Evolution and diversity of biomineralized columnar architecture in early Cambrian phosphatic-shelled brachiopods

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Abstract Biologically-controlled mineralization producing organic-inorganic composites (hard skeletons) by metazoan biomineralizers has been an evolutionary innovation since the earliest Cambrian. Among them, linguliform brachiopods are one of the key invertebrates that secrete calcium phosphate minerals to build their skeletons. One of the most distinct shell structures is the organo-phosphatic cylindrical column exclusive to phosphatic-shelled brachiopods, including both crown and stem groups. However, the complexity, diversity and biomineralization processes of these microscopic columns are far from clear in brachiopod ancestors. Here, exquisitely well-preserved columnar shell ultrastructures are reported for the first time in the earliest eoobolids. The hierarchical shell architectures, epithelial cell moulds, and the shape and size of cylindrical columns are scrutinised in *Latusobolus xiaoyangbaensis* gen. et sp. nov. and *Eoobolus acutulus* sp. nov from the Cambrian Series 2

Shuijingtuo Formation of South China. The secretion and construction of the stacked

sandwich model of columnar architecture, which played a significant role in the evolution of

linguliforms, is highly biologically controlled and organic-matrix mediated. Furthermore, a

continuous transformation of anatomic features resulting from the growth of columnar shells

is revealed between Eoobolidae, Lingulellotretidae and Acrotretida, shedding new light on

the evolutionary growth and adaptive innovation of biomineralized columnar architecture

among early phosphatic-shelled brachiopods during the Cambrian explosion.

Introduction

The early Cambrian witnessed a great burst in diversity of animal body plans and

biomineralized shell architectures around half a billion years ago (Briggs, 2015; Erwin, 2015,

2020; Budd and Jackson, 2016; Murdock, 2020; Yun et al., 2021; Zhang and Shu, 2021;

Zhang et al., 2021c). The novel processes of biologically-controlled mineralization producing

organic-inorganic composites (hard skeletons), in complex animals had played a vital role in

the survival and fitness of early clades (Balthasar and Cusack, 2015; Cuif et al., 2010; Li et

al., 2022; Skovsted et al., 2008; Yun et al., 2022), and in turn built the fundamental blocks of

complex marine ecosystems (Bicknell and Paterson, 2018; Buatois et al., 2020; Chen et al.,

2022; Zhang et al., 2010, 2020). Since the early Cambrian to the present day, this adaptive

evolution has been demonstrated and continuously preserved in brachiopods, one of the key

members of the Cambrian Evolutionary Fauna (Carlson, 2016; Harper et al., 2021, 2017;

Sepkoski, 1984). Among them, the phosphatic-shelled brachiopods are some of the most

common animals in early Cambrian faunas (Harper et al., 2017). With a high fidelity of

preservation and high abundance of biomineralized shells in the fossil record, their

morphological disparity, diversity of shell structures and growth patterns, together with

ecological complexity are preserved in great detail (Chen et al., 2021; Claybourn et al., 2020;

Topper et al., 2018; Zhang, 2018; Z. L. Zhang et al., 2020a, 2021b).

Studying the processes by which organisms form biomineral materials has been the

focus at the interface between earth and life sciences (Leadbeater and Riding, 1986;

Simonet Roda, 2021). Brachiopods are unique animals in having the ability to secrete two

different minerals, calcium phosphate and calcium carbonate, making them the ideal group

to further explore the processes of biomineralization (Simonet Roda et al., 2021; Williams,

1977). Hard tissues composed of calcium phosphate with an organic matrix are also largely

present in vertebrates, which have remarkably shaped the evolutionary trajectory of life on

Earth (Ruben and Bennett, 1987; Wood and Zhuravlev, 2012). The origin of phosphate

biomineralization in the evolutionary distant invertebrate brachiopods and vertebrates is still

a big mystery in animal evolution (Luo et al., 2015; Neary et al., 2011; Simonet Roda, 2021).

Thus, more research is needed in order to understand how to relate hierarchical structure to

function in the very early examples of calcium phosphate-based biomineralization processes

(Kallaste et al., 2004; Weiner, 2008). It is noteworthy that South China has been considered

as one of centres for the origination and early dispersal of phosphatic-shelled brachiopods

(Zhang et al., 2021a). It provides a great opportunity to explore the unique biomineralization

process and consequent adaptive evolution of their earliest representatives during the

Cambrian radiation.

The shell-forming process of brachiopods, although critical to understanding their poorly

resolved phylogeny and early evolution, has long been problematic(Carlson, 2016; Cusack

and Williams, 2007; Holmer et al., 2008; Murdock, 2020; Streng et al., 2007; Temereva,

2022; Williams et al., 2004; Williams and Cusack, 1999). The epithelial cells of outer mantle

lobes have been considered to be responsible for the shell ornamentation and fabrics of

brachiopods (Williams and Cusack, 1999). However, the biologically controlled process of

brachiopod shell secretion at the cellular level is still unclear, although organic substrates are

observed to be available for biomineral deposition during mantle activity (Williams, 1977).

Extensive studies have been conducted on living and fossil shells, but most of them are

focused on articulated or carbonate-shelled representatives (Cusack et al., 2010;

Griesshaber et al., 2007; Simonet Roda et al., 2021; Simonet Roda, 2021; Ye et al., 2021).

By contrast, the linguliform brachiopods with shells composed of an organic matrix and

apatite minerals that show extremely intricate architectures and permit exquisite preservation,

are less studied. The shell structural complexity and diversity, especially of their fossil

representatives require further investigations (Butler et al., 2015; Cusack et al., 1999; Streng

et al., 2007; Williams and Holmer, 1992; Zhang et al., 2017). The building of shells by

microscopic cylindrical columns is a unique feature, which is restricted to the phosphatic-

shelled brachiopods and their assumed ancestors (Butler et al., 2015; Holmer et al., 2002,

2008; Holmer, 1989; Skovsted and Holmer, 2003; Williams and Holmer, 2002). This type of

columnar shell was previously believed to be exclusively restricted to micromorphic

acrotretide brachiopods, a group which demonstrate more complex hierarchical architectures

and graded structures compared to simple lamella shell structure in older lingulides (Holmer,

1989; Williams and Holmer, 1992). Further studies however reveal that diverse columnar

shell architectures occur in other brachiopod groups, including stem group taxa, such as

Mickwitzia, *Setatella* and *Micrina* (Butler et al., 2015; Holmer et al., 2008; Skovsted et al.,

2010; Skovsted and Holmer, 2003; Williams and Holmer, 2002), the lingulellotretid

Lingulellotreta (Holmer et al., 2008), the eoobolid *Eoobolus* (Zhang et al., 2021a), and the

enigmatic *Bistramia* (Holmer et al., 2008). However, the columnar architectures among the

oldest linguliforms and their evolutionary variations have not been studied in detail. Different

conditions of fossil preservation with varied taphonomic histories has compounded this issue

as different depositional environments, pore-water geochemistry, and subsequent diagenetic

and tectonic alteration often obscure the finder details of shell structures (Butler et al., 2015;

Holmer et al., 2008; Streng et al., 2007; Ushatinskaya and Korovnikov, 2014; Zhang, 2018).

For the first time, exquisitely well-preserved columnar shell structures are described here

from the oldest known eoobolid brachiopods. *Latusobolus xiaoyangbaensis* gen. et sp. nov.

and *Eoobolus acutulus* sp. nov. are reported, based on new specimens from the Cambrian

Series 2, Shuijingtuo Formation of southern Shaanxi and western Hubei in South China. In

this study, the shell architectures, epithelial cell moulds, and the shape and size of cylindrical

columns are examined, shedding new light on our understanding of the architecture intricacy,

biomineralization process and evolutionary fitness of early phosphatic-shelled brachiopods.

Results

Systematic palaeontology

Brachiopoda Duméril, 1806

Linguliformea Williams, Carlson, Brunton, Holmer and Popov, 1996

Lingulata Gorjansky and Popov, 1985

Lingulida Waagen, 1885

Linguloidea, Menke, 1828

Eoobolidae, Holmer, Popov and Wrona, 1996

Genus **Latusobolus** Zhang, Zhang and Holmer gen. nov.

Type species. *Latusobolus xiaoyangbaensis* sp. nov., here designated.

Etymology. From the Latin '*latus*' (wide) with the ending '*obolus*', to indicate the

transversely oval outline of both ventral and dorsal valves, morphologically similar to *Obolus*.

The gender is masculine.

Diagnosis. For a full description and discussion of *Latusobolus* gen. nov., refer to

Appendix 1.

Latusobolus xiaoyangbaensis Zhang, Zhang and Holmer sp. nov.

Figure 1 and ***Appendix 2—figures 1-4***, ***Appendix 3—table 1***.

Etymology. After the occurrence at the Xiaoyangba section in southern Shaanxi, China.

Type material. Holotype, ELI-XYB S5-1 BR09 (***Appendix 2—figure 1M–P***), ventral valve, and paratype, ELI-XYB S4-2 BO11 (***Appendix 2—figure 2M–P***), dorsal valve, from Cambrian Series 2 Shuijingtuo Formation at the Xiaoyangba section (Zhang et al., 2021a) near Xiaoyang Village in Zhenba County, southern Shaanxi Province, China.

Diagnosis. As for the genus.

Description. For a full description and discussion of *Latusobolus xiaoyangbaensis* gen. et sp. nov., refer to ***Appendix 1***.

Genus ***Eoobolus*** Matthew, 1902

Type species. *Obolus* (*Eoobolus*) *triparilis* Matthew, 1902 (selected by Rowell, 1965)

Diagnosis. See Holmer et al. (p. 41) (Holmer et al., 1996).

Eoobolus acutulus Zhang, Zhang and Holmer sp. nov.

Figure 2 and ***Appendix 2—figures 5–7***, ***Appendix 3—table 2***.

Etymology. From the Latin '*acutulus*' (somewhat pointed), to indicate the slightly acuminate ventral valve with an acute apical angle. The gender is masculine.

Type material. Holotype, ELI-AJH S05 BT11 (***Appendix 2—figure 5E–H***), ventral valve, and paratype, ELI-AJH S05 1-5-07 (***Appendix 2—figure 5M***), dorsal valve, from Cambrian Series 2 Shuijingtuo Formation at the Aijiahe section (Z. L. Zhang et al., 2016) near Aijiahe Village in Zigui County, north-western Hubei Province, China.

152   **Diagnosis**. For a full description and discussion of *Eoobolus acutulus* sp. nov., refer to

153   *Appendix 1*.

Discussion

Diversity of linguliform brachiopod shells

Although the supposed living fossil *Lingula* has long been considered to virtually lack

morphological evolutionary changes (Schopf, 1984), more recent studies have shown that

lingulide brachiopods have experienced dramatic modifications in many aspects (Liang et al.,

2023), including internal organs (Zhang et al., 2008), life mode (Topper et al., 2015), shell

structures (Cusack et al., 1999), and even genome (Goto et al., 2022; Luo et al., 2015). The

complexity and diversity of linguliform shell architecture was increasingly recognised in the

pioneering study of Cusack, Williams and Holmer (Cusack et al., 1999; Holmer, 1989;

Williams and Cusack, 1999; Williams and Holmer, 1992). Moreover, such complex

architectures have a wide distribution in closely related brachiopod groups when they made

their first appearance at the beginning of the Cambrian. In connection with an ongoing

comprehensive scrutiny of well-preserved linguliform shell ultrastructures from the lower

Cambrian limestones of South China, their complexity and diversity hidden in the

conservative oval shape is becoming more and more intriguing. However, compared to their

ancestral representatives, the shell structures in living lingulides are relatively simple,

revealing profound modifications during their long evolutionary history (Cusack et al., 1999;

Holmer et al., 2008; Williams, 1977; Williams and Cusack, 1999).

In general, the shells of organo-phosphatic brachiopods are stratiform, composed of an

outer periostracum and inner rhythmically-disposed succession of biomineralized lamellae or

laminae (Holmer, 1989; Williams, 1977). The organic periostracum, serving as a rheological

coat to the underlying shell, is scarcely fossilized. However, its wrinkling and vesicular

features have largely been preserved as superficial imprints (pits, pustules, fila, grooves,

ridges, rugellae, drapes, reticulate networks and spines) on the surface of the primary layer,

and are important characters in understanding brachiopod phylogeny (Holmer 1989; Holmer

et al. 1996; Cusack *et al.* 1999). By contrast, the structures preserved in the secondary layer

have been characterised in fossil and living groups by three ancient fabrics –columnar,

baculate, laminated – all of which persists in living shells except for the columnar fabric

(Cusack et al., 1999).

The primary layer commonly consists of heavily biomineralized compact laminae

composed of apatite granules, with a thickness from 2 µm to 20 µm (Williams, 1977)

(***Figures 1B, C*** and ***2G; Appendix 2—figure 4J***). Usually, the concentric growth lines are

evenly distributed on the surface of the primary layer of the post-metamorphic shell. By

contrast, the surface ornamentation tends to be unevenly distributed, demonstrating a strong

phylogenetic differentiation. Superficial pustules are one of the most distinct patterns and are

readily recognised in one of the oldest brachiopod groups, the Eoobolidae (Holmer et al.,

1996). The pustules are roughly circular in outline, composed of apatite aggregates, and

range from 2 µm to 20 µm in diameter (Zhang, 2018) (***Figure 1G***). Such pustules are also

found on early obolid, zhanatellid and acrotheloid shells with relatively wide size variations

from 5 µm to 30 µm (Cusack et al., 1999; Zhang, 2018). Although different in size, the similar

pattern may indicate the same secretion regime that originated as vesicles during the very

early stages in periostracum secretion (Cusack et al., 1999). The thickness of the underlying

secondary layer varies greatly in different brachiopod groups, depending on the shell

component and fabric type. Columnar, baculate and laminated fabrics are incorporated into

the basic lamination component to form the diverse stratiform successions of the secondary

shell layer.

The fossil record reveals that the columnar shell structure is generally preserved in most

early linguliforms (Holmer et al., 2008; Streng et al., 2007; Williams, 1977; Zhang, 2018; Z. L.

Zhang et al., 2021a, 2020a). It is a multi-stacked sandwich architecture (multi-columnar units

are stacked in a vertical direction), which is developed in the earliest linguliform *Eoobolus*

with a relatively simple architecture as an early developmental stage (Zhang et al., 2021a).

Each stacked sandwich columnar unit consists of numerous columns disposed orthogonally

between a pair of compact stratiform lamellae (Cusack et al., 1999; Holmer, 1989). One to

three stacked sandwich columnar units with short orthogonal columns can also be found in

Latusobolus xiaoyangbaensis gen. et sp. nov. (***Figure 1***) and *E. incipiens* (***Figure 4A***), while

the number of stacked sandwich units increased in later eoobolids (***Figure 2***), obolids and

lingulellotretids (***Figure 4B–D***). Eventually, the acrotretides developed a more complex

columnar architecture with multiple stacked sandwich units (***Figure 4F–J***). Moreover, the

columnar shell structures are also found in stem group brachiopods, e.g., *Mickwitzia*,

Setatella and *Micrina*, but with different column size and numbers of laminae (Butler et al.,

2015; Holmer et al., 2002, 2008; Skovsted et al., 2010; Williams and Holmer, 2002). It is

assumed here that the columnar architecture may be plesiomorphic character in linguliform

brachiopods, inherited from stem group brachiopods.

219   Biomineralization process of organo-phosphatic columnar architecture

Metazoans are known for secretion of very different types of biominerals through the

process of biological mineralization, This linking of living soft organic tissues with solid earth

minerals is a process that has changed the nature of Earth's fossil archive (Addadi and

Weiner, 2014; Lowenstam and Weiner, 1989; Simonet Roda, 2021; Wood and Zhuravlev,

2012). Because of the fine quality of phosphate biomineralization in linguliforms (Cusack et

al., 1999; Williams and Cusack, 1999), they can have exquisitely finely preserved shell

ultrastructures (***Figures 3*** and ***4***), including epithelium cell moulds (***Figure 3C*** and ***F–I)***. This

permits us to reconstruct the biomineralization process of their apatitic cylindrical columns

and address key questions about how these hierarchical structures relate to mechanical

functions. Although the biomineralization process of living brachiopods at the cellular level is

not well known, biochemical experiments (Cusack et al., 1999, 1992; Lévêque et al., 2004;

Williams and Cusack, 1999) have revealed the possibility that the biologically-controlled,

organic matrix mediated extracellular mineralization during brachiopod shell secretion. This

process can be compared to the hard tissue forming process of mollusc shells and

vertebrate teeth (Golub, 2011; Neary et al., 2011; Simonet Roda, 2021).

Many polygonal structures (*Figure 3C* and *F*), preserved on the internal surface of

successively alternating laminae (*Figure 3G–I*), have generally been considered to

represent the moulds of epithelial cells (McClean, 1988). The average size of epithelial cells

is 20 µm, ranging from 5 to 30 µm. The rheological vesicle environment may cause the

shape variation of epithelia, while different secreting rate could result in different sizes.

Generally, smaller epithelial cells had higher secretion rates (Z. L. Zhang et al., 2016). As

the active secretion activity of outer epithelial cells, they can be easily embedded in the

newly formed inner and outer bounding surfaces of shell laminae (McClean, 1988; Winrow

and Sutton, 2012), which left shallow grooves between epithelial margins as intercellular

boundaries *(Figure 3F*). Epithelial cells are preserved as moulds – presumably by the

phosphatization of smooth organic sheets (*Figure 4G* and *H*) – that had been secreted

relatively slow by the outer plasmalemma of the outer mantle (Cusack et al., 1999).

These active cells are responsible for the secretion of linguliform periostracum with

rheomorphic wrinkling features (Cusack et al., 1999), which left typical microtopography on

the external surface of the underlying primary layer (*Figure 3A* and *B*). On the other hand,

they also secreted linguliform biominerals in a rheological extracellular environment. It is

inferred from living *Lingula* as apatite growing from an amorphous calcium phosphate

precursor, which forms the basic crystals of apatite around 5 nm in diameter (Lévêque et al.,

2004; Williams and Cusack, 1999). These nanoscale crystals were packaged into apatite

granules with an average size of 100 nm (*Figures 1I* and *2H*; *Appendix 2—figure 4K*),

which acted as the fundamental component building up the hierarchical stratiform shells,

including the primary layer and secondary layer (*Figure 5*). The apatite granules were

probably coated or saturated with organic compounds to form granule aggregations or

clusters (spherular mosaics) as irregular spherules or rods of about 500 nm (*Figures 1I* and

2H; Appendix 2—figures 4J, *7J* and *K*), usually leaving gaps between the aggregation

boundaries in fossils after the degradation of organic counterparts (*Figure 2H; Appendix*

2—figure 4K). These apatite spherules are aggregated in the planar orientation as compact

thin lamella less than 4 μm in thickness. Several thin lamellae are closely compacted to form

the primary layer (*Figure 1B* and *C*; *Appendix 2—figure 4I* and *J*). On the other hand,

similar nanometre scale networks of spherules are aggregated and organised as orthogonal

columns perpendicular to a pair of stratiform lamella surfaces, forming one columnar unit.

Multi-columnar units are stacked in a vertical direction from the exterior to interior to form the

secondary layer, applying a stacked sandwich model (*Figure 5*), which differs from the layer

cake model. A very thin gap, commonly less than 1 μm, between each pair of stacked

sandwich columnar units is obvious in well preserved specimens (*Figures 1C*, *I*, *2C*, *F* and

4D; Appendix 2—figures 4I and *7J*). This is likely indicating an organic membrane acting

as an extracellular matrix with functions of a template guiding mineral nucleation (Addadi

and Weiner, 2014; Cusack et al., 1999; Lévêque et al., 2004). This is also supported by the

almost symmetrical nature of the columnar architecture, revealing a homogeneous organic

substrate responsible for the succeeding rhythmic sequence. Although, newly secreted

columnar units may succeed the older one unconformably with overlap (*Figures 2C*, *3I* and

4D) and be involved in lateral changes of column size (*Figures 2K* and *4C*), it is supposed

to reflect intracellular deviation with the same secretory cycle of the outer mantle as a whole

as in living lingulids (Cusack et al., 1999; Williams et al., 1992). The secreting and building

processes of the early Cambrian phosphatic-shelled brachiopod columnar shells, secreted

by the underlying outer epithelium cells of the mantle lobe, are illustrated in *Figure 5*.

It is worth noting that, on well preserved specimens nanoscale openings are permeated

on the terminal ends of the orthogonal columns (*Figures 1C, 4E* and *H*) and surfaces of

stratiform lamellae (*Figures 1H, 2G*, *3D*, *E*, *4G* and **I**). The openings are rounded with a

mean diameter of 600 nm. Observation through natural fractures of shells, show that some

canals can be traced continuously through several columnar sequences (*Figures 1D*, *2F*

and *4E*), while in poorly preserved fossils, they are filled with secondarily phosphatised

spherules or mosaics (*Figures 1I* and *2G*), occasionally leaving random gaps (*Figures 1D*,

2F and *4I*). These canals are very likely the empty space left by degraded organic material,

which is confirmed by different taphonomic processed that preserve canals in

contemporaneous early Cambrian Burgess Shale–type fossil Lagerstätte in South China

(Duan et al., 2021). The regular arrangement of the canal systems and closely related

columns reveal that the process is organic matrix-mediated. Furthermore, the even

disposition of the organic matrix in columns indicate the existence of rheological and central

areas, on which biologically controlled biomineralization took place. It reveals that the

nucleation, growth and aggregation of the deposited amorphous calcium phosphate are

directed by the same group of epithelial cells (Cusack et al., 1999; Pérez-Huerta et al., 2018;

Simonet Roda, 2021; Weiner and Dove, 2003). In the stacked sandwich columnar units, the

empty chambers between each columns(*Figure 4F* and *J*) would be originally filled with

glycosaminoglycans (GAGs) as in living *Lingula* (Cusack et al., 1999; Williams et al., 1994).

These chambers are often filled with coarse spherular mosaics when being secondarily

phosphatised and consequently they become indistinguishable from the columns, paired

stratiform lamellae and organic membrane (*Figures 1D*, *2E*, *F* and *4I*). At the posterior

margin of mature shells, especially the ventral pseudointerarea where the vertical

component of the growth vector becomes increasingly important in ventral valves, the short

columns are succeeded by relatively taller columns, resulting in the change from two stacked

sandwich columnar units into one unit during growth anteriorly (*Figures 2C*, *K*, *3I* and *4D*).

This may demonstrate allometric growth of shell (Cusack et al., 1999).

The most intriguing and enigmatic phenomenon of skeletal biomineralization is the

evolutionary selection of calcium carbonate and calcium phosphate in invertebrates and

vertebrates, respectively (Lévêque et al., 2004; Luo et al., 2015). However, the Brachiopoda

is a unique phylum that utilises both minerals. The appearance of apatite as a shell

biomineral dates back as far as Cambrian Age 2 for stem group brachiopods (Skovsted et al.,

2015; Topper et al., 2013; Ushatinskaya, 2002) and persists to the present in living

linguliforms (Carlson, 2016). Calcium phosphate can build a relatively less soluble skeletal

component as compared with calcium carbonate shells, but with the disadvantage of a

greater energetic and physiologic cost (Wood and Zhuravlev, 2012). The acquisition of this

specific biomineral in phosphatic-shelled brachiopods has been considered an ecological

consequence of the globally elevated phosphorous levels during phosphogenic event in the

calcite seas with a low Mg:Ca ratio and/or high CO_2 pressure (Balthasar and Cusack, 2015;

Brasier, 1990; Cook and Shergold, 1984; Wood and Zhuravlev, 2012). In such situations,

linguliforms were able to utilise the sufficient phosphorous in ambient waters, unlike

brachiopod ancestors that possessed an unmineralized shell coated with detrital grains, like

Yuganotheca found in the Chengjiang Lagerstätte (Zhang et al., 2014). The acquisition of a

calcium phosphatic shell may have been an evolutionary response of prey to an escalation

of predation pressure during the Cambrian explosion of metazoans (Cook and Shergold,

1984; Wood and Zhuravlev, 2012). Consequently, organic-rich biomineral composites of

linguliform brachiopod shells possessed innovative mechanical functions, providing

competitive superiority and adaptation on Cambrian soft substrates as well as reducing

susceptibility to predation (Wood and Zhuravlev, 2012). Despite the physiological cost of

calcium phosphate biomineralization and subsequent reduction in phosphorus levels post

Cambrian period, linguliforms have retained their phosphate shells during dramatical

oscillations of seawater chemistry and temperature over 520 million years.

Given the long history of this subphylum, the possession of a phosphatic shell likely has

numerous advantages. The innovative columnar architecture can mechanically increase the

thickness and strength of the shell by the presence of numerous, stacked thinner laminae,

comparable with the laminated fabric seen in obolids (Cusack et al., 1999; Z. F. Zhang et al.,

2016). Furthermore, the stacked sandwich columns also increases the strength, flexibility

and the ability to resist crack propagation by filling the space between the stratiform lamellae

with organic material, comparable with the baculate fabric (Lévêque et al., 2004; Merkel et

al., 2009). Thus, the stacked sandwich model of the columnar architecture possesses a

greater advantage of mechanical functions and adaptation with a superior combination of

strength, durability and flexibility in laminated and baculate fabrics, resembling the

colonnaded and reinforced concrete often used in urban construction. New data from

nuclear magnetic resonance spectroscopy and X-ray diffraction reveals that apatite in

brachiopod shells is highly ordered and thermodynamically stable crystalline and it is more

robust in the extremes of moisture, ambient osmotic potential and temperature, unlike the

poorly ordered crystal of vertebrate bone (Neary et al., 2011). This type of more efficient and

economical shell may also have been responsible for the early diversity of major linguiform

brachiopods during the Cambrian explosion, resulting in this group becoming a significant

component of the Cambrian Evolutionary Fauna (Bassett et al., 1999; Sepkoski, 1984;

Zhang et al., 2008, 2020; Z. L. Zhang et al., 2021b).

Evolution of stacked sandwich columnar architecture in early brachiopod

clades

Evolutionary transformations have repeatedly modified the organo-phosphatic

architecture consisting of various aggregates of spherular apatite, held together by a

scaffolding of glycosaminoglycan complexes, fibrous proteinaceous struts and chitinous

platforms, in linguiform brachiopod shells since the early Cambrian (Cusack et al., 1999). As

one of the oldest forms of brachiopod shell architectures, the columnar shell has previously

been regarded as an unique character of acrotretide brachiopods (Cusack et al., 1999;

Holmer, 1989). However, recent discoveries of columnar shell structures in a diversity of

early Cambrian stem group brachiopods have revealed that the same biomineralization

strategy is utilised much more widely than previously thought (Butler et al., 2015; Holmer et

al., 1996, 2008; Skovsted et al., 2010; Streng et al., 2007; Ushatinskaya and Korovnikov,

2014; Zhang, 2018; Zhang et al., 2021a). This highlights the need for a better understanding

of the origin and adaptive modification of stacked sandwich columnar architectures in early

lophophorate evolution.

The Eoobolidae is presently considered to be the oldest known linguliform brachiopods

(Zhang et al., 2021a). The biomineralized orthogonal columns in *Eoobolus incipiens* from

lower Cambrian Stage 3 probably represents an early and simple shell structure type with a

poorly developed secondary columnar layer (***Figure 4A***). The columns are relatively small

with a mean diameter of 1.8 μm ranging from 0.6 to 3.0 μm, and a mean height of 4.1 μm

(***Figure 6***; ***Appendix 3—table 3***). The complete secondary layer is only composed of two to

three stacked sandwich columnar units, resulting in a shell thickness of about 30 μm. Such a

simple shell structure is also developed in the slightly younger *Latusobolus xiaoyangbaensis*

gen. et sp. nov. (***Figure 1***), but with a slightly taller column of about 6.2 μm. From Cambrian

Age 4, the number of stacked sandwich columnar unit increases rapidly in the Eoobolidae,

growing to as many as 10 stacked sandwich columnar units in *Eoobolus acutulus* sp. nov.

(***Figure 2***), and as many as 10 in *Eoobolus*? aff. *priscus*, and the shell thickness also

increases to 50 μm (Streng et al., 2007). However, the size of individual columns keeps

within a stable range, around 4 μm in height and 2 μm in diameter. Compared with

Eoobolidae, the Lingulellotretidae demonstrates a more developed columnar shell, which

has a relatively larger number of stacked sandwich columnar units (up to 20), effectively

increasing the shell thickness to 70 μm. The column size is very similar in both

Lingulellotreta malongensis and *L. ergalievi* (***Figure 4B*** and ***D***), and matches that of

contemporaneous *Eoobolus*. But the slightly younger *L. ergalievi* has more columnar units,

resulting in a thicker shell than *L. malongensis*.

Among all early Cambrian linguliforms with columnar architectures, the acrotretides have

developed the most complex shell structure (***Figure 4F–J***). The fossil record in the

Cambrian unveiled a clear pattern of the increasing growth (regarding both the diameter and

height of columns and the number of stacked sandwich columnar units) of columnar

architecture in acrotretides: from a very simple type in *Palaeotreta shannanensis* like that of

E. incipiens to slightly more developed in *Palaeotreta zhujiahensis* like that of *L.*

malongensis (Z. L. Zhang et al., 2020b), and to the most developed *Eohadrotreta*

zhenbaensis onward (**Figure 6**). The diameter of single orthogonal columns increase about

two times in acrotretides compared to eoobolids, whereas the general height of column

increases to 10 µm in *Eohadrotreta zhenbaensis* and to 29 µm in *Hadrotreta primaeva*,

which is about 10 times as high as seen in *Eoobolus variabilis*. Furthermore, the number of

columnar units has also increased to about 30, collectively increasing the shell thickness to a

maximum value of more than 300 µm in *Eohadrotreta*.

The homology of the columnar architecture in early linguliforms outlines a clear picture,

most likely representing a continuous transformation between the Lingulida (Eoobolidae and

Lingulellotretidae) and Acrotretida. Two groups that represent a major component of early

Cambrian benthic communities (Chen et al., 2021; Claybourn et al., 2020; Topper et al.,

2015; Zhang et al., 2008; Zhang, 2018; Z. L. Zhang et al., 2020a). The possible occurrence

of this shell architecture within the family Obolidae cannot be discounted, as detailed

information on the possible columnar shell structures in early Cambrian representatives such

as, *Kyrshabaktella* and *Experilingula* are poorly known (Cusack et al., 1999; Streng et al.,

2007; Williams, 1977). The small size and simple pattern of stacked sandwich columnar

architectures remain stable in the Eoobolidae, and this stability likely limits both the shell

thickness and overall body size, with ventral and dorsal valves of the family remaining below

6 mm until Miaolingian (middle Cambrian). Stacked sandwich columnar architectures are a

character state of the *Lingulellotreta* shell structure as well; however more columnar units

are developed that slightly increases the shell thickness and subsequently species of

Lingulellotreta reach body sizes twice that of Eoobolidae (Zhang et al., 2007). A continuous

transformation of anatomic features can be deduced from the evolutionary growth of

columnar shells between the two clades. Firstly, the orthogonal columns are markedly

developed at the pseudointerarea area of ventral valve of *Lingulellotreta* (**Figure 4B**),

resulting in a greater elevation of the pseudointerarea above the shell floor. It leaves a large

amount of space for the posterior extension of the digestive system, which is well protected

by the covering mineralized ventral pseudointerarea. This is supported by the discovery of a

curved gut under the pseudointerarea of *Lingulellotreta malongensis* in the Chengjiang

Lagerstätte (Zhang et al., 2007). Secondly, with the continuous growth of the ventral

pseudointerarea, the opening obolide-like pedicle groove is sealed, resulting in a unique

pedicle opening exclusively observed in the lingulellotretid brachiopods. Thus, the early

pedicle-protruding opening between the ventral and dorsal valves of the Linguloidea is

transformed into a new body plan where the pedicle opening is restricted in the ventral valve

of lingulellotretids. It supports the scenario that the columnar architecture is monophyletic in

at least Linguloidea, and that the slightly younger lingulellotretid columnar architecture was

derived from an *Eoobolus*-like ancestor during late Cambrian Age 3.

In another evolutionary direction, acrotretide brachiopods fully utilise the columnar

architecture including the derived camerate fabric as shell structure in the whole clade

(Streng and Holmer, 2006). The similarity and gradually evolutionary transformations of shell

structure from simple forms in lingulides to complex forms in acrotretides suggests the

stacked sandwich columnar architecture did not evolve independently in actrotretides. In

terms of the derived camerate fabric, more mineralized material is utilised compared to its

precursor, the columnar shell structure (Streng et al., 2007). The column size, including

height and the number of stacked sandwich columnar units uniformly increase to about 10

times greater in acrotretides than in *E. incipiens and L. xiaoyangbaensis*, since late

Cambrian Age 3 (**Figure 6**). Despite the increase in size of the columns and the number of

stacked sandwich columnar units, the whole body of acrotretides is restricted to only

millimetre size (Holmer, 1989; Williams et al., 2000). A continuous transformation of

anatomic features and shell structure functions can be deduced from the evolutionary growth

of columnar shells in early acrotretides. Firstly, the stacked sandwich columns were

markedly developed at the pseudointerarea area of the ventral valve, resulting in a greater

elevation of the pseudointerarea above the valve floor, compared to that of *Lingulellotreta*.

Secondly, two transformations subsequently changed an obolid-like flat ventral valve

(*Palaeotreta shannanensis*) to a cap shape valve (*P. zhujiahensis*) (Z. L. Zhang et al.,

2020b), and a conical shape (*Eohadrotreta zhenbaensis*) (Zhang et al., 2018), and

eventually to a tubular shape (*Acrotreta*) (Holmer and Popov, 1994). During this evolution,

dorsal valves constantly kept a flat shape. The obolid-like ventral pseudointerarea changed

from orthocline to catacline and eventually to procline with strongly reduced propareas, while

the ventral muscular system moved towards the elevated posterior floor, resulting in the

formation of a new apical process (Popov, 1992; Zhang et al., 2018). With the increasing

growth of the stacked sandwich columnar shells, the thick organo-phosphatic shell with

increased strength might have provided more mechanical support to the conical or tubular

valve in a turbulent environment.

Based on the similar evolutionary trajectory, regarding the increasing growth of the shell

in Lingulellotretidae and Acrotretida, the formation of their ventral pedicle foramens is very

likely homologous, modified from an obolid-like pedicle groove between the two valves.

Furthermore, the similarity and continuity in the increasing number and size of the

orthogonal columns suggest that columnar architecture is a plesiomorphic character in

Linguloidea and Acrotretida. However, the phylogenetic puzzle of whether the columnar

architecture is paraphyletic with the baculate fabric in Linguliformea, or even in Lingulata

hangs on two pieces of important fossil evidence. The shell structure of the earliest

linguliform brachiopods on a global scale needs to be comprehensively investigated based

on more well-preserved fossils. Moreover, more extensive scrutinization of the shell

architecture and composition in widely distributed early Cambrian stem group brachiopods is

required to conclusively resolve their phylogenetic relationship with linguliforms. The

complex shell in stem group taxa *Setatella* and *Mickwitzia*, that are younger and have more

advanced columnar shell features than *Eoobolus incipiens*, might reveal the plesiomorphic

state of the columnar architecture in Linguliformea (Butler et al., 2015; Holmer et al., 2008;

Skovsted and Holmer, 2003; Williams and Holmer, 2002). But this assumption depends on

future work and whether the columnar shell structure is preserved in older ancestors and

other stem group taxa. In another scenario, baculate, laminated and columnar architectures

might have originated independently from an unmineralized ancestor like the agglutinated

Yuganotheca during the early Cambrian (Cusack et al., 1999; Zhang et al., 2014).

Regardless of what scenario is true, the origin of the innovative columnar architecture with a

stacked sandwich model has played a significant role in the evolution of linguliform

brachiopods. The evolutionary diversity of shell architectures would match the general

increase in diversity of phosphatic-shelled brachiopods during the Cambrian explosion.

Among them, the micromorphic acrotretides demonstrate the superb application of the

columnar architecture combined with its innovative conical shape and possible exploitation

of secondary tiering niches (Topper et al., 2015; Wang et al., 2012; Zhang et al., 2008; Z. L.

Zhang et al., 2018, 2021b). The fitness of the diminutive body size of acrotretides is likely a

trade-off between the increasing metabolic demand of phosphate biomineralization after the

Cambrian phosphogenic event and the increased chance of evolutionary survival and

adaptation by producing a high mechanical skeleton for protection in the shallow water

environment (Cook and Shergold, 1984; Garbelli et al., 2017; Lévêque et al., 2004; Neary et

al., 2011; Simonet Roda, 2021; Wood and Zhuravlev, 2012).Their relatively large

surface/volume ratio mechanically requires strong support from the composition of stacked

sandwich columnar architecture and possibly a relatively lower density of the shell by

organic biomineralized material for the secondary tiering life. Such adaptive innovations may

account for the flourish of phosphatic-shelled acrotretides in the latter half of the Cambrian,

continuing to the Great Ordovician Biodiversification Event, thriving and playing an important

role in marine benthic communities for more than 100 million years.

Material and Methods

The brachiopod material studied here was collected from the Cambrian Series 2 Xihaoping

Member of the Dengying Formation and the Shuijingtuo Formation at the Xiaoyangba

section of southern Shaanxi (Zhang et al., 2021a), and the Shuijingtuo Formation at the

Aijiahe section and Wangjiaping section of western Hubei (Z. L. Zhang et al., 2016). All

specimens are recovered through maceration of limestones by acetic acid (~10%) in lab,

and deposited in the Early Life Institute (ELI), Northwest University, China. Selected

specimens were coated and studied further using Fei Quanta 400-FEG SEM at Northwest

University, Zeiss Supra 35 VP field emission at Uppsala University and JEOL JSM 7100F-

FESEM at Macquarie University. Measurements of length, width and angle of different parts

of *Latusobolus xiaoyangbaensis* gen. et sp. nov. and *Eoobolus acutulus* sp. nov. are

performed on SEM images of well-preserved specimens by TpsDig2 v. 2.16. Measurements

of diameter and height of orthogonal columns and thickness of different shell layers are

performed on SEM images of available adult specimens from this study and previously

published literatures by TpsDig2 v. 2.16. Raw data is provided in ***Appendix 3-tables 1-3***.

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Competing interests. The authors declare no competing interests.

Additional files

Supplementary files

Supplementary file 1. Appendix 1-Systematic Palaeontology.

Supplementary file 2. Appendix 2-figures 1-7.

Supplementary file 3. Appendix 3-table 1.

Supplementary file 4. Appendix 3-table 2.

Supplementary file 5. Appendix 3-table 3.

Data availability

All data are available in the main text and the supplementary materials.

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【Figure captions】

Figure 1. Shell architecture of *Latusobolus xiaoyangbaensis* gen. et sp. nov. from

the Cambrian Series 2 Shuijingtuo Formation in southern Shaanxi, South China. **A-C**,

ELI-XYB S5-1 BS01. **A**, cross section of a ventral lateral margin, note post-

metamorphosis pustules by arrow, primary layer 1 and stacked sandwich columnar

units 2-7, box indicates area in **B**. **B**, enlarged view of **A**, showing top primary-

underlying secondary layer boundary by dashed line. **C,** enlargement of shell layers

1-4 of **A**, note organic canals of columns (arrows), organic layer (tailed arrows)

between two stratiform lamellae by dashed lines. **D**, poorly phosphatised columns of

ventral valve, note organic canals by tailed arrows, ELI-XYB S5-1 BR06. **E**, columns

of dorsal valve, ELI-XYB S5-1 BS17. **F**, cross section of a ventral lateral margin,

show developed post-metamorphosis pustules, box indicates the area in **G**, ELI-XYB

S4-2 BO06. **G**, enlarged primary layer pustules and underlying secondary layer

columns. **H**, one unit of dorsal stacked columnar architecture with the exfoliation of

top primary layer, noting column canals on the stratiform lamella surface by arrows,

ELI-XYB S4-2 BO08. **I**, apatite spherules of granule aggregations of ventral

columnar shell structure, note granule rods by arrow and thin gap left by the

degradation of organic counterparts by tailed arrow, ELI-XYB S4-2 BO06. Scale bars:

A, 50 μm; **B**, **E**, **G**, 20 μm; **C**, **H**, 10 μm; **D**, 5 μm; **F**, 100 μm; **I**, 1 μm.

Figure 2. Shell architecture of ventral *Eoobolus acutulus* sp. nov. from the Cambrian

Series 2 Shuijingtuo Formation in Three Gorges areas, South China. **A-C**, ELI-AJH

S05 BT12. **A**, cross section of a ventral lateral margin, note primary layer 1 and

stacked sandwich columnar units 2-11, box indicates area in **B**. **B**, enlarged view of

A. **C**, enlarged view of **B**, show thin gap left by the degradation of organic

counterparts (tailed arrow) between two stratiform lamellae by dashed lines, the

fusion point of two columnar units by arrow. **D-F**, ELI-AJH 8-2-3 BT02. **D**, cross

section of shell margin, box indicates area in **E**. **E**, different preservation condition of

columnar architecture. **F**, poorly phosphatised columns, note the opening of organic

canals along the organic membrane by arrows, and space between two stratiform

lamellae by tailed arrows. **G-H**, ELI-AJH 8-2-3 BT03. **G**, note organic canals on the

cross section and surface of stratiform lamella by arrows, and partly exfoliated

primary layer by tailed arrow. **H**, magnified columns composed of granule spherules

with canal in **G** by arrow. **I**, cross section of shell margin, box indicates area in (**J**),

ELI-AJH 8-2-3 BT04. **J**, enlarged short columns. **K**, imbricated columnar architecture

(arrows), ELI-AJH S05 BT12. Scale bars: **A**, **E**, 50 μm; **B**, **I**, 20 μm; **C**, 5 μm; **D**, 200

μm; **F**, **G**, **J**, **K**, 10 μm; **H**, 1 μm.

Figure 3. Shell ornamentation, ultrastructure and epithelial cell moulds of Cambrian

Series 2 brachiopods. **A**, post-metamorphic pustules of *Latusobolus*

xiaoyangbaensis gen. et sp. nov., ELI-XYB S4-3 AU11. **B-D**, *Eoobolus acutulus* sp.

nov. **B**, metamorphic hemispherical pits, ELI-AJH 8-2-2 Lin01. **C**, epithelial cell

moulds, note column openings on layer surfaces beneath by arrows, ELI-AJH S05

N31. **D**, enlarged column openings on layer surface, ELI-WJP 7 CE05. **E-I**,

Eohadrotreta zhenbaensis. **E**, partly broken columns, note organic canals by arrows,

ELI-AJH F36. **F**, polygonal epithelial cell moulds on valve floor, dash lines indicate

margin of one epithelial cell, ELI-WJP 6 R79. **G-I**, ELI-AJH Acro 053. **G**, epithelial

cell moulds on dorsal median septum with columns between. **H**, enlarged view of **G**,

note rudiment of columns by tailed arrows and one column on epithelial cell margin

by arrow. **I,** epithelial cell moulds on stratiform lamella surfaces of successive three

stacked sandwich columnar units developed on cardinal muscle areas with columns

between (arrow). Scale bars: **A**, **D**, 50 µm; **B**, **E**, **F**, **H**, 10 µm; **C**, **G**, **I**, 20 µm.

Figure 4. Biomineralized columnar architecture of Cambrian Series 2 brachiopods. **A**,

Eoobolus incipiens, AJXM-267.5 DT12. **B-D**, *Lingulellotreta ergalievi*, ELI-AJH 8-2-3

CI11. **B**, cross section of shell margin, box indicates area in **C**, note primary layer 1

and stacked sandwich columnar units 2-22, and raised pseudointerarea (tailed

arrow). **C**, enlarged view of thin organic layer (tailed arrow) between two stratiform

lamellae by dashed lines, the fusion of two stacked columnar units into one by arrow.

D, imbricated growth pattern of stacked columnar units. **E**, *Palaeotreta zhujiahensis*,

note column openings (arrow) on eight successive columnar units by tailed arrows,

ELI-AJH 8-2-1 AE09. **F-J**, *Eohadrotreta zhenbaensis*. **F**, relatively taller columns (ca.

20 µm), ELI-AJH 8-2-1 acro16. **G**, apatite spherules of granule aggregations in one

columnar unit, note column openings (arrows) on both stratiform lamella surfaces,

ELI-AJH S05 E18. **H**, cross section show column openings on 4 successive units by

tailed arrows, ELI-WJP 7 AB98. **I**, poorly phosphatised columns (arrows), note

openings of organic canals on surface of stratiform lamella by tailed arrows, ELI-AJH

S05 I76. **J**, stacked columnar units in imbricated pattern, ELI-WJP 6 R47. Scale bars:

A, **E**, **I**, 10 µm; **B**, 100 µm; **C**, **F**, 20 µm; **G**, 2 µm; **H**, 5 µm; **D**, **J**, 50 µm.

Figure 5. Biomineralization process of typical columnar architecture by the stacked

sandwich model of phosphatic-shelled brachiopods. Abbreviation: D=Diameter;

H=height; T=Thickness. (modified from (Williams and Holmer, 1992; Z. L. Zhang et

al., 2016))

Figure 6. The evolution of stacked sandwich columnar architecture in early

Eoobolidae *Eoobolus incipiens*, *Latusobolus xiaoyangbaensis* gen. et sp. nov.,

Eoobolus acutulus sp. nov., *Eoobolus variabilis*, *Eoobolus*? aff. *priscus*,

Lingulellotretidae *Lingulellotreta malongensis*, *Lingulellotreta ergelievi*, Acrotretida

Palaeotreta shannanensis, *Palaeotreta zhujiahensis*, *Eoohadrotreta zhenbaensis*,

Hadrotreta primaeva, *Canthylotreta crista* and stem group *Setatella significans*. The

height and diameter data of column are from the literatures (Skovsted and Holmer,

2003; Streng et al., 2007; Streng and Holmer, 2006; Ushatinskaya and Korovnikov,

2014; Z. L. Zhang et al., 2016, 2020a, 2020b).

periostracum
primary layer
secondary columnar layer
mantle

metamorphosis pits
post-metamorphosis pustules
periostracum
primary layer
secondary columnar layer
fold or drape system
organic membranes
mantle
mantle epithelial cell
connective tissue

primary layer lamellae / laminae
T=2-20 µm

thin lamella
T<1 µm

apatite spherule of granule aggregations
(~500 nm)

T=~1 µm
D=~2 µm
H=20 µm
D=~500 nm
T=~1 µm
orthogonal column

stacked sandwich columnar unit
(columnar shell structure)