DIATOMS: BIOLOGY AND APPLICATIONS SERIES

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Chapter 3

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Chapter 4

Figure 4.1 Morphology of an araphid diatom (a), Staurosira construens var. venter (scale bar: 2 μm) and a raphid diatom (b), Navicula radiosa (scale bar: 10 μm), diatom on valve view. The raphe (indicated by a white arrow) runs through the whole valve and is a primary structure for adhesion to surfaces and moving. (c) A close-up of the raphe (indicated by a white arrow) is shown (scale bar: 1 <u>µm). (SEM images downloaded from diatoms.org</u> <u>with permission [4.107] [4.120] [4.143].)</u>

Figure 4.2 Schematic of a benthic biofilm. Algae, predominantly pennate biofilms together with bacteria, protists, and fungi, are embedded within a protective matrix of extracellular polymeric substances (EPS). A benthic biofilm experiences high variability of spatial and temporal gradients of environmental factors such as inorganic nutrients, dissolved organic matter (DOM), and light. (Figure from Sabater *et al.* [4.129]. Reprinted under CC-BY license.)

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Chapter 5

Figure 5.1 The effect of incubation time on Stauroneis response times in a mixed culture of Craticula cuspidata and Stauroneis phoenicenteron. This graph displays the average direction change response times for Stauroneis phoenicenteron cells in the presence of ca. 9:1 ratio of live C. cuspidata: S. phoenicenteron cells. S. phoenicenteron cells were isolated and washed from culture and incubated together with *C. cuspidata* cells (C/S) in a 9:1 *C. cuspidata:S. phoenicenteron* ratio. Cells were then irradiated at their leading end with high irradiance (ca. $10^4 \mu mol/m^2$ s) 1s pulses of blue (470 nm) light, and observed to determine the time until they changed direction (response time). Response times significantly increased almost 2-fold from the initial incubation interval (0-10 min) within 20-30 min ($30\pm 2 \mu m/s$ and $57\pm 7 \mu m/s$ respectively, *P*=0.003). Graphs represent the mean values of response times ± 1 SE. For comparison, unirradiated Stauroneis cell response times were $155\pm 11 \mu m/s$. Error bars represent ± 1 SE.

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the presence of dead *P. viridis* (Dead Pinn) that had been killed by immersing the *P. viridis* cells for 30 sec in 95% ethanol prior to rinsing the *P. viridis* with distilled water and fresh diatom medium. The *S. phoenicenteron* cells in the presence of living *P. viridis* showed a significant increase in response time, even though they were exposed to the same medium as the isolated *S. phoenicenteron*. while those in the presence of dead *P. viridis* showed no such increase in response time. Error bars represent ± 1 SE.

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Figure 6.3 Top: Change in environmental driver (pressure) dependent on the presence of a "stabilizing biofilm" or under "regular erosion." Bottom: Low-temperature scanning electron micrograph of MPB biofilm structure. (Image: Irvine Davidson, University of St Andrews.)

Chapter 7

Figure 7.1 Proposed model of the control of vertical migration by sediment-inhabiting benthic pennate diatoms, as responding to main directional environmental stimulus, light and gravity. The figure illustrates the variation with the time of day of the diatom biomass at the sediment surface on samples kept in the dark (closed circles) and exposed to constant low light (150 μ mol m-² s-¹) during the subjective low tide period (open circles) (for more details, see [7.18]). Example of a day when the low tide takes place during the middle of the day. The gray horizontal plots represent the strength (bar thickness) of photo- and geotaxis, of negative and positive signal, along the day. (1) <u>Upward migration starts before the beginning of</u> the light period, driven by negative geotaxis. (2) Negative geotaxis ceases roughly at the time expected for start of the low tide light period: if no light is available at surface, the diatoms stop migrating upwards and the incipient biofilms start to disaggregate, due to random cell movement or weak positive geotaxis. (3) During light exposure, cell movements are controlled mainly by phototaxis, either positive (under low light intensities) or <u>negative (under high light intensities). In the</u> particular case of the data in the figure, positive phototaxis dominates, as samples were exposed to low intensity. (4) Anticipating the end of the light period, geotaxis becomes dominant over phototaxis, as cells begin to migrate downwards without any changes in incident illumination. Vertical gray areas represent periods of darkness. Vertical white area represents the period of light exposure (150 µmol m^{-2} s⁻¹) of the light exposed samples.

Chapter 8

<u>Figures 8.1–8.13 LM images of *Eunotia* taxa in valve view (Figures 8.1–8.8) and girdle view, ventral-side up (Figures 8.9–8.11) to show variation in morphology and raphe shape and location (arrow</u>

in some images). (Figure 8.1 [8.28]) E. bilunaris (Ehrenb.) Souza - raphe recurved almost 180°. (Figure 8.2) E. serra Ehrenb. - raphe on valve face with slight curve toward apex. (Figure 8.3 [8.28]) E. <u>areniverma Furey, Lowe et Johansen</u> – raphe follows margin of apex from ventral to dorsal margin, and up onto dorsal mantle. (Figure 8.4 [8.28]). *E. pectinalis* var. *ventricosa* (Ehrenb.) Grunow (E. pectinalis var. ventralis (Ehrenb.) Hust. = synonym). (Figure 8.5) *E. bigibba* Kütz. - raphe with slight recurve. (Figure 8.6) *E. incisa* Smith ex. <u>Gregory – raphe (not visible) only present on valve</u> mantle (see Figure 8.17). (Figure 8.7) E. muscicola Krasske - raphe (not visible) on the valve face with slight recurve (see Figure 8.20). (Figure 8.8) Close up of valve apex and raphe of *E. bilunaris* in Figure 8.1. (Figure 8.9) *E. bigibba* and (Figure 8.10) *E. tetraodon* Ehrenb. - ventral-girdle view. Depth of valve only permits part of raphe to be in focus (on one plane) at a time. (Figure 8.11) Unknown valve in ventral-girdle view. Raphe almost all in focus. (Figure 8.12) Epiphytic cells of E. bilunaris (E) standing up on end from the mucilaginous sheath(s) of the cyanobacterium *Hapalosiphon* Nägeli ex Bornet et Flahault (Image credit R.L. Lowe). (Figure 8.13) SEM image of *Eunotia* on a bryophyte. For all images except for Figure 8.8: black scale bar = $10 \mu m$. Figure 8.8: white scale <u>bar = 5 µm. (Figures 8.1, 8.3, 8.4 originally</u> published in Furey et al. [8.28] www.schweizerbart.de/journals/bibl diatom). Figures 8.14-8.22 SEM images of Eunotia taxa to show variation in morphology, along with external and internal raphe (R) shape and location on the

valve face and valve mantle, helictoglossa (h),

shape, location, and internal expression of the rimoportula (r), and external expression of the rimoportula pore (rp). (Figure 8.14 [8.28]) E. <u>areniverma – raphe follows margin of apex from</u> ventral to dorsal margin, and up onto dorsal mantle. (Figure 8.15 [8.28]) E. areniverma internal view of apex. Rimoportula located mid apex. (Figure 8.16 [8.28]) E. pectinalis var. ventricosa - external view showing path of raphe from mantle onto valve face with slight recurve. (Figure 8.17 [8.28]) E. incisca - raphe located completely on valve mantle. External expression of rimoportula. (Figure 8.18) E. bigibba - curve of raphe onto valve face with slight recurve. (Figure 8.19) *E. serra* – raphe on valve face with slight curve toward apex. (Figure 8.20) E. muscicola curve of raphe onto valve face with a slight recurve and (Figure 8.21) internal view of valve apex with rimoportula located close to the helictoglossa. (Figure 8.22) *E. serra* – internal view of valve apex with rimoportula located closer to dorsal margin. Scale bars as shown. (Figures 8.14–8.17 originally published in Furey et al. [8.28] www.schweizerbart.de/journals/bibl diatom). [8.38] [8.53] [8.66], though their position could be derived if a more complex raphe system became reduced (discussed by Kociolek [8.44] and Siver and Wolfe [8.69]). Movement in eunotioid diatoms with their short raphe system, typically described as slightly $(< 2 \mu m/sec [8.31])$ or weakly motile (2 to 4 $\mu m/sec$ [8.31]) (see eunotioid taxa in DONA [8.21] - Furey [8.26]), contrasts with that of more motile forms like naviculoid, nitzschioid, or surirelioid diatoms with more extensive raphe systems, typically described as moderately to highly motile. Examination of motility in *Eunotia* species may

provide unique insight into motility in diatoms, especially for diatoms with more complex raphe systems. A search for the terms "motile" or "move" in florae focused on *Eunotia* (e.g., [8.20] [8.28] [8.48] [8.51]) and in >40 manuscripts with descriptions of *Eunotia* taxa new to science revealed little to no mention of motility. This chapter focuses on motility in the diatom genus *Eunotia*, but does not cover cellular or biomechanical details around the mechanisms of movement, as other chapters in this book discuss these aspects at length.

Figures 8.23–8.27 Schematic representation of some of the movement types for *Eunotia*. (Figure 8.23) Schematic of forward movement – apical displacement where cells tilt slightly so the anterior ends of the valves remain in contact with the surface and the posterior ends become slightly raised (**schematic modeled after Palmer [8.60] plate vi. fig. 2, and Bertrand [8.6] fig. 1**). (Figure 8.24) Valve in girdle view. (Each raphe branch labeled after Bertrand [8.6] (**see also Harbich [8.30]**). Black arrow following the line of the raphes on B to C apices represents diagonal line of direction, where the raphe on the C end becomes active. (Figure 8.25) Black horizontal arrow represents diagonal line of direction. Bidirectional arrow shows transition between raphes involved in forward motion (**schematic modeled after Bertrand [8.6], fig. 1**). (Figure 8.26) Schematic of a vertical polar pivot which can return a cell in girdle view, dorsal-side down (a) to ventral-side down (b,c) where a cell can then continue forward movement (c) (**schematic modeled after Palmer [8.60], plate vi. fig. 2, and Bertrand ([8.6], fig. 5]**).

(Figure 8.27) Schematic of a horizontal, polar pivot (a,b) to show direction of raphe activity (straight arrows, A and B). Note the cell is depicted ventral side up so the raphe branches are visible (rather than dorsal side up as the movement occurs). Curved arrows show direction of rotation. Dot represents the pivot point. (**Modified after images from Harbich [8.30]). See additional schematics in Bertand [8.5].

Chapter 9

Figure 9.1 Gliding cells of *Nitzschia sigmoidea* with stalked epiphytes of *Pseudostaurosira parasitica*: a single epiphyte in connective view (a), in valve view (b), and two epiphytes attached to the same frustule (c), which can be seen in movement [9.39].

Figure 9.2 Cells of *Nitzschia sigmoidea* with adnate epiphytes of *Fallacia helensis* (OM): single epiphyte in connective view and host in valve view (a), four epiphytes in valve view and host in connective view (b), which can be seen in movement [9.43].

Figure 9.3 Cells of *Nitzschia sigmoidea* with many epiphytes (OM): epiphytes of *Amphora copulata* on a still gliding host (a), epiphytes of *Amphora copulata* on a dividing host (b), epiphytes of *Amphora copulata* and *Pseudostaurosira parasitica* on the same host (c), which can be seen in movement [9.41].

Figure 9.4 Focus on epiphytes of *Nitzschia* sigmoidea (SEM). Adnate, *Fallacia helensis*, and stalked, *Pseudostaurosira parasitica*, epiphytes on the same host (a), two cells of *Pseudostaurosira parasitica* still associated after division (b), two superimposed cells of *Fallacia helensis* after division and a third single one attached on the edge of the frustule (c), apex of a cell of *Pseudostaurosira parasitica* with the mucilaginous pad secreted for adhesion (d), individual of *Amphora copulata* (e) and internal view of one valve of *Amphora* sp., probably *Amphora copulata* var. *epiphytica* Round & Kyung Lee, considering the almost circular areolae on the ventral side (f). Scale bars indicate 10 µm, except in 9.4d (=5 µm).

Figure 9.5 Variations in the specific composition of epiphytes on *Nitzschia sigmoidea* between two sampling sites located on two connected rivers (upand downstream sites), expressed as the occurrence of three epidiatomic species on frustules of *N. sigmoidea*. In fact, through the observation of fresh material, *N. sigmoidea* could not be strictly distinguished from the close species *N. vermicularis*. Both species were present in each site (see [9.44] for details).

Figure 9.6 Sigmoid frustules of *Nitzschia sigmoidea* (NSIO) and *Gyrosigma attenuatum* (GYAT) (OM, H₂O₂ treated material). Two species co-occurring in rivers samples with similar abundance, valve length and motility. However, *Gyrosigma attenuatum* was never seen with epiphytes.

Chapter 10

Figure 10.1 Drawing adapted from O.F. Müller (1783, translated in [10.54]), who was the first to characterize *Bacillaria* colonies. Examples 1 through 8 show the various states of expansion and contraction (dynamic phenotypes) of colonies.

<u>Figure 10.2 *Bacillaria* close-up images of single</u> <u>cells using scanning electron microscopy (SEM). (a)</u> a whole valve seen from the inside. (b) close up of the same, middle section. The horizontal slit is the raphe. It lacks a central node. (c) Tip of the inside of a valve. (d) Middle section of a valve, exterior view. Note the raphe is a slit through the whole valve. (e): External view of the tip of a valve [10.33]. (Reprinted with permission of Amgueddfa Cymru, National Museum Wales.)

Figure 10.3 Demonstration of the image tracking procedure. (a) definition of tracked feature (white ellipse within a cell). (b) labeled (numbered) cells with relative measurements provided in red. The determined coordinates refer to a target in the middle of the template. The target can be moved and placed on the apex of the diatom being tracked. Then the coordinates of the apex are captured. This position is indicated in Figure 10.3a by a mark and a vertical line. Image scale: 38.36 µm per cm, or 0.325 µm per pixel. Scale bars 50 µm.

Figure 10.4 A diagram showing the five points on a sample cell (two ends, midpoint of the transverse line, and edges of the cell).

Figure 10.5 Point A is on the edge (vertical direction). The gradient direction is normal to the edge. Points B and C are in gradient directions. So, point A is checked with point B and C to see if it forms a local maximum. If so, it is considered for the next stage, otherwise, it is suppressed (set to zero). The result is a binary image with "thin edges."

<u>Figure 10.6 Diagram showing an example of</u> <u>hysteresis thresholding and the labeled edge</u> <u>relative to the "sure edge" threshold (V_{max}).</u> <u>Figure 10.7 An example of feature identification</u> <u>performance for the Watershed Segmentation</u> <u>algorithm (left, red boundary) and Canny Edge</u> <u>Detection algorithm (right, white boundary). Image</u> <u>scale: 38.36 µm per cm, or 0.325 µm per pixel.</u> <u>Scale bars 50 µm.</u>

Figure 10.8 An example of feature identification training (purple rectangles) for the deep learning approach on a single set of cells. Notice the resolution of the colony. An example of correct performance is shown in Figure 10.5. Image scale: 38.36 µm per cm, or 0.325 µm per pixel. Scale bar 50 µm.

<u>Figure 10.9 Rank-order analysis of bounding box</u> (cell) sizes (area) across the dataset. The area is measured in pixels squared. Image scale: 38.36 µm per cm, or 0.325 µm per pixel.

<u>Figure 10.10 Rank-order analysis of height (blue)</u> and width (red) of bounding boxes (cells) across the dataset. The area is measured in pixels squared. Image scale: 38.36 µm per cm, or 0.325 µm per pixel.

Figure 10.11 (Top) location of centroids in normalized coordinate space in selected dataset for static analysis. (Bottom) First two principal components from PCA analysis (PC1 represents horizontal position, while PC2 represents vertical position) of coordinates representing the *x*,*y* position for all four edges of each bounding box using the selected datasets for static analysis. Image scale: 38.36 µm per cm, or 0.325 µm per pixel. Figure 10.12 An example of feature identification optimization procedures implemented in DeepLabv3. GRAY: no optimization applied, RED: Optimization #1, BLUE: Optimization #2. Given an initial number of training frames (y-axis), the nonoptimized procedure (originally detected) will yield a certain number of boxes (x-axis). Applying various optimization procedures generally leads to a decreased number of boxes per frame for both low and high numbers of boxes.

Figure 10.13 Four examples of how the identified features map to two different images (a, c, e, g) of a *Bacillaria* colony. Points (b, d, f, h) represent the centroids for all bounding boxes identified in images a, c, e, and g, respectively. Image scale: 38.36 µm per cm, or 0.325 µm per pixel. Image scale: 38.36 µm per cm, or 0.325 µm per pixel. Scale bars 50 µm.

Figure 10.14 Three examples of how images of a Bacillaria colony are converted into a skeleton image. (Top Row) light microscopy images, (Middle Row) thin skeletonization based on a procedure implemented in GIMP, (Bottom Row) thick skeleton based on a procedure implemented in GIMP. Image scale: 38.36 μm per cm, or 0.325 μm per pixel. Image scale: 38.36 μm per cm, or 0.325 μm per pixel. Scale bars 50 μm.

Figure 10.15 Examples of relative movement of cells in a sample colony. (a) Comparisons between changes of position for cell #2 relative to cell #1 (red) and cell #3 relative to cell #2 (blue). (b) Comparisons between changes of position (red) and changes of velocity (blue) for cell #2 relative to cell #1. (c) a phase diagram of the data shown in b,