

Diatom identification in the face of changing species concepts and evidence of phenotypic plasticity

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ABSTRACT – Although it is often suggested that diatom wall morphology is faithfully replicated at each cell division, it is also well known that the average cell size of a diatom population usually decreases as cells proliferate. Comparisons between the two valves of a single frustule may also reveal morphological differences, indicating that valve ontogenetic processes are susceptible to modification. This paper will discuss the different factors affecting valve morphology in diatoms and some of the implications for ecological and palaeo-ecological studies using diatoms. It will also consider some of the problems of evaluating variation between clonal populations, and the influence of findings from molecular biology and reproductive studies on the interpretation of morphology and species concepts.

KEYWORDS: *species concepts, environmental variability, diatom life history, clonal populations, diatom identification*

INTRODUCTION

Diatoms are well known for their regularly patterned siliceous cell walls (frustules) and the apparent fidelity with which wall construction is replicated at each cell division, yet it is also known that morphological changes can occur as cell size decreases, following successive vegetative divisions. In addition, examples of heterovalvy, when the two halves (valves) of a single frustule differ morphologically (apart from the monoraphid diatoms, which are inherently heterovalvate), can be found in the literature (Stoermer, 1967; Jordan *et al.*, 1991; Teubner, 1995; Meyer & Håkansson, 1996; McBride & Edgar, 1998; Werner & Smol, 2006), indicating that ontogenetic processes during wall formation may be modified by external factors. The existence of contrasting linkage and separation (or intercalary and terminal) valves in some chain-forming diatoms, summer and winter stages in others, or the formation of resting spores in neritic species, also indicates that a single genotype can have more than one morphological expression (Fryxell, 1978; Crawford, 1979; Fryxell & Prasad, 1990; Bentley *et al.*, 2012). Although results of molecular studies of diatoms are challenging some traditional interpretations of diatom relationships, as well as species concepts (e.g. Mann, 1999; Mann *et al.*, 2004; Kaczmarska *et al.*, 2013), for most ecological and palaeo-ecological studies, taxon identification still rests heavily on the morphology of the siliceous walls under light microscopy. Furthermore, inferring ecological conditions from diatom assemblages relies on known correlations between species occurrences and environment, assuming consistent identification of the taxa. It is therefore important to know to what extent morphology is consistent within a taxon, what changes may occur during size reduction and what effects environment can have on the morphology of a particular taxon.

This paper will consider some of the issues raised by changes in diatom species concepts based on reproductive and molecular studies on the one hand, and challenges to traditional species definitions from evidence of phenotypic plasticity (the ability of an organism to change its phenotype in response to environmental changes) on the other. What does this mean for palaeontologists

and palaeo-ecologists who must rely on morphological evidence for taxon identification? It will also consider whether subtler influences affect the interpretation of morphological evidence. Does the fact that diatom assemblages in any one sample (representing a short time-scale, days or weeks) are usually represented by clonal populations (at least in contemporary samples), each comprising many cells of identical genotype and very similar morphology, affect how the observer interprets the significance of morphological variation between populations at different sites? (Fossil assemblages can represent much longer time periods of accumulation, which may then include a range of populations and morphologies, but whether this inevitably results in wider species concepts is uncertain.) In addition, to what extent do developments in one research area impinge on another, subtly shifting paradigms and concepts by inference rather than by evidence?

DIATOM WALL FORMATION AND LIFE HISTORIES

One of the distinctive characteristics of diatoms is their cell wall construction and mode of formation. With few exceptions, all diatom cell walls essentially comprise two larger siliceous components (valves) with a series of linking elements (girdle bands) (Fig. 1), which are external to the protoplast, although variously anchored to the latter when alive (Round *et al.*, 1990; Cox, 2011). Following mitosis, new valves are generated in membrane-bound vesicles inside the daughter cells, being released to the exterior on completion (Fig. 2). Thus the valves of a single cell are not formed simultaneously, but each cell comprises an older (epi-) and a younger (hypo-) valve (Fig. 1). Girdle band formation occurs over the life of a cell, bands being added sequentially to the hypovalve until just before the next mitosis (Pickett-Heaps *et al.*, 1990; Round *et al.*, 1990; Cox, 2012). Once valves are formed and released their size is fixed; cell volume increases only by the addition of more girdle bands and expansion of the cell in the perivalvar direction. As the bands are also siliceous, there is little flexibility in the diatom frustule, constraining the space in which new valves can be formed after cell division. Depending on the thickness and rigidity of the cell wall (particularly in the

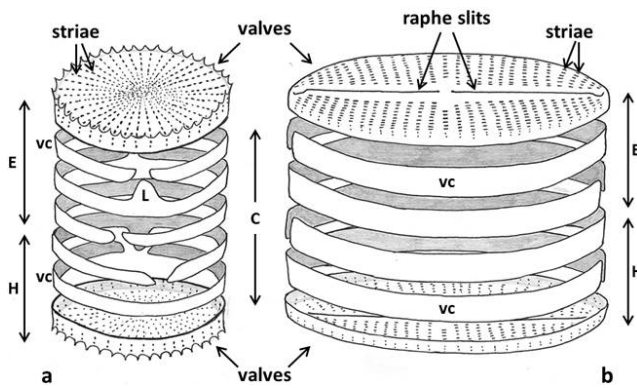


Fig. 1. Diagrams showing the basic cell wall structure of (a) a radial centric and (b) a raphid pennate diatom. Each frustule comprises two thecae, an older epitheca (E) and younger hypotheca (H), each comprising a valve with a series of girdle bands (cingulum, C). Rows of pores (striae) are radially oriented on centric diatoms (a), about a longitudinal axis of symmetry in pennate (b) diatoms. Raphid diatoms are characterized by the presence of paired raphe slits through the valve, enabling the diatom to move over surfaces. Girdle bands are often split rings, sometimes with a tongue-like extension (ligula, L) that inserts between the ends of the adjacent band. The band nearest the valve is usually designated the valvocopula (vc). Spines may be present around the periphery of a valve.

girdle region), new valves may therefore be more or less markedly smaller than the parent valves (Crawford, 1981).

Although there are some diatoms that show no size reduction over time (Rose & Cox, 2013), in most cases, a sexual reproductive phase followed by cell expansion (auxosporulation) (Figs 3–4), is necessary to prevent populations dying out as size reduction proceeds (Round *et al.*, 1990; Mann, 2011; Kaczmarska *et al.*, 2013). After expansion within some sort of lightly protected structure (perizonium, incunabulum, properizonium), new valves are laid down and vegetative division proceeds again. The first-formed (initial) valves are usually somewhat atypical, in part at least because they are not formed within the constraints of a

normal frustule, juxtaposed to a sibling valve. Thus, initial valves of radial centric diatoms may be hemispherical, while those of pennate diatoms have a convex rather than flattened valve face and some may be inflated at the centre (Cohn *et al.*, 1989; Assmy *et al.*, 2006), reflecting the perizonium shape. It has also been suggested that valve features, such as stria arrangements, may be atypical because full cytoskeletal control of morphogenesis, which determines the positioning of some wall structures, is not re-established until the first vegetative division (Pickett-Heaps *et al.*, 1990; Schmid 1994, Cox, 2002).

CONSISTENCY IN DIATOM STRUCTURE

While cell size and shape are subject to variation over the life cycle of particular diatom species, within any taxon the component parts of a frustule (processes [Fig. 5a], spines [Fig. 5b, c], girdle bands [Fig. 5c–e], striae [Fig. 5a, b, e–h], raphe [Fig. 5f–h]) (Fig. 1) are always formed in the same way and, therefore, have a consistent ultrastructure (Cox, 2011, 2012). Thus, species within a genus will be expected to share the same type of pore structure, e.g., poroid with cribrate occlusions, poroid with hymenate occlusions, loculate with external cribrate occlusions (Fig. 5e), loculate with internal hymenate occlusions (Round *et al.*, 1990; Cox, 2011), although some genera do not meet this requirement because they have been only partially revised (e.g. *Navicula* Bory sensu lato, *Triceratium* Ehrenberg sensu lato). Cingulum construction is similarly consistent within taxa, e.g. bands are with or without pores that are in turn occluded by cribra or hymenes; bands are with or without septa, are complete or split, etc. (Pocock & Cox, 1982; Round *et al.*, 1990; Cox, 2011 and references therein). Other fundamental characteristics of the cell wall show comparable consistency within taxa. Features associated with particular functions, e.g. apical pore fields through which polysaccharide pads or stalks are secreted, or raphe slits associated with motility, are also generally unchanged throughout the life cycle. On the other hand, stria arrangements may be more susceptible to modification, spacing varying within a certain range, as cell dimensions change. Thus, stria density within a clone will gradually increase with decreasing cell length, then

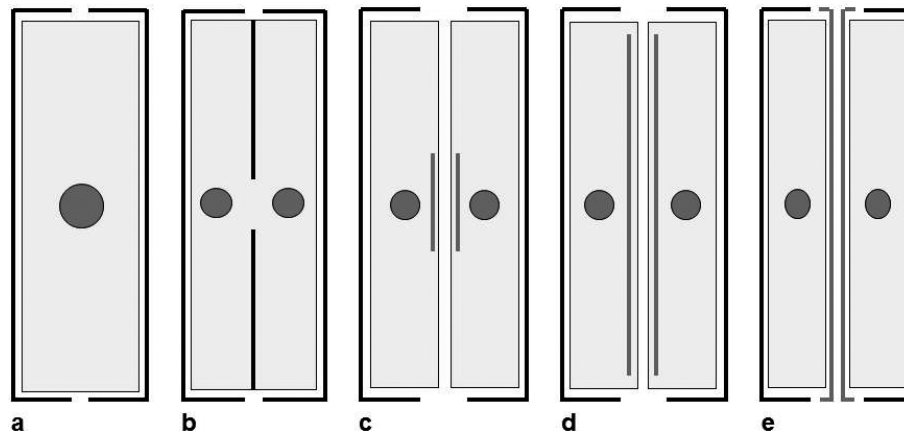


Fig. 2. Diagram showing process of vegetative cell division and formation of new valves. (a) vegetative cell; (b) cytokinesis proceeding after mitosis; (c) formation of new valves beginning at centre; (d) new valve formation continuing; (e) new valves completed and released to exterior. Thick black outer lines represent the valves (in girdle view), paler thick lines show the forming hypovalves, which are laid down within a membrane-bound vesicle inside the daughter cells, only released to the exterior after completion. Nuclei represented by grey circles.

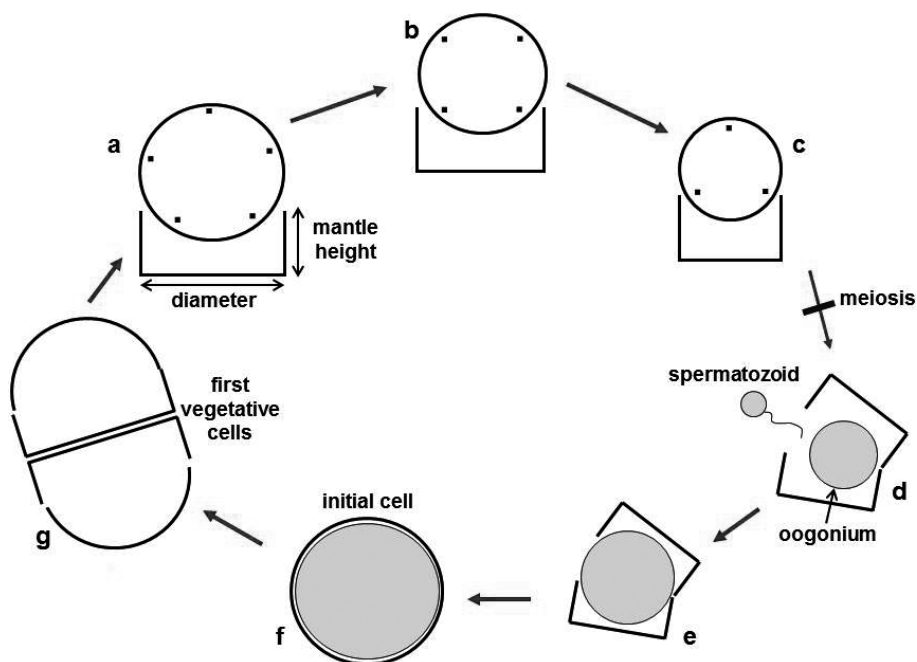


Fig. 3. Diagram summarizing typical life cycle of an oogamous (usually centric) diatom. (a–c) Valve diameter decreases over a series of vegetative divisions, peripheral processes (shown as dots) may decrease in number with decreasing diameter, while diameter: mantle height ratio changes. (d) Meiosis occurs to produce oogonium and flagellate spermatozoa. (e–f) Following gametic fusion, zygote (auxospore) expands to maximum size and (f) deposits new walls externally. (g) Initial cells are often more or less spherical, first vegetative cells are often heterovalvate with an initial valve and a ‘normal’ vegetative valve. Non-silicified stages are shown with a thin line, siliceous walls by thick black lines. Timing of meiosis indicated by arrow with transverse bar.

revert to a minimum before gradually increasing again, the cycle then repeating as cells continue to decrease in length (Cox, 1983); striae are packed slightly closer together before size restrictions preclude further packing and the number of striae in 10 μ m is reduced. Stria densities vary because units of fixed dimensions (pores) are being ‘fitted into’ changing areas. Similarly, if processes at the circumference of a round valve must have a minimum distance between them, their number per valve will be decreased if that valve decreases significantly in size (diameter and circumference) (Fig. 3).

Whereas electron microscopy (particularly SEM) reveals the 3D structure of diatom cell walls (Fig. 5a–h), this cannot always be determined with light microscopy (LM). Similar morphologies (in LM) could be generated from contrasting ultrastructural elements, e.g. longitudinal lines may mark the edge of internal alveolar openings (in *Pinnularia* Ehrenberg and *Caloneis* Cleve), longitudinal canals (in *Neidium* Pfitzer) or aligned gaps within striae (in *Neidiopsis* Lange-Bertalot & Metzeltin) (Bahls, 2014). Similar morphologies under LM can also be generated by contrasting morphogenetic processes, e.g. stauros v. fascia (Cox, 2001, 2002). It is also increasingly clear that shape and symmetry are not infallible guides to relationships at generic (Cox, 1979, 1982) or higher taxonomic levels (Cox, 2002; Kooistra *et al.*, 2003; Alverson *et al.*, 2006).

INTRINSIC CHANGES IN DIATOM MORPHOLOGY

I refer to changes that are a function of the life cycle of the diatom, unaffected by changing environment, as intrinsic. The effect of size reduction on diatom cell shape varies with the shape and

symmetry of the taxon (taxon = any group of organisms considered a unit), as well as the depth and rigidity of the girdle region. Diatoms with circular valve faces (radial centrics) (Figs 1, 5a–c) do not change shape in valve view, although the cell proportions in girdle view may change because the cell diameter decreases but mantle depth does not (Fig. 3). This is most striking in diatoms such as *Aulacoseira* Thwaites and *Melosira* C. Agardh, in which valve diameter in large cells may be two or three times that of the smallest cells, seen most clearly when auxospores develop still associated with the original filament of small cells (Crawford, 1974, figs 5, 6). In multipolar diatoms the angularity of the valves may change, with angles becoming less acute as maximum diameter decreases. There is also some evidence that angles may be lost (e.g. *Hydrosera* Wallich – Cox, 2013, figs 2–7).

Although valve shape does not change in many radial centric diatoms, there can be changes in the relative proportions of central and peripheral areas. Thus, in some species, the size of marginal alveoli in *Cyclotella* (Kützing) Brébisson may remain approximately the same, but if valve diameter decreases, these occupy a larger proportion of the valve face in smaller than larger specimens (Prasad & Nienow, 2006, figs 35–39). In other centrics the number of marginal processes may decrease with decreasing cell size (Fig. 3), if a minimum distance must be maintained between adjacent processes (Theriot & Stoermer, 1981; Theriot, 1988; Beszteri *et al.*, 2005).

In so-called bipolar centric and pennate diatoms, which both have a longer and a shorter valve axis (Fig. 5d–h), size reduction is usually accompanied by a shift in valve proportions, length decreasing more than width; i.e. size reduction is allometric

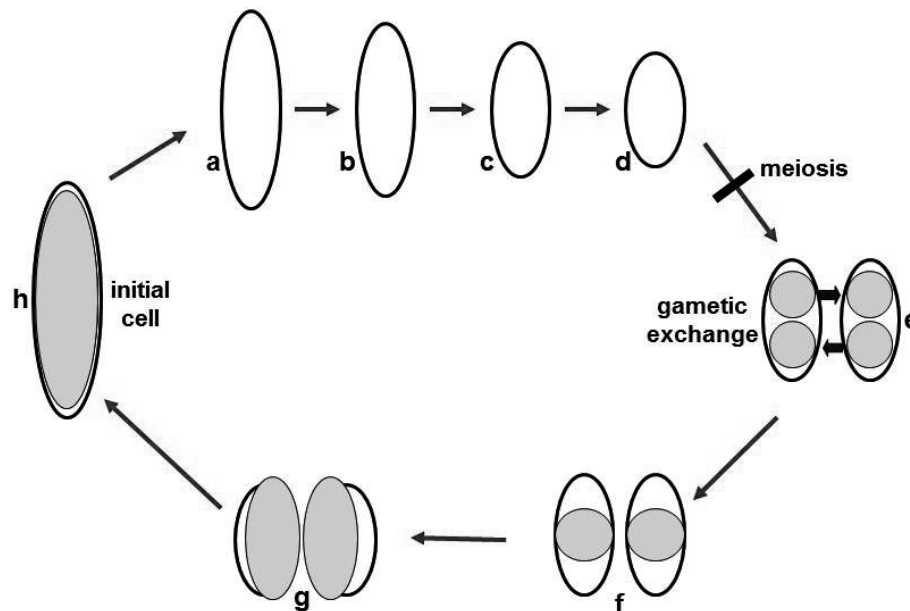


Fig. 4. Diagram summarizing the typical life cycle of an isogamous (usually pennate) diatom. (a–d) Valve length decreases over a series of vegetative divisions although valve width may remain more or less constant. Shape of valve therefore changes. Meiosis occurs when compatible cells pair, after which, (e–f) there is an exchange of gametes and fusion. The auxospore expands to its maximum size outside the confines of the parent valves (g) before (h) laying down new valves externally. Non-silicified stages are shown with a thin line, siliceous walls by thick black lines. Timing of meiosis indicated by arrow with transverse bar.

(Fig. 4) (Cox, 1985, 1986; Amato *et al.*, 2005; D’Alelio *et al.*, 2009). There may also be changes in the shape of the valve apices, which may become more rounded in some taxa, but in other cases, apical shape is maintained or even exaggerated (e.g. Geitler, 1932; Hustedt, 1955; Cox, 1986). Length reduction may also modify the proportional relationships of valve features, e.g. raphe length in *Berkeleya rutilans* (Trentepohl) Grunow remains more or less constant, but central area length decreases with decreasing valve length thereby changing raphe length: central area length ratio (Cox, 1975; Lobban, 1984). Meanwhile, in *Donkinia* Ralfs in Pritchard the central inflexion of the sigmoid raphe becomes sharper as valve length decreases (Cox, 1983). It is therefore impossible to make a simple generalization about shape changes with size reduction, or to be certain of the particular morphological changes within a taxon (species, variety) without following it for its entire life cycle, or at least being able to link the extremes of the cell size range (Hostetter & Hoshaw, 1972; Steinman & Sheath, 1984; Cox, 1985; Rose & Cox in press). This also has implications when comparing different-sized specimens in samples from different locations, e.g. because of the interrelationship between raphe inflexion and valve length in *Donkina lata* E. J. Cox, different-sized cells looked rather different, although culture studies showed that they were part of same spectrum (Cox, 1983).

Legendre polynomial shape analysis has been used (Stoermer & Ladewski, 1982; Stoermer *et al.*, 1986) to compare presumed conspecific specimens from different samples/locations to assess the likelihood of conspecificity, or to explore variation within a single assemblage (Theriot & Ladewski, 1986). It has also been applied to taxa which contain diverse forms (Rhode *et al.*, 2001; Pappas & Stoermer, 2003) to help evaluate the taxonomic status of different shape groups. Continuity in shape variation is

interpreted as indicating that the specimens represent stages within the life cycle of a single species, whereas discontinuous shape variation may indicate the existence of discrete species (Pappas & Stoermer, 2003). More recently, other morphometric studies have shown that shapes of apices and central areas in raphid diatoms are more informative than traditional size measures with respect to identity in morphologically similar taxa (Vesela *et al.*, 2009).

PHENOTYPIC RESPONSES TO EXTRINSIC FACTORS

The intermittent occurrence of so-called Janus or heterovalvate diatom frustules (where the two valves in a single frustule are morphologically different) reveals the effect of changing environment on single genotypes (Stoermer, 1967; Jordan *et al.*, 1991; Teubner, 1995; Meyer & Håkansson, 1996; McBride & Edgar, 1998; Werner & Smol, 2006); i.e. diatoms are able to modify their wall morphology to a greater or lesser extent under different conditions. In the case of *Gomphonema* (McBride & Edgar, 1998) the striae on one valve are much more closely spaced than on the other, while in *Mastogloia* Thwaites (Stoermer, 1967) the stria pattern was sufficiently different that the valves would be assigned to different species (*Mastogloia grevillei* W. Smith and *M. elliptica* var. *dansei* [Thwaites ex W. Smith] Cleve). The recognition of spring and winter forms in *Proboscia* Sundström and *Eucampia antarctica* (Castracane) Mangin indicates that different morphologies correlate with seasons (Fryxell, 1991, 1994; Jordan *et al.*, 1991), while *Chaetoceros dichaeata* Ehrenberg produces cells with shorter setae in sea ice than in the plankton, and the setae are bent inside the filament rather than turning outwards (Ligowski *et al.*, 2012). Meanwhile, the production of linkage and separation valves in some filamentous diatoms, e.g. *Aulacoseira*, *Skeletonema* Greville, *Eucampia* Ehrenberg, *Cymatosira* Grunow

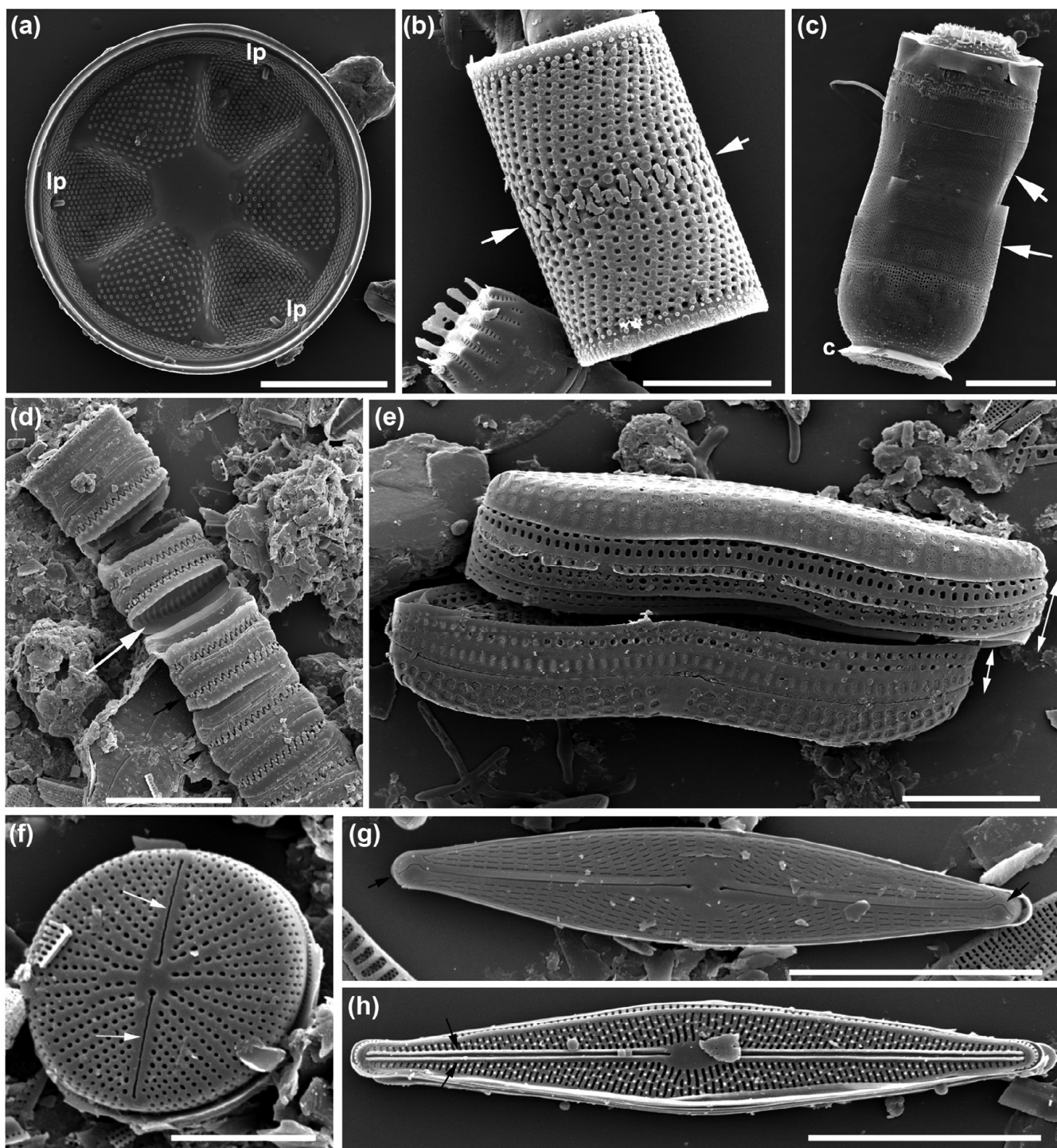


Fig. 5. Scanning electron micrographs of diatoms to illustrate some of the morphological features. (a) Internal view of whole valve of *Actinoptychus*, showing valve undulation, radially arranged striae and labiate processes (lp) near the valve margin in alternate sectors. (b) External view of two valves of *Aulacoseira*, linked by spines (arrow). Pores form spiral striae on the valve mantle. (c) External view of two pairs of cells of *Melosira*, held together by shared girdle bands (arrows). Valve faces are defined by a corona (c) and have irregular spines at the centre. (d) External view of a chain of *Pseudostaurosira* cells linked by interlocking spines (black arrows) on sibling valves. Note that some cells have partially opened and the internal structure of the striae is visible (white arrow). (e) External view of two sibling cells of *Achnanthes*, recent products of mitosis. Note the bands associated with each daughter cell (double-headed arrows) and the cribrate occlusions to the pores in the striae and on the bands. (f) External view of a valve of *Cavinula*, showing the straight central raphe slits (arrows) and the radial striae comprising round areolae. (g) External view of a cell of *Navicula*, showing the central raphe slits with hooked polar fissures (arrows), and radiate striae formed of lineate areolae. (h) External view of *Brachysira* showing central raphe slits flanked by siliceous ribs (arrows) and with a ridge around the valve margin, at the junction of the valve face and mantle. Small knobs of silica are irregularly distributed over the valve face between the radiate striae. Scale bars 5 μm (b, d, f) and 10 μm (a, c, e, g, h).

(Fryxell, 1978; Crawford, 1979; Fryxell & Prasad, 1990), indicates another phenotypic response to the environment, allowing the diatoms to modulate their position in the water column as filament length changes. However, although light regimes have been implicated in the production of linkage valves in *Eucampia* (Fryxell & Prasad, 1990) and computer simulations suggest that light regime could control the type of spine production in *Aulacoseira* (Bentley *et al.*, 2012), there have been no controlled experiments to determine precisely how environmental conditions control the switch from one type of valve production to another.

Studies on populations of single species across a range of environmental conditions can reveal correlations between morphology and environment that may have indicator value. Thus, Cortese & Gersonde (2007) found that valves of *Fragilariopsis kerguelensis* (O'Meara) Hustedt were longer and slimmer with more costae in 5µm in warm than cold waters, and suggested that this could be a useful proxy for sea surface temperature. Meanwhile, based on analysis of three sediment cores from different parts of the Southern Ocean, Shukla *et al.* (2013) suggested that average valve size of this diatom reflects nutrient availability, particularly iron, and could reveal the effects of climate and nutrient cycling in the ocean.

Experimental studies allow cells to be grown under a range of known conditions and can therefore reveal how factors such as salinity and temperature modify morphology (Schultz, 1971; Schmid, 1976; Fryxell, 1988; Syvertsen, 1987; Hausmann & Lotter, 2001; Balzano *et al.*, 2011; Trobajo *et al.*, 2011). With such information, it then becomes possible to link particular morphs to particular environmental conditions causally (Schmid, 1976; Syvertsen, 1987; Fryxell, 1988; Hausmann & Lotter, 2001), rather than relying on correlations alone (Hürlimann & Straub, 1991; Theriot *et al.*, 1988; Cortese & Gersonde, 2007). Morphology can then be used as a proxy for environmental factors, and applied to past climatic and oceanic change reconstruction (Crosta, 2009; Cortese *et al.*, 2012; Shukla *et al.*, 2013). More dramatic phenotypic plasticity is seen in a few diatoms, e.g. *Diademsis gallica* W. Smith, in which the raphe slits are variously reduced or even 'lost' (i.e. completely infilled) when grown in culture, and linking spines are developed around the valve face margin (Granetti, 1977; Cox, 2006). Unfortunately experimental studies have been conducted on only a tiny proportion of diatoms, so we have little idea of the extent to which morphology could be modified in the majority of taxa, or of the potential environmental signals reflected in morphology.

INFERENCES FROM OTHER CONTEMPORARY DIATOM RESEARCH

Molecular studies

The application of molecular techniques to diatoms has revealed genetic diversity, based on sequence differences in targeted genes, within and between clones of traditionally defined taxa, which has in turn stimulated the search for correlated morphological variation. Investigations of *Skeletonema costatum* Greville *sensu lato* (Medlin *et al.*, 1991; Sarno *et al.*, 2005; Zingone *et al.*, 2005) resulted in the recognition of five new species, the molecular differences being supported by morphological differences. However, variability in two taxa (*S. dohrnii* Sarno & Kooistra and *S. marinoi* Sarno & Zingone) subsequently showed that these could not be separated on morphological grounds (Kooistra *et al.*, 2008).

Investigations of *Pseudonitzschia* H. Peragallo have also led to a proliferation of new species, discriminated on molecular and morphological grounds, probably stimulated by the potential toxicity of some of those species (Lundholm *et al.*, 2012). In evaluating the significance of sequence differences, some workers suggest that divergence thresholds could be used to determine whether taxa warrant separation at different taxonomic levels (Coleman, 2009; MacGillivray & Kaczmarek, 2011, 2014), but while defined degrees of difference might be logically attractive, they are intrinsically arbitrary.

Reproductive studies

Work on the *Sellaphora pupula* Kützing complex and some other benthic freshwater diatoms (Mann, 1984, 1988, 1989) revealed that within some apparently well-known species, subtle morphological differences reflected breeding patterns. Individual diatoms would only mate with those with the same morphology (assuming both were below the size threshold for sexualization) to produce viable offspring. Building on this work, Mann (1999) argued that reproductively distinguishable strains should be recognized as biological species, eventually formally creating new species for the different taxa (Mann *et al.*, 2004). Further work has revealed that not only can subtle morphological differences indicate different breeding populations, but also that within obligately allogamous populations of diatoms (i.e. where different strains must pair to reproduce), there is discrimination into mating types (Chepurnov *et al.*, 2004; Mann & Chepurnov, 2005; Vanormelingen *et al.*, 2008).

Biogeography

Whereas many diatom species were once thought to be cosmopolitan, based on the assumption that micro-organisms could be easily transported ('everything is everywhere'), plus the use of European floras to identify taxa from around the world, resulting in force-fitting taxa (Tyler, 1996), closer examination of specimens from diverse sites has revealed that they are often morphologically distinguishable (Zidarova *et al.*, 2010). Endemism has also been shown for some diatoms (Theriot *et al.*, 2006; Vyverman *et al.*, 2010); others have distinct geographical ranges (Kooistra *et al.*, 2008) and taxa that were once thought to be bipolar are being reassessed (e.g. Van de Vijver *et al.*, 2005). Where molecular analyses have been employed, biogeographical patterns have been revealed for some apparently widespread taxa (Pouličková *et al.*, 2010; Kermarrec *et al.*, 2013), yet widely separated clones of others may share the same genetic identity (Trobajo *et al.*, 2009; Evans *et al.*, 2008, 2009). In other words, individual taxa behave in different ways; some have been able to spread widely, others are much more restricted in their distribution.

DISCUSSION

Molecular, reproductive and biogeographical studies on diatoms have generated many new taxa that are morphologically more finely discriminated, and almost invariably recognized at the species level. Meanwhile, many diatomists still work primarily with morphology and light microscopy. Yet even here, narrower species concepts are being applied, assuming that subtle morphological discontinuities indicate species differences. In other words, authors are inferring that, if small morphological differences between other taxa can reflect the existence of breeding barriers

(Droop, 1995; Droop *et al.*, 2000), molecular differences can warrant species-level recognition, or taxa can be geographically restricted, the same must apply to their specimens. Yet they are concluding this merely by inference; they are doing so without independent supporting evidence from molecular, reproductive behaviour or biogeographical studies. Yet on the other hand, it is clear from life history and experimental studies that diatom wall morphology can be significantly modified such that genetically identical individuals (or pre-gametangial and their corresponding post-initial cells) could be identified as different species (Schultz, 1971; Cox, 1985; Trobajo *et al.*, 2006; Rose & Cox in press).

Studies into the effects of salinity on diatom morphology demonstrate most clearly the flexibility in diatom morphogenesis (and its outcome) (Schultz, 1971; Trobajo *et al.*, 2004, 2011; Balzano *et al.*, 2011). Responses are species-specific (Trobajo *et al.*, 2011) and, in some cases, discriminating between supposed bioindicators becomes problematic (Trobajo *et al.*, 2004). The effects of temperature on diatom morphology have also been clearly demonstrated for a number of marine taxa, e.g. *Thalassiosira* species (Syvertsen, 1979; Fryxell, 1988). It is, however, impossible to predict, *a priori*, the morphological range of a particular taxon. Shape analysis can be helpful in determining (*a posteriori*) whether or not specimens could be part of the same taxon (Stoermer & Ladewski, 1982; Stoermer *et al.*, 1986; Theriot & Ladewski, 1986; Rhode *et al.*, 2001; Pappas & Stoermer, 2003), but perhaps focusing on other morphometric features, such as shapes of central areas and valve apices (Veselá *et al.*, 2009), could improve our understanding of the morphological range of other taxa. It may also provide more reliable taxonomic criteria for species separation and identification.

It is clear that there is considerable genetic diversity within the diatoms, and it is also clear that many previously considered widespread taxa should probably be more finely discriminated, and that finer discrimination would in turn provide useful ecological or biogeographical information. However, there is a danger that every newly discovered variant is being treated as a new species (even if cryptic or pseudo-cryptic); this ignores the fact that genetic diversity is an intrinsic property of any species, allowing for its ability to adapt to different or changing environments and thereby survive.

Diatoms are primarily clonal

Because diatoms reproduce asexually for most of their life cycle, clonal populations can develop in which a single genotype is represented by many cells, which are likely to show very little morphological variability (other than size-related) at any one time and site under the same environmental conditions. (Fossil populations may show greater variation because they represent accumulations of cells that may have grown under different conditions and may also represent different clonal populations.) When populations from different sites are compared, because each is represented by many, very similar, individuals, it may then be easy to over-emphasize the significance of any differences, forgetting that the individuals at each site do not necessarily represent different genotypes. In other words the differences between two populations, each comprising a single genotype, are no more significant than the differences between two individuals, one of each genotype, representing those populations.

In order to obtain sufficient DNA, most molecular studies on diatoms use cultured strains, which again will comprise many cells with the same morphology. Therefore the morphological comparisons will probably involve many individuals in a sample, which similarly may tend to encourage over-emphasis on the significance of any morphological differences observed between those clones. Thus, where sequence differences have been paralleled by morphological variation, the inference has usually been that these represent taxa that warrant (formal) recognition (Medlin *et al.*, 1991; Sarno *et al.*, 2005; Lundholm *et al.*, 2012) and a proliferation of species, cryptic species and pseudo-cryptic species has resulted. But what do these really tell us? Very few studies have investigated the morphological variability of these clones under different conditions; the genetic sequence is linked to a single morphological expression, much as traditional typological taxonomy links the name to a specimen. Furthermore, some of these fine morphological criteria are now being shown to be more variable and therefore unreliable indicators of taxon identity (Kooistra *et al.*, 2008).

Species definitions v. ecological interpretation

While, in broad terms, the taxonomist seeks to describe and define taxonomic units, using a variety of tools to help discriminate those units, the ecologist is usually seeking to understand how those units respond to, or provide information about, their environment. The question is, to what extent do the units defined by the taxonomist provide consistent information about their biology or ecology? If the discovery of genetic diversity within taxa is leading to a proliferation of names, yet there is little or no physiological or ecological difference between them, how should an ecologist treat them? Is there any relevance (or sense) in giving each genotype a different name? If different morphological forms of the same taxon are given different specific epithets, will the unwitting ecologist record a floristic change if both are encountered at different times, or in different places, rather than understanding that a difference in environment has been reflected in contrasting morphology? Clearly recording that different morphs have been observed provides ecological information, but what is the most appropriate way to do this?

Previously (Cox, 1997), I suggested that if the taxonomic unit 'species' is used only where there is evidence of breeding barriers between populations, 'variety' could be used for within-species populations that are ecologically and morphologically distinct, whereas cases of phenotypic plasticity within species be designated as forms. The latter can then be used to convey ecological information, whether that be salinity, temperature or some other factor. Since we still do not understand patterns of relationship or diversification for the majority of diatoms, and demonstrating sexual compatibility or incompatibility for even a reasonable proportion of diatoms would be a monumental task, it is probably unreasonable to expect a biological species concept (Mann, 1999) to be used for most diatoms. However, expanding the use of varietal and form designations is more practicable, and would help convey ecological and biogeographical information about morphologically similar taxa, where the application of different specific epithets may obscure that. Nevertheless, discrimination of different varieties and forms (not just the use of a species name '*sensu lato*') remains necessary if such ecological information is to be conveyed.

CONCLUSIONS

The worldwide distribution, ecological diversity and distinctive wall characteristics of diatoms have all contributed to their use as contemporary ecological indicators and also, given their extensive fossil record because of the resistant nature of their silica walls, as palaeo-indicators. But for any bioindicator system to work effectively, ecological information must be unequivocally linked to reliably identifiable taxonomic units. Traditionally, diatoms have been identified based on the morphology of their silica cell walls under light microscopy. While light microscopy remains the technique of choice for much routine investigation, electron microscopy has revealed both the intricacies of wall construction and also subtle variation within and between classical species. Molecular biology is also uncovering unsuspected genetic diversity within supposedly well-known diatom species, which is being interpreted as evidence of the existence of unrecognized species, cryptic or pseudo-cryptic species. (There is an apparent reluctance by diatomists to use classical infraspecific categories, such as subspecies, variety or form.) But the ecological ranges and tolerances of these new taxa are often unknown and thus their relevance to environmental/palaeoenvironmental research is unclear.

One problem seems to be the continued view of diatoms as static expressions of their genotype, rather than as organisms that can potentially modify their wall morphology in response to their environment. Experimental studies can help us understand the possibilities of, and constraints on, phenotypic responses to particular environmental factors. In turn, this can aid the ecological interpretation of morphological assemblages. Is there real species turnover, or are the same species modifying their morphology in response to environmental change? Reviving the use of forma when morphological change is a response to environment would allow ecological information to be recorded effectively, without suggesting that totally different species are involved. Meanwhile, although they may be challenging and time-consuming, experimental studies into key ecological indicators, in conjunction with genetic studies, are an important way to improve our understanding of species response and adaptation to environment, and thereby to evolutionary processes.

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