory method of the shell microstructure whenever possible, but SEM is not always available to all brachiopodologists (as pointed out by Motchurova-Dekova et al. 2008) and microstructural analysis performed in the present paper exposes that conventional light microscopy can be a suitable equipment to reveal the main microstructural patterns in proper serial sections. Thus, high resolution microphotographs of acetate peels were taken using a Nikon CFI60 600POL microscope. The fibres of the secondary layer were quantified as close as possible to the maximum shell width, as recommended by previous authors (Radulovic et al. 2007; cf. Manceñido and Motchurova-Dekova 2010). All specimens figured and acetate peels are deposited in the collections of the Earth and Environmental Sciences Department of the University of Alicante (DCTMA), the Department of Geodynamics, Stratigraphy and Palaeontology of the Complutense University of Madrid (DPUCM), and the Department of Palaeontology of the University of Granada (DPUGR).

With reference to the ribbing pattern criterion, Vörös (2010) defined four ornamentation categories in Mesozoic rhynchonellides: 1. smooth or capillate; 2. costate or antidichotomous; 3. strongly costate; and 4. spinose. Partly based on this categorization, we adopted the following groupings: 1. smooth or capillate with paucicostate shells, mainly restricted to the anterior margin; 2. strong multicostate or fully ribbed shells, 3. spinose shells.

Regarding the crural types, nomenclature herein followed is that proposed by Manceñido et al. (2007), later followed by Manceñido and Motchurova-Dekova (2010) considering four crural groups: ensimergal, raducal, septifal, and arcual. Concerning the shell microstructure on the secondary layer, the duality eurinoid/leptinoid (coarse/finer fibres respectively) established, among others, by Radulovic et al. (2007), Manceñido et al. (2007, in *Treatise*, v. 6), and Manceñido and Motchurova-Dekova (2010), is considered.

Several critical bioevents were plotted against the rhynchonellide diversity curve obtained using the global dataset of the Paleodb. This diversity dynamics curve was processed to genus-level by means of a stage-level resolution. Biostratigraphical data of each taxon were taken into account to evaluate the range and distribution of each morphogroup. Environmental epioceanic/epicontinental duality was also considered, when available, in the interpretation of the graphics to better constrain the environmental objective of the research (see environmental occurrence data in the supplementary matrix for habitat preferences of each taxon; columns Q and R).

The binary (presence/absence) matrix, built up exclusively considering morpho-structural criteria, i.e. the crural type, microstructure, and ribbing criteria, was evaluated through hierarchical cluster analysis with the UPGMA algorithm using the Jaccard similarity coefficient. Independently, binary values were also considered to plotting the environmental preferences of each rhynchonellide morphogroup (columns Q and R in the supplementary matrix) against their biostratigraphic range. All statistical analyses were carried out by means of the PAST software package (Hammer et al. 2001).

3. Crural architecture plus shell microstructure.

Significance of type of crura as the main endoskeletal structure with most important taxonomic value has a long-way trajectory. Traditional systematics of rhynchonellides is based upon this character earlier in the morphological interpretations of Rothpletz (1886), Wisniewska (1932), and Muir-Wood (1934), subsequently followed by the comprehensive works of Ager (1956b-1967), Cooper (1959), Makridin (1964), Dagys (1968), Ager et al. (1972), Manceñido (2000), Manceñido and Owen (2001), Savage et al. (2002), Manceñido and Motchurova-Dekova (2010), Schreiber et al. (2014), among many others (for a recent revision on this issue, see e.g. two latter quoted works).

Over the last years, a more accurate taxonomical hierarchy has been achieved by means the exploration of the fibrous microstructure pattern on the secondary layer of the shell, simultaneously applied in correspondence with types of crura. In this sense, an extensive background (e.g. Motchurova-Dekova and Simon 2007; cf. Manceñido et al. 2007; Radulovic et al. 2007; Motchurova-Dekova et al. 2008, 2009; Manceñido and Motchurova-Dekova 2010; Simon et al. 2018) has improved the suprageneric taxonomy, also revealing major evolutionary lineages within rhynchonellides. Thus, an eurinoid coarse fibres pattern has been revealed corresponding with the arcual and septifal crural groups, contrasting with a finer leptinoid pattern which matches with the ensimergal and raducal crural groups (Manceñido and Motchurova-Dekova 2010).

As results of this previous extensive background, a starting point dataset has been built up herein. A total number of 107 genera has been considered, compiling information from 44 genera recorded in the Western Tethys Ocean, arranged into the Superfamilies Rhynchonelloidea, Hemithiridoidea, Rhynchotetradoidea, Dimerelloidea, Pugnacoidea, Wellerelloidea, and Norelloidea, taking previous comprehensive research of Manceñido and Motchurova-Dekova (2010) as a keystone, as well as data from some other authors revealing microstructure criteria either directly in descriptions or by observing the illustrations (e.g. Tomašových 2006; Dulai et al. 2008, Motchurova-Dekova et al. 2009, Simon et al. 2018, Berrocal-Casero et al. 2020). In all these

genera, a positive correspondence between microstructure pattern and expected crural types was revealed (see supplementary material).

Additionally, new data testing this matching have been analysed in 23 Mesozoic-Cainozoic genera (29 species) recorded in the peri-Iberian platforms-basins system (Table 1), accomplishing therefore a profitable post-Palaeozoic database for further deductions (see supplementary material). These taxa are encompassed into 8 families: Wellerellidae (*Cirpa briseis*, *Cirpa* sp. aff. *latifrons*, *Salgirella albertii*), Tetrarhynchiidae (*Gibbirhynchia curviceps*, *Quadratirhynchia tenuata*), Rhynchonellidae (*Cuneirhynchia cartieri*, *C. dalmasi*, *C. rastuens* s. *Homoeorhynchia meridionalis*, *Choffatirhynchia vasconcellosi*, *Rhynchonelloidea goyi*), Prionorhynchiidae (*Prionorhynchia guembeli*, *P. quinqueplicata*, *P. polyptycha*); Cyclothyrididae (*Albusirhynchia vorosi*, *A. jorali*, *Globirhynchia subobsoleta*), Basiliolidae (*Apringia atlaeformis*, *Soaresirhynchia bouchardi*, *Pseudogibbirhynchia jurensis*, *Mondegia atla*, *Fortunella decipiens*, *Aphelesia bipartita*), Erymnariidae (*Septocrurella microcephala*), and Norellidae (*Nannirhynchia pillula*, *Atychorhynchia falsiorigo*, *Praemonticlarella distercica*, *Monticlarella triloboides*). The remaining 35 genera in the matrix are tentatively arranged in basis of their suprageneric taxonomic attribution widely substantiated from different conterminous taxa from their respective lineages.

All the peri-Iberian Basiliolidae herein analysed reveal a eurinoid microstructure pattern (Fig. 1). Their coarse fibres show a non-ovoidal cross-section outline and range from 45 to 75 μm width and 30 to 50 μm thickness. Crural developments in the studied Basiliolidae are represented by the hamiform type (figs. 1 A, B; 2C) in the Lower Jurassic representatives (*Apringia atlaeformis*, *Pseudogibbirhynchia jurensis*, and *Soaresirhynchia bouchardi*) and by the falciform type in the Middle Jurassic (*Mondegia atla*), Cretaceous (*Fortunella decipiens*), and Cainozoic (*Aphelesia bipartita*) counterparts (Fig. 1, C-E). Other taxon having an eurinoid pattern has been observed in Erymnariidae (*Septocrurella microcephala*), presenting coarse fibres with a mean of 70 μm wide and 45 μm thick, and septiform crura, thus satisfying the predicted pattern of the septifal crural group (Fig. 1F).

The remaining species revealing a eurinoid microstructure pattern are arranged into Wellerellidae (*Cirpa briseis*, *Cirpa* sp. aff. *latifrons*, and *Salgirella albertii*). They present rhomboidal to subquadrate fibres in cross section, ranging from 60-70 μm wide and 35-40 μm thick, also corresponding with hamiform crural developments (Fig. 2). Taxa with arcual crural developments (*Nannirhynchia pillula*, *Monticlarella triloboides*, and *Atycorhynchia falsiorigo*) evidence slightly stacked and finer fibres (50-60 μm wide and around 20 μm thick) but also fitting the eurinoid pattern, with pseudo-rhombic fibres and rather homogeneous structure (Fig. 3). This eurinoid pattern is much more evident in *Praemonticlarella distercica*, also with arcual crural architecture (Fig. 3B).

By analysing the dataset matrix, the correspondence eurinoid/septifal can be extended to all the Basiliolidae, Erymnariidae, and Wellerellidae representatives as well as the eurinoid/arcual relationship can be broadened to Norellidae, as foreseen to some extent by Manceñido and Motchurova-Dekova (2010) and references therein.

On the other hand, a striking positive correspondence is detected between the leptinoid microstructure pattern and the raducal crural group (Fig. 4). It is substantiated in all the Jurassic families and genera studied with radiform development, i.e. Tetrarhynchiidae (*Gibbirhynchia curviceps*), Rhynchonellidae (*Cuneirhynchia cartieri*, *C. dalmasi*, *C. rastuensis*, *Homoeorhynchia meridionalis*, and *Choffatirhynchia vasconcellosi*), Prionorhynchiidae (*Prionorhynchia quinqueplicata*, *P. guembeli* and *P. polyptycha*) and Cyclothyrididae (*Alebusirhynchia jorali*, *A. vorosi*, *Globirhynchia subobsoleta*). These taxa present finer halberd-like fibres in *Cuneirhynchia*, *Homoeorhynchia*, and *Prionorhynchia* (Fig. 4 A-F, L); subrectangular and finest in *Gibbirhynchia* (Fig. 4 G); being more irregular and heterogeneous in *Alebusirhynchia* and *Choffatirhynchia* (Fig. 4 I-K). The fibres size varies from 8-45 μm wide (exceptionally 55 μm in a few *Cuneirhynchia* cross sections) and 3-10 μm thick (except for some *Cuneirhynchia* fibres as well, with 15 μm thick).

The correlation leptinoid/raducal is also corroborated in all the Tetrarhynchidae, Rhynchonellidae, Prionorhynchiidae, and Cyclothyrididae representatives present in the dataset matrix, as also foreseen by Manceñido and Motchurova-Dekova (2010) and references therein.

4. Ribbing pattern reflected in the diverse crural-microstructure lineages of rhynchonellides

In his seminal work about smooth Mediterranean brachiopods, Vörös (2005) portrayed the evolutionary lineages of several smooth Norelloidea and Basiliolidae rhynchonellides, in part taking into account their internal architecture. The analysis now performed estimates the ornamentation pattern considering the Western Tethys post-Palaeozoic rhynchonellide database (see supplementary material), bearing in mind the crural/microstructure equivalence previously deduced in each evolutionary lineage. Considering the different ribbing pattern categories aforementioned in the materials and methods section, the dominant pattern in the diverse groups based on the internal structure is evaluated, resulting the following combinations (figs. 1-4):

1. Multicostate or fully strong-ribbed taxa are the abundant “ordinary-looking” post-Palaeozoic rhynchonellides. This ornamentation category is mainly distributed in the Western Tethys in two major groups according to their internal body plan:

1a. The raducal (unusually ensimegal)/leptinoid is the most numerous ribbed group (Fig. 4). Their representatives are arranged in several supra-generic positions: the better represented is Tetrarhynchiidae, with 23 genera (herein depicted with *Gibbirhynchia*). Superfamily Hemithiridoidea is also frequent in this class, emphasising 8 ribbed genera into the Family Cyclothyrididae (herein *Alebusirhynchia*). The Family Rhynchonellidae is typified by 6 ribbed genera (*Choffatirhynchia* and *Cuneirhynchia* herein). Further multicostate families such as Prionorhynchiidae (*Prionorhynchia* herein) or Peregrinellidae are less diverse.

1b. The septifal (exceptionally arcual)/eurinoid group is less abundant. Ribbed taxa are mainly arranged into the subfamilies Cirpinae (Family Welleriidae), Lacunosellinae (Superfamily Pugnacoidea), and, to a lesser extent, into the Subfamily Pamirorhynchiinae (Family Basiliolidae;

such as *Pseudogibbirhynchia* herein) and the Family Erymnariidae. The specimens illustrated in this work belonging to this group are mainly arranged in *Cirpa*, *Salgirella*, and *Pseudogibbirhynchia* (Fig. 2). Interestingly, some genera included in this group show a typical “bifurcate” ribbing pattern, more infrequent in the previous group. The ribbed representatives with arcual developments are arranged into Praemonticlarellinae, herein represented by *Praemonticlarella* (Fig. 3).

2. Spinose forms are completely embodied into the Family Acanthothirididae, all with raducal/leptinoid internal pattern.

3. Among the smooth, paucicostate, and capillate pattern there are different internal body plans, but major groups are included in Norelloidea and Basiliolidae:

3a. Smooth or paucicostate ribbing pattern with septifal/eurinoid structure is conspicuous in the Family Basiliolidae, showing falciform, subfalciform, and hamiform crura, and mainly arranged into the Subfamily Basiliolinae (13 genera). This Subfamily is typified by *Apringia*, *Soaresirhynchia*, *Mondegia*, and *Fortunella* in this work (Fig. 1), also analysing Aphelesiinae (*Aphelesia*) and Pamirorhynchiinae representatives. Completing the septifal/eurinoid group there are three poorly ribbed genera belonging to Erymnariidae with septiform crura (*Septocrurella* herein).

3b. The paucicostate and smooth ornamentation is also recurrent in the rhynchonellides with arcual/eurinoid internal architecture. They are all arranged in Norelloidea (*Nannirhynchia*, *Monticlarella*, and *Atycorhynchia* herein, Fig. 3), mainly in the Subfamily Norellinae, as reflected in the matrix dataset.

3c. Leptinoid microstructure pattern is minority among the smooth (paucicostate) morphotypes. Five genera belonging to the Family Rhynchonellidae show this type of ornamentation with raducal/leptinoid structure (*Homoeorhynchia* herein, Fig. 4L) and three genera into Dimerelloidea fulfil the smooth leptinoid pattern with ensimergal crural development.

In summary, ribbing pattern is variable in almost all the groups defined by their crura-microstructure features, reinforcing its secondary role for phylogenetic relationships. Therefore, it seems to be an important feature in adaptive strategies, as will be discussed in the next paragraphs.

5. Discussion. Testing twofold hypothesis

5.1. Main post-Palaeozoic rhynchonellide morphogroups in the Western Tethys

Cluster multivariate analysis performed in the matrix dataset of Western Tethys rhynchonellides expose a clear grouping on the basis of microstructure pattern (eurinoid/leptinoid) as primary criterion (Fig. 5). The subsequent subordinate division takes the ribbing pattern as more relevant factor, being the crural group the last criterion of arranging.

These criteria are marked in the nodes of the cluster (Fig. 5) and apparently accomplish a discernible taxonomical component as primary factor for splitting morphogroups, i.e., leptinoid vs. eurinoid microstructure in turn related with the crural type, acquiring the ribbing pattern a decisive role in the clustering. However, all features globally considered acquire an environmental significance as discussed below, without diminishing its phylogenetic meaning.

In the eurinoid group, the phylogenetic and systematic relationship is evident, grouping on the basis of the similarity degree, the arcuiform Norellidae interrelated with the septifal crural architecture of the Erymnariidae (septiform) and Basiliolidae (hami/falciform). However, as previously mentioned, cluster distance gives more significance to the ribbing pattern than the crural group. Thus, several taxa diverge from the phylogenetic structure and we can observe ribbed eurinoid hamiform Basiliolidae, such as *Pseudogibbirhynchia*, *Stolmorhynchia*, and *Orbirhynchia* clustered together with ribbed eurinoid hamiform Welleridae such as *Cirpa*, *Salgirella*, or *Calcirhynchia*. Hence, this clustering releases an interesting systematic discussion. For instance, the so far assumed systematic arrangement of *Pseudogibbirhynchia* into Basiliolidae was obviously based on the type species *Pseudogibbirhynchia moorei* (Davidson), which is the only paucicostate species assigned to this genus. This species is very scarce and presumably endemic from the South-

England basins, since the occurrences attributed to this taxon outside its type area (Thévenin 1908; Comas-Rengifo and Goy 1975; Alméras 1996; Alméras and Becaud 2002; Baeza-Carratalá 2013) are tentative and/or ambiguous. On the other hand, the occurrences assigned to *Pseudogibbirhynchia* in the Upper Toarcian-Aalenian from Spain, Portugal, and the Alps (García Joral 1993; García Joral and Goy 2009; Andrade 2006; Andrade et al. 2016) are fully costate (frequently dichotomous) forms that can be considered closer to representatives of Welleridae, such as *Cirpa*, as resulted in our clustering analysis. It is therefore plausible postulating a phyletic relationship between *Pseudogibbirhynchia* and *Cirpa*, as envisaged by Ager (1962), and consequently, a systematic revision of *Pseudogibbirhynchia* and other ribbed Basiliolidae such as *Orbirhynchia*, both currently in the Subfamily Pamirorhynchinae would be desirable.

5.2. Morphogroups linked to the key post-Palaeozoic turnover and extinction events

By analyzing the global diversity curve of rhynchonellides obtained using the global dataset of the Paleodb, several events of extinction and diversification in the overall history of the group can be individualized (Fig. 6). We can distinguish four major post-Palaeozoic crises, entailing a considerable diversity decrease followed by subsequent noticeable recovery episodes: The Rhaetian, linked to the end-Triassic extinction (ETE) events; the Early Toarcian; the Oxfordian; and the Maastrichtian/Danian boundary. Several subordinate episodes involving loss of diversity, less conspicuous in the rhynchonellide groups, are also detected (e.g. Cenomanian-Turonian boundary or Messinian events in the Fig. 6). These secondary episodes took place just when brachiopods diversity was very low, even in the background conditions, and no relevant inferences can be obtained from their analysis. On the other hand, previous authors (e.g. Miller and Foote 2009) have confirmed that environmental perturbations during mass extinctions and their macroevolutionary signatures were manifested differently in epicontinental and epi-oceanic habitats (*sensu* Hallam 1971; Olóriz 2000). Connecting this aspect with the morphogroups resulting from the figure 5, allowing to organize the 9 groups previously determined by the cluster, arranging them into 4 stocks

with palaeoenvironmental significance. For the sake of the accuracy of the analysis, only those morphogroups with relative abundance related to the extinction/diversification events have been considered, being the remaining groups too scarce to allow reliable interpretations. These groups are: 1. eurinoid arcuiform smooth; 2. eurinoid septifal smooth; 3. eurinoid septifal ribbed; and 4. leptinoid raducal ribbed. The chronostratigraphic distribution of the studied genera representing these four morphogroups is exposed in the Fig. 7, distinguishing epioceanic and epicontinental occurrences, and the main extinction and diversification events since the Early Triassic.

The eurinoid pattern of microstructure occurs in three of these morphogroups, all consistent with habitats related with pre- or post-stressed environmental conditions. The smooth eurinoid arcuiform morphogroup has been interpreted as constituted by deeper epioceanic taxa (cf. Vörös 2005). However, if we interrelate this morphogroup with the stratigraphical distribution and the epicontinental/epioceanic occurrences of the taxa analysed from the Western Tethys post-Palaeozoic database (Fig. 7), it can be deduced also for this group a resilient pattern in epicontinental platforms around extinction episodes as well as a pioneer post-extinction occurrence after three events, the early Toarcian, the Callovian/Oxfordian and the Cenomanian/Turonian extinctions. Likewise, it can be observed that the smooth eurinoid septifal morphogroup is mainly constituted by deeper epioceanic inhabitants, which conducted opportunistic post-crisis epicontinental colonization; while the distribution of the eurinoid falciform ribbed morphogroup is consistent with epicontinental resilient taxa.

Finally, the leptinoid, raducal, and ribbed forms, seems to be linked to epicontinental specialist taxa with secondary adaptations, depending on ecological factors such as the substrate type, trophic source, hydro-dynamics, or reproductive adaptations. This specialistic component is noticed also for the leptinoid ensimergal representatives (not represented in the figure 7). Thus, several dimerelloids such as *Peregrinella*, *Sulcistrostra*, and *Rhynchonellina* have been related with cold seeps or chemosynthesis-based ecosystems (Campbell and Bottjer 1995; Sandy 1995, 2010; Dulai 2002; Peckmann et al. 2007, 2013; Kiel et al. 2014; Sandy and Peckmann 2016; Palfy et al. 2017, among

others); in the same way, the raducal spinose Acanthothyrids, have been often related with pararecifal environments (e.g. Childs 1969) and *Cryptopora* could be adapted to deeper (up to abyssal) environments (Surlyk and Johansen 1984). The role of the leptinoid raducal smooth or paucicostate morphogroup is more difficult to analyse because of their scarcity, but it is noticeable its record just prior to the first occurrence of the leptinoid raducal ribbed forms in epicontinental areas, both in the Triassic and in the Paleogene.

With this analysis, a matching seems to appear between major post-Palaeozoic rhynchonellide turnover events after ecological crises and a specific body plan, consisting in smooth/eurinoid, arcual (Norellidae) or falciform and derived (mainly Basiliolidae) genera, otherwise typical inhabitants of epioceanic areas. Conversely, those ribbed leptinoid, raducal (some ensimergal) types do not reveal such a patent distribution pattern around critical events, but they are the main components of the assemblages in the diversification events in epicontinental environments. On the other hand, ribbed falciform (and derived) eurinoid types, recurrently recorded both in epicontinental and epioceanic environments, seem to be more resilient to these events (Fig. 8).

One of these a priori deeper epioceanic lineages (Norelloidea) was previously presumed as responsible of the rhynchonellide diversity recovery after the End-Permian and End-Triassic (ETE) crises (Manceñido and Owen 2001; Vörös 2005). These authors hypothesized that minute Norelloidea could lead the post-Palaeozoic radiation from paedomorphic survivors of this family, performing the recovery of brachiopod lineages after the Permian-Triassic biotic catastrophe. According to the rhynchonellide global diversity curve (Fig. 6), from the minimal richness values reached in the End-Permian extinction, when other 5 rhynchonelliform orders were extinct (Curry and Brunton 2007), the first rhynchonellide re-diversification pulse in the Western Tethys was recorded in the Anisian, probably conducted by smooth, arcuiform, eurinoid, *Norella* and *Holcorhynchella* (Vörös 2005; Gradinaru and Gaetani 2019) arranged into the Norelloidea lineage (Fig. 7). *Holcorhynchella* did not evolve beyond the Middle Triassic, but the epioceanic *Norella* lasted up to the End-Triassic crisis, in the Rhaetian. Thus, Norellidae (and most of the resulting

morphogroups herein) apparently are consistent with a pattern of declining diversity in the ETE (at least in the westernmost Tethyan areas) whose onset was detected also in other fossil groups at or around the Norian/Rhaetian boundary by several previous authors (e.g. Rigo et al. 2020).

The succeeding record of this lineage is detected during the main Early Jurassic diversification event after the End-Triassic crisis (Sinemurian) due to the occurrence of the small-sized arcual, eurinoid, and paucicostate *Pisirhynchia* in epioceanic habitats (Vörös 2005; Vörös and Dulai 2007), together with the minute Pliensbachian-Bathonian *Nannirhynchia* (arcual, eurinoid, and mostly paucicostate species). This last genus has been repeatedly associated to relatively deep and oxygen-depleted environments (Ager 1967; Alméras et al. 1989, 1995; Andrade 2006).

As can be deduced from the figures 6 and 7, rhynchonellides suffered a Late Jurassic-Lower Cretaceous debacle with development of unfavourable conditions for brachiopod settlement in the Western Tethys as a whole, specially noticed in the epioceanic Mediterranean habitats by the so-called “age of radiolarites” (Vörös 2005; Colás and García Joral 2012) from the Callovian onwards. The only Norellidae genus recorded at this time is *Monticlarrella*, which seemingly was capable of surviving this crisis by colonizing the epicontinental platforms of the northern margin of Western Tethys, just like other epioceanic brachiopods such as nucleatids (Colás and García Joral 2012).

Later representatives of this smooth lineage are very scarce up to their extant counterparts, e.g. *Abyssorhynchia*, *Neorhynchia*. The lineage probably survived in deeper epioceanic habitats establishing steady communities in deep faunal reservoirs (Vörös 2005; Thuy et al. 2014). This can be corroborated by the occasional record of this lineage in deep epioceanic environments, as typified by *Atycorhynchia* in deep Subbetic grabens around the Early Toarcian Mass Extinction Event (ETMEE) crisis (Baeza-Carratalá et al. 2017), which has revealed a eurinoid arcuiform smooth morphotype (Fig. 3).

On the other hand, a virtually uninterrupted trajectory can be detected in the smooth falciform (and related subfalciform, hamiform) Basiliolidae (Vörös 2005), also with eurinoid microstructure. This evolutionary lineage, clearly represented early in the Jurassic by the hamiform *Apringia*-group

(*Apringia*, *Serratapringia*, *Megapringia*) probably started with the Triassic representatives of *Austriellula*. It should be mentioned that *Austriellula* was arranged with uncertainty in Norellinae in the revised Treatise (Manceñido et al. 2002), but subsequent authors (Vörös 2005; Dulai et al. 2008; Baeza-Carratalá et al. 2017) preferred considering this taxon as the baseline of the evolutionary lineage *Austriellula* (Triassic)-*Fortunella* (Early Cretaceous). First records of *Austriellula* (Carnian-Norian) are reported in the maximum of the Triassic rhynchonellide diversification and persisted until the End-Triassic extinction event (except for the tentative attribution to it of Renz's Toarcian material from Western Greece by Manceñido (1993), with inaccessible internal features).

The following record in this lineage is typified by the epioceanic *Apringia* group in the main speciation recovery event that took place in the Sinemurian (Luhai 2003; Vörös and Dulai 2007; Baeza-Carratalá 2013), leading to the subsequent heyday of the brachiopod Mediterranean smooth fauna in the Pliensbachian (Vörös 2005, 2009; Vörös and Dulai 2007) and not exceeding the Early Toarcian Mass Extinction Event (ETMEE).

After the ETMEE, monospecific beds bearing the prolific *Soaresirhynchia bouchardi* are recorded throughout the Western Tethys Ocean (e.g. Alméras 1994; García Joral and Goy 2000; Graziano et al. 2006; García Joral et al. 2011; Baeza-Carratalá 2013; Baeza-Carratalá et al. 2017). This Basiliolidae species is characterized by a smooth to paucicostate shell, moderately small, hamiform crura, and eurinoic macrostructure (Fig. 1). The post-ETMEE pattern of recovery led by *S. bouchardi* in the Serpentinum Chronozone (Elegantulum Subchronozone) of the Early Toarcian, occupying vacant ecospace, is distinctive of an opportunistic strategy in the repopulation interval of the aforementioned biotic crisis (e.g. García Joral and Goy 2000; Gahr 2005; García Joral et al. 2011; Baeza-Carratalá et al. 2017). This turnover episode has been considered as the most important post-Palaeozoic renewal interval in the brachiopod fauna in the Western Tethys (García Joral et al. 2011; Andrade et al. 2016) subsequently allowing, in the stabilization of the background conditions, the settlement of the leptinoid raducal morphotypes (i.e. *Homoeorhynchia*, *Choffatirhynchia*).

The subsequent taxon recorded in this evolutionary lineage is *Mondegia*, also showing smooth, eurinoid, and falciform morphotypes (Andrade 2006; Baeza-Carratalá et al. 2011; Fig. 1 herein). In the mainly epiocenic Mediterranean bioprovince, the onset of the first post-ETMEE diversification episode was conducted by smooth brachiopods (Vörös 2005), with this maximum in diversity dated in the Early Bajocian with the pervasive occurrence of *Mondegia atla* in most of the epiocenic basins (e.g. Oppel 1863; Ferrari 1962; Vörös 1997; Vörös and Dulai 2007; Baeza-Carratalá et al. 2011, 2014) preceding the rhynchonellide bloom in the Bajocian-Callovian interval (Fig. 7). *Mondegia* was also recorded in the Sauzei Chronozone (Lower Bajocian) in relatively deep areas of the epicontinental NW-European bioprovince (Lusitanian basin), subsequently to the dominance of the Basiliolidae (*Soaresirhynchia*) and Norellidae (*Nannirhynchia*), thus conducting these smooth eurinoid morphotypes the diversification episode in the Valaisian-lowermost Bajocian of this area (Andrade 2006). The Bajocian-Callovian interval represents the maximum diversity peak in the post-Palaeozoic history of rhynchonellides, highly contributing the deep-epiocenic smooth taxa to reaching this bloom (cf. Vörös 2005), after which *Mondegia* became extinct.

After the Late Jurassic crisis, the smooth eurinoid/falciform lineage continued in the Jurassic-Cretaceous transition with the genus *Fortunella*. This taxon appeared and diversified after recovery of environmental stress conditions, marked (similarly to the ETMEE) by a negative isotopic oxygen excursion (Curry and Bruntorn 2007). This genus, just like some other aforementioned Basiliolidae taxa, also reveals an opportunistic strategy, showing high morphologic plasticity, being widely diversified from the Oxfordian to the Aptian, and attaining a nearly cosmopolitan distribution, from East-Spain to the Alps and from Algeria to the Russian Platform (Sulser and Calzada 1991). In this sense, *Fortunella* could play a decisive role in the colonization of vacant ecospace in this Upper Jurassic-Cretaceous rhynchonellide decline interval, since it was considered by Vörös et al. (2018) as pioneer taxon in the repopulation phases after crucial perturbation events, such as the Early Cretaceous Weissert anoxic event (Upper Valanginian), when monospecific levels of *Fortunella* were recorded recolonizing intra-Tethyan basins (Vörös et al. 2018).

The record of this lineage is virtually disrupted up to the Turonian times (Vörös 2005) when, once again, after the Bonarelli extinction event, the phylogenetic line of the extant *Basiliola* seems to emerge (figs. 6, 7) leading to a diversification phase of this morphogroup which lasted up to the Maastrichtian. The renowned Cretaceous-Paleogene (K-Pg) event also affected brachiopod fauna (Curry and Brunton 2007, Dulai et al. 2008, Vörös et al. 2019) leading to a very low rhynchonellide diversity. Once again, Basilioloidea representatives occurred as opportunistic brachiopod fauna after K-Pg event. Thus, Dulai et al. (2008) recorded numerous individuals of *Basiliocostella* (an eurinoid falciform and weakly ribbed morphotype) forming monospecific assemblages in the Danian.

Similar situation occurred with the Neogene counterparts of this lineage. The smooth Mesozoic Basiliolidae lineage depicted by Vörös (2005) continued with the Neogene *Aphelesia*. This taxon is profusely recorded in the Upper Miocene-Lower Pliocene of the Mediterranean basins. In the westernmost peri-Iberian basins (Mediterranean Atlantic seaway), *Aphelesia* is recorded in the Late Tortonian, and re-emerged again in the earliest Pliocene (Bittner and Martinell 2001, Bittner and Moissette 2003, García-Ramos 2004, Toscano-Grande et al. 2010; Giannetti et al. 2018) when the background conditions after the Messinian Mediterranean Salinity Crisis were re-established (Giannetti et al. 2018).

The extant representative of this lineage (e.g. *Basiliola*, *Basiliolella*) are typical inhabitants of cryptic or deeper habitats (Zezina 1994, 2010; Vörös 2005). If we consider the present-day as a declining episode in the rhynchonellides diversity, we can postulate that the smooth eurinoid arcual (or falciform) body plans of the Norellidae and Basiliolidae respectively withstand singular environmental conditions (caves, crevices, bathyal habitats) expecting future better conditions or less competitive pressure on the platforms, constituting “reservoirs” for potential diversification, since they are the clear dominant rhynchonellide fauna: 14 of the total 19 extant genera are arranged into Norellidae and Basiliolidae, and a great number of them are smooth and micromorphic forms,

especially in the extant arcual Norelloidea. The same occurs to species-level: 24 of 38 total extant rhynchonellide species are arranged in these body plans.

The eurinoid/septifal ribbed morphogroup is consistent with an epicontinental resilient pattern in the Early and Middle Jurassic. *Cirpa* seems to be the only big-sized rhynchonellide that survives until the last levels prior to the Early Toarcian crisis (García Joral et al. 2018), and *Pseudogibbirhynchia* is the typical representative of the impoverished brachiopod communities that are frequent in the Late Toarcian and the Early Aalenian in epicontinental areas (García Joral and Goy 2009; Andrade et al. 2016). Some representatives of this group have even been considered as disaster-taxa flourishing during Cretaceous OAEs or platform drowning events (Graziano and Taddei-Ruggiero 2008). Thus, *Orbirhynchia* blooms coincided with the Early Aptian Selli OAE and *Costerymnaria* beds are recorded concurring with both the Early Cenomanian and Late Campanian drowning episodes. This lineage seems to become extinct at the Cretaceous-Paleogene boundary.

The leptinoid raducal group is present in epicontinental environments when “normal” conditions prevail in the Jurassic and the Cretaceous, with maximum of diversity in the Pliensbachian, the Bajocian-Bathonian, the Aptian to Cenomanian and the Santonian to Maastrichtian stages, and is scarce or absent during crisis events.

5.3. Functional adaptive body plan

As can be deduced from the previous analysis, there is a correspondence between the rhynchonellide morphogroup consisting of smooth eurinoid, mainly with arcual or septifal crura and crucial episodes in the post-Palaeozoic evolutionary history of rhynchonellides. Thus, the smooth eurinoid arcual structure is the most recurrent pattern just in the pre-extinction boundary intervals, but also is regularly recorded in the post-extinction recoveries, thus contributing to the main diversification events in both the epiocceanic and epicontinental areas (Fig. 8). In the same way, the smooth eurinoid septifal structure seems to be the most effective pattern to conduct the

recolonization and diversification events from the deeper seas, when the environmental conditions have been re-established in shallow-water environments after ecological perturbations, as opportunistic or pioneer taxa indicative of the onset of repopulation episodes in both epicontinental and epi-oceanic platforms (figs. 7, 8). Finally, the leptinoid microstructure with ribbed raducal morphotype, is characteristic of the diversification climax, mainly in epicontinental areas (Fig. 8), and does not seem to play an important role in these critical bioevents.

Although genetic was probably the most determinant factor on the shell microstructure, the eurinoid type seemingly requires more metabolic energy and oxygenated habitats to construct coarser fibres than the leptinoid pattern (cf. Simon et al. 2018), and cannot be excluded that its iterative occurrence after extinction may have been a consequence of heterochrony processes (heterochronic “Lazarus effect” *sensu* Flessa and Jablonsky 1983; Jablonsky 1986; Vörös 2018). In this sense, Manceñido and Motchurova-Dekova (2016) considered that eurinoid microstructure could represent an “apomorphic stage” of development. Among the eurinoid morphogroups, all Basiliolidae and Norellidae develop a simple spirulophous lophophore, but the falciform (and derived) crural architecture of Basiliolidae may represent a more developed type of crura than those raduliform or arcual types developed in other rhynchonellide clades. The septifal crural developments entail a better attachment and consequently a less effective ciliate retractable lophophore (cf. Vörös et al. 2015). The proliferation of taxa with this better-attached lophophore can be consistent just when the record of the eurinoid septifal group increase, i.e. in the background conditions in the epi-oceanic environments and the onset of the repopulation intervals, since in such environmental contexts with abundant resources, redirecting flow regime through retractile lophophore is not essential for filter-feeding.

However, while the epicontinental environments are constrained by adverse conditions or oxygen-depleted habitats, the smooth eurinoid falciform group is mostly restricted to the deeper habitats (Fig. 8) and the eurinoid pattern (with higher metabolic and oxygen requirements) is only represented in the epicontinental settings by small smooth eurinoid rhynchonellides which

developed relatively smaller-type arcuiform cruras (e.g. *Norella*, *Nannirhynchia*). These arcual crura provide a lesser-attached lophophore, thus achieving major surface of retractable ciliate lophophore, more efficient for filtering in less advantageous conditions. Likewise, spinuliform representatives of this arcual crural group were invoked as capable to separate the two coils of the spirolophous lophophore anteriorly (cf. Schreiber et al. 2013, p. 561), enhancing the flow regime within some mantle cavities. These adaptive strategies could lead to Norelloidea and related groups to tolerate the short-term impoverished conditions of the pre-extinction intervals.

Additionally, it is inferred that the small, smooth or paucicostate Erymniariids, with septifal crura and eurinoid microstructure (e.g. *Septocrurella*) could only develop a schizolophous lophophore stage (Manceñido and Motchurova-Dekova 2010, p. 209). Speculatively, the simple physiology and the conservative external morphotype of Erymniariids let them to play the role of a resilient stock diversified in epiocenic deeper refugia.

Conversely, the leptinoid pattern associated to raducal (or ensimergal) crura, and mainly ribbed shell, shows an uneven distribution unlinked to critical events, suggesting, as previously deduced, a more specialized pattern. This type of crura and microstructure require less metabolic and environmental restrictions. The more primitive crural architecture (mainly raducal) supports a simple spirolophous lophophore whose versatility, together with the less-resources consuming microstructure, makes this morphogroup consistent with a strategy that promotes specialization based on trophic efficacy. As previously mentioned, the diversification of this leptinoid group seems to be related with niche partitioning and adaptations to different substrate types or even to chemosynthesis processes.

According to the raised hypothesis linking the morphogroups distribution (epicontinental/epiocenic) around the extinction and posterior diversification events, and taking into account the fixation of lophophore and the metabolic requirements to generate the different microstructure, the eurinoid ribbed falciform morphogroup, whose distribution can be consistent with a resilient pattern both in epiocenic and epicontinental environments, does not fulfil the proposed model. As

postulated above, the lower versatility in terms of filter-feeding efficiency conditioned by falciform crura does not accomplish the requirements to be resilient in epicontinental areas during adverse conditions. The development of the ornamentation on this morphogroup could then come into play in these shallower platforms. These rhynchonellides develop strong angular ribs in the shell to better restrict the coarse particles potentially transferred into the mantle cavity (e.g. Rudwick 1964; Fürsich and Hurst 1974). On the other hand, lophophore tends to occupy all the available space within the shell. Consequently, a strong ribbed shell, apart from playing the role as a defence against shell-crushing predators (cf. Vörös 2010), may be efficient for both functions, the restriction of coarse sediment from entering the mantle cavity and the amplifying of the lophophore surface, which, in turn, improves the filter-feeding efficiency, allowing inhalant currents with a minor gaping of the shell (cf. Fürsich and Hurst 1974). This could be an alternative to the development of a retractile lophophore, enabling the resilience in a disfavoured environmental situation.

A particular case after the well-known K-Pg event recovery can lead to masking the true pattern of post-extinction recolonization carried out by the eurinoid morphogroups. After a barren interval just after the extinction boundary, the record of rhynchonellides in some epicontinental settings immediately starts in the Danian with raducal leptinoids, consisting of micromorphic species belonging to the smooth *Cryptopora* and the ribbed *Cretirhynchia* (e.g. Johansen 1989, Hakansson and Thomsen, 1999). However, *Cretirhynchia* is a raducal leptinoid genus which is also widely recorded in the Campanian-Maastrichtian background conditions prior the extinction interval (Johansen 1989). This genus commonly occurred in the assemblages from the Cenomanian-Turonian event onwards. Thus, it cannot be considered as opportunistic taxon, but an epicontinental resilient suffering miniaturization as a solution to the environmental deterioration and the shortage of resources. Otherwise, *Cryptopora* is a peculiar raducal leptinoid genus seemingly adapted to deeper environments (Surlyk and Johansen 1984). Seemingly, the epiocceanic eurinoid and falciform *Basiliocostella* could have found an effective substitute in this depth-adapted micromorphic leptinoid forms to recolonize some platforms.

Accordingly, we suggest that the diversity dynamics of the smooth rhynchonellides stock can be globally affected by post-Palaeozoic biotic crises, since diversity turnovers after crisis events are documented to a genus (or species) level, but the body plan (smooth, eurinoid, septifal/arcual) persisted: they were the typical inhabitants of the deeper epioceanic environments (Vörös 2005) who are the last to withdraw the epicontinental platforms (Norellidae, Erymnaridae?) when the environmental conditions were not propitious, and the pioneers and opportunistic taxa when the conditions are re-established in the aftermath of the biotic crises (Norellidae, Basiliolidae).

6. Conclusions

An updated Western Tethys post-Palaeozoic rhynchonellide database has been compiled to appraise conclusive inter-relationship coupling the crural architecture, microstructure of the secondary layer of the shell, and the ribbing pattern of 101 genera, testing both previous available data and new observations empirically evaluated in several taxa from the peri-Iberian basins. This analysis allows to differentiate four main rhynchonellide morphogroups: smooth eurinoid arcuiform, smooth eurinoid septifal, ribbed eurinoid septifal, and ribbed leptinoid raducal).

The changing relative occurrence of these morphogroups in response to the key extinction and diversification events recorded in the overall post-Palaeozoic history of rhynchonellides reveals a recurrent pattern with an adaptive significance beyond the phylogenetic meaning of the grouping:

The smooth, eurinoid, arcual morphogroup is a distinctive inhabitant of deeper epioceanic environments as well as the last to withdraw from the epicontinental platforms when the environmental conditions are adverse and, together with the smooth (epioceanic), eurinoid, septifal morphogroup played the role of pioneers and opportunistic taxa conducting repopulation of epicontinental habitats when the conditions were re-established after biotic crises. The ribbed, eurinoid, falciform morphogroup seems to be the more resilient stock in relation to ecological crises. Finally, the ribbed, leptinoid, raducal group mainly represents the faunal core of epicontinental communities when normal conditions predominate, evidencing an uneven

distribution unlinked to critical events, thus suggesting a more specialist pattern with a diversification related to a secondary differentiation.

The relationship between morpho-structural features defining morphogroups and their predominance or scarcity in substantial ecological crises or replacement events is probably controlled by the availability and utilization of resources. In the progressively deteriorating conditions of extinction events, the more effective and versatile forms, represented by smooth eurinoid arcual morphologies, otherwise resilient in deep environments, are dominant in both epioceanic and epicontinental habitats. Severe ecological perturbations in epicontinental seas could be only short-term withstood by the ribbed eurinoid falciform morph group, as the increasing of the filter-feeding efficiency of the lophophore is postulated for this group. When resources become abundant after crises, the repopulation is conducted mainly by smooth eurinoid septifal morphologies with opportunistic strategies, together with the aforesaid versatile smooth arcual group. When diversity increases towards normal background levels, the leptinoid raducal ribbed morphotypes are benefited due to their trophic efficiency, with economic structures and more specialized strategies.

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Table and Figure captions

Table 1. Peri-Iberian post-Palaeozoic rhynchonellides integrated in the analysis with internal and external features description. See repository acronyms in the material and methods section.

Fig. 1. Representative post-Palaeozoic rhynchonellid species from the peri-Iberian basins, combining eurinoid microstructure with septifal crural architecture and smooth ribbing pattern. **A.** *Apringia atlaeformis*, specimen O4.2. TS3, Pliensbachian, westernmost Subbetic area. A1, A2. Serial sections at 2.60 and 3.20 mm respectively from the apex showing hamiform crural development. A3, A4. Microphotographs of the secondary layer of the shell at the same sections respectively (2.60 and 3.20 mm) showing eurinoid microstructure. A5. Anterior view of the same specimen. **B.** *Soaresirhynchia bouchardi*, Early Toarcian of the Iberian Range. B1. Serial section performed at 5.00 mm from the apex on the specimen VO0.2 showing hamiform crura. B2, B3.

Microstructure of the secondary layer of the shell in a section performed at 1.70 mm from the apex (B2) in the Esc.Bo1 specimen, and at 1.30 mm from the apex in the PT.ES.1 specimen (B3). B4. Dorsal view of the SC2a.bo.1 specimen. **C.** *Mondegia atla*, Early Bajocian, Western Subbetic. C1. Serial section performed at 3.80 mm from the apex in the specimen LSR-Map-sec.1, showing incipient falciform crura. C2, C3. Microphotographs on the same specimen showing eurinoid microstructure at 1.80 and 4.60 mm from the apex respectively. C4. Dorsal and anterior views of specimen LSR.A-Aap6. **D.** *Fortunella decipiens*, specimen XA28.276, Barremian, Prebetic domain. D1. Serial section performed at 3.00 mm from the apex, showing falciform crura. D2. Detailed view of the shell in the same section showing eurinoid microstructure. D3. Dorsal and anterior view of the same specimen. **E.** *Aphelesia bipartita*, Late Tortonian, Gaudin post-orogenic Betic basin. E1. Serial section performed at 2.20 mm from the apex in the specimen AL6.AB9, showing an incipient stage of the falciform crura. E2. Dorsal and anterior view of the AL1.AB1 specimen (note that shell microstructure is not exposed because of the poor preservation of the material). **F.** *Septocrurella microcephala*, Early Bajocian, Subbetic area. F1. Serial section performed on the specimen O.7.3.Ts.1 at 2.10 mm from the apex showing septiform crura. F2. Microphotograph of the same specimen in a section at 1.80 mm from the apex showing eurinoid microstructure. F3. Anterior view of the specimen O.7.3.Ts.5.

Fig. 2. Representative post-Palaeozoic rhynchonellid species from the peri-Iberian basins, combining eurinoid microstructure with septifal crural architecture and multicostate ribbing pattern. A, B, and D derive from the Pliensbachian from the westernmost Subbetic area, and C from the Early Toarcian from the Iberian Range **A.** *Cirpa briseis*. A1. Serial section performed on the specimen CCC.3 at 2.60 mm from the apex showing fused hinge plates which evolve onwards to hamiform crura. A2. Dorsal view of the CCA.8.Cb.1 specimen. **B.** *Cirpa* sp. aff. *latifrons*. B1. Microphotograph of the I.12.T26(26).1 specimen in a section at 2.60 mm from the apex showing eurinoid microstructure. B2. Dorsal view of the same specimen. **C.** *Pseudogibbirhynchia jurensis*,

1.AR.103.1 specimen. C1. Serial section at 2.50 mm from the apex showing hamiform crura. C2. Dorsal and anterior views of the same specimen. C3. Eurinoid microstructure pattern revealed in the previous section (2.50 mm). **D. *Salgirella albertii***. D1. Serial section at 3.30 mm from the apex performed in the I12.T26(26).2 specimen, showing hamiform crura. D2. Dorsal view of O8.20.T5.1 specimen. D3. Microphotograph of the I.12.T26(26).2 specimen in a section at 2.30 mm from the apex showing eurinoid microstructure.

Fig. 3. Representative post-Palaeozoic rhychonellid species from the peri-Iberian basins, combining eurinoid microstructure with arcual crural architecture. **A. *Atycorhynchia falsiorigo***, Early Toarcian, Subbetic area. A1. Serial section performed on the specimen CE.28b.Rh1 (paratype) at 3.20 mm from the apex showing arcuiform crura. A2. Microphotograph of the same specimen in a section at 3.50 mm from the apex showing eurinoid microstructure. A3. Dorsal view of the specimen CE.28.Rh1 (holotype). **B. *Præmonticlarella distercica***, Aalenian, Iberian Range. B1. Serial section performed at 2.10 mm from the apex on the specimen FZ.68.17 (paratype) showing arcual development. B2. Microphotograph of the same specimen in a section at 1.10 mm from the apex showing wide and coarse eurinoid microstructure. B3. Dorsal and anterior views of specimen FZ.74.3 (paratype). **C. *Nannirhynchia pillula***, Late Pliensbachian, Subbetic area. C1. Serial section performed on the specimen O4.1.T1.7 at 0.80 mm from the apex showing arcuiform crura. C2. Microphotograph of the same specimen in a section at 1.40 mm from the apex showing eurinoid microstructure. C3. Anterior view of O8.23.6 specimen. **D. *Monticlarella triloboides***, specimen TO3 R51-9.1, Middle Oxfordian, Iberian Range. D1. Partial view of serial section at 3.20 mm from the apex showing arcuiform crura. D2. Detailed microphotograph of the same specimen in a section at 1.40 mm from the apex showing eurinoid microstructure. D3. Dorsal and anterior views of the same specimen.

Fig. 4. Representative post-Palaeozoic rhynchonellid species from the peri-Iberian basins, combining leptinoid microstructure with a raducal crural architecture and multicostate ribbing pattern. A-H derive from the Pliensbachian from the westernmost Subbetic area and J-K from the Sinemurian of the same Subbetic Domain. I, L derive from the Toarcian from the Iberian Range. **A.** *Cuneirhynchia cartieri*, specimen Cu.Ca1. A1. Section at 1.60 mm from the apex showing initial stage of raduliform crura. A2. Dorsal view of the same specimen. **B.** *Cuneirhynchia rastuensis*, specimen LL.A.Cu.1, B1. Section at 3.90 mm from the apex showing leptinoid microstructure of the shell. B2. Dorsal view of the same specimen. **C.** *Cuneirhynchia dalmasi*, specimen Cu.Da1. C1. Section at 3.40 mm from the apex showing leptinoid microstructure. C2. Dorsal view of the same specimen. **D.** *Prionorhynchia polyptycha*, specimen O.5.B4. D1. Section at 3.20 mm from the apex showing raduliform crura. D2. Anterior view of the same specimen. **E.** *Prionorhynchia quinqueplicata*, specimen O.9.2.1. E1. Section at 4.70 mm from the apex showing leptinoid microstructure. E2. Dorsal view of the same specimen. **F.** *Prionorhynchia guembeli*, specimen I.12.26c. F1. section at 5.10 mm from the apex showing leptinoid microstructure. F2. Anterior view of the same specimen. **G-H.** *Gibbirhynchia curviceps*, specimen O.8.20.3. G1. Section at 3.10 mm from the apex showing raduliform crura. G2. Dorsal view of the same specimen. H1. Section at 5.00 mm from the apex showing leptinoid microstructure of the shell. **I.** *Choffatirhynchia vasconcellosi*, specimen AF.11.701, paratype. I1. Section at 3.60 mm from the apex showing raduliform crura. I2. Dorsal and anterior views of the same specimen. I3. section at 2.50 mm from the apex showing leptinoid microstructure. **J.** *Alebusirhynchia jorali*, specimen O.12.10.2, paratype. J1. Section at 5.30 mm from the apex showing raduliform crura. J2. Dorsal view of the same specimen. **K.** *Alebusirhynchia vorosi*, specimen SG1.ANP7. K1. section at 3.20 mm from the apex showing leptinoid microstructure. K2. Dorsal and anterior views of the holotype of this species (specimen SG1.ANP1). **L.** *Homoeorhynchia meridionalis*, specimen VO.345. L1. Section at 3.50 mm from the apex showing raduliform crura. L2. Dorsal and anterior views of the same specimen. L3. Section at 2.10 mm from the apex showing leptinoid microstructure.

Fig. 5. Phenogram of the rhynchonellide dataset studied based on a cluster analysis performed with the Jaccard similarity index. The nodes in this multivariate approach are understood as the most significant internal and external diagnostic features herein used to link/discriminate among genera (shell microstructure, ornamentation, type of crura).

Fig. 6. Global diversity curve of post-Palaeozoic Western Tethys rhynchonellides (from PaleoDB, <http://paleobiodb.org/>, slightly modified after addition of several new recent genera) remarking the main critical ecological crises and extinction events and the major post-Palaeozoic diversification episodes.

Fig. 7. Stratigraphic range distribution and diversity of the four rhynchonellide morphogroups analysed in relation to the main extinction and diversification events of the post-Palaeozoic rhynchonellide record. Ranges are differentiated after the occurrences of the genera in epicontinental (left) or epiocceanic (right) environments. When genera occur in both environments, they are represented on both sides of the logs.

Fig. 8. Schematic model explaining the spatial distribution pattern (epicontinental vs. epiocceanic environments) of post-Palaeozoic rhynchonellide morphogroups (1 to 4) in relation to ecological crises or critical bioevents taking into account the different phases around mass extinction events.

Taxa	Author/year	Repository/specimen	Family	Age	Locality	Crural type
<i>Cirpa briseis</i>	(Gemmellaro, 1874)	DCTMA/CCC.3	Wellerellidae	Pliensbachian	Subbetic	hamifor
<i>Cirpa</i> sp. aff. <i>latifrons</i>	-	DCTMA/I.12.T26(26).1	Wellerellidae	Pliensbachian	Subbetic	hamifor
<i>Salgirella albertii</i>	(Oppel, 1861)	DCTMA/I12.T26(26).2	Wellerellidae	Pliensbachian	Subbetic	hamifor
<i>Gibbirhynchia curviceps</i>	(Quenstedt, 1852)	DCTMA/O.8.20.3	Tetrahynchidae	Pliensbachian	Subbetic	raduca
<i>Quadratirhynchia attenuata</i>	(Dubar, 1931)	DPUCM/AR.7.102	Tetrahynchidae	E. Toarcian	Iberian Range	raduca
<i>Cuneirhynchia cartieri</i>	(Oppel, 1861)	DCTMA/Cu.Ca1	Rhynchonellidae	Pliensbachian	Subbetic	raduca
<i>Cuneirhynchia dalmasi</i>	(Dumortier, 1869)	DCTMA/Cu.Da1	Rhynchonellidae	Pliensbachian	Subbetic	raduca
<i>Cuneirhynchia rastuensis</i>	(Benigni, 1978)	DCTMA/LL.A.Cu.1	Rhynchonellidae	Pliensbachian	Subbetic	raduca
<i>Homoeorhynchia meridionalis</i>	(Deslongchamps, 1863)	DPUCM/VO.3.45	Rhynchonellidae	E. Toarcian	Iberian Range	raduca
<i>Choffatirhynchia vasconcellosi</i>	(Choffat in Dubar, 1931)	DPUCM/AR.11.501	Rhynchonellidae	E. Toarcian	Iberian Range	raduca
<i>Rhynchonelloidea goyi</i>	García Joral, 1982	DPUCM/RE.6	Rhynchonellidae	L. Toarcian	Iberian Range	raduca
<i>Prionorhynchia guembeli</i>	(Oppel, 1861)	DCTMA/I.12.26c	Prionorhynchiidae	Pliensbachian	Subbetic	raduca
<i>Prionorhynchia quinqueplicata</i>	(Zieten, 1832)	DCTMA/O.9.2.1	Prionorhynchiidae	Pliensbachian	Subbetic	raduca
<i>Prionorhynchia polyptycha</i>	(Oppel, 1861)	DCTMA/O.5.34c	Prionorhynchiidae	Pliensbachian	Subbetic	raduca
<i>Globirhynchia subobsoleta</i>	(Davidson, 1852)	DPUCM/RE.1.36	Cyclothyrididae	Aalenian	Iberian Range	raduca
<i>Alebusirhynchia jorali</i>	Baeza-Carratalá, 2011	DCTMA/O.12.10.2	Cyclothyrididae	L. Sinemurian	Subbetic	raduca
<i>Alebusirhynchia vorosi</i>	Baeza-Carratalá, Dulai & Sandoval, 2018	DCTMA/SGI.ANP7	Cyclothyrididae	L. Sinemurian	Subbetic	raduca
<i>Apringia atlaeformis</i>	(Böse, 1897)	DCTMA/O4.2.TS3 DPUCM/VO.0.2	Basiliolidae	Pliensbachian	Subbetic	hamifor
<i>Soaresirhynchia bouchardi</i>	(Davidson, 1852)	DCTMA/Esc.Bo1 DCTMA/PT.ES.1	Basiliolidae	E. Toarcian	Iberian Range	hamifor
<i>Mondegia atla</i>	(Oppel, 1863)	DCTMA/LSR-Map-sec.1	Basiliolidae	E. Bajocian	Subbetic	falcifor
<i>Pseudogibbirhynchia jurensis</i>	(Quenstedt, 1858)	DPUCM/1.AR.103.1	Basiliolidae	E. Toarcian	Iberian Range	hamifor
<i>Aphelesia bipartita</i>	(Brocchi, 1814)	DCTMA/AL6.AB9	Basiliolidae	L. Tortonian	post-orogenic Betic basins	falcifor
<i>Fortunella decipiens</i>	(D'Orbigny, 1847)	DPUGR/XA28.276	Basiliolidae	Barremian	Prebetic	falcifor
<i>Septocrurella microcephala</i>	(Parona, 1896)	DCTMA/O.7.3.Ts.1	Erynnariidae	E. Bajocian	Subbetic	septifor
<i>Praemonticlarella distercica</i>	García Joral, 1993	DPUCM/FZ.68.17	Norellidae	Aalenian	Iberian Range	arcual
<i>Atychorhynchia falsiorigo</i>	Baeza-Carratalá, Reolid & García Joral, 2017	DCTMA/CE.28b.Rh1	Norellidae	E. Toarcian	Subbetic	arcual
<i>Monticlarella triloboides</i>	(Quenstedt, 1858)	DPUCM/TO3 R51-9.1	Norellidae	M. Oxfordian	Iberian Range	arcual
<i>Nannirhynchia pillula</i>	(Schlosser in Böse & Schlosser, 1900)	DCTMA/O4.1.T1.7	Norellidae	L. Pliensbachian	Subbetic	arcual

Linking Western Tethyan Rhynchonellide morphogroups to the key post-Palaeozoic extinction and turnover events

José Francisco Baeza-Carratalá^{a*} and Fernando García Joral^b

Highlights:

Rhynchonellides are arranged coupling shell-microstructure, crura type and costation

Singular morphogroups are linked to key post-Palaeozoic crisis/diversification events

A recurrent pattern with adaptive meaning beyond a phylogenetic meaning is revealed

Adaptive response to extinction events differ in epicontinental/epioceanic seas

Resources availability or filter-feeding efficiency regulated distribution of groups

Journal Pre-proof

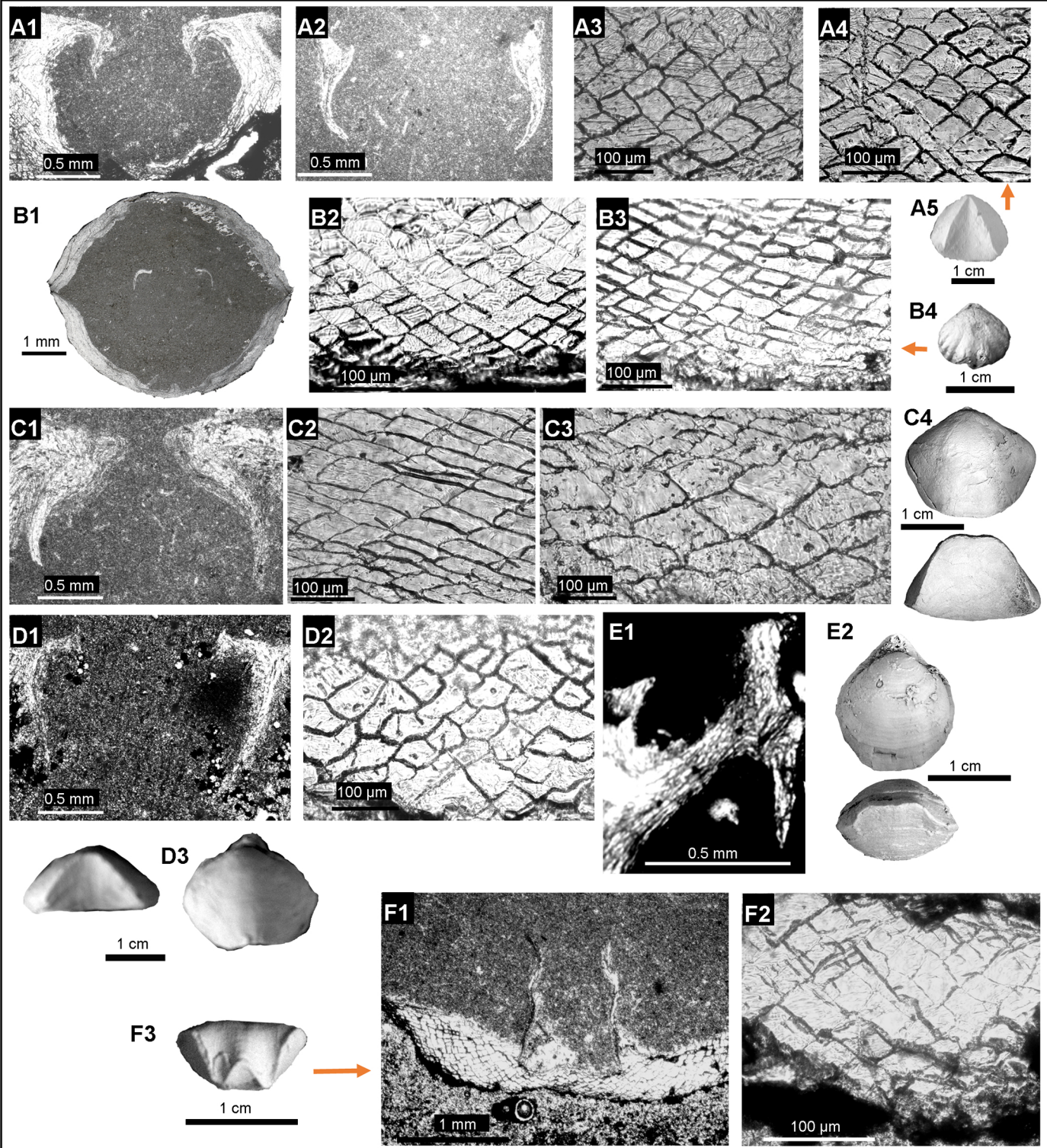


Figure 1

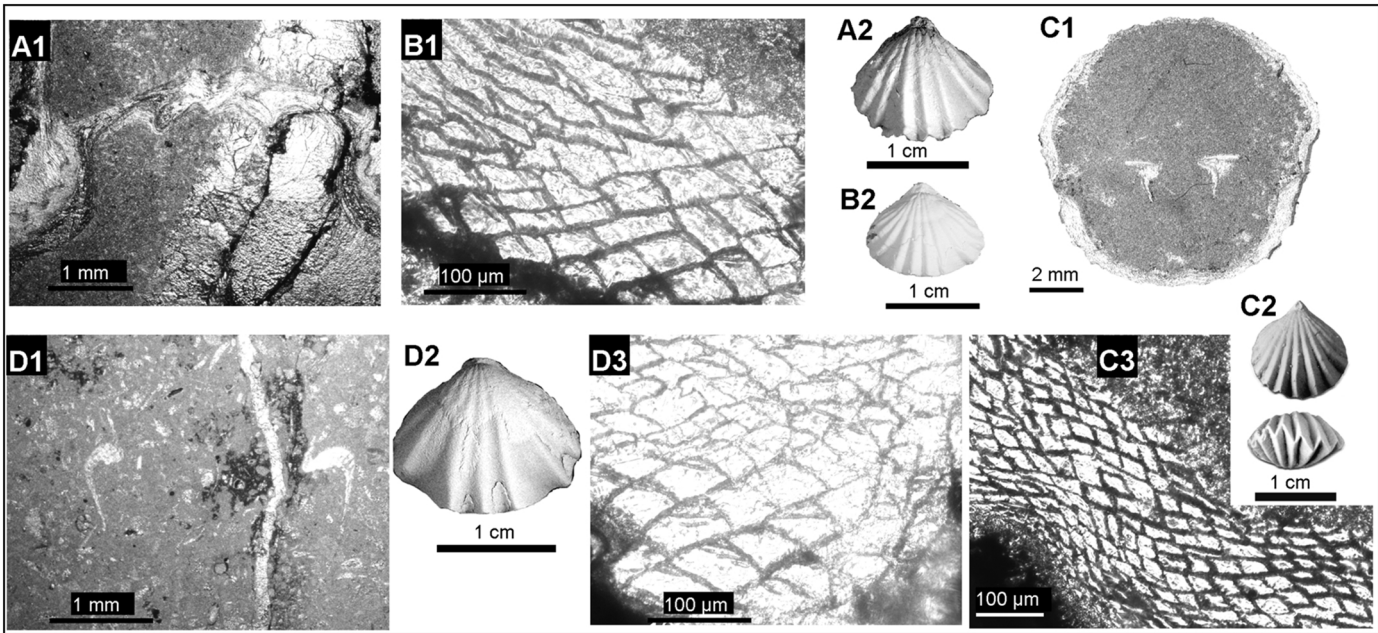


Figure 2

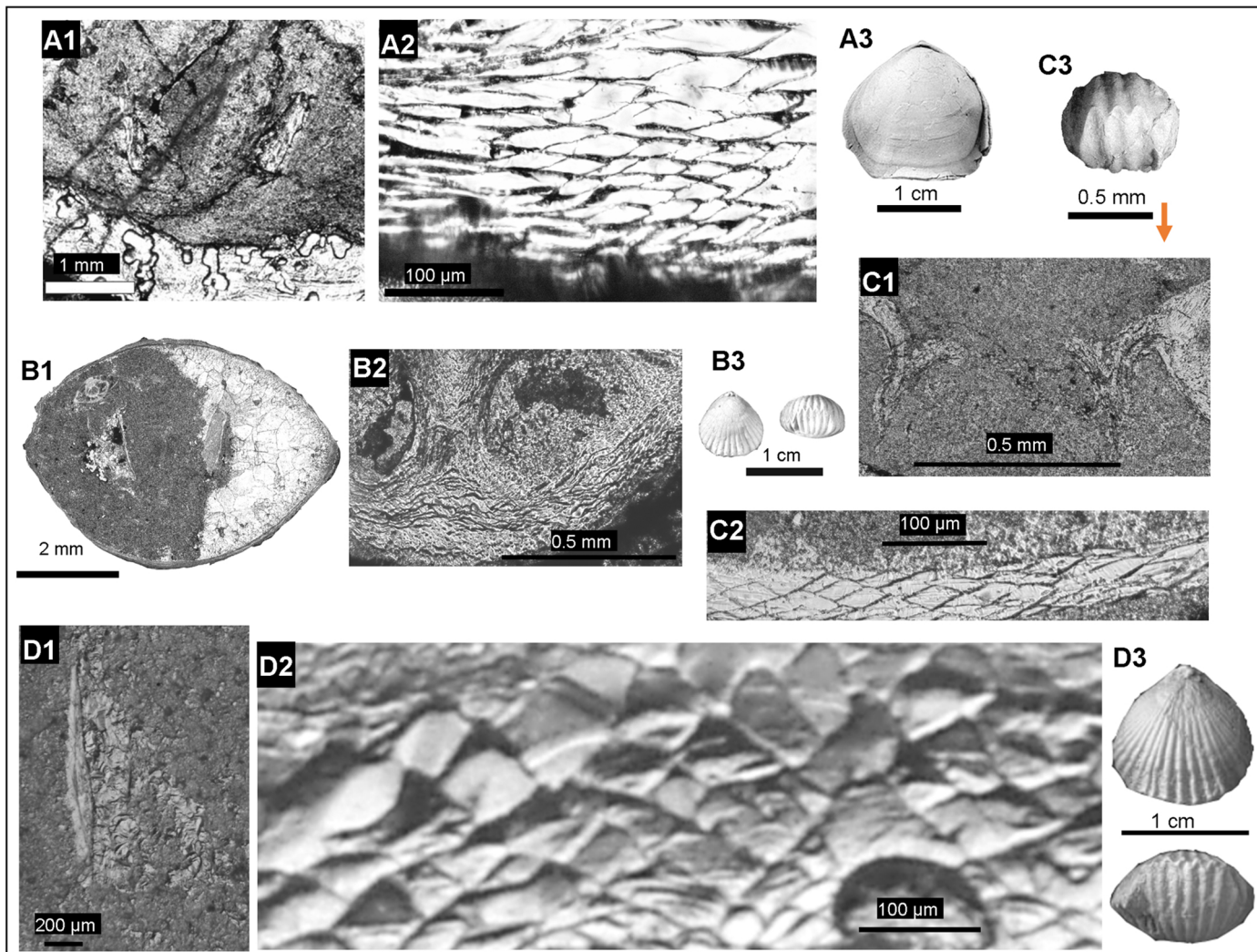


Figure 3

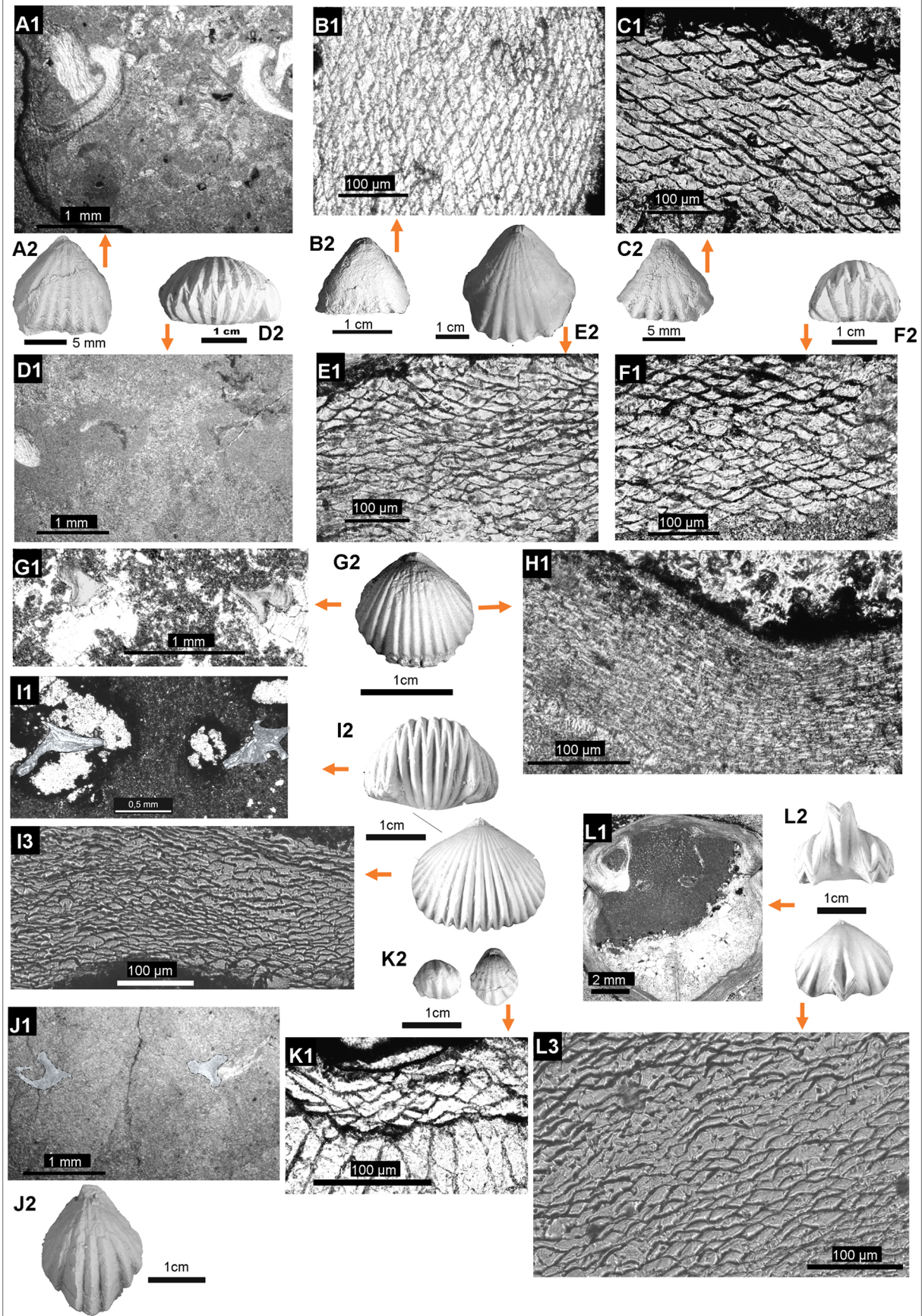


Figure 4

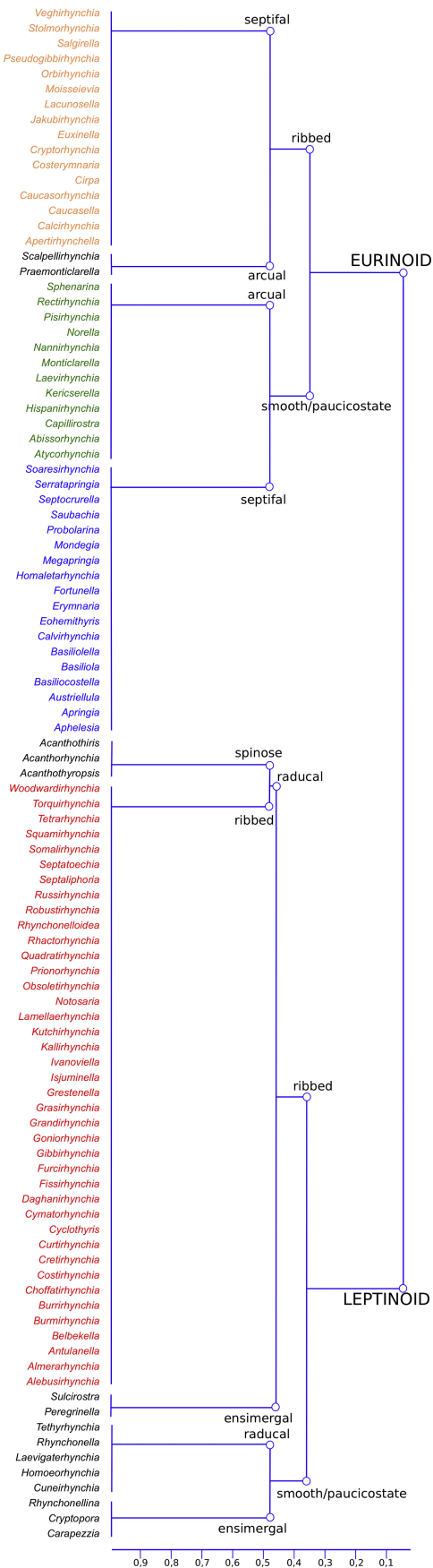


Figure 5

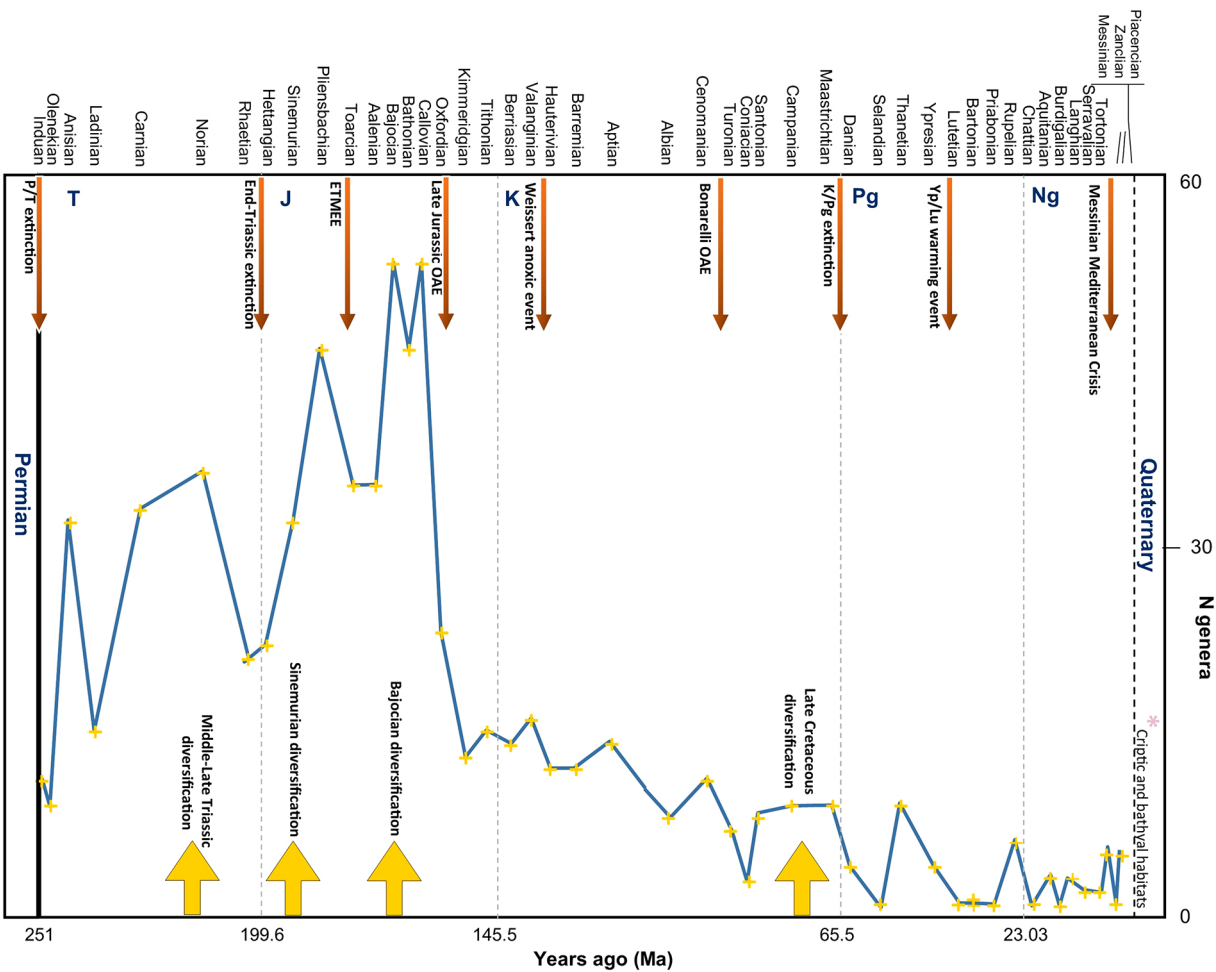
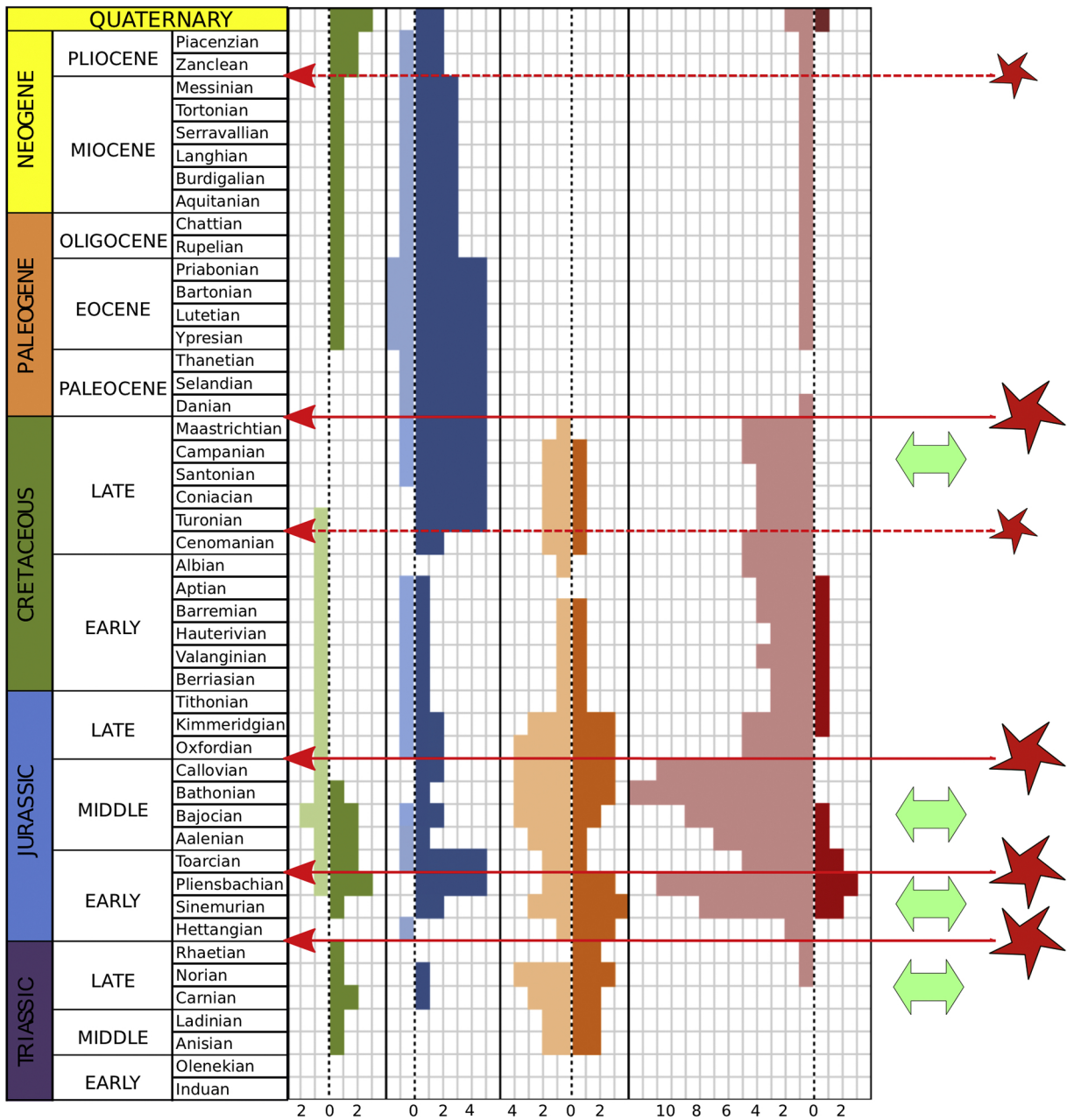


Figure 6



Diversification Event

Extinction Event

①	②	③	④
ARCUIFORM	SEPTIFAL		RADUCAL
SMOOTH/PAUCICOSTATE		RIBBED	
EURINOID			LEPTINOID

number of genera

Epicontinental | Epicoeanic/
deep

Figure 7

Epicontinental environment

Epiocceanic and deeper environments

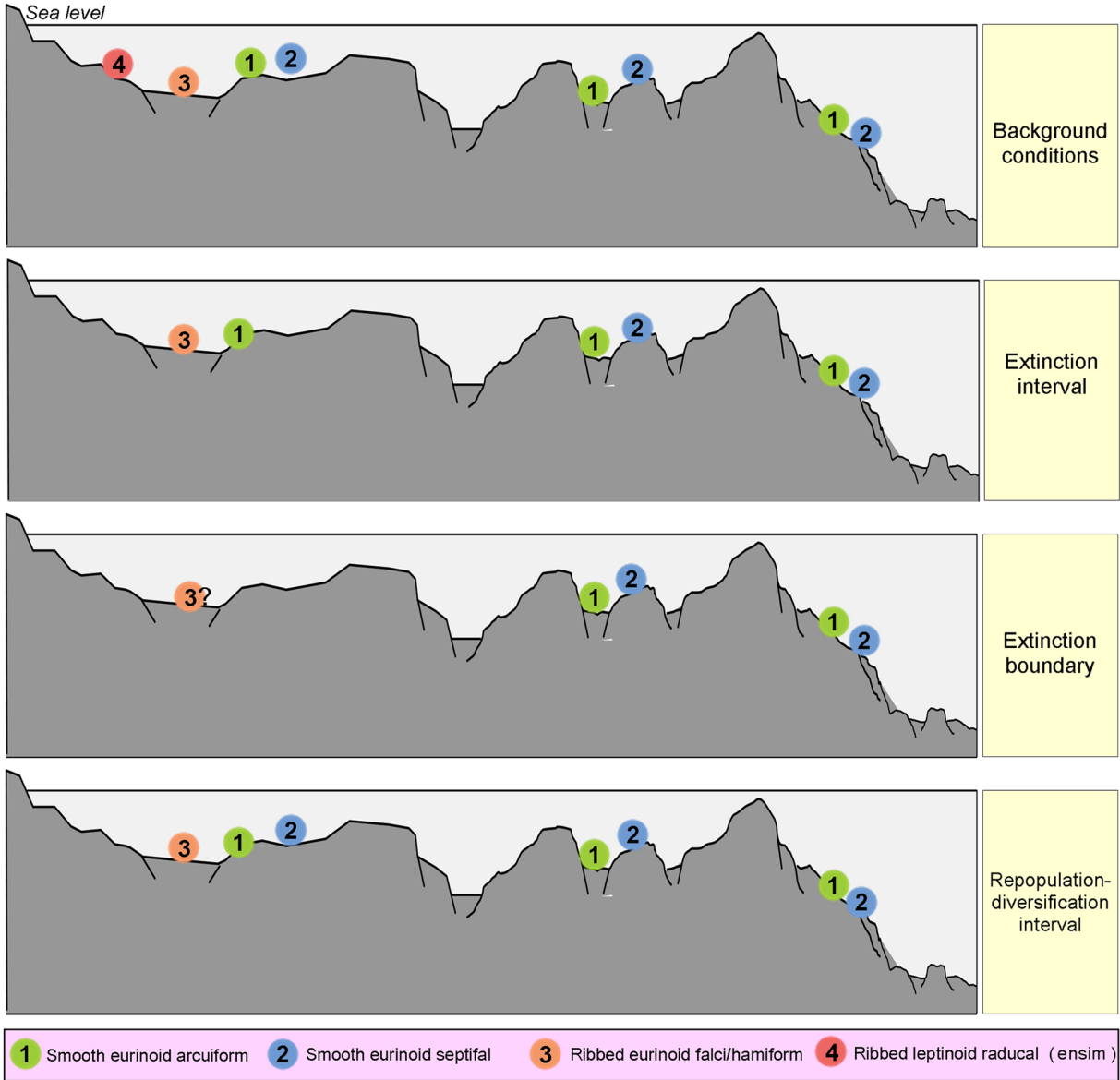


Figure 8