

Reverse engineering the euglenoid movement

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Euglenids exhibit an unconventional motility strategy amongst unicellular eukaryotes, consisting of large-amplitude highly concerted deformations of the entire body (euglenoid movement or metaboly). A plastic cell envelope called pellicle mediates these deformations. Unlike ciliary or flagellar motility, the biophysics of this mode is not well understood, including its efficiency and molecular machinery. We quantitatively examine video recordings of four euglenids executing such motions with statistical learning methods. This analysis reveals strokes of high uniformity in shape and pace. We then interpret the observations in the light of a theory for the pellicle kinematics, providing a precise understanding of the link between local actuation by pellicle shear and shape control. We systematically understand common observations, such as the helical conformations of the pellicle, and identify previously unnoticed features of metaboly. While two of our euglenids execute their stroke at constant body volume, the other two exhibit deviations of about 20% from their average volume, challenging current models of low Reynolds number locomotion. We find that the active pellicle shear deformations causing shape changes can reach 340%, and estimate the velocity of the molecular motors. Moreover, we find that metaboly accomplishes locomotion at hydrodynamic efficiencies comparable to those of ciliates and flagellates. Our results suggest new quantitative experiments, provide insight into the evolutionary history of euglenids, and suggest that the pellicle may serve as a model for engineered active surfaces with applications in microfluidics.

microswimmers | self-propulsion | stroke kinematics | active soft matter

Unicellular microorganisms have developed effective ways of locomotion in a fluid, overcoming fundamental physical constraints such as the time reversibility of low Reynolds number (Re) hydrodynamics (1). Amongst eukaryotes, most species swim beating cilia or flagella. Yet, through a long evolutionary history, some protists have developed unconventional functional strategies, accomplished by highly diverse subcellular structures (2). A notable example is the euglenoid movement, or metaboly, executed by some species of euglenids (3). This peculiar motility mode is characterized by elegantly concerted, large-amplitude distortions of the entire cell with frequencies of about $f \approx 0.1$ Hz (4). Euglenids have attracted the attention of scientists since the earliest days of microscopy, when van Leeuwenhoek referred to them in 1674 as microscopic motile “animalcules” that were green in the middle, which challenged the classification of organisms into animals and plants (5). More recently, metaboly has inspired models for artificial microswimmers (6), although even its locomotory function remains unclear. In contrast with flagellar or ciliary motion, the euglenoid movement has not undergone close biophysical scrutiny, and fundamental questions remain open, including a precise understanding of the actuation mechanism leading to the body shape changes, or its hydrodynamic efficiency. Because all euglenids move primarily with their flagella, it is difficult to discern the role of the body distortions in the observed motion.

Euglenids are abundant in a wide range of aquatic environments and, with typical sizes from tens to hundreds of micrometers, are easily observed by optical microscopy. They display a striking diversity in terms of morphology, nutrition (phagotrophs, phototrophs, osmotrophs), and motility, making of euglenids an excellent system to study evolution (7, 8). As other protists, they cannot resort to cell

specialization to accomplish vital functions. Instead, they encapsulate all the functionality required for survival in a single cell, with an elaborate machinery revealed by ultrastructural studies. Euglenids are enclosed by a striated surface with two opposing poles called pellicle. The pellicle, the major trait of morphological diversity in the group, is a cortical complex consisting of the plasma membrane, a set of interlocking proteinaceous strips, microtubules, and tubular cisternae of endoplasmic reticulum arranged along the strips (3). The ability of euglenids to undergo metaboly has been correlated with the morphology of the pellicle. Flexible pellicles possess a large number of articulated strips (a few tens) often wrapped helically around the cell, whereas rigid pellicles can have as few as four fused strips arranged either helically or along the meridians of the cell. Some euglenids exploit their body flexibility to merely round up, elongate, or gently bend. Here, we focus on the most extreme oscillatory motions displayed by some species.

Detailed observations of *Euglena fusca*, possessing a pellicle decorated by easily traceable particles, have shown that shape changes are accompanied by sliding between adjacent pellicle strips (9, 10); as the shape changes, the pellicle strips retain their length and width, but change their helicity. This mechanism is consistent with the commonly observed twisting motions around the long axis of the cell accompanying metaboly. These pellicle rearrangements are thought to be driven by the sliding of microtubule complexes relative to the proteinaceous strips and controlled by calcium sequestration in the tubular cisternae of endoplasmic reticulum. Although the molecular motor has not been identified, the active motions are thought to be fueled by ATP (11). Further experimental interrogation of metaboly with modern biophysical techniques is still lacking.

The euglenoid movement confronts us with far-reaching biological and biophysical questions. One such issue is the competitive advantage of metaboly. Although metaboly is an oddity in microscopic locomotion, it has persisted through natural selection coexisting within individuals with flagellar motility in a significant part of the phylogenetic tree of euglenids. Another intriguing aspect of metaboly is the operation of the distributed machine that elegantly and robustly executes the body motions, and that may inspire man-made active envelopes. As a first step, our goal here is to understand the kinematics and hydrodynamics of the euglenoid movement. For this purpose, we quantitatively analyze movies of metaboly recorded by others, representative of common observations by protistologists and physical scientists interested in motility (4). We examine these recordings with statistical learning methods. We then propose a model for the pellicle kinematics, which establishes a precise connection between distributed actuation and shape changes. Finally, we assess

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by numerical simulations the hydrodynamic features of the euglenoid movement.

Quantitative Observations

Data and Methods. We start from video recordings of four photosynthetic specimens, labelled euglenid #1 through #4, three of which are of the genus *Eutreptiella*, and the fourth belongs to either a *Euglena* or a *Eutreptiella* species. We assume that these euglenids remain essentially axisymmetric as they undergo metabolism, in agreement with their general description (12) and with detailed scrutiny of the movies studied here (see Fig. S1). Therefore, they can be described by their generating curve in the symmetry plane.

From the viewpoint of low Re locomotion, a swimming stroke is a closed path in shape space, which needs to be nonreciprocal to accomplish net motion. We next propose a method to identify and parametrize such a path for further quantitative examination (see Fig. 1). In recent years, statistical learning methods have been increasingly used to understand stereotyped animal behavior; see, e.g., related studies on the motility of *Caenorhabditis elegans* (13, 14). Because the stroke is expected to be a nonlinear manifold of dimension one, here we resort to a nonlinear dimensionality reduction technique called Isomap (15). This method identifies nonlinear correlations of high-dimensional data points (here snapshots of the cells describing their shape) by embedding them in low dimensions in a geometry-preserving manner.

First (Fig. 1A), we gather the collection of shapes adopted by each euglenid by segmenting and aligning the video frames. We represent numerically the shape of each frame by fitting a spline curve to the boundary of the aligned segmentation. Second (Fig. 1B), we identify the geometric structure underlying the set of shapes (*i*) with Isomap. We embed the shapes in 2D (*ii*) to examine the nature of the path described by the euglenids in shape space. Because the shape ensemble is essentially a one-dimensional loop, we further embed the shapes in a periodic 1D segment labelled by τ (*iii*), where the frames are ordered by shape

proximity. With the 1D embedding, we parametrize the geometric stroke smoothly (*iv*). This procedure filters some of the noise while closely following the data; see Fig. S2A and B. The resolution of the result is limited by optical artifacts and images out of focus, which bias the segmentation, and possibly by slight departures from axisymmetry. Yet, the quality of the numerical stroke is sufficient for quantitative examination (see Movie S1).

Because our method is based on geometric similarity, τ measures arc length in shape space rather than physical time. The hydrodynamic efficiency depends on the relative rate at which different parts of the stroke are executed (the net displacement after one cycle does not). To inspect the pace along the stroke, we obtain from the data the relation between τ and normalized physical time t .

Results. We find that nonlinear dimensionality reduction reveals vividly the signature of low Re swimming for each of the four euglenids, with well-defined nonreciprocal paths in shape space over the many cycles captured in the movies (Fig. 2A and Fig. S3A). Different specimens perform geometrically distinct strokes (Fig. S2C), and, for a given specimen, each stroke is executed multiple times with the same pace; i.e., the relation between τ and t is nearly the same over several strokes (Fig. 2B and Fig. S3D). With the numerical stroke at hand, we first examine elementary features such as the surface area $S(t)$ and the volume $V(t)$ enclosed by the pellicle. We express volume in nondimensional terms as the reduced volume $v = 6\sqrt{\pi}V/S^{3/2}$, the ratio between the cell volume and the volume of a sphere with the same surface area. Thus, v ranges between 0 to 1. We find that the surface area of the four euglenids remains nearly constant along the stroke, with deviations below 5% (Fig. S3C). This observation is consistent with the hypothesis that pellicle deformation is mediated by relative sliding of pellicle strips, which retain their length and width (9). Such a deformation is called simple shear in continuum mechanics, and preserves area locally. The small area deviations may be due to the systematic errors in

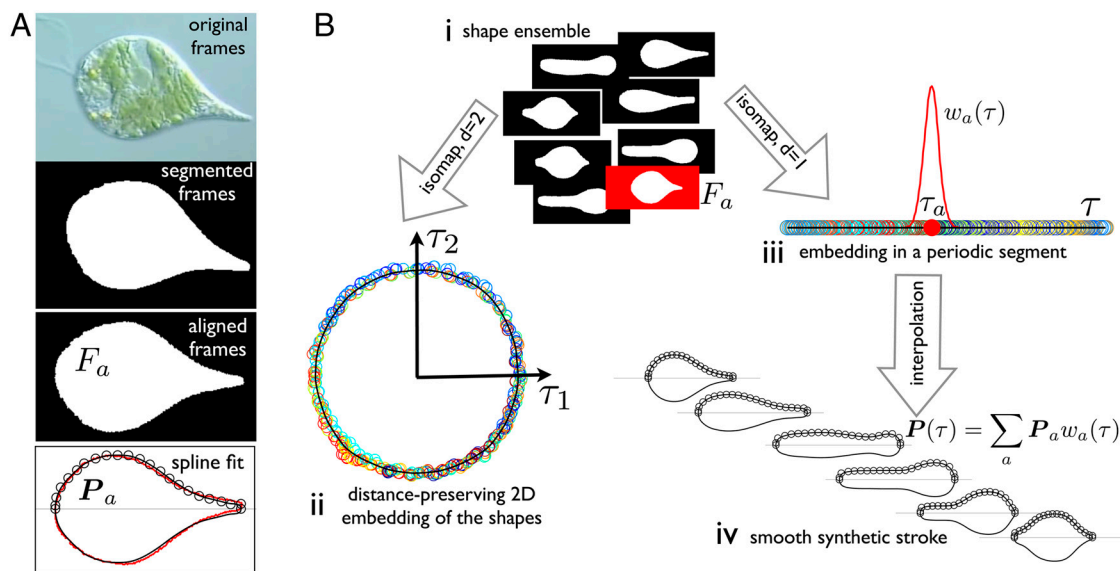


Fig. 1. Quantitative analysis of the movies: method. (A) The frames are segmented and aligned to obtain images of F_a containing information about the shape alone, devoid of translation, rotation, or textures. A B-spline curve, given by its control polygon P_a (black circles), is fitted to the boundary of F_a and is a generating curve of the axisymmetric representation of the pellicle. Original frame image courtesy of Richard E. Triemer. (B) The segmented and aligned frames (*i*), representative of the shapes adopted by the cell, are embedded in low dimensions by a nonlinear dimensionality reduction technique called Isomap (15). The algorithm maps each frame to a low-dimensional point (circles in *ii* and *iii*, color labeling chronological order), so that the intrinsic distance between frames (shape dissimilarity) is preserved as much as possible by the low-dimensional representation. The embedding in the plane (τ_1, τ_2) (*ii*), shows that the stroke is a closed nonreciprocal path in shape space. Consequently, it can be most compactly described by embedding the frames in a periodic 1D segment (*iii*), from which we parametrize the stroke as a function of τ by interpolation (*iv*) with smooth basis functions $w_a(\tau)$. At any given τ , the synthetic stroke is a weighted average of the curves fitting video frames whose 1D embedding is in the vicinity of τ . The parameter τ is not proportional to physical time, ignored by the manifold learning algorithm, but rather to arc length in shape space. See *SI Text*, section *Video Processing and Parametrization of the Stroke*, for technical details.

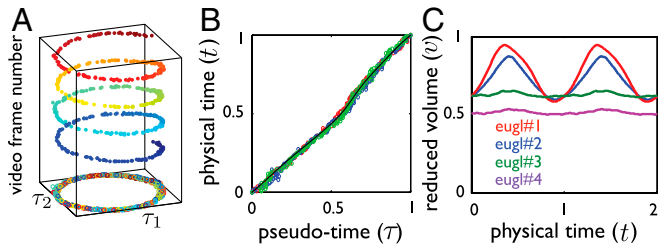


Fig. 2. Quantitative analysis of the movies: results. (A) Two-dimensional embedding of a stroke plotted against video frame number, showing the shape and pace uniformity of the stroke over several realizations. Isomap identifies similar shapes from different realizations (here about four cycles) and yields a single geometric stroke. (B) Pseudo-time parameter τ against normalized physical time t during three full strokes represented in different colors, showing that path in shape space is traveled with a well-defined pace (here, euglenid #3). The black line shows the fit used to reparameterize time. (C) Reduced volume, v , as a function of time during two strokes. We distinguish two distinct motility styles: a volume-changing style for euglenids #1 and #2, and a volume-preserving style for euglenids #3 and #4. See Fig. S3 for the full data.

the data acquisition mentioned earlier, or to a slight area deformability of the pellicle. Strikingly, the body volume behaves very differently for euglenids #1 and #2 and for euglenids #3 and #4; see Fig. 2C. The former exhibit reduced volume deviations of around 20% and average reduced volumes of around 0.75, while the latter show essentially constant reduced volume of smaller magnitude. The volume-changing euglenids necessarily flow significant amounts of fluid in and out of their body quite rapidly during the stroke. The contracting vacuole of euglenids, connected to an invagination called reservoir located at one end of the cell, is known to regulate water, although generally at longer time scales. The motion of the internal organelles during metaboly, clearly seen in the movies, suggests that the volume variations are accommodated by a distributed fluid permeation rather than by localized jets.

Pellicle Kinematics

Theory. The information about the kinematics of the pellicle obtained above is incomplete, because shape alone does not report on the motion of material particles; specifically, no quantitative data is available about the tangential motions within the symmetry plane and in the azimuthal direction. A careful inspection of some of the movies reveals that, while changing shape, the cell body twists nonuniformly around the symmetry axis. We next describe a theory for the pellicle kinematics, which allows us to recover the full velocity field of the pellicle and also provides a precise understanding of the concerted actuation leading to metaboly.

The key assumption in this theory, as discussed earlier, is that the pellicle deforms by simple shear γ along the strips (Fig. 3A). Consequently, the pellicle deformation is locally area preserving, and we can label the material parallels of the axisymmetric pellicle with a body coordinate λ , measuring the fraction of total area delimited by a given parallel (see Fig. S4). By time differentiation of the generating curve at fixed λ , we can compute the pellicle velocity within the symmetry plane.

We next recover the azimuthal motions and the pellicle shear. Consider an arbitrary but fixed conformation of the pellicle, given by its generating curve $[r_0(\lambda)$ and $z_0(\lambda)]$ and by the orientation of the pellicle strips, which could be obtained by *in vivo* imaging. This orientation is given mathematically by the vector field \mathbf{s}_0 along the strips, or by the angle the strips form with the surface parallels $\alpha_0(\lambda)$ (Fig. 3A). We take this conformation as a reference. The strain of the pellicle surface relative to the reference configuration can be characterized at each point by the Cauchy–Green deformation tensor \mathbf{C} (16), which by axisymmetry only depends on the body coordinate λ . From the actuation viewpoint,

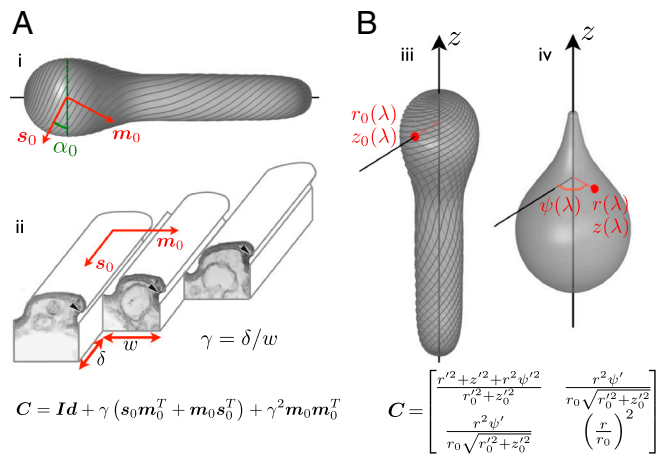


Fig. 3. Pellicle kinematics: theory. (A) Surface strain (the 2×2 matrix \mathbf{C}) is derived assuming simple shear along the strips, γ , acting on a reference pellicle. The reference pellicle conformation (i,iii) is defined by a shape and a pellicle conformation, given by the tangent vector fields \mathbf{s}_0 and \mathbf{m}_0 , along the strips and perpendicular to them, or by the angle α_0 between the pellicle strips and the surface parallels. (ii) Ultrastructure of the pellicle (transmission electron micrograph from ref. 7, Copyright 2001 The Society for the Study of Evolution), and depiction of the sliding between adjacent strips causing the pellicle shear γ . (B) Surface strain, \mathbf{C} , is now derived by comparing a reference configuration, (iii), given by $r_0(\lambda)$ and $z_0(\lambda)$, where λ is the body coordinate, and a deformed configuration (iv), given by $r(\lambda)$, $z(\lambda)$ and the azimuthal displacement relative to a fixed direction $\psi(\lambda)$. The red point denotes a material particle attached to the pellicle, and (\prime) denotes differentiation with respect to λ . Matching the microstructural (A) and the shape-derived (B) expressions for the strain, we find equations relating pellicle shear, pellicle orientation, shape changes, and azimuthal motions.

we obtain a first expression for \mathbf{C} , depending on the reference pellicle orientation and the pellicle shear; Fig. 3A. On the other hand, comparing the reference state with a new configuration, given by $r(\lambda)$, $z(\lambda)$, and the azimuthal displacement $\psi(\lambda)$, we obtain a second expression for \mathbf{C} ; Fig. 3B. Matching the two expressions, we find the relation between the global conformation (including shape and pellicle helicity) and the local actuation; see *SI Text*, section *Pellicle Kinematics*, for details and Fig. S5 for a validation.

Here, we have obtained from the movies a trajectory of shapes, $r(\lambda, t)$ and $z(\lambda, t)$. The theory then provides expressions for the associated azimuthal displacement $\psi(\lambda, t)$ and the pellicle shear $\gamma(\lambda, t)$, from which in turn we evaluate the full velocity field and the conformation of the deformed pellicle strips along the stroke. The theory can also be used to evaluate the deformed pellicle shape and helicity resulting from a prescribed pellicle shear $\gamma(\lambda, t)$ acting on a reference conformation, although we do not explore here the design of strokes.

The theory imposes a constraint on the shapes achievable from a given reference pellicle; namely, $r/r_0 \geq |\sin \alpha_0|$. Consider a reference pellicle with strips parallel to the meridians; i.e., $\alpha_0(\lambda) = \pi/2$. Then the shape at instant t is accessible from the reference pellicle if $r(\lambda, t) \geq r_0(\lambda)$, consistent with the fact that thin parts of flexible pellicles display strips with almost no helicity. We can access all the shapes of the stroke by defining $r_0(\lambda)$ as the smallest radius adopted by the material parallel λ , and integrate $z_0(\lambda)$ from local area preservation. This procedure defines an optimal reference pellicle for each euglenid, which achieves the observed shapes with minimal pellicle shear at every point; we adopt this reference pellicle in the absence of detailed imaging.

Results. Fig. 4 summarizes the application of our theory to the four euglenids, highlighting the link between distributed actuation, pellicle conformation, and shape. The reconstructed pellicle shears $\gamma(\lambda, t)$ of the four euglenids exhibit distinct patterns, but

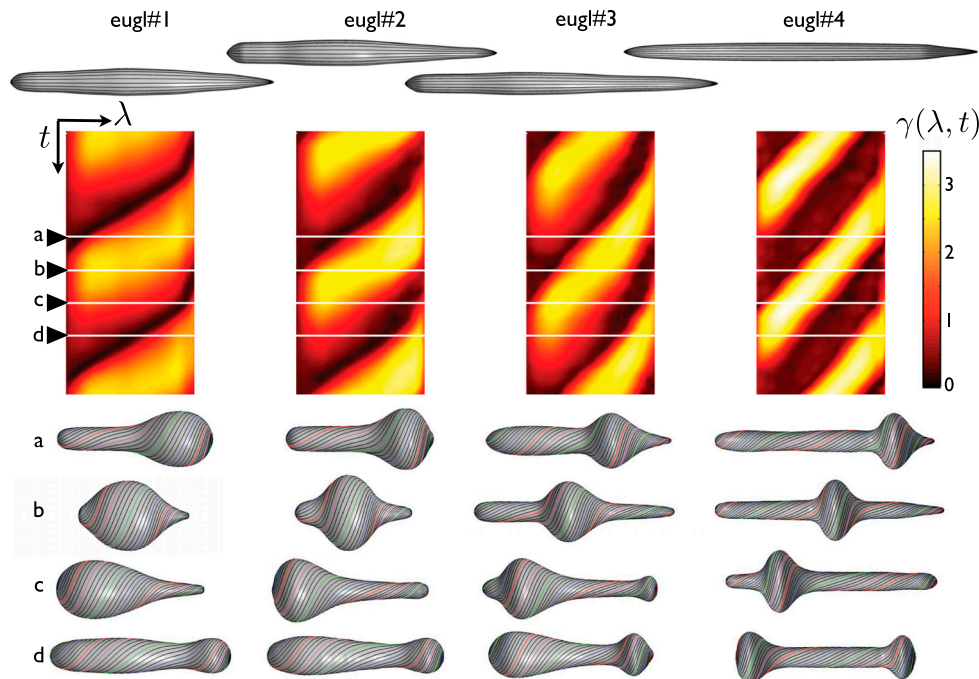


Fig. 4. Stroke kinematics: relationship between the local actuation and shape changes. Reference conformation of the pellicle for each euglenid (top), exhibiting very small reduced volumes. Map of the pellicle shear recovered from the observations as a function of time and body coordinate $\gamma(\lambda, t)$ (center), shown for two full synthetic strokes. The pellicle shear distributions acting on the reference configurations produce the shapes and pellicle conformations shown below.

also common features. A bump of high pellicle shear near the head of the cell (snapshot *a*) travels toward the tail (snapshots *b* and *c*). This traveling shear bump reflects in the shape as a bulge that moves from head to tail, reminiscent of peristaltic movements. When it reaches the tail, the shear bump decays while a new one appears at the head (snapshot *d*) before the cycle closes. Euglenid #1 exhibits the bluntest pattern and shape trajectory, and only reaches pellicle shears of 250%. In contrast, euglenid #4 executes a remarkably sharp and regular pattern of pellicle shears, reaching 340%, with a corresponding well-defined bulge that slides along a cylindrical body of nearly constant radius. Some species of osmotrophic genera, such as *Astasia* and *Distigma*, exhibit a similar swimming style to the photosynthetic euglenid #4. With an estimated pellicle width of $w \approx 250$ nm, we predict a sliding displacement between adjacent strips of up to $\delta \approx 850$ nm for this euglenid. By differentiating the map of $\gamma(\lambda, t)$ in Fig. 4, we find that $\dot{\gamma}$ reaches $20f$ Hz, leading to a maximum sliding velocity of $\dot{\delta} = 20fw \approx 500$ nm/s, consistent with the velocities of molecular motors along microtubules (17, 18). The similarity between our pellicle reconstructions and observations of flexible pellicles, with a clear correlation between radius and pellicle orientation, is remarkable.

Hydrodynamics of the Stroke

Combining the movie processing and the theory for the pellicle kinematics, we compute the surface velocity during the strokes, and analyze the hydrodynamics by placing the model euglenids in a Newtonian fluid. With a typical size of 50 μm and a stroke frequency $f \approx 0.1$ Hz, we estimate $\text{Re} \approx 10^{-4}$. We solve the Stokes equations with a boundary integral method, and the linear velocity along z and the azimuthal spin of the cell are calculated from the self-propulsion condition (19); see *SI Text*, section *Hydrodynamics*, for details. We assume that volume changes are accommodated by a uniform normal permeation velocity. The fluid velocity at the pellicle is then the surface velocity plus the permeation velocity, which is not very significant even for the volume changing euglenids #1 and #2 (see Fig. S6).

Table 1 summarizes the kinematic features and swimming performance of the four euglenids (see Table S1 for additional data). We evaluate the swimming performance with the linear displacement in one stroke U , measured in units of body length $2R = \sqrt{S/\pi}$, and with Lighthill's efficiency Eff_L . This conventional notion of efficiency is defined as the ratio between the power needed to drag the swimmer at the average velocity $W = U/T$ and the average power exerted by the swimmer on the surrounding fluid during the stroke, P^{out} . Because our swimmers change shape continuously, we consider $\text{Eff}_L^{\text{out}} = 6\pi\eta RW^2/P^{\text{out}}$, where η is the fluid viscosity. We find that all euglenids move by a few tenths of a body length in one stroke, and exhibit efficiencies in the order of 1%. With the typical frequencies of metaboly, the swimming speeds are very low, $W \approx 1\text{--}2$ $\mu\text{m/s}$, between one and two orders of magnitude slower than during flagellar locomotion. Remarkably, the hydrodynamic efficiency is comparable to that estimated for ciliates and flagellates (20). We find that euglenid #1 is the slowest and least efficient swimmer, followed by #2 and #3, with comparable performance, and finally #4 is the best, with a displacement of 40% of the body length per stroke and $\text{Eff}_L^{\text{out}} = 2.0\%$. The table suggests that swimming performance is correlated with maximum pellicle shear. The best swimmer is a volume-preserving specimen with the lowest reduced volume,

Table 1. Summary of the kinematic features and swimming performance of the four strokes

Euglenid	$v_{\text{min}} v v_{\text{max}}$	γ_{max}	U	$\text{Eff}_L^{\text{out}}$, %	$\text{Eff}_L^{\text{out+inn}}$, %
#1	0.58 0.76 0.95	2.5	0.16	0.71	0.24
#2	0.60 0.72 0.87	3.0	0.27	1.6	0.57
#3	0.61 0.63 0.65	3.0	0.25	1.6	0.48
#4	0.50 0.51 0.53	3.4	0.40	2.0	0.76

U denotes the net displacement in one stroke, in units of body length $2R = \sqrt{S/\pi}$ (the diameter of a sphere with the surface area of the swimmer). Eff_L denotes the Lighthill efficiency, accounting for either the outer, or both the outer and inner fluids. \bar{v} denotes the average reduced volume during the stroke.

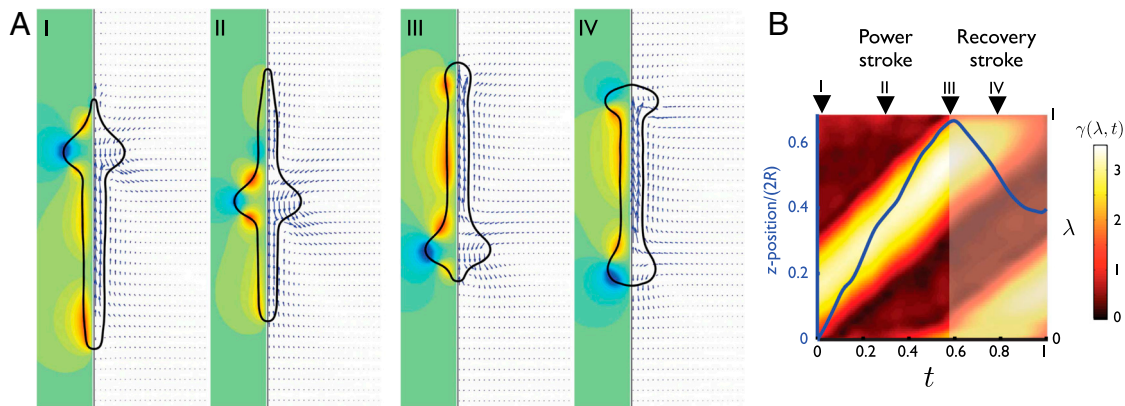


Fig. 5. Stroke hydrodynamics of euglenid #4. (A) Flow pattern around the pellicle at selected instants along the stroke. The fluid velocity field in the symmetry plane is indicated by blue arrows, and the azimuthal component is shown with isocontours, where green is zero, blue negative, and red positive. (B) Relationship between actuation and forward motion, where the vertical axis is the body coordinate λ for the color map of $\gamma(\lambda, t)$ and z position (average between the head and the tail) in units of body length for the curve in blue. Both the actuation pattern and the forward motion clearly delineate a power and a recovery phase in the stroke.

although there is no clear trend in this respect. See [Movie S2](#) for a visual depiction of the stroke kinematics and the resulting linear and angular motion for the four euglenids.

The hydrodynamics in the symmetry plane are completely decoupled from the azimuthal hydrodynamics, by the linearity and symmetry of the equations. Thus, the azimuthal energy dissipation in the fluid is a toll on the swimmer efficiency resulting from the mechanism used by the pellicle to change shape. It accounts for about 20% of the outer fluid dissipation (see [Table S1](#)). Lighthill's efficiency disregards the internal energy losses, passive or metabolic, known to be dominant in other microorganisms (21). Without attempting to evaluate these, we now view the swimmer as an isolated active surface embedded in a fluid, and consider the inner fluid dissipation as well. Remarkably, the inner dissipation is very large, about three times the outer counterpart, resulting in efficiencies about three times smaller.

Fig. 5 illustrates the fluid flow patterns around euglenid #4 and correlates these with the forward displacement during the stroke. Similar results are obtained for the other euglenids. We can clearly distinguish between a power and a recovery phase. During the power stroke, a bulge slides down the slender body, producing a characteristic flow pattern that propels the swimmer forward monotonically. During the recovery stroke, the swimmer moves backward and the bulge at the tail disappears at the expense of a growing bulge at the head. This requires a considerable forward rearrangement of fluid and pellicle mass, clearly noticeable in the flow pattern. The large internal streaming during this phase agrees with the motion of the organelles observed in movies of metabolic euglenids. While the energy dissipation in the outer fluid is quite uniformly distributed along the stroke for all four euglenids, the average dissipation power of the inner fluid is three to four times larger during the recovery phase than during the power phase. Because optimal strokes with respect to a dissipation metric exhibit uniform dissipation rate (22), this shows that the observed strokes do not minimize internal dissipation, despite that being the largest contribution. Fig. 5B summarizes the correspondence between the distributed actuation mechanism and the resulting forward motion during the power and recovery phases.

Discussion

Our study suggests many new questions. A number of quantitative experiments could test and complement our theory. A detailed kinematical picture of the pellicle during metaboly can be obtained *in vivo* by light microscopy under ultraviolet illumination (23), while particle image velocimetry (24) can resolve the

hydrodynamics and shed light on the inflow and outflow management of volume-changing euglenids. Our observation about the volume variations challenges the predominant conceptual models of swimming at low Re (6, 25), although swimmers permeating fluid across a fixed body shape have been analyzed (26). It is not clear if volume changes are passive, as a consequence of the pellicle deformation, or if these two phenomena are regulated in concert. Experiments on extracted cell models suggest that the euglenoid movement is mediated by a spatiotemporally non-uniform calcium sequestration, which triggers the pellicle activity (11). This mechanism could osmotically drive water permeation, which may enhance the exchange of nutrients and wastes. It is known that photosynthetic euglenids can acquire nutrients by osmotrophy.

Having established that metaboly is a competent motility mode, the evolutionary benefit of such a large biological investment remains unclear, when coexisting with flagella, an alternative strategy of demonstrated success. Current evolutionary theories suggest that the earliest ancestors, rigid and feeding on small bacteria, developed a flexible pellicle to engulf larger eukaryotic prey. Photosynthetic species evolved from phagotrophs after engulfing green algae, and some of these lost the pellicle flexibility by strip fusing (8). Presumably, the pellicle flexibility may have found a secondary utility, besides mediating in eukaryovory, and may have persisted because of this in photosynthetic euglenids such as those studied here, as well as in some osmotrophs. The benefit of metaboly may be related to the stirring of the cytosol or the surrounding medium, or to locomotion. Interestingly, many euglenids live in interstitial, soil-rich, or eutrophic water, suggesting that metaboly may be particularly well-suited to move in "difficult" conditions such as confined environments, or in media of complex rheology, known to affect flagellar efficiency (20, 27). This hypothesis is supported by a marine photosynthetic species, lacking emergent flagella but exhibiting strong metaboly, known to migrate vertically through the sand of the benthic zone in diurnal cycles (28). Thus, the heterogeneity of the biophysical traits of metaboly could complement the morphological and molecular data used in current phylogenetic studies of euglenids, and provide new clues about their evolutionary history.

Besides complementing experimental observations to obtain a complete kinematical picture of metaboly, the theory for the pellicle deformation provides the background for further biophysical studies. It is possible to optimize the pellicle shear to achieve maximal efficiency, and investigate the role of volume constraints or different dissipation metrics. Similar studies have brought new insight for cilia (29), flagella (30, 31), and model swimmers

undergoing full body distortions (22, 32, 33). Furthermore, the theory can be used to compute the resisting forces, arising from the internal and external friction and from the pellicle bending elasticity, that need to be overcome by metabolic activity.

The pellicle suggests itself as an appealing design concept for an active material (34). The local actuation of the pellicle produces a local strain incompatible with the current geometry, which is only realizable by further curving and producing shape changes (35). Similar concepts of material-as-machine relying on martensitic materials have been put forth (36, 37), and actually operate in biological organisms such as the tail sheath of bacteriophage T4 (38). Viewed as an active surface, the pellicle exhibits an interesting balance between shape flexibility and controllability, which explains why metaboly is such an elegantly concerted motion as compared, e.g., with the amoeboid movement. The pellicle has a single degree of freedom available for the local surface deformation, the pellicle shear, which nevertheless is sufficient to realize a wide family of shapes. Observations show that flexible euglenids can perform bending and torsional maneuvers of remarkable agility, including sharp turns. A general theory considering nonuniform pellicle shears in the azimuthal direction may delimitate the possibilities of this shape actuation concept, which may find applications in artificial microswimmers or peristaltic micropumps.

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Conclusions

We have provided a detailed biophysical analysis of the euglenoid movement, a widely appreciated but poorly understood motility mode of euglenids. We have developed a statistical learning methodology to analyze strokes and applied it to four movies recorded by others. We have established the shape and pace uniformity of the strokes, and have parametrized the motions in a computer model. Some euglenids exhibit very large variations of their body volume during the stroke, while others keep their volume constant. A kinematical theory, assuming that the cell surface deforms by simple shear along pellicle strips, provides a precise link between distributed actuation and shape changes, explains common observations about the pellicle helicity, and provides the input for hydrodynamics. Metaboly is found to be a slow but efficient motility mode in water, which enhances stirring of the cytosol and the surrounding fluid, and may be advantageous in granular or complex media.

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Supporting Information

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SI Text

Video Processing and Parametrization of the Stroke. The procedure outlined here assumes that the specimens adopt axisymmetric shapes, which agrees well with the detailed inspection of the videos; see Fig. S1. Additionally, the movies suggest that the axis of symmetry of the euglenid remains roughly parallel to the focal plane. Fully automated segmentation procedures require video inputs of high quality. The movies processed here include frames slightly out of focus, are nonuniformly illuminated, and have featured backgrounds. Consequently, our segmentation strongly relies on manual operations. We first enhance the edges of the cell by blurring the surroundings with Adobe Photoshop. This procedure also detaches the flagella from the cell body and allows us to complete small body parts lying outside of the field of view. We then segment the resulting images with Matlab and ImageJ, and keep the best result. All other computations in the paper are carried out with in-house Matlab codes. The segmented frames are aligned (translated and rotated) with Principal Component Analysis (PCA). The boundary pixels are tagged, and split in two by the symmetry axis detected by PCA. One of the halves is reflected to yield a larger set of points sampling the axisymmetric surface. We then fit to these points a B-spline curve (1) for each frame, expressed as

$$c_a(u) = \sum_{I=1}^N B_I(u) \{r_I, z_I\}_a,$$

where $B_I(u)$ are the B-spline basis functions referred to a reference interval $u \in [0, 1]$ and $\{r_I, z_I\}_a$ are cylindrical coordinates of the control points associated with the a th frame. We use cubic B-splines with tangents along the r direction at the end points. The geometry can be adequately described with 20 knot intervals ($N = 23$ control points). However, for accurate resolution of the fluid flow, we consider 30 knot intervals (see section *Hydrodynamics*). The video analysis is carefully checked frame by frame by visual inspection, and the procedure is iterated if needed. Movie S1 illustrates that this method provides a low-resolution, yet quantitative, representation of the strokes.

We find low-dimensional embeddings of the frames with Iso-map (2), using 30 neighbors to approximate geodesic distances. The results are quite insensitive to this parameter in the range 20 to 50. This algorithm embeds each frame into a point τ_a in a periodic segment scaled to $[0, 1]$. This segment represents a geometric pseudo-time. We use this embedding to parameterize in pseudo-time the generating curve of the euglenid surface during the stroke as

$$c(u, \tau) = \sum_{a=1}^n w_a(\tau) c_a(u) = \sum_{I=1}^N B_I(u) \sum_{a=1}^n w_a(\tau) \{r_I, z_I\}_a,$$

where n is the number of frames, and $w_a(\tau)$ are smooth meshfree maximum-entropy basis functions (3). See Fig. S2 for a discussion on the selection of their support size. Fig. S3A shows the two-dimensional embeddings of the four euglenids, where each frame is mapped to a point $\tau_a \in \mathbb{R}^2$, together with the reconstruction of the stroke path given by

$$\tau(\tau) = \sum_{a=1}^n w_a(\tau) \tau_a.$$

To estimate the relation between τ and physical time, we use the fact that the movies are shot at constant frame-rate Δt . For each frame associated to the time instant $(a-1)\Delta t$, we find the corresponding pseudo-instant by minimizing $|\tau(\tau) - \tau_a|^2$ with respect to τ . We then identify several distinct strokes and plot the resulting pairs of pseudo-time and physical time, normalized in the interval $[0, 1]$ (see Fig. S3D). We fit the one-to-one relation $t = g(\tau)$ with a spline shown in black, and reparameterize the reconstructed stroke in time as $\tilde{c}(u, t) = c(u, g^{-1}(t))$, which, in an abuse of notation, we simply write as $c(u, t)$.

Pellicle Kinematics. Axisymmetric surface and generating curve. Let $c(u, t) = \{r(u, t), z(u, t)\}$ denote the generating curve of the axisymmetric surface Γ acquired from the data as detailed above. We denote its speed by $a = \sqrt{r'^2 + z'^2}$, where $()'$ denotes partial differentiation with respect to the space coordinate. A unit normal is given by $\mathbf{n} = 1/a \{-z', r'\}$. We orient the surface so that \mathbf{n} is an outer normal. The element of area of the axisymmetric surface can be expressed as $dS = 2\pi r du$. The surface area and volume of the axisymmetric surface can be computed as

$$S = 2\pi \int_0^1 ar du,$$

$$V = \frac{1}{3} \int_{\Gamma} \mathbf{x} \cdot \mathbf{n} dS = \frac{2\pi}{3} \int_0^1 (-z'r + r'z)r du.$$

If the motion of the pellicle is locally area preserving with constant total area S , then the parameter

$$\lambda = f_t(u) = \frac{1}{S} \int_0^u 2\pi r(v, t) a(v, t) dv \quad [\text{S1}]$$

measuring the fraction of the total area delimited by u , labels material parallels of the pellicle surface during the motion. See Fig. S4 for an illustration. By reparameterizing the generating curves with this body coordinate, we obtain a Lagrangian description (i.e., tracking material particles) of the pellicle up to azimuthal motions, $\tilde{c}(\lambda, t) = \{\hat{r}(\lambda, t), \hat{z}(\lambda, t)\} = c(f_t^{-1}(\lambda), t)$. We assume from this point on that the surface is parametrized by λ , and, with an abuse of notation, we denote the reparameterized curve by $c(\lambda, t) = \{r(\lambda, t), z(\lambda, t)\}$.

Deformation of the surface. The reference pellicle configuration is given by

$$\mathbf{x}_0(\lambda, \theta) = \{r_0(\lambda) \cos \theta, r_0(\lambda) \sin \theta, z_0(\lambda)\}, \quad [\text{S2}]$$

where the body coordinate $\lambda \in [0, 1]$, and the azimuthal angle $\theta \in [0, 2\pi]$ label material particles. The full prescription of the pellicle Lagrangian motion along the stroke is given by the known functions $r(\lambda, t)$ and $z(\lambda, t)$, together with the unknown azimuthal displacement $\psi(\lambda, t)$

$$\mathbf{x}(\lambda, \theta, t) = \left\{ \begin{array}{l} r(\lambda, t) \cos[\theta + \psi(\lambda, t)] \\ r(\lambda, t) \sin[\theta + \psi(\lambda, t)] \\ z(\lambda, t) \end{array} \right\}. \quad [\text{S3}]$$

The natural basis vectors of the tangent plane to the deformed configuration can be expressed in Cartesian coordinates as (4)

$$\mathbf{x}_\lambda = \begin{Bmatrix} r' \cos(\theta + \psi) - r\psi' \sin(\theta + \psi) \\ r' \sin(\theta + \psi) + r\psi' \cos(\theta + \psi) \\ z' \end{Bmatrix},$$

$$\mathbf{x}_\theta = \begin{Bmatrix} -r \sin(\theta + \psi) \\ r \cos(\theta + \psi) \\ 0 \end{Bmatrix}.$$

By axisymmetry, the local geometry does not depend on the azimuthal position, and therefore without loss of generality we can particularize the expressions at $\theta + \psi = 0$. Consequently, the metric tensor in the natural basis, $g_{ij} = \mathbf{x}_i \cdot \mathbf{x}_j$, takes the form

$$\mathbf{g} = \begin{bmatrix} a^2 + r^2(\psi')^2 & r^2\psi' \\ r^2\psi' & r^2 \end{bmatrix}.$$

The components of this tensor are also those of the Cauchy–Green deformation tensor \mathbf{C} expressed in the natural basis of the reference configuration given by $\mathbf{x}_{0\lambda}$ and $\mathbf{x}_{0\theta}$ (5). Again, particularizing the expressions at $\theta = 0$ without loss of generality, these vectors can be expressed as

$$\mathbf{x}_{0\lambda} = \begin{Bmatrix} r'_0 \\ 0 \\ z'_0 \end{Bmatrix} = a_0 \mathbf{e}_{0\lambda}, \quad \mathbf{x}_{0\theta} = \begin{Bmatrix} 0 \\ r_0 \\ 0 \end{Bmatrix} = r_0 \mathbf{e}_{0\theta},$$

where $\mathbf{e}_{0\lambda}$ and $\mathbf{e}_{0\theta}$ denote tangent unit vectors along meridians and parallels of the axisymmetric surface. By a change of basis, we find the components of \mathbf{C} in the orthonormal basis $\{\mathbf{e}_{0\lambda}, \mathbf{e}_{0\theta}\}$; see Fig. 3B in the main text.

Pellicle deformation. The tangent vector fields along and perpendicular to the strips can be expressed as

$$\mathbf{s}_0 = \sin \alpha_0 \mathbf{e}_{0\lambda} + \cos \alpha_0 \mathbf{e}_{0\theta}, \quad \mathbf{m}_0 = -\cos \alpha_0 \mathbf{e}_{0\lambda} + \sin \alpha_0 \mathbf{e}_{0\theta},$$

where $\alpha_0(\lambda)$ is the angle of the pellicle strips with the surface parallels in the reference configuration. Under the assumption that the local deformation is the result of simple shear along the pellicle strips, the local deformation gradient takes the form

$\mathbf{F} = \mathbf{R} \left(\mathbf{Id} + \gamma \mathbf{s}_0 \mathbf{m}_0^T \right)$, where \mathbf{R} is an orthogonal matrix. This local deformation is area preserving by construction. The Cauchy–Green deformation tensor takes the form

$$\mathbf{C} = \mathbf{F}^T \mathbf{F} = \mathbf{Id} + \gamma (\mathbf{s}_0 \mathbf{m}_0^T + \mathbf{m}_0 \mathbf{s}_0^T) + \gamma^2 \mathbf{m}_0 \mathbf{m}_0^T = \begin{bmatrix} 1 - 2\gamma \sin \alpha_0 \cos \alpha_0 + \gamma^2 \cos^2 \alpha_0 & \gamma (\sin^2 \alpha_0 - \cos^2 \alpha_0) - \gamma^2 \sin \alpha_0 \cos \alpha_0 \\ \gamma (\sin^2 \alpha_0 - \cos^2 \alpha_0) - \gamma^2 \sin \alpha_0 \cos \alpha_0 & 1 + 2\gamma \sin \alpha_0 \cos \alpha_0 + \gamma^2 \sin^2 \alpha_0 \end{bmatrix} \cdot \kappa$$

By equating this microstructural deformation tensor (Fig. 3A) with the deformation derived from the shape changes (Fig. 3B), we obtain three equations:

$$(a^2 + r^2 \psi'^2)/a_0^2 = 1 - 2\gamma \sin \alpha_0 \cos \alpha_0 + \gamma^2 \cos^2 \alpha_0 \quad [\text{S4}]$$

$$(r^2 \psi')/(r_0 a_0) = \gamma (\sin^2 \alpha_0 - \cos^2 \alpha_0) - \gamma^2 \sin \alpha_0 \cos \alpha_0 \quad [\text{S5}]$$

$$(r/r_0)^2 = 1 + 2\gamma \sin \alpha_0 \cos \alpha_0 + \gamma^2 \sin^2 \alpha_0 \quad [\text{S6}]$$

This set of algebraic and differential equations relates the reference pellicle conformation (r_0, z_0, α_0) , the deformed configuration (r, z, ψ) , and the pellicle shear relative to the reference state (γ).

Assuming (r_0, z_0, α_0) and (r, z) are data, the equations above allow us to determine ψ and γ . The pellicle shear follows from Eq. S6:

$$\sin^2 \alpha_0 \gamma^2 + 2 \sin \alpha_0 \cos \alpha_0 \gamma + 1 - (r/r_0)^2 = 0,$$

which has real solutions if the discriminant $\Delta = 4 \sin^2 \alpha_0 [(r/r_0)^2 - \sin^2 \alpha_0]$ is nonnegative, i.e., if

$$r/r_0 \geq |\sin \alpha_0|.$$

If $\Delta \geq 0$, the pellicle shear is

$$\gamma = (-\cos \alpha_0 \pm \sqrt{(r/r_0)^2 - \sin^2 \alpha_0}) / \sin \alpha_0. \quad [\text{S7}]$$

The solution with a plus sign is the one with physical meaning, because the pellicle shear vanishes when the deformed and the reference configurations coincide. Then, Eq. S5 provides an ordinary

differential equation (ODE) for ψ , or alternatively, combining Eqs. S5 and S6, we find

$$\frac{r^2}{a_0 r_0} \psi' = \frac{\cos \alpha_0}{\sin \alpha_0} \left[1 - \left(\frac{r}{r_0} \right)^2 \right] + \gamma, \quad [\text{S8}]$$

which is easily integrated to find ψ . The initial condition for the ODE sets the surface rotation around the symmetry axis, and is determined by the hydrodynamics. For definiteness, we choose $\psi(0)$ such that $\int_{\Gamma} \psi r dS = 0$. By a direct calculation, plugging Eqs. S7 and S8 into Eq. S4, we find $[(ar)^2 - (a_0 r_0)^2]/(a_0 r)^2 = 0$, a compatibility condition identically satisfied if the motion is area preserving.

If a reference configuration with pellicle strips parallel to the meridians is considered, then the pellicle shear is $\gamma = \sqrt{(r/r_0)^2 - 1}$. This expression clearly shows that the larger r_0 , the smaller γ . This is why the choice of reference pellicle given in the main text minimizes the pellicle shear point-wise.

Validation of the theory. We use the experimental observations in ref. 6 to examine the validity of the proposed theory; see Fig. S5. In ref. 6, two images showing different pellicle conformations of a specimen of *Euglena fusca* are reported. This species has a pellicle decorated with particles, making it easy to capture the pellicle conformation with light microscopy. We consider the left image as the reference conformation, and to simplify the analysis assume its pellicle strips are exactly meridians of the surface. We acquire from the image on the right r and z , and compute the pellicle shear and the azimuthal displacement required to accommodate this shape change. The resulting pellicle conformation agrees very well with the observation. This deformation mobilizes a maximum pellicle shear of 150%, which means that pellicle strips slide relative to each other as much as 1.5 times their width.

Definition of a canonical pellicle conformation. By requiring that $\alpha_0 = \pi/2$, the pellicle equations are solvable along the stroke

if $r(\lambda, t) \geq r_0(\lambda)$ for all $t \in [0, 1]$. We define $r_0(\lambda) = \min_t r(\lambda, t)$. To find $z_0(\lambda)$, we recall the definition of λ to write

$$\lambda S = 2\pi \int_0^\lambda \sqrt{r_0'^2(\nu) + z_0'^2(\nu)} r_0(\nu) d\nu.$$

Differentiating this expression with respect to λ , we find

$$z_0'(\lambda) = \sqrt{\left(\frac{S}{2\pi r_0(\lambda)}\right)^2 - r_0'(\lambda)}, \quad [\text{S9}]$$

which is easily integrated to find $z_0(\lambda)$.

Pellicle velocity. Recalling Eq. S3 and the axisymmetry of the kinematics, the velocity field of the pellicle in the $x-z$ plane is

$$\dot{\mathbf{x}} = \begin{Bmatrix} \dot{r} \\ r\dot{\psi} \\ \dot{z} \end{Bmatrix} = v_n^s \mathbf{n} + v_\lambda \mathbf{e}_\lambda + v_\theta \mathbf{e}_\theta,$$

where the normal velocity due to shape changes is given by $v_n^s = \mathbf{n} \cdot \dot{\mathbf{x}} = (-z'\dot{r} + r'\dot{z})/a$, the tangential velocity along generating curves (meridians) is given by $v_\lambda = \mathbf{e}_\lambda \cdot \dot{\mathbf{x}} = (r'\dot{r} + z'\dot{z})/a$ with $\mathbf{e}_\lambda = 1/a\{r', 0, z'\}$, and the tangential velocity along the parallels is given by $v_\theta = \mathbf{e}_\theta \cdot \dot{\mathbf{x}} = r\dot{\psi}$ with $\mathbf{e}_\theta = \{0, 1, 0\}$. By $(\dot{\quad})$ we denote differentiation with respect to time. See Fig. S6 for an illustration.

Hydrodynamics. For the volume-changing euglenids, $\dot{V} = \int_\Gamma v_n^s dS \neq 0$. In order to satisfy the balance of mass of the internal fluid, assumed to be incompressible, the total normal fluid velocity needs to include a permeation contribution v_n^p satisfying $\int_\Gamma v_n^p dS = -\dot{V}$. The total fluid velocity at the pellicle surface is then

$$\mathbf{v}^{\text{out}} = (v_n^s + v_n^p) \mathbf{n} + v_\lambda \mathbf{e}_\lambda + (v_\theta + \omega r) \mathbf{e}_\theta + V_c \mathbf{e}_z,$$

where ω and V_c are (constant and a priori unknown) rigid body rotational and translational velocities. The simplest possibility for the permeation normal velocity is a uniform field given by $v_n^p = -\dot{V}/S$, but any choice of the kind $v_n^p(u, t) = -\dot{V}h(u, t)$, where $h(u, t)$ is a profile with unit integral over the surface, is possible.

The interior part of the swimmer can be either ignored or assumed to be a fluid with the same characteristics of the outer fluid. In this second option, the swimmer is seen as an isolated active surface embedded in a fluid, and the interior fluid satisfies Stokes equations with Dirichlet data given by $\mathbf{v}^{\text{inn}} = \mathbf{v}^{\text{out}}$.

We use a boundary integral formulation for the low Reynolds number (Re) hydrodynamics of a swimmer immersed in a quiescent (Newtonian) fluid (7). Given a swimmer occupying the region Ω with boundary Γ , the velocity of the fluid at any point \mathbf{x} in space satisfies the equations

$$v_i(\mathbf{x}) + \int_\Gamma T_{ijk}(\mathbf{x}-\mathbf{y}) v_j^{\text{inn}}(\mathbf{y}) n_k(\mathbf{y}) d\Gamma_y = \int_\Gamma G_{ij}(\mathbf{x}-\mathbf{y}) f_j^{\text{inn}}(\mathbf{y}) d\Gamma_y \quad \mathbf{x} \in \Omega \quad [\text{S10}]$$

$$v_i(\mathbf{x}) - \int_\Gamma T_{ijk}(\mathbf{x}-\mathbf{y}) v_j^{\text{out}}(\mathbf{y}) n_k(\mathbf{y}) d\Gamma_y = \int_\Gamma G_{ij}(\mathbf{x}-\mathbf{y}) f_j^{\text{out}}(\mathbf{y}) d\Gamma_y \quad \mathbf{x} \in R^3 \setminus \Omega, \quad [\text{S11}]$$

where summation is implied over repeated indices, \mathbf{v} is the velocity in the fluid, $\mathbf{f}^{\text{inn}}(\mathbf{y})$ and $\mathbf{f}^{\text{out}}(\mathbf{y})$ are the inner and outer tractions at $\mathbf{y} \in \Gamma$ defined as $\mathbf{f}^{\text{inn}}(\mathbf{y}) = \boldsymbol{\sigma}^{\text{inn}}(\mathbf{y}) \mathbf{n}(\mathbf{y})$ and $\mathbf{f}^{\text{out}}(\mathbf{y}) = -\boldsymbol{\sigma}^{\text{out}}(\mathbf{y}) \mathbf{n}(\mathbf{y})$, where $\mathbf{n}(\mathbf{y})$ is the outer unit normal to $\partial\Omega$ at the point \mathbf{y} , $\boldsymbol{\sigma}$ is the Cauchy stress in the fluid, and

$$G_{ij}(\mathbf{r}) = \frac{1}{8\pi\nu} \left(\frac{\delta_{ij}}{|\mathbf{r}|} + \frac{r_i r_j}{|\mathbf{r}|^3} \right) \quad T_{ijk}(\mathbf{r}) = -\frac{6}{8\pi} \left(\frac{r_i r_j r_k}{|\mathbf{r}|^5} \right)$$

are the free space Green functions of Stokes flow. Letting \mathbf{x} approach Γ in Eqs. S10 and S11 above, considering the integrals in the principal value sense and exploiting the symmetry of the domain, we obtain two Fredholm integral equations of the first kind for the unknown tractions \mathbf{f}^{inn} and \mathbf{f}^{out} , which we solve numerically via a Galerkin boundary element method (8).

Ignoring the inner behavior of the swimmer is equivalent to setting the Dirichlet boundary conditions for the inner velocity \mathbf{v}^{inn} to zero. In this case, the inner traction \mathbf{f}^{inn} is also identically zero, and the interior part of the swimmer does not contribute to the motion, nor to the dissipation.

The a priori unknown rigid body translational and rotational velocities V_c and ω of the swimmer are obtained by imposing the conditions of self-propulsion, which, consistently with the low Re regime considered here, reduce to annihilating the total viscous force and torque. By axisymmetry, the only two nontrivial equations are

$$\mathbf{e}_z \cdot \int_\Gamma (\mathbf{f}^{\text{inn}}(\mathbf{y}) + \mathbf{f}^{\text{out}}(\mathbf{y})) d\Gamma_y = 0, \\ \mathbf{e}_z \cdot \int_\Gamma (\mathbf{y} - \mathbf{o}) \times (\mathbf{f}^{\text{inn}}(\mathbf{y}) + \mathbf{f}^{\text{out}}(\mathbf{y})) d\Gamma_y = 0,$$

where \mathbf{o} is a point on the axis of symmetry of the swimmer. The power dissipated through the flow induced in the inner and outer fluid are given, respectively, by

$$W^{\text{inn}}(t) = \int_\Gamma \mathbf{f}^{\text{inn}}(\mathbf{y}, t) \cdot \mathbf{v}(\mathbf{y}, t) d\Gamma_y, \\ W^{\text{out}}(t) = \int_\Gamma \mathbf{f}^{\text{out}}(\mathbf{y}, t) \cdot \mathbf{v}(\mathbf{y}, t) d\Gamma_y,$$

from which we obtain the average power exerted by the swimmer on the surrounding fluid P^{out} and the average ambient fluid dissipation $P^{\text{inn+out}}$ when considering the swimmer as an isolated active surface embedded in the fluid:

$$P^{\text{out}} = \frac{1}{T} \int_0^T W^{\text{out}}(t) dt, \\ P^{\text{inn+out}} = \frac{1}{T} \int_0^T (W^{\text{inn}}(t) + W^{\text{out}}(t)) dt.$$

Here, T denotes the duration of one stroke.

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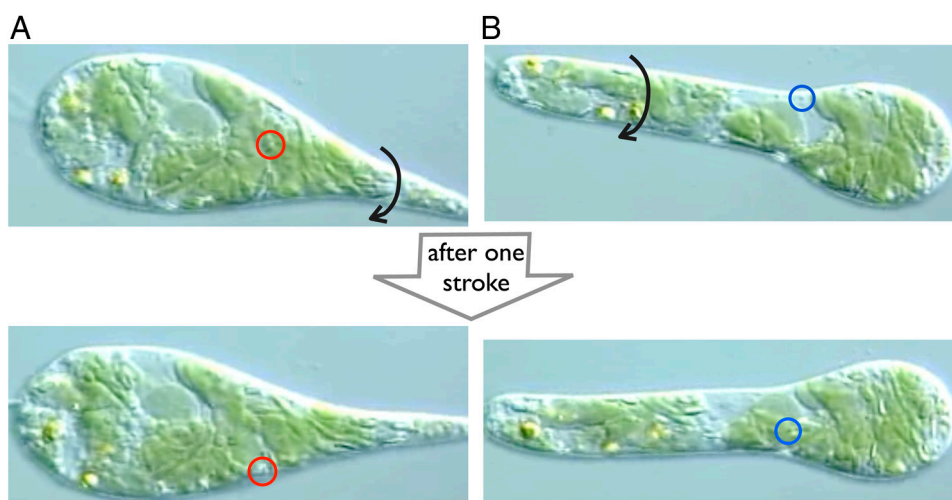


Fig. S1. Evidence for near-axisymmetric cell shape. In some movies, particles attached to the pellicle are clearly noticeable. By tracking these markers and comparing snapshots one stroke apart, (A and B), we conclude that the cell has rotated about its axis by a significant amount. If the cell body was not axisymmetric, the shapes would differ significantly, and Isomap would show distinct paths for each realization of the stroke. However, this is not the case; see Fig. S3A. Frame images courtesy of Richard E. Triemer.

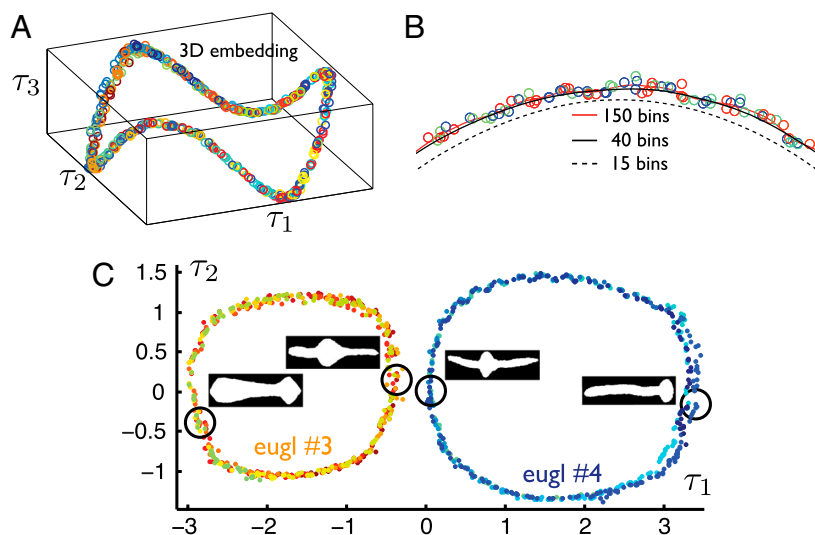


Fig. S2. (A) Embedding of the frames in 3D for euglenid #3, showing that the specimen indeed follows a closed curve in shape space, and that the 2D picture is not the projection of a more complex trajectory. (B) Effect of the width of the basis functions on the reconstructed stroke, here visualized on the 2D embedding. The maximum entropy basis functions (3) used here satisfy $w_a(\tau) \propto \exp[-(n\tau)^2]$, where n is the number of bins used to subdivide the interval $[0, 1)$ and the typical support size of the basis functions is $\sim 1/n$. If n is too large, the reconstructed stroke follows spurious details caused by errors in the data acquisition, whereas if n is too small, then the stroke is excessively smoothed out and shrunk, resulting in less sharply reconstructed motions. We find an optimal compromise with $n = 40$. (C) Joint embedding of the volume-preserving euglenids #3 and #4, showing that the strokes are geometrically distinct, with nearby shapes in the middle of the power stroke.

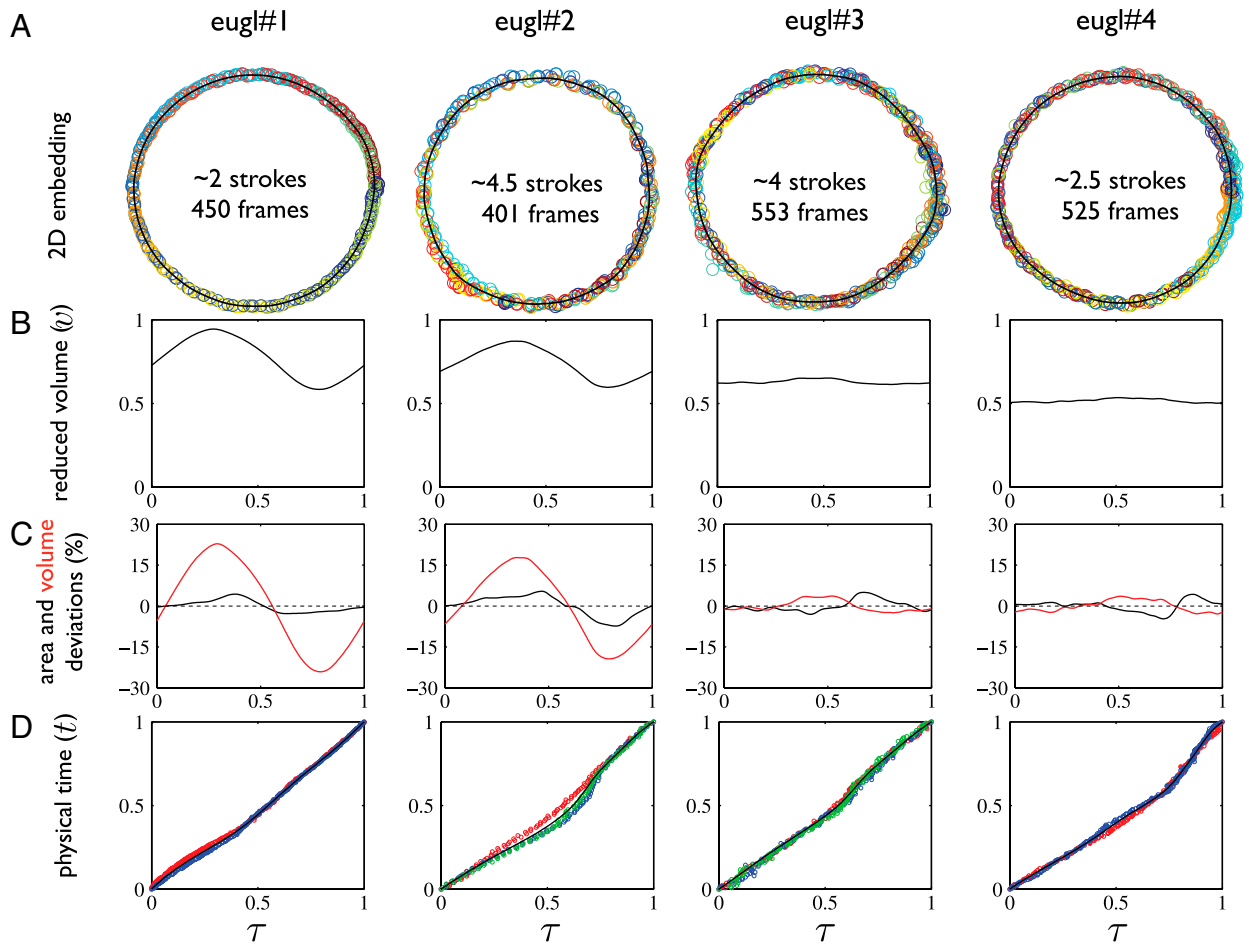


Fig. S3. Quantitative analysis of four video recordings: results. (A) Two-dimensional embedding of the frames, where the points are colored (from dark blue to red) chronologically from the video recording. The solid black line shows the smooth reconstruction of the stroke path in this embedding. (B) Reduced volume v along the reconstructed stroke. (C) Area and reduced volume deviations relative to the average along the stroke. (D) Relationship between the scaled physical time (t) and arc-length in shape space (pseudo-time τ) during two (euglenids #1 and 4) or three (euglenids #2 and 3) strokes. Data from different strokes are represented in different colors. The spline fit to the data is shown in black.

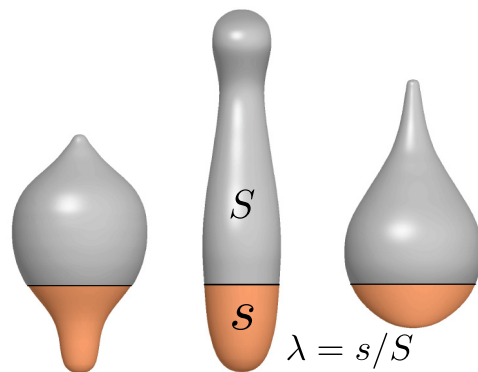


Fig. S4. Definition of the body coordinate λ , labeling material parallels. Under the assumption of local area preservation, the total surface area of the pellicle S remains constant during the stroke. At each instant, consider the region delimited by a parallel such that its surface area is a constant s (shown in orange), and define the ratio $\lambda = s/S (=0.25$ in the figure). By the local area preservation, this parameter labels material parallels along the stroke, shown in black in the figure.

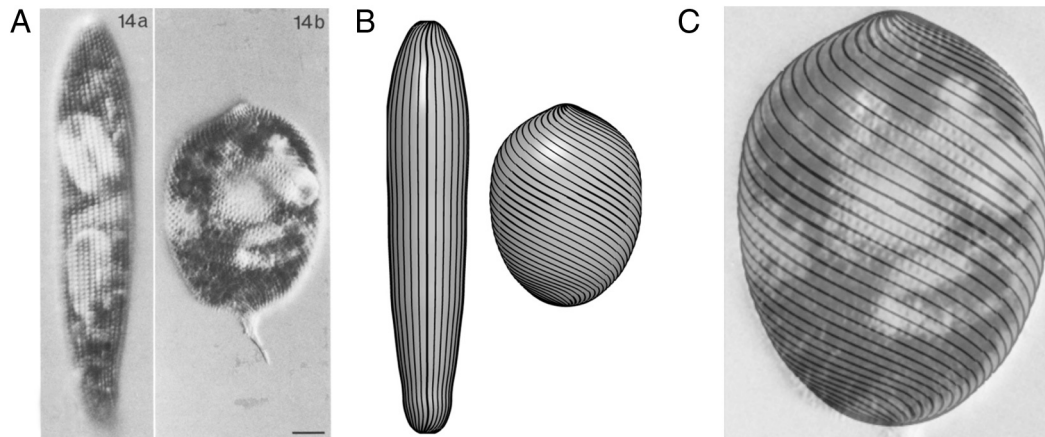


Fig. 55. Validation of the theory on the pellicle kinematics with the experiments in ref. 6 with kind permission from Springer Science and Business Media (Copyright 1985 Springer-Verlag). (A) experimental observations, (B) configurations computed with our model, and (C) overlay of computed and observed configurations confirming the accuracy of the model.

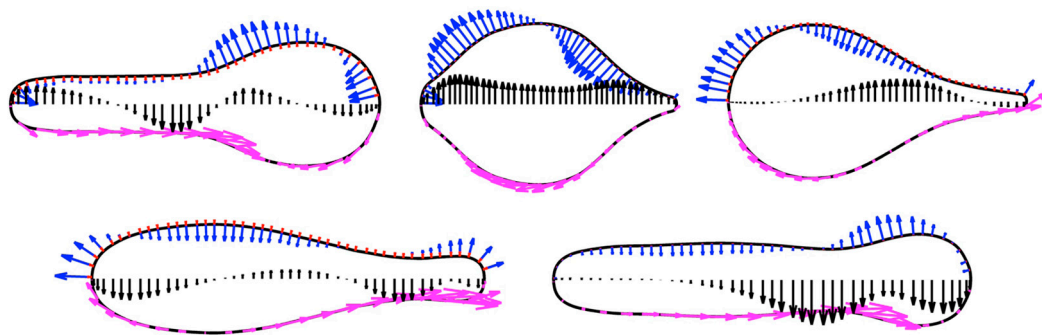
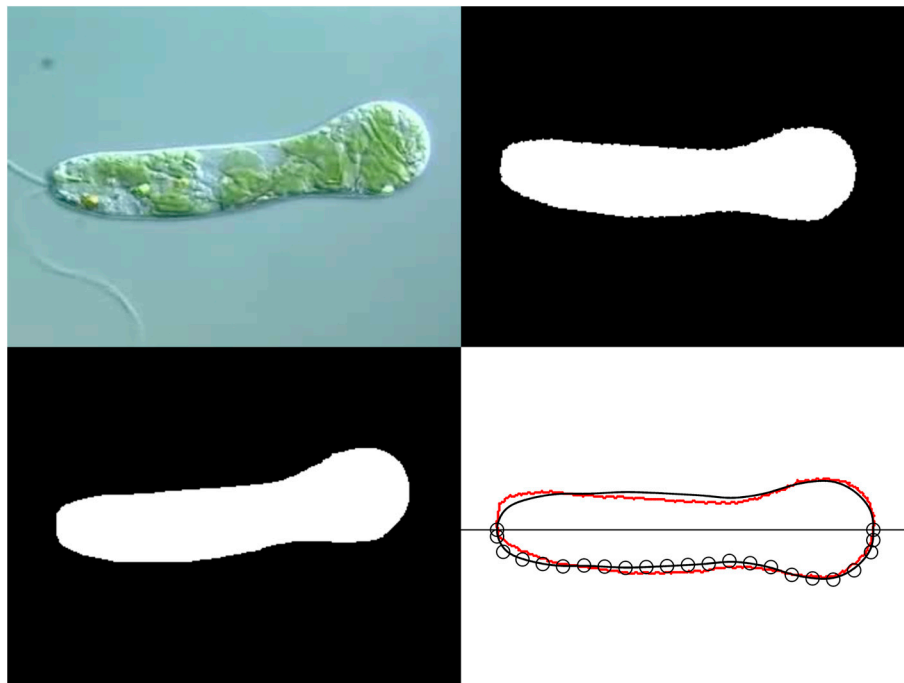
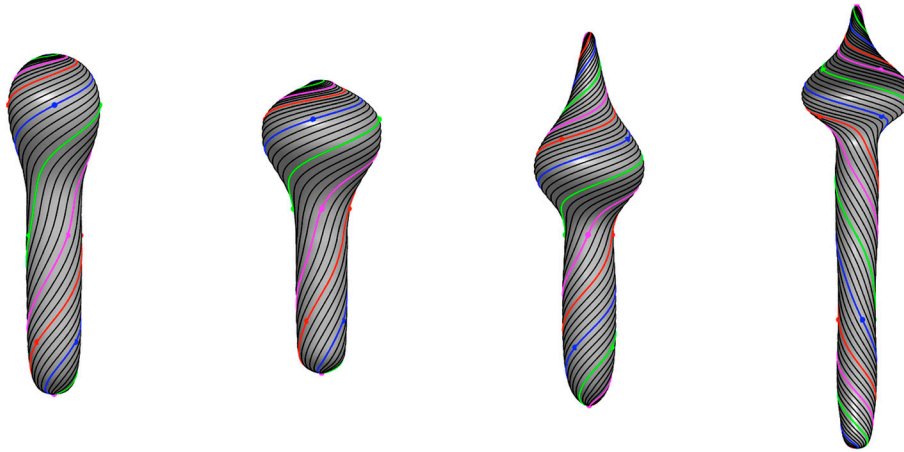


Fig. 56. Velocity of the pellicle surface of euglenid #1 along the stroke at five selected instants, computed as described in section *Pellicle velocity*. All snapshots and velocity components are represented with the same arrow scale. The normal velocities due to shape changes v_n^s and to permeation v_n^p are shown in blue and red, respectively. The tangential velocity along meridians v_λ is shown in magenta, and the azimuthal tangential velocity v_θ is shown in black at a pellicle meridian aligned with the symmetry axis.



Movie S1. Processing of the video recording of euglenid #1. Original frames (Upper Left), segmented frames (Lower Left), aligned segmentation (Upper Right), and B-spline fit to the boundary (Lower Right). The red dots represent the boundary pixels, the black curve is the B-spline fit, and the circles represent the control points of the B-spline. Contains frames from a movie by Richard E. Triemer.

[Movie S1 \(MOV\)](#)



Movie S2. The four model euglenids executing metachronal side by side. The net linear and angular motion along the symmetry axis is determined by the hydrodynamical analysis. Material markers, shown on selected pellicle strips in color, help appreciate the net angular motion. The four models have been scaled to have the same surface area.

[Movie S2 \(MOV\)](#)

Table S1. Summary of the kinematic features and swimming performance of the four strokes

Euglenid	U	$\Delta\theta/(2\pi)$	$P_{r,z}^{\text{out}}/\bar{P}$	$P_{\theta}^{\text{out}}/\bar{P}$	$\text{Eff}_L^{\text{out}}, \%$	$P_{r,z}^{\text{out+inn}}/\bar{P}$	$P_{\theta}^{\text{out+inn}}/\bar{P}$	$\text{Eff}_L^{\text{out+inn}}, \%$
# 1	0.16	0.0070	11	3.2	0.71	39	3.6	0.24
# 2	0.27	0.055	15	3.8	1.6	47	4.5	0.57
# 3	0.25	0.086	12	3.5	1.6	46	4.0	0.48
# 4	0.40	0.20	25	6.0	2.0	76	7.3	0.76

U denotes the net linear displacement in one stroke, in units of body length $2R = \sqrt{S/\pi}$, defined as the diameter of the sphere with the same surface area of the pellicle. $\Delta\theta$ denotes the net angular displacement after one stroke. Eff_L denotes the Lighthill efficiency, accounting for the outer fluid, or the outer and inner fluids. The power is nondimensionalized by $\bar{P} = 6\pi\eta R^3/T^2$, where T is the duration of one stroke. The different components of the dissipation power are reported, inner or outer, and azimuthal or within the symmetry plane.