

1 **Commentary**

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4 **Silicon drives the evolution of complex crystal morphology in calcifying algae**

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6 Quote:

7 *“The formation of intricately shaped coccoliths from rudimentary calcite crystals requires*

8 *the presence of silicon. This novel insight brings us closer to understand the evolution of*

9 *morphological diversity in algae that have shaped planet Earth”*

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26 **Main text**

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28 Coccolithophores are oceanic microalgae that have influenced the global climate for millions
29 of years because of their ability to calcify (e.g. Monteiro *et al.* 2016). Their life cycle is
30 haplo-diplontic with significant differences in the structure and morphology of the calcium
31 carbonate plates (coccoliths) between haploid and diploid life-cycle stages (e.g. De Vries *et*
32 *al.* 2021, Frada *et al.* 2019, de Vargas *et al.* 2007). Whereas coccoliths of haploid life-cycle
33 stages (Holococcoliths – HOLs) are uniform in shape and size, diploid stages are
34 characterized by intricately-shaped coccoliths (Heterococcoliths – HETs) of almost infinite
35 morphology. As HOLs seem to be formed differently and only appear in the fossil record
36 about 30 Mya after the first HETs, it has been suggested that holococcolith-formation
37 represents an independent process of calcification, evolving after the emergence of HETs
38 (e.g. Bown *et al.* 2004, De Vargas *et al.* 2007). Yet, in this issue of New Phytologist, Langer
39 *et al.* (2021; pp. ABC-XYZ) have challenged this view by carefully analysing the process of
40 holococcolith-formation. Combining state-of-the-art microscopy tailored to preserve all
41 subcellular structures, and experiments to reveal the role of silicon in the process of
42 calcification, they show that HOLs are formed in intracellular compartments similar to HETs
43 and that silicon is only required for the formation of intricately shaped coccoliths. These
44 results suggest that HOLs might represent an ancestral form of calcification and that the
45 ability to use silicon in the process of calcification evolved later and is responsible for the
46 synthesis of the elaborately shaped HETs.

47

48 Calcification is the most characteristic feature of coccolithophores, which belong to the group
49 of prymnesiophytes and diverged from their non-calcifying ancestors approximately 310 Mya
50 (e.g. Liu *et al.* 2010). There are over 250 known species of coccolithophores in sunlit oceans,
51 contributing up to 10% of annual marine primary production (e.g. Poulton *et al.* 2007). Some
52 species including *Emiliana huxleyi* are so productive that their blooms can be seen from
53 space (Fig. 1). Despite their significance for the global carbon cycle, most studies so far have
54 only focussed on a limited number of diploid coccolithophores with the best studied likely to
55 be *E. huxleyi* (e.g. Gal *et al.* 2018, Read *et al.* 2013). Intricately shaped coccoliths allow
56 easier identification of diploid species, which is possibly why haploid life-cycle stages, many
57 of which do not calcify (e.g. von Dassow *et al.* 2012), have largely been neglected, biasing
58 our current knowledge on coccolithophore biology and evolution.

59

60 How and when the life phase transitions occur is not well known for most coccolithophore
61 species although several drivers have been identified for *E. huxleyi* (e.g. Frada *et al.* 2019).
62 For instance, viral infections can cause a switch from the diploid to the haploid life-cycle
63 phase to increase survival rates in response to a virus infection (Frada *et al.* 2008). Typically,
64 though, ploidy and the proliferation of life cycle-stages are decoupled, similar to macroalgae
65 and some plants where gametes develop independent life-cycle stages that reproduce
66 asexually, i.e. gametophytes (e.g. Coelho & Cock 2020, Taylor *et al.* 2005). Thus, both life-
67 cycle stages are exposed to evolutionary forces and therefore might even speciate
68 independently. Generally, it can be assumed that haplo-diplontic life cycles are better at
69 exploring the adaptive landscape of a species because of the larger allelic diversity. Indeed,
70 there is some evidence that different oceanic environments appear to select for different life-
71 cycle stages of coccolithophores (e.g. De Vries *et al.* 2021), however, our knowledge of the
72 adaptive benefits is still very limited. Nevertheless, it is likely that calcification, which
73 underpins the formation of distinct phenotypes, is under selection and therefore the molecular
74 machinery driving it. Depending on the species, basic calcium carbonate crystals (calcite) can
75 transform into nanopatterned and elaborate coccoliths of seemingly infinite shape and form.
76 HETs are formed inside a specialized Golgi-derived vesicle (e.g. Brownlee *et al.* 2015).
77 Before they are extruded, they are formed by an unknown mechanism that controls crystal
78 morphology and the overall shape and form of HETs. Together they form the coccosphere,
79 which can include various appendages and in which the cell resides. In contrast to well-
80 studied HETs, HOLs have received little attention, but their crystals resemble the typical
81 rhombohedral geometry of inorganic calcite (e.g. Young *et al.* 1999). Furthermore, the
82 morphological diversity of HOLs is much more constrained.

83

84 In 2016, the same laboratory at the Marine Biological Association (MBA) in Plymouth, UK,
85 discovered that calcifying coccolithophores have something in common with their silicifying
86 cousins: diatoms (Durak *et al.* 2016). However, diatom shells are made of silica and therefore
87 thought to represent a distinct mechanism of biomineralization. This concept was challenged
88 by the discovery of silicon transporters (SITs) in calcifying diploid coccolithophores (Durak
89 *et al.* 2016). Some of these species even appear to have an obligatory requirement for Si,
90 similar to diatoms. Although the cellular mechanism by which Si contributes to the process of
91 calcification is still unknown, studies in other organisms have suggested that silica might be
92 essential for the formation of ordered calcite crystals as seen in HETs but not in HOLs (e.g.
93 Gal *et al.* 2012).

94

95 The study by Langer *et al.* (2021) has tested the hypothesis that HOLs represent an ancestral
96 state of calcification, which is contradictory to the fossil record. As support for their
97 hypothesis, they combined knowledge on the role of silicon for the formation of HETs and
98 applied advanced microscopy to re-assess the calcification processes in HOLs. By using
99 scanning electron microscopy in combination with high pressure freezing and freeze
100 substitutions to preserve both inorganic and organic structures, it was possible for the first
101 time to reveal that HOLs are formed inside the cells in vesicles similar to the synthesis of
102 HETs. Langer *et al.* (2021) argue that this result provides first evidence for the presence of a
103 last common ancestor that was capable of producing both HOLs and HETs as they share not
104 only the same chemical process of calcification but also the same cell biology required to
105 produce calcite crystals. Thus, Langer *et al.* (2021) have provided an evolutionary link
106 between both modes of calcification (Fig. 2). As HOLs are structurally more simplistic, it
107 suggests that they have evolved first, which was already postulated a few years ago by Frada
108 *et al.* (2019). To identify why the additional complexity observed in HETs evolved later,
109 Langer *et al.* (2021) drew on their insights into the role of silicon for the formation of
110 complex calcite crystal morphology. Remarkably, they found that HOLs do not require Si for
111 crystal formation. They also discovered the presence of rhombohedral HOL crystals in
112 diploid coccolithophores after replacing Si in the growth medium by germanium. These
113 results suggest that Si is required for the synthesis of different crystal shapes as both life-
114 cycle stages develop rudimentary rhombohedral crystals but intricately shaped coccoliths are
115 only formed with the help of Si (Fig. 2).

116

117 Although these results seem to have resolved a long-standing paradigm in the evolution of
118 calcification in microalgae (e.g. Bown *et al.* 2004, De Vargas *et al.* 2007), they raise
119 interesting questions. For instance, not all diploid coccolithophores with HETs require Si
120 during formation including the model species *E. huxleyi* (Durak *et al.* 2016). Furthermore,
121 although the requirement for Si explains why there is complex calcite crystal morphology, it
122 does not explain the almost infinite morphological diversity of coccospheres. I argue that
123 answers to these questions can be found by applying evolutionary theory to silicon and
124 calcium carbonate metabolism. Although phylogenetics has been applied to reveal
125 relationships between individual genes involved in biomineralization, the field will benefit
126 from revealing how the evolutionary forces of mutation, selection, genetic drift and gene flow
127 shaped the genetic and morphological diversity of biomineralizing microalgae. Langer *et al.*

128 (2021), speculate that high concentrations of Si in the surface oceans about 250 Mya were
129 driving the evolution of HETs. A subsequent decline of Si due to the rise of diatoms might
130 have caused the loss of an obligate Si requirement at least in some species such as *E. huxlei*
131 and therefore provided a fitness advantage under lower Si concentrations. Thus, they argue
132 that changes in the environment selected for the evolution of complex crystal morphology in
133 calcifying algae.

134

135 Combining molecular markers and fossils from the geological record of coccolithophores
136 with demographic inference such as coalescence theory (e.g. Rosenberg & Nordborg 2002),
137 which provides a view backwards in time, will provide evidence as to whether environmental
138 change (e.g. Si concentrations) coincides with the point where gene genealogies (e.g. silicon
139 transporters) come together ('coalesce'). Furthermore, identifying signals of selection will
140 inform biochemical studies because they reveal which genes and functional domains likely
141 contribute to the evolution of morphological diversity, which potentially is the outcome of
142 diversifying selection. As mutational and demographic models are available for
143 coccolithophores (Krasovec *et al.* 2020, Bendif *et al.* 2019), I consider this an exciting
144 avenue for providing further insights into what drives the evolution of complex crystal
145 morphology in calcifying algae. If extended to other biomineralizers, it might even reveal a
146 unifying concept on which the apparently distinctive processes of calcification and
147 silicification coalesce.

148

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150

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155

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232

233 **Figure legends**

234

235 **Figure 1.** NERC Earth Observation Data Acquisition and Analysis Service (NEODAAS)
236 enhanced-colour view of waters around the British Isles in June 2019 (Median composite
237 between 17th and 23rd of June). Coccolithophores are blue-white and non-calcifying
238 phytoplankton (e.g. diatoms) are from dark to light red. The colour gradients (blue to white;
239 dark to light red) reflect the cell-density gradients of phytoplankton populations in the surface
240 layers with increasing densities from darker to lighter colours. Sediment is yellow.

241

242 **Figure 2.** Conceptual model based on Langer *et al.* (2021) describing the evolution of
243 complex crystal morphology in calcifying algae over the past ca. 350 million years. Anc =
244 non-calcified ancestor of coccolithophores; HOL = coccolithophores with holococcoliths;
245 HET = coccolithophores with heterococcoliths. Si = silicon.