DIATOM GLIDING MOTILITY

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Figure 2.6 In Figures 2.6a and 2.6c the orientation angle corresponding to the diatoms whose trajectories were previously investigated are shown (diatoms #2 and #3 in a and c, respectively). In Figure 2.6e the orientation angle of another diatom (diatom #4) is shown. Orientation angle is the angle between the major axis of the diatom and the x-axis. Figures (b), (d), and (f) are overlay images of the diatom boundary for different frames of the data shown in (a), (c), and (e), respectively. White pixels show the stationary points in diatom boundary, while colors indicate change in boundary location. An extra Gaussian smoothing was included in the image processing algorithm for the diatom #4, whose orientation angle is shown in Figure 2.6e. Otherwise, it was not possible to extract orientation data for diatom #4.

Figure 2.7 (a) The diatom in Figure 2.6d is rotated around the arc center of the white pixel region in Figure 2.6d; (b) the consecutive intersections of the major axis of the diatoms during rotation are shown as yellow dots. In (b), while the green dots at the centerlines are close to one another, they would coincide exactly if the rotation were precisely around the center of the diatom.

Figure A1 Displacement histograms of the image of the stationary polystyrene particle situated in the micromachined channel; (a) and (b) show x and y displacement histograms, respectively. Dots represent microscopy measurement while continuous lines are the Gaussian fits to the measurements. Two Gaussian terms are used for fitting total displacement data. The particle was imaged at 821 fps.

Figure A2 Fast Fourier Transform (FFT) of the x and y coordinates of the centroid location for the recordings made with the stationary polystyrene particle and a diatom; (a) and (c) are the FFTs of the x centroid data and (b) and (d) are the FFTs of y centroid data of the polystyrene particle and diatom #3, respectively. Prior to the FFT the centroid location mean was subtracted from the data.

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Figure A4 Zoomed in views of the large displacements in the representative trajectories given in Figure A3. The plots (a), (b), and (c) show the centroids of diatoms #1, #2, and #3, respectively.

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μm). (SEM images downloaded from diatoms.org with permission [4.107] [4.120] [4.143].)

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.)

Figure 4.3 A summary of some factors and gradients affecting diatom adhesion and motility on intertidal sediment. While adhering to the surface, cells sense their wettability through the production of intracellular nitric oxide (NO), which in turn mediates EPS production. Pennates generally have a preference for hydrophobic surfaces for attachment. Shear force or water flow affects substrate attachment with weakly attached cells easily dislodged by stronger shear. Once attached, the gliding capability and reversal control are mediated by the availability of extracellular and intracellular Ca^{2+}, respectively. As light only penetrates a few mm on the sediment, the substrate is divided into the photic and aphotic zone. Cells undergo diel vertical migration to the photic zone to photosynthesize on the surface with a taxa-specific temporal rhythm, thus leading to micro-niche segregation. Nutrient concentrations vary with depth, and motile raphids take up dissolved silicate (dSi) and phosphate (dP) on the aphotic zone as there are higher concentrations of these mineral nutrients with depth. The aphotic

zone also provides a protected and stable environment for vegetative and sexual reproduction. (Figure modified and redrawn from Saburova *et al.* [4.132] and adapted from Fenchel [4.59]. Reprinted with permission.)

Figure 4.4 (a) Scanning electron microscopy (SEM) images of EPS or mucilage trails and strands from the motile raphid diatom, *Navicula* sp. (scale bar: 5 µm). EPS trails were observed to be either straight or curved. (SEM images from Chen *et al.* [4.26]. (Reprinted under CC-BY license); (b) Adhesion of *Nitzschia palea* towards surfaces with different surface wettabilities (scale bar: 15 µm). Cells adhered more to hydrophobic surfaces after 4 hours. Insets are duplicate experiments. (Optical microscopy images from Laviale *et al.* Reprinted (adapted) with permission from Laviale *et al.* Copyright 2019 American Chemical Society [4.91]).

Figure 4.5 (a) A representative cell trajectory of *Seminavis robusta* showing the run-reverse motility of cells from high cell density experiments. (Cell track replotted from data from Bondoc *et al.* [4.15]). (b) The circular run-reverse gliding technique observed in *Nitzschia communis* under isotropic environmental conditions. Cells form arc-like runs with constant speed, reverse, and continue the arc-like runs in the opposite clock direction. Filled squares signifies the starting point of the cells. (Figure from Gutiérrez-Medina *et al.* [4.68]. Reprinted with permission.)

Figure 4.6 Motility of the model raphid pennate *Seminavis robusta* towards different stimuli across its life cycle. For every mitotic division, cells undergo size reduction until they reach a sexual

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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 μmol/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 ± 2 μm/s and 57 ± 7 μm/s respectively, $P=0.003$). Graphs represent the mean values of response times ± 1 SE. For comparison, unirradiated Stauroneis cell response times were 155 ± 11 μm/s. Error bars represent ± 1 SE.

Figure 5.2 The effect of *Craticula cuspidata* or *Pinnularia viridis* presence on *Stauroneis phoenicenteron* response times. This graph displays the average direction change response times for *S. phoenicenteron* cells alone on slide chamber (Control). On a mixed slide chamber in a group by themselves (Isolated) or in a slide chamber in the presence of a high ratio of live *C. cuspidata* or *P. viridis* cells to *S. phoenicenteron* cells. Cells were then irradiated at their leading end with high irradiance 1s pulses of blue light, and observed to determine the time until they changed direction (response time). The presence of either *C. cuspidata* or *P. viridis* caused significant increases ($P < 0.03$) in cell response time from either control or isolated *S. phoenicenteron*. Error bars represent ± 1 SE.

Figure 5.3 The effect of culture medium on *Stauroneis phoenicenteron* response times. This graph displays the average direction change response times for *S. phoenicenteron* cells placed in the presence of culture medium in which either

S. phoenicenteron, Craticula cuspidata, or a mixture of *S. phoenicenteron* and *C. cuspidata* cells had been growing. Samples of the media were extracted, then briefly centrifuged in a microcentrifuge to remove any contaminating cells or cell debris. The *S. phoenicenteron* cells were then immersed into the desired medium, then isolated onto a slide chamber, allowed to incubate, then irradiated at their leading end with high irradiance 1s pulses of blue light, and observed to determine the time until they changed direction (response time). There was no significant difference between any of the treatment groups. Error bars represent ± 1 SE.

Figure 5.4 The effect of *Pinnularia viridis* on *Stauroneis phoenicenteron* response times. This graph displays the average direction change response times for *S. phoenicenteron* cells placed in two different areas of a cell chamber with differing proximities to *P. viridis*. This graph displays the average direction change response times for unirradiated *S. phoenicenteron* (Unirradiated), *S. phoenicenteron* irradiated in a slide chamber by themselves (Control), and *S. phoenicenteron* placed in a two-area slide chamber in which the entire chamber was subject to the same medium. Cells were irradiated at their leading end with high irradiance 1s pulses of blue light, and observed to determine the time until they changed direction (response time). The *S. phoenicenteron* in the two-area chamber were either in an area by themselves (Isolated), or in an area of the chamber where they were in close proximity to a large number of *P. viridis* (Live Pinn). In some trials the *S. phoenicenteron* were placed in

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.

Figure 5.5 Effect of _Craticula cuspidata_ fixation method on repression of _Stauroneis phoenicenteron_ direction change response time. _S. phoenicenteron_ cells isolated and washed from culture were incubated together with _C. cuspidata_ cells in a VALAP-spaced cell chamber where some _Stauroneis_ were in contact with _Craticula_ (grey bars) while other _Stauroneis_ (isolated, solid bars) had no _Craticula_ in near proximity. Stauroneis cells were then irradiated at their leading end with high irradiance flashes of blue light, and observed to determine the time until they changed direction (response time). _Stauroneis_ in a mixed region with live _Craticula_ had a significantly greater response time than _Stauroneis_ in an isolated area ($P = 0.01$). This increase in response time remained when _S. phoenicenteron_ were in the presence of _C. cuspidata_ cells killed in ethanol or acetone fixation alone, but no significant difference in response time was observed when _C. cuspidata_ were killed with a 50:50 acetone:ethanol mixture. The isolated _S. phoenicenteron_ in the chamber also showed increased response times in the treatments containing ethanol or acetone fixed _C. cuspidata_.

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 µmol 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) Upward migration starts before the beginning of 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 negative (under high light intensities). In the 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

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

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. areniverma* Furey, Lowe et Johansen – 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. Gregory – raphe (not visible) only present on valve 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 µm. Figure 8.8: white scale bar = 5 µm. (Figures 8.1, 8.3, 8.4 originally 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. areniverma* – raphe follows margin of apex from 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 µm/sec [8.31]) or weakly motile (2 to 4 µm/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]**).

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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."

Figure 10.6 Diagram showing an example of hysteresis thresholding and the labeled edge relative to the "sure edge" threshold (V_{max}).

Figure 10.7 An example of feature identification performance for the Watershed Segmentation algorithm (left, red boundary) and Canny Edge Detection algorithm (right, white boundary). Image scale: 38.36 μm per cm, or 0.325 μm per pixel. Scale bars 50 μm.

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.

Figure 10.9 Rank-order analysis of bounding box (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.

Figure 10.10 Rank-order analysis of height (blue) 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 non-optimized 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,