



## Evolutionary Transformations of Sporoderm Ultrastructure in Certain Monophyletic Lineages of Higher Plants

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### ABSTRACT

Data of transmission electron microscopy can be successfully used to reveal evolutionary transformations, which occurred in pollen or spores during the existence of monophyletic lineages of higher plants. On the basis of such data, one can discover gradual transitions in groups, which include new forms, that at the first glance seem to be totally dissimilar to the predecessors. One can trace sequences of morphological transitions within lineages from the earliest to latest members of a given group and connect the first and last members, which do not share any common features. Hidden transformations are traceable in groups, which seemed unchanged during the entire period of their existence. Parallelisms can be revealed between unrelated groups of higher plants not only in final forms, but also in succession of transformations. The successful application of this approach has been exemplified in lycopodioidaleans, peltasperms, equisetaleans, and the Cheirolepidiaceae. Particularly promising seem groups that have a long geological history and survive until now.

**Keywords:** sporoderm ultrastructure, evolutionary transformations, peltasperms, lycopodioidaleans, horsetails, Cheirolepidiaceae

### РЕЗЮМЕ

**Завьялова Н.Е.** Эволюционные преобразования в ультраструктуре спородермы в некоторых монофилетических группах высших растений. Данные трансмиссионной электронной микроскопии могут с успехом применяться для выявления эволюционных преобразований в ультраструктуре пыльцевых зерен или спор, происходивших в монофилетических линиях высших растений. На основе таких данных можно находить постепенные переходы в тех случаях, когда новые и совершенно не похожие на предшественников формы возникали в геологической летописи на первый взгляд внезапно. Можно проследить последовательные морфологические преобразования в ряду от самых ранних к самым поздним представителям определенной группы, что удастся даже если между первыми и последними членами цепи вообще не наблюдается сходных признаков. Скрытые преобразования удастся проследить в группах, казалось бы оставшихся практически неизменными в течение всего периода их существования. Могут быть выявлены параллелизмы между неродственными группами высших растений, касающиеся не только отдельных стадий, но и последовательностей преобразований, которые к этим стадиям привели. В качестве примеров рассмотрены лигноптерисовые и пельтаспермовые семенные папоротники, хвощи, хейролепидиевые. Особенно перспективными представляются группы, имеющие длительную геологическую историю и дожившие до настоящего времени.

**Ключевые слова:** ультраструктура спородермы, эволюционные преобразования, пельтаспермовые, лигноптеридиевые, хвощи, хейролепидиевые

In my perception, the morphology and ultrastructure of fossil pollen and spores are such an interesting subject that even little discoveries of novelties are worth of spent efforts. It is even truer nowadays, when few scientists continue dealing with the ultrastructure of fossil exines. Quite often, when I am curious to learn about the inner structure of a given type of fossil spores or pollen, a search in the published literature does not help, and the only way to get this information is to study it with TEM myself. The current knowledge on the sporoderm is very limited and the situation will hardly change to the best within my lifetime. However, already obtained data invite to generalize over

them, in spite of the fact that they are always incomplete and often incomparable. There are "palynological puzzles" that have attracted the attention of pollen morphologists for a long time (Chaloner 2013). Some of such puzzles seem less unsolvable than others. For example, this is probably the case of pollen and spores of some monophyletic groups of higher plants and transformations that took place in their morphology and sporoderm ultrastructure in course of the evolution. Using such groups as preferable objects of ultrastructural studies of sporoderms could be very prospecting in the following aspects:

- It often seems that new forms appeared in some long-living groups of higher plants suddenly and that they were completely dissimilar to earlier forms of the group. Ultrastructural studies can show that morphological transitions did exist and gradational evolution did take place even in such cases.
- Moreover, we can confidently trace a succession of morphological transformations from the earliest to latest members of a given group. As a result, one can reveal a connection between the first and the last links of the lineage, even in the case when the latest links have lost any similarity to the earliest links.
- We can prove that morphological evolution did take place in groups which appeared unchanged during the entire period of their existence.
- We can discover parallelisms between unrelated groups not only in final evolutionary stages, but also in chains of morphological transformations, which led to them.

Below these aspects are considered in several examples, such as peltasperms, lyginopteridaleans, equisetaleans, and the Cheirolepidiaceae.

### A transition between such dissimilar types as saccate and non-saccate pollen: the Peltaspermaceae

The Permian/Triassic family of seed ferns Peltaspermaceae is an example, where a morphological transition was revealed between apparently dissimilar earlier and later forms. The family is very diverse in pollen morphology by comparison to other gymnosperm families. Peltasperm pollen organs are known to contain the following pollen types: bisaccate striate *Protobaploxypinus*, bisaccate non-striate *Vesicaspora* and *Falcisporites*, non-saccate striate *Vittatina*, and non-saccate boat-shaped *Cycadopites* (Gomankov 1986, Zavialova & Karasev 2015). Of interest is that nearly all of these pollen types occur not only in peltasperms. The distribution of these types in peltasperms through time is also peculiar. All, but *Cycadopites* and *Falcisporites*, are known from peltasperm pollen organs from Permian deposits. *Falcisporites* associates with peltasperms both in Permian and Early Triassic deposits. Late Triassic members of the family are characterized only by *Cycadopites* pollen type. By contrast to pollen grains, pollen organs of early and late peltasperms are very similar to each other. The Late Paleozoic fashion on striate and saccate pollen grains, which is known in many unrelated gymnosperm groups, was replaced in the Mesozoic by a fashion on boat-shaped monosulcate non-saccate pollen grains. Peltasperms had their fling to both fashions. In sum, their pollen types were very diverse during the Permian, but were represented by the only one and completely different pollen type at the end of the Triassic.

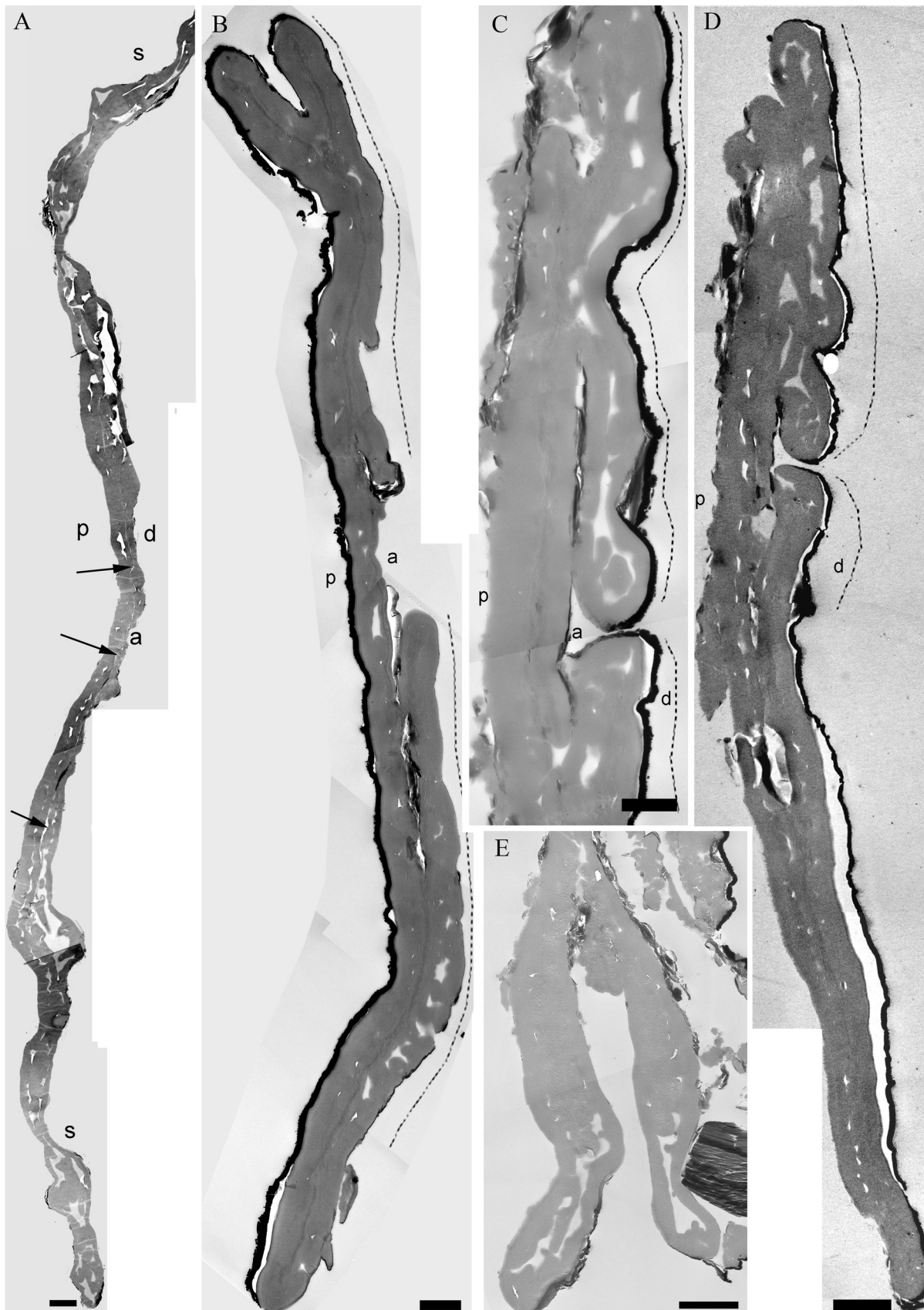
Until now, the rich diversity of pollen types in the Permian peltasperms has remained unexplainable; and no transitions have been known towards non-saccate monosulcate pollen of late peltasperms. However, it seems that now we can better understand the sudden appearance in the Triassic of a non-saccate pollen type, which replaced the earlier diversity of pollen types, most of which were saccate. Studies of pollen of Permian (Zavialova et al. 2001) and Triassic (Zavialova & Van Konijnenburg-van Cittert 2011) peltasperms allowed Zavialova & Van Konijnenburg-van

Cittert (2011) to hypothesize how a transition could have taken place from earlier saccate pollen to later non-saccate pollen. They studied pollen grains from pollen organs of *Antevsia zeilleri* (Nathorst) Harris 1937 from the Rhaetian of Germany and found that areas of the ectexine that surround the sulcus are thicker than the ectexine of other regions and even slightly resemble sacchi by the ultrastructure. Sacchi in non-striate saccate pollen grains of *Vesicaspora* type, which seems the most suitable candidate for this transformation among Permian pollen types of peltasperms (Fig. 1A), could have reduced up to the extensions surrounding the distal sulcus, as it was observed in ultrathin sections of pollen grains of *A. zeilleri* (Fig. 1B–1E). Although a non-saccate pollen type is known in Permian peltasperms (*Vittatina*), its granulate exine is much more different from that of pollen of *Antevsia* than the exine of *Vesicaspora*. Among the three saccate types of Permian peltasperms, *Vesicaspora* was chosen for this reconstruction, because this pollen is not striate; as to *Falcisporites*, it still waits for a thorough ultrastructural study, which would be better carried out on an in situ material.

A similar morphological transformation was earlier suggested for pollen grains of *Ginkgo biloba* as a transition to them from saccate pollen of a hypothetical ancestor (Sahashi & Ueno 1986). Therefore, the morphological transition which was reconstructed for pollen of early and late peltasperms is also an example of a parallelism to an analogous transformation that was reconstructed in a different group of gymnosperms.

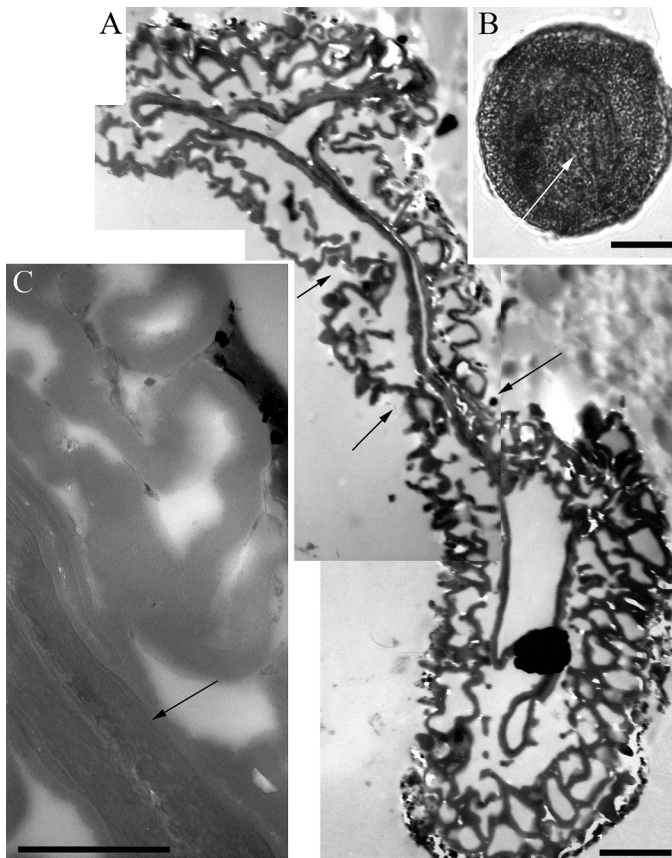
### Development of a distal aperture in primitive prepollen: the Lyginopteridales

The seed fern order Lyginopteridales is primitive as far as concerns the general morphology of their prepollen. Prepollen grains found in the majority of their pollen organs are assignable to dispersed genera *Cyclogranisporites* or *Punctatisporites* (Balme, 1995). They are sub-circular, granulate/punctate, with a trilete proximal scar, and indistinguishable from spores in dispersed state. The ultrastructure of prepollen grains extracted from *Crossotheca hughesiana* also confirms the primitiveness of the order: the only aperture is the proximal scar, and the ultrastructure of the sporoderm is homogeneous (Taylor & Taylor 1987; fig. 7). Both features are intrinsic to spores rather than pollen. Orlova et al. (2009) studied the fine morphology of prepollen grains associated with the lyginopteridalean *Telan-giopsis nonnae* Orlova & Zavialova 2009. By general morphology, these prepollen grains also would be defined as spores if found in dispersed state (Fig. 2B). However, their ectexine is alveolate, and the endexine is lamellate – both features are clearly gymnospermous (Fig. 2C). In addition, although no clearly shaped distal aperture was observed under LM and SEM, ultrathin sections show thinned areas in the exine on both sides of the sporoderm (proximal and distal; Fig. 2A). That means that not only a proximal scar was observed, but also the first step towards the distal aperture (Orlova et al. 2009). The ultrastructure of these prepollen grains is more advanced than their general morphology; it reveals a shift from the spore towards pollen state, that is yet undetectable at the level of general morphology.



**Figure 1** Ultrathin sections of pollen of *Vesicaspora* sp. and *Antersia zeilleri* (Nathorst) Harris 1937 (illustrations are reproduced from Zavialova 1998 and Zavialova & Van Konijnenburg-van Cittert 2011). A. Pollen of *Vesicaspora* sp., composite longitudinal section through the body and sacci, proximal face to the left, arrows point to the obliterated gametophyte hollow. B–E. Pollen of *A. zeilleri*. B. Composite section shows thinned ectexine in the apertural region and extensions of the exine bordering the aperture, proximal face to the left. C. Enlargement of D. D. Composite section of a pollen grain, proximal exine shows regular alveolae, distal-equatorial extensions surround the aperture. E. Section of a pollen grain compressed in lateral position, distal face to the bottom of the figure. Scale bars (A, D, E) 1  $\mu$ m, (B, C) 0.67  $\mu$ m; (a) aperture, (d) distal face, (p) proximal face, (s) saccus, intermittent line indicates the position of extensions surrounding the aperture





**Figure 2** Prepollen grains of *Telangiopsis nonnae* Orlova & Zavialova 2009 (illustrations are reproduced from Orlova et al. 2009, pl. VIII). A. Composite image of an ultrathin section, thinned areas are present on both sides of the exine (arrows). B. General view of a prepollen grain, arrow indicates one of three rays of a closed proximal scar, LM. C. Enlargement of the compressed exine, arrow points on the endexine. Scale bar (A) 1.67  $\mu\text{m}$ , (B) 10  $\mu\text{m}$ , (C) 0.5  $\mu\text{m}$

### A succession of transformations connects dissimilar morphological types: the Equisetales

The next example demonstrates a succession of morphological transformations, which connect totally dissimilar first and last members of a group. It was shown that apertural features in spores are usually clearly visible in suitable sections of mature exospores, and are quite different in spores of filicopsids, homosporous and heterosporous lycopsids, and living sphenopsids, and, therefore, they can be successfully used to differentiate between taxa of higher rank of spore-bearing plants (Tryon & Lugardon 1990). This premise allowed Lugardon & Brousmuche-Delcambre (1994) and Grauvogel-Stamm & Lugardon (2009) to reveal evolutionary transformations in the spore morphology and ultrastructure within the Equisetales.

Spores of modern *Equisetum* have an indistinct small oval aperture. The exospore consists of only two layers (outer and inner exospores), of nearly equal and uniform thickness (Fig. 3e). The outer exospore thins abruptly in the apertural region and forms a pronounced hollow delimiting the exact site of the aperture. The exospore is devoid of any fold in the apertural region, but is inwardly reinforced with a thick, biconvex, lens-shaped mass of sporopollenin, the so-called "sub-apertural obturator", which

is attached below the thin apertural wall. It is a unique ultrastructural feature, which is only known in spores of horsetails. The extra-exospore layer (episore) is present as a thin envelope (Tryon & Lugardon, 1990). Lugardon studied the sporoderm ultrastructure of in situ *Calamospora* from Carboniferous sphenophytes (Lugardon & Brousmuche-Delcambre, 1994). The spores are subcircular, with a trilete proximal scar. In the apertural area, the exospore is thickened and consists of three layers: the lower one is more electron-dense, thin, and continuous; the middle one shows a complex stratification with sheets, canals and cavities; and the outer one is much thicker than the inner layer, consists of amorphous sporopollenin and nearly lacks any structural features (Fig. 3a). The apertural fold is strongly protruding. No extra-exospore layers are present. This type of sporoderm ultrastructure differs drastically from that of modern horsetails and, paradoxically, show many similar features to certain ferns, particularly to the Ophioglossales (Lugardon & Brousmuche-Delcambre, 1994).

It should be pointed out that the latest members of the chain share no common features with the earliest members. To solve this paradox, Lugardon and his co-authors studied spores of horsetails of Triassic and Jurassic ages. They discovered transitional steps in the sporoderm ultrastructure and documented the appearance and further development of features, which later became specific of spores of modern horsetails. The Triassic spores are very close to the Carboniferous spores by the apertural features and sporoderm ultrastructure, but show a tendency of the apertural slit to open at the base and the apertural fold to become less protruding (Fig. 3b). Sporopollenin globules fill a widened area of the apertural fold (Fig. 3c); this filling represents the precursor of the obturator of *Equisetum* spores. An extra-exosporial structure also appeared (extra-exosporial layer is lacking in the Carboniferous spores and is present in modern spores). The Jurassic member shows a further step in the evolutionary modifications (Fig. 3d). These spores seem to be alete and bear short elaters. Although they still shows an apertural fold, the exospore is not thickened in this area anymore and does not show the foliated substructure which characterizes the Carboniferous and Triassic spores. The apertural fold is broadly open and occupied by a mass of sporopollenin that unambiguously corresponds to the obturator. The only primitive vestiges are a slightly developed apertural fold and a few small cavities along its both sides (Grauvogel-Stamm & Lugardon 2009).

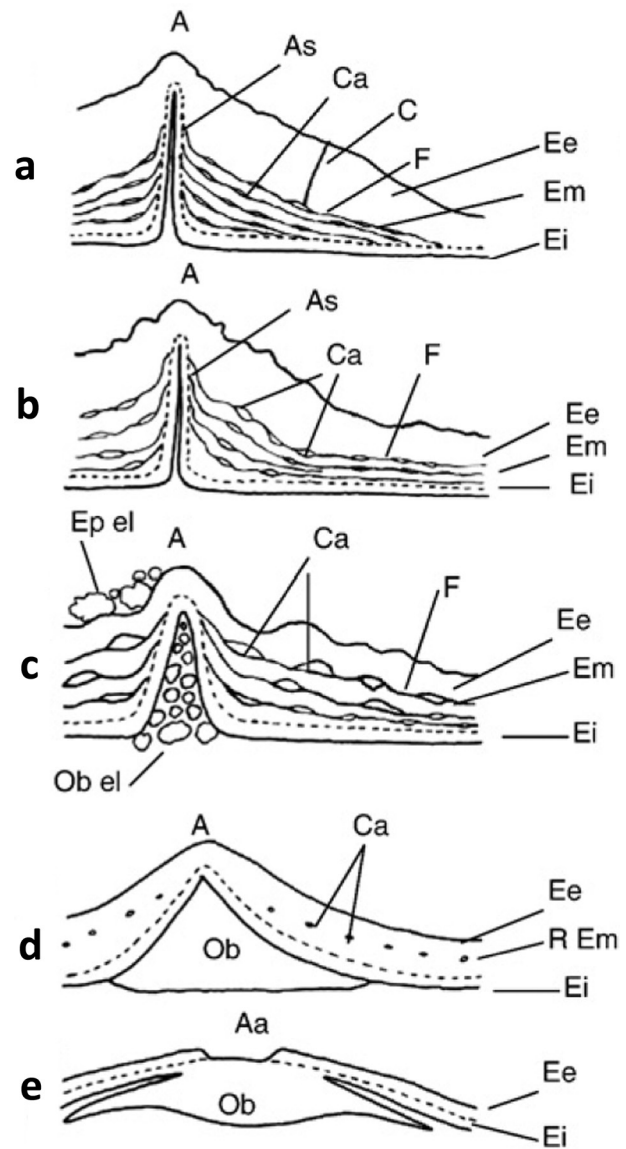
This is an example of how a study of spore morphology and ultrastructure can shed light on the gradational evolution in a plant group, which in the absence of ultrastructural data appeared sudden and interrupted. The above examples about lyginopteridalean prepollen and peltasperm pollen showed only one morphological transformation in each group, whereas the study of horsetail spores discovered several successive transformations. One can connect the first and the last links of chain, even in the case when the latest links have lost any similarity to the earliest links (Fig. 3a, 3e).

### Hidden ultrastructural transformations in seemingly unchanged pollen: the Cheirolepidiaceae

By means of ultrastructural studies, one can reveal that morphological evolution of pollen or spores did take place even in groups, in which they appeared unchanged during the entire period of the existence of a given group. This will be shown in the Cheirolepidiaceae family, which is a long-living group of gymnosperms. Macrofossil records show that the Cheirolepidiaceae appeared as early as in the Late Triassic and became extinct at the end of the Cretaceous (Watson 1988). According to data on dispersed pollen, they appeared in the Late Triassic, existed during the Jurassic and Cretaceous, and, possibly, not for long survived into the Paleogene (Petrosianz & Bondarenko 1983, Barreda et al. 2012). At the first glance, their pollen morphology is very stable. Watson (1988) noted that *Classopollis* pollen type is perhaps the only character that all members of this family shared. However, the fact that the general morphology of pollen seems unchanged does not necessarily mean that no changes happened to the sporoderm ultrastructure. On the contrary, the above example of fossil sphenopsids shows that Carboniferous and Triassic spores of the same morphological type differ in the ultrastructure (Grauvogel-Stamm & Lugardon 2009).

Pollen grains of *Classopollis* were among the first fossil pollen types ever observed with TEM (Pettitt & Chaloner 1964); later, they also have been repeatedly studied with help of TEM (see references in Pocock et al. 1990, Kedves 1994, Zavalova et al. 2010ab). Since the already accumulated information on the exine ultrastructure of *Classopollis* is relatively voluminous, this pollen is a suitable model object for tracing transformations in the inner structure of the exine in course of the evolution of the parent group. Pollen grains of *Classopollis* are spheroidal, with a small trilete proximal scar, a distal cryptopore, a rimula (= subequatorial circum-polar canal), and an equatorial girdle. The ectexine comprises a double tectum and an infratectum; a foot layer is probably present in some pools of specimens, but lacking in others (Lugardon 1985, Kedves 1994). Several variants of the endexine structure are known, but no directional changes in the endexine are revealed. Even pollen grains from the same sample are shown to have slightly different endexine, supposedly due to slightly different ontogenetic stages, and no endexine, supposedly due to preservation (e.g., Zavalova et al. 2010a). The most famous feature of this pollen type is its ectexine that resembles columellate ectexine of angiosperms and is a common example of angiospermization.

For the aim of the present paper, I focus on the ectexine, since several variants of the infratectum arrangement have been observed in this sublayer and these variants seem to alternate through geological time (Fig. 4). The earliest pollen grains of *Classopollis* show a rather peculiar granular infratectum. Large granules are mostly arranged in one row, being sandwiched between overlying and underlying layers. This type was revealed in Rhaetian and some Jurassic pollen grains (Fig. 4B, 4C): in *Classopollis meyerana*, *C. torosus*, and *Geopollis zwołinskai* from the Rhaetian of England (Zavalova et al. 2010) and *Classopollis harrisii* from the Rhaetian-Liassic of France (Médus, 1977).



**Figure 3** Evolution of the apertural ultrastructure in the Equisetales (figure is redrawn from Grauvogel-Stamm & Lugardon 2009: fig. 1).

a: *Calamites* spore of the *Calamospora* type, Carboniferous; b: *Schizoneura* – *Echinostachys*, Triassic; c: *Equisetites* – *Equisetostachys*, Triassic; d: *Equisetum columnare*, Jurassic; e: *Equisetum* sp., present-day.

A = apertural fold; Aa = apertural area; As = apertural slit; Ee = external layer of the exospore; Ei = inner layer of the exospore; F = sheet; C = canal; Ca = cavity; Ob = sub-apertural obturator; Ob el = element of the sub-apertural obturator; Ep el = epispore elements. See Grauvogel-Stamm & Lugardon (2009, 2010) for a more detailed legend

Pollen grains with such an infratectum were later replaced by pollen with an infratectum that was composed of more elongated elements (Fig. 4D); it more and more resembled the columellate infratectum of angiosperm pollen. The structural elements are firmly attached to the overlying layer and can freely hang towards the underlying layer or touch it (Fig. 4E). Pollen grains with such an infratectum were produced during the Jurassic and Early Cretaceous. This transition (from large granules mostly arranged in one row to columella-like elements) is easy to imagine. Some Jurassic *Classopollis* show alternating columellate-like and



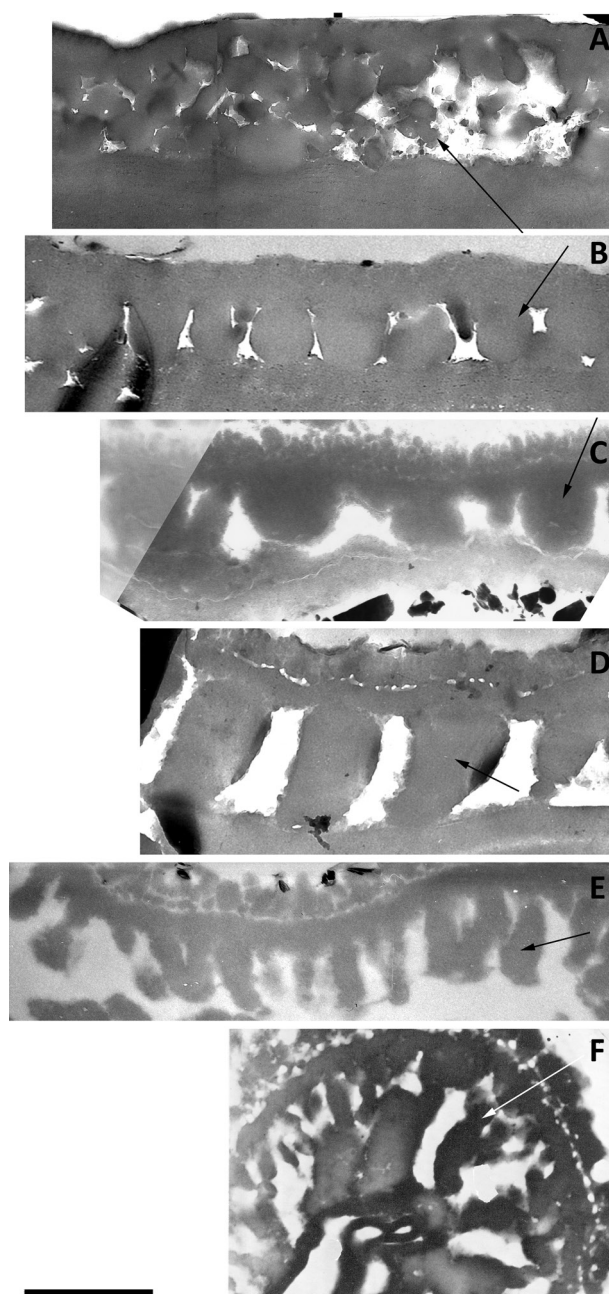
granular elements (e.g. *C. meyerana* from the Callovian? of Kazakhstan<sup>1</sup>, Tekleva & Krassilov 2009; and *C. torosus* from the Oxfordian of England, Rowley & Srivastava 1986). However, some other Jurassic and Cretaceous pollen show an infratectum of only columellate-like elements (*C. torosus* from the Callovian of Siberia, Zavialova 2003; *C. torosus* from the Barremian of England, Taylor & Alvin 1984; and *C. martinottii* from the Albian-Cenomanian?<sup>2</sup> of Lebanon, Zavialova et al. 2010b). Only *C. torosus* from England does not fit to this scheme, showing a columellate-like ectexine already in the Rhaetian–Liassic (Pettitt & Chaloner 1964, pl. I, 1, fig. 1), but I think that additional sections (only one section was shown in the paper) probably would have revealed granules alternating with columellate-like elements.

The third type demonstrates branching columellate-like elements (Fig. 4F); it was found in the Late Cretaceous *Classoidites glandis* Van Ameron 1965 (Médus 1977, Kedves 1994). Therefore, we can hypothesize rather confidently that the earlier type of the infratectum (one row of large granules) transformed into a columellate-like infratectum in more advanced members; and this infratectum finally evolved in the branching-columellate type in late members of the group.

*Classopollis* belongs to the morphological group Circumpolles, which included also other pollen types and was rather diverse during the Late Triassic. Apart from *Classopollis*, no pollen types of this group have ever been found in situ in cheirolepidiaceae cones (as well as in cones of other plant groups). However, in addition to the morphological similarity between *Classopollis* and other members of Circumpolles, there are some more indirect indices implying that these pollen grains were produced by members of the Cheirolepidiaceae as well. They have been found in the same beds that contain dispersed cheirolepidiaceae cuticles as well as conifer amber and wood. These indices allow one to treat early Circumpolles in the analysis as possible cheirolepidiaceae pollen with a sufficient degree of certainty (Roghi et al. 2006). If we

<sup>1</sup> Krassilov et al. (1997) found these pollen grains in guts of a fossil insect from the Mikhailovka (= Aulie) locality of the Karatau Mountain Range in Kazakhstan. Although they mentioned a Late Jurassic age of the enclosing deposits, I state a Callovian with a question mark in the present paper, since Krassilov et al. (1997) repeated the information from Doludenko & Orlovskaya (1976), who provided paleobotanical and palynological dating. At those times, Soviet paleontologists considered the Callovian as a part of the Upper Jurassic, and while Doludenko & Orlovskaya (1976) stated that the age is a Late Jurassic, they wrote that the list of pollen and spores they discussed and fossil plants they studied suggested that the age of these lacustrine marls was most probably Callovian.

<sup>2</sup> Dr. Dejax (Muséum national d'histoire naturelle, Paris; personal communication; Azar et al. 2011) believes that the palynological assemblage with *C. martinottii* is much older in age than an Albian–Cenomanian. He suspects a Barremian or Aptian age on the basis of photos of the members of the assemblage (Zavialova et al. 2010b) and his knowledge about the stratigraphy of the region, but refrains from any conclusion as far as he did not study the assemblage himself. Therefore, I cite the original dating made in our study by Sophia Smirnova (Zavialova et al. 2010b), but add a question mark. The probable older age of these Lebanon deposits does not influence the proposed reconstruction of the ectexine evolution, though an unequivocal understanding of the age of these deposits is highly desirable.



**Figure 4** Transformations of the infratectum in members of Circumpolles. Early members of Circumpolles (A: *Duplicisporites granulatus*, Carnian, Italy) had an infratectum of numerous small irregularly distributed granules. Since the Rhaetian an infratectum appeared that consists of large granules arranged predominantly in one row (B: *Classopollis torosus*, Rhaetian, England and C: *Classopollis meyerana*, Callovian?, Kazakhstan). This infratectum gradually transformed during the Jurassic into a columellate-like infratectum (D: *Classopollis torosus*, Callovian, Siberia and E: *Classopollis martinottii*, Albian–Cenomanian?, Lebanon). In the Late Cretaceous, the columellate-like infratectum gave rise to an infratectum with branching columellate-like elements (F: *Classoidites glandis*, Turonian, France). Scale bar 1  $\mu$ m. Arrows point on elements of the infratectum. The illustrations are reproduced from Zavialova & Roghi 2005 (A); Zavialova et al. 2010a (B); Tekleva & Krassilov 2009 (C); Zavialova 2003 (D); Zavialova et al. 2010b (E); Médus 1977 (F)

assume that early members of Circumpolles were pollen grains of early members of the Cheirolepidiaceae, then the infratectum of one row of large granules can be derived from the infratectum of numerous irregularly distributed small granules (as, for example, in pollen of *Duplicisporites granulatus*, Fig. 4A; Zavialova & Roghi 2005).

This scheme differs from reconstructions, which were earlier proposed for the origin and further morphological evolution of *Classopollis*. De Jersey (1973) hypothesized a transformation from the Early Triassic *Grebespora concentrica* Jansonius 1962, through Late Triassic *Discisporites psilatus* (de Jersey) Reine et al. 1988, to *Classopollis meyerana*. This idea seems unconvincing both considering the conceivability of the structural transformation and keeping in mind the possible botanical affinity. Indeed, it was later abandoned by its author and *Discisporites psilatus* was re-interpreted as a lycopsid spore (Reine et al. 1988). Scheuring (1976) proposed a transition from *Triadispora* to *Classopollis*, based on data of pollen morphology and keeping in mind the idea that the Cheirolepidiaceae could have originated from the Voltziales (the Majoniaceae) as was supposed by Clement-Westerhof & Van Konijnenburg-van Cittert (1991). The family Majoniaceae is characterized by the *Lueckisporites* pollen type, which represents, similarly to *Triadispora*, pollen grains with two protosacci and a small trilete scar. This scheme leaves no place for early Circumpolles. The transition from *Triadispora* or *Lueckisporites* to *Classopollis* also implies a transformation from a protosaccate structure of the ectexine. Such a transformation would lead to an ectexine with, at least occasionally, branching units, which is not the case of Triassic and Jurassic members of *Classopollis*. The scheme proposed by Schering (1976) is beautiful and agrees with the hypothesis made on macrofossil material. However, it belongs to reconstructions that appear the more convincing the less data are available. Indeed, it would appear better substantiated, if we were aware only about the infratectum with branching units (Fig. 4F).

*Classopollis*, with its complicated general morphology, apertures of several types, and columellate-like ultrastructure, is a classical example of angiospermization. Possible parallelisms to angiosperm pollen include the transformations of the ectexine ultrastructure, which are similar to transformations that possibly took place in angiosperms. This concerns the development of columellate exine from granular exine. Moreover, the occurrence of branching columellae is known in such advanced angiosperms as the Compositae, similarly to the latest cheirolepidiacean pollen that also shows branching columellate-like elements of the ectexine. Now it is shown that parallelisms with angiosperms concern not only final stages, but also a chain of similar transformations that could have occurred in these two groups.

In spite of the fact that pollen grains of the Cheirolepidiaceae have been studied with TEM much more often than pollen of most other groups of fossil gymnosperms, further studies are needed. In particular, it would be a long-awaited piece of luck if any pollen grains of early Circumpolles are finally found in pollen organs and become available for a TEM study. Such a study would shed light on the first steps in the evolution of the family. Speaking about late members, the latest material that is so far studied with TEM is dated to the Turonian: branching columellate-like elements were revealed in the ectexine (only one section was published, Médus 1977). Information about post-Turonian members would be very interesting to better substantiate

the occurrence of branching elements. It is also not excluded that additional transformations will be detected. Thus, although pollen grains of *Classopollis* from deposits of different ages show very low variations in dimensions, much larger dispersed pollen grains of *Classopollis* have been recently found in Danian deposits (Barreda et al. 2012). This discrepancy in dimensions is perhaps an indirect index that the latest members of the genus had differences in their exine ultrastructure as well.

## Conclusions

To conclude, TEM studies of pollen or spores of members of monophyletic groups of higher plants can reveal gradual changes in the morphology and ultrastructure of their pollen or spores. Without TEM data, such changes remain hidden from the scientist and it erroneously seems that a given new morphological type originated suddenly and discretely. It is important to apply this approach to monophyletic groups, then no problems will appear with correct homologization of particular sporoderm layers and structures. The approach is very suitable for groups which existed during a long geological time, and, particularly, which managed to survive until nowadays. They are very promising objects, especially because they allow one to extrapolate on fossil members the information about the sporoderm ontogenesis of their modern descendants.

As such a group, the Cycadales seems a very suitable object, which unfortunately so far lacks enough information for the analysis. The group appeared in the Triassic and has survived until nowadays (Taylor et al. 2009). Currently available data show that the elongated perpendicular alveolae of the ectexine, which are so characteristic of pollen of modern cycads, existed at least as early as in the Jurassic (Zavialova & Van Konijnenburg-van Cittert, submitted), or, perhaps, even earlier (Klavins et al. 2003). This seems to be the case of evolutionary stability. However, the problem of the aperture is waited for additional information. There is a possibility that some extinct cycads produced inaperturate pollen grains (Tekleva et al. 2007; Zavialova & Van Konijnenburg-van Cittert 2012), unlike other known fossil cycads and all modern cycads (Audran & Masure 1977, Meyer 1977, Tekleva et al. 2007). In addition, the aperture of fossil cycads differs from that of modern cycads by the ultrastructure (Zavialova & Van Konijnenburg-van Cittert, submitted). At the current state of knowledge, what we can discern is a very peculiar diversity in the apertural condition among cycads, but there are not enough data to order this diversity and to reveal any directed changes during the existence of the group.

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