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Abstract
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Oscillatory Flows Induced by Microorganisms Swimming in Two Dimensions

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We present the first time-resolved measurements of the oscillatory velocity field induced by swimming unicellular microorganisms. Confinement of the green alga C. reinhardtii in stabilized thin liquid films allows simultaneous tracking of cells and tracer particles. The measured velocity field reveals complex time-dependent flow structures, and scales inversely with distance. The instantaneous mechanical power generated by the cells is measured from the velocity fields and peaks at 15 fW. The dissipation per cycle is more than 4 times what steady swimming would require.

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Motile single cells use a variety of strategies for locomotion in low Reynolds number environments. Some methods produce roughly steady swimming (e.g., rotation of helical flagella in E. coli), while others produce oscillatory cell motion by beating flagella in a periodic manner [1]. Understanding the hydrodynamics of swimming microorganisms can give insight into complex biological processes such as predator-prey interactions and flagellar mechanics [2,3]. A variety of mathematical models have been proposed to describe interactions between swimming cells and their environment [1,4,5]. However, experimental studies of velocity fields are rare [6], and oscillatory near-field hydrodynamics have not been investigated experimentally despite their importance [7,8].

The flows induced by swimming organisms are responsible for biogenic mixing [9], and microorganisms in particular enhance the transport properties of suspensions [5,10]. Active particles and microorganisms also affect the rheological properties of suspensions, where the effective viscosity is increased in the case of “pullers” (e.g., Chlamydomonas reinhardtii) and decreased for “pushers” (e.g., E. coli) [11,12]. Hydrodynamic interactions leading to coherent motion have been studied for concentrated bacterial cells [13], but not for algal cells.

In many of these phenomena, the oscillatory swimming motion of organisms may be important. Furthermore, the concentration of swimmers may be sufficiently high that organisms are within several radii of one another, close enough that near-field effects should be considered. Understanding the resulting time-dependent flow fields is required for the interpretation of interactions between swimming cells and their environments.

We present the first time-resolved measurements of the flow field around oscillatory swimming microorganisms (C. reinhardtii). This is accomplished through simultaneous tracking of swimming cells and passive tracer particles within quasi-two-dimensional (quasi-2D) thin liquid films, where direct optical access and long observation times are achieved. These measurements reveal complex near-field flow structures that evolve throughout the beat cycle, and provide a richer picture than time-averaged flow fields and simplified models. The instantaneous mechanical power transferred by flagella to the fluid is measured via the viscous dissipation. These observations carry important implications for the interpretation and modeling of transport processes, locomotion, and flagellar mechanics.

C. reinhardtii is a single-celled swimming alga with a 7–10 μm diameter body [14]. It typifies many biflagellated microorganisms with two anterior 10–12 μm long flagella that beat at 50–60 Hz propelling the cell in an oscillatory manner at a mean speed of 100–200 μm/s (see video [15]). The flagella, actuated along their lengths by dynein motors, take on asymmetric conformations during the power and recovery strokes to overcome the time reversibility of low Reynolds number flows.

Wild-type C. reinhardtii (cc1690) is grown in minimal media (M1) on a light cycle (16 hr bright/8 hr dark) to ensure uniform size (radius R ≈ 3.5 μm) and motility [14]. An adjustable wire-frame device [12] is used to stretch a 2 μl droplet of the cell suspension into a square film with side length L = 6 mm and measured thickness h = 15 ± 2 μm. A trace amount of surfactant (Tween 20) stabilizes the film without harming the cells. Polystyrene microspheres (1 μm diam., Thermo Scientific), passivated with bovine serum albumin to prevent adhesion to flagella [16], are incorporated with the cells prior to stretching the film.

The device is mounted on an upright microscope (40 × 0.75 NA objective) where suspensions are observed under bright field illumination with red light (>610 nm) to prevent phototaxis [17]. A high-speed digital camera captures the motion over ≈3000 frames (50 or 500 fps). Cells and tracer particles are segmented from one another by size and tracked using a predictive algorithm [18].

The thin liquid film ensures that the algae are coincident with the focal plane of the objective and prevents distortion.
from out-of-focus particles. This allows detailed observation of the cells (mean speed $U_0 = 134 \, \mu m/s$) for up to about 8 s at 50 fps. Using high-speed imaging (500 fps), oscillations of the cell body, $U(t)$, become evident [Fig. 1 (inset)]. The peak forward velocity is 4 times the mean value and can be negative during the recovery stroke. The probability density function (PDF) of the beat frequency $f$ for 170 cell tracks is shown in Fig. 1. The oscillations have a fairly narrow distribution with a mean frequency $\bar{f} = 53 \pm 5$ Hz corresponding to a beat cycle period $T = 1/\bar{f} = 18.9$ ms.

By simultaneously tracking swimming cells and passive tracer particles (see video [15]), we measure the velocity field induced by the cells by translating and rotating the instantaneous tracer particle measurements to a common coordinate system based on cellular orientation. Swimmer trajectory segments with large curvature or irregular beat frequencies are excluded [19]. The beat-cycle averaged velocity field is measured by imaging the cells at a frame rate comparable to the cell beat frequency (50 fps), thus capturing the net motion over a cycle. The velocity field resulting from 560 cell tracks is shown in Fig. 2(a), where the swimmer motion is to the right and solid lines are instantaneous streamlines [20].

While the general shape of the flow field has some qualitative similarities to a force dipole (“stresslet”) approximation [1], several unexpected features are apparent [6]. The location of the hyperbolic stagnation point far away from the anterior of the cell is surprising, since it typically is thought to be located between the centers of drag (body) and thrust (flagella). Two strong vortices are visible lateral to the organism (associated with the flagella), while two weaker vortices appear far from the cell body beyond the separatrix of the hyperbolic point.

The fluid-air interfaces of the liquid film are nearly stress-free, which minimizes velocity gradients transverse to the film. This effect along with comparability of the film thickness and cell size ($h/2R \sim 2$) creates a quasi-2D environment [21], where we expect longer-range hydrodynamic disturbances compared to 3D flows. Fluid velocity magnitude normalized by the mean swimmer speed, $u/U_0$, where $u = |\mathbf{u}|$, is shown in Fig. 2(b) as a function of radial distance from the organism, $r/R$, in various directions. Toward the posterior, fluid speed scales nearly as $u \sim r^{-1}$ up to 20 cell radii away, similar to a force dipole in 2D [22], with slight deviations from $u \sim r^{-1}$ likely due to minor 3D effects. These results are consistent with similar measurements in 3D that show the expected $u \sim r^{-2}$ scaling [6]. In the lateral direction, the velocity magnitude passes through a local minimum when traversing the parabolic stagnation points (vortex), before recovering $u \sim r^{-1}$ scaling in the far field. Similarly, in front of the cell, fluid...
speed decreases rapidly near the hyperbolic stagnation point 7–8 radii from the swimmer.

The time-averaged velocity field in Fig. 2(a) shows the limitations of some current swimmer models, as also noted in [6], and raises several important questions. For example, how does the velocity field evolve throughout the oscillatory flagellar beat cycle? Using high-speed imaging (500 fps), we measure the instantaneous swimmer phase, and identify tracer particles at corresponding times in the beat cycle. Velocity fields (resulting from 170 cell tracks) are constructed from tracer velocities at each phase of the oscillation with resolution \( T = 15 \) s.

A time series of the velocity field during the beat cycle is shown in Fig. 3 (see video [15]). Insets show swimmer speed and phase (lower left) and approximate flagellar position (lower right). At the beginning of the power stroke [Fig. 3(a)], the velocity field is neatly divided into four symmetric quadrants with the hyperbolic stagnation point located slightly forward from the body, in line with conventional force dipole swimmer models [1]. As the flagella move toward the posterior and the power stroke peaks, the vortices lateral to the organism strengthen [Fig. 3(b)], then shift across the body to the anterior side [Fig. 3(c)–3(e)]. After the power stroke ends and the recovery stroke begins, the flagella extend out in front of the organism, and the cell velocity becomes negative. The flow shown in Fig. 3(f) is qualitatively reversed from Fig. 3(a), changing the sign of the dipole. The instantaneous flow field generated by an oscillatory swimmer such as \( C. \) reinhardtii is complex and highly time dependent.

In Stokes flows, all of the mechanical energy generated by a swimmer for locomotion is rapidly dissipated by the fluid. The power transferred to the fluid by the organism is calculated from the velocity field gradient through the viscous dissipation \( P = \int 2\mu (\nabla \cdot u)^2 \, dA \), where \( \mu \) is the fluid viscosity, \( \frac{1}{2} \left( \frac{\partial u}{\partial r} + \left( \frac{\partial u}{\partial x} \right)^T \right) \) is the rate of strain tensor, and \( dA \) is a differential area element of the film [15]. The instantaneous mechanical power output, \( P_{osc}(t) \), is calculated from the velocity fields [Fig. 3] throughout the beat cycle and shown in Fig. 4(a). The peak power output (\( \approx 15 \) fW) occurs during the power stroke and corresponds to the maximum instantaneous speed of the cell body. A secondary local maximum also occurs at the peak speed of the recovery stroke [see Fig. 4(b) (inset)]. Because of the
measurable of the velocity field throughout the flagellar beat cycle, revealing complex underlying flow structures that evolve in time. From these measurements, we demonstrate that the mechanical power dissipated by these swimming microorganisms is more than 4 times what the mean velocity field predicts. This has important consequences regarding the input of chemical energy required to power oscillatory swimming of flagellated cells.

We thank R.E. Goldstein, K. Drescher, E. Lauga, and M.D. Graham for helpful discussions, as well as B. Boyes for technical assistance. This work was supported by NSF Grant No. DMR-0803153.

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The mechanical power scales with the cell body speed \( U \) as \( P \sim U^2 \) with deviations at small \( U \) due to the finite accuracy of measuring tracer velocities [Fig. 4(b)]. This behavior might be expected for drag-based thrust at low Reynolds number, since the forces scale linearly with the body speed \( F \sim U \), and the power output of the organism is \( P \sim FU \sim U^2 \).

This work represents an important step toward understanding the locomotion of swimming microorganisms and collective behaviors for which hydrodynamics are responsible. Measurements of the time-averaged velocity field around \( C. reinhardtii \) in quasi-2D thin liquid films demonstrate the predicted \( u \sim r^{-1} \) scaling of the fluid velocity with detailed resolution of the near-field flow features. Also, high-speed imaging allows for phase-resolved measurements of the velocity field throughout the flagellar beat cycle, revealing complex underlying flow structures that evolve in time. From these measurements, we demonstrate that the mechanical power dissipated by these swimming microorganisms is more than 4 times what the mean velocity field predicts. This has important consequences regarding the input of chemical energy required to power oscillatory swimming of flagellated cells.

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