

Swimming dynamics and efficiency in chain diatom colonies

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Diatom chains are cohesive assemblies of unicellular microorganisms typically found in still and fresh water [1]. While some species are passively transported by ambient currents and settle due to the weight of their dense silica shells, others use various strategies to move or self-propel [2]. One species in particular, called *Bacillaria paxillifer*, forms colonies of stacked rectangular cells that slide along each other while remaining parallel (cf. Fig. 1 a). This unique collective motion leads to beautiful and nontrivial trajectories at the colony scale (cf. Fig. 1 b-c). Using a numerical method developed to simulate rigid bodies with kinematic constraints and hydrodynamic interactions in Stokes flows [3], we show that the swimming speed of such chains changes non-monotonically with the sliding delay between adjacent cells. The swimming efficiency, as a function of sliding delay between cells, shows multiple local maxima, which contrasts with the behavior typically observed in flagellate microorganisms [4]. In addition, the optimal cell aspect ratio for swimming found with our simulation matches those observed in real diatoms.

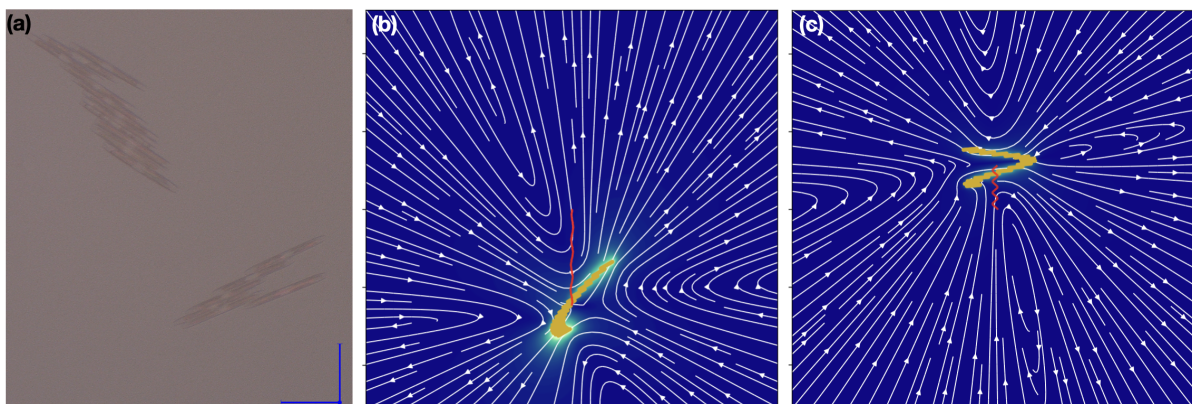


Figure 1. a) Microscope view of two *Bacillaria paxillifer* colonies. The scale bar is $50 \mu\text{m}$. b-c) Simulated flow fields around two colonies with different conformations. The red line represents the trajectory of the center of mass of the colony.

Références

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