Swimming, Feeding and Inversion of Multicellular Choanoflagellate Sheets

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The recent discovery of the striking sheet-like multicellular choanoflagellate species *Choanoeca* flexa that dynamically interconverts between two hemispherical forms of opposite orientation raises fundamental questions in cell and evolutionary biology, as choanoflagellates are the closest living relatives of animals. It similarly motivates questions in fluid and solid mechanics concerning the differential swimming speeds in the two states and the mechanism of curvature inversion triggered by changes in the geometry of microvilli emanating from each cell. Here we develop fluid dynamical and mechanical models to address these observations and show that they capture the main features of the swimming, feeding, and inversion of C. flexa colonies.

Some of the most fascinating processes in the developmental biology of complex multicellular organisms involve radical changes in geometry or topology. From the folding of tissues during gastrulation [1] to the formation of hollow spaces in plants [2], these processes generally involve coordinated cell shape changes, cellular division, migration and apoptosis, and formation of an extracellular matrix (ECM). It has become clear through multiple strands of research that evolutionary precedents for these processes exist in some of the simplest multicellular organisms such as green algae [3, 4] and choanoflagellates [5], the latter being the closest living relatives of animals. Named for their funnel-shaped collar of microvilli that facilitates filter feeding from the flows driven by their beating flagellum, choanoflagellates serve as model organisms for the study of the evolution of multicellularity.

While well-known multicellular choanoflagellates exist as linear chains or "rosettes" [6] held together by an ECM [7], a new species named *Choanoeca flexa* was recently discovered [8] with an unusual sheet-like geometry (Fig. 1) in which hundreds of cells adhere to each other by the tips of their microvilli, without an ECM [9]. The sheets can exist in two forms with opposite curvature, one with flagella pointing towards the center of curvature ["flagin"] with a relatively large spacing between cells, and another with the opposite arrangement ["flag-out"] with more tightly-packed cells. Transformations from *flag-in* to flag-out can be triggered by darkness, and occur in \sim 10 s. Compared to the *flag-out* form, the *flag-in* state has limited motility and is better suited to filter-feeding. It was conjectured [8] that the darkness-induced transition to the more motile form is a type of photokinesis.

As a first step toward understanding principles that govern the behavior of such a novel organism as C. flexa,

we analyze two models for these shape-shifting structures. First, the fluid mechanics are studied by representating the cell raft as a collection of spheres distributed on a hemispherical surface, with nearby point forces to represent the action of flagella. Such a model has been used to describe the motility of small sheet-like multicellular assembles such as the alga *Gonium* [10]. The motility and filtering flow through these rafts as a function of cell spacing and curvature explain the observed properties of *C. flexa*. Second, abstracting the complex elastic interactions between cells to the simplest connectivity, we show that a model based on linear elasticity at



FIG. 1. The multicellular choanoflagellate *Choanoeca flexa*. Top views of (a) *flag-in* and (b) *flage-out* states at times relative to removal of light. (c) Close-up of the collar connections in the two states. (d) Electron micrograph showing round white cell bodies connected by microvilli. Adapted from [8].



FIG. 2. Models for *C. flexa.* (a) Cell body and flagellar force in the *flag-in* state. (b,c) Mechanical model of interconnecting microvilli in rafts; cells (green spheres, not to scale) are at the vertices of a geodesic icosahedron. Blue arrows indicate flagella forces, red segments represent microvilli, blue dots the microvilli tips, and blue lines the collar-collar interface. (b) Connectivity of the whole raft. (c) Connections between two cells. Effect of curvature 1/R on (d) geometry of raft and (e) swimming speed U (blue) and flow rate Q (red) passing through S_f at constant. (f,g) Cross section of the disturbance flows $\tilde{\mathbf{u}}_d$ and \mathbf{u}_f around the a raft ($\Phi = 0.31, R = 8.61$) in the reciprocal problems for calculating U and Q.

the microscale produces bistability on the colony scale.

Fluid mechanics of feeding and swimming – The cells in a C. flexa raft are ellipsoidal, with major and minor axes $a \sim 4 \,\mu\text{m}$ and $b \sim 3 \,\mu\text{m}$, with a single flagellum of length $2L \sim 24.5 \,\mu\text{m}$ and radius $r \sim 0.5 \,\mu\text{m}$ beating with amplitude $d \sim 2.3 \,\mu\text{m}$ and frequency $f \sim 44 \,\text{Hz}$ [11], sending bending waves away from the body. A cell swims with flagellum and collar rearward; the body and flagellum comprise a "pusher" force dipole. From resistive force theory [13] we estimate the flagellar propulsive force to be $F \sim 2L \left(\zeta_{\perp} - \zeta_{\parallel} \right) (1 - \beta) f \lambda \sim 8 \,\mathrm{pN}$, where β is a function of the wave geometry, $\lambda \sim 12 \,\mu \text{m}$ is the wavelengths along the direction of the bending wave [11], ζ_{\perp} and ζ_{\parallel} are transverse and longitudinal drag coefficients, $\zeta_{\perp} \sim 2\zeta_{\parallel} \sim 4\pi\mu/\ln(2L/r)$, with μ the fluid viscosity. These features motivate a computational model in which N identical cells in a raft have a spherical body of radius a and a point force $F\hat{\mathbf{n}}_i$ acting on the fluid a distance L from the sphere center, oriented along the vector $\hat{\mathbf{n}}_i$ that represents the collar axis [Fig. 2(a,b)]. An idealization of a curved raft involves placing those spheres on a connected subset of the vertices of a *geodesic* icosahedron (one whose vertices lie on a spherical surface) of radius $\rho \gg a$; the area fraction Φ of the sheet occupied by cells scales as $\Phi \sim N(a/\rho)^2$. The pentagonal neighborhoods within the geodesic icosahedron serve as topological defects that allow for smooth large-scale surface curvature [14]. Importantly, confocal imaging of C. flexa colonies shows that a significant fraction (~ 0.25) of the cellular neighborhoods defined by the microvilli connections are pentagonal [11], and earlier work on C. perplexa [9] also found non-hexagonal packing. We use the geodesic icosahedron $\{3,5+\}_{(3,0)}$ in standard notation [15], with 92 total vertices, and take patches with N = 58 for computational tractability. The vectors $\hat{\mathbf{n}}_i$ point towards (away from) the icosahedron center in the flaq-in (flaq-out) forms (Fig. 2(a)). A deformation of the sheet to a new radius ρ' , at fixed Φ , requires the new polar angle θ'_i of a cell with respect to the central axis of the sheet be related to its original angle θ_i via ${\rho'}^2 (1 - \cos \theta_i') = \rho^2 (1 - \cos \theta_i)$. We define the scaled force offset length $\ell = L/a \sim 3$ and sheet radius $R = \rho'/a \gtrsim 6$, and take R > 0 in the flag-in state.

Images of many colonies of *C. flexa* [11] show that the packing fraction in the *flag-out* state $\Phi_{out} = 0.47 \pm 0.06$, considerably less than both the maximum packing frac-

tion $\Phi_{\text{max}} = \pi \sqrt{3}/6 \simeq 0.907$ for a hexagonal array of spheres in a plane, and the estimated maximum packing fraction $\tilde{\Phi}_{\text{max}} \simeq 0.83$ for circles on a sphere [16]. The packing fraction in the *flag-in* state is $\Phi_{\text{in}} = 0.34 \pm 0.03$, and we use the extremes $\Phi_{\text{out}} = 0.53$ and $\Phi_{\text{in}} = 0.31$ as representative values to explore the consequences of the differences between the two forms.

Consider first an isolated force-free spherical cell at the origin moving at velocity $U_s \hat{\mathbf{e}}_x$ with point force $-\mathbf{F} = -F\hat{\mathbf{e}}_x$ at $-L\hat{\mathbf{e}}_x$ acting on the fluid and its reaction force \mathbf{F} acting on the cell. The cell experiences Stokes drag $-\zeta_s U_s \hat{\mathbf{e}}_x$, where $\zeta_s = 6\pi\mu a$, and a disturbance drag $-D\hat{\mathbf{e}}_x$ arising from the disturbance flow created by the point force. By the reciprocal theorem [17], the disturbance drag is $D = \mathbf{F} \cdot \tilde{\mathbf{u}}_d (-L\hat{\mathbf{e}}_x)/\hat{U}$, where $\tilde{\mathbf{u}}_d(\mathbf{r})$ is the disturbance flow created when the cell is dragged along $\hat{\mathbf{e}}_x$ with unit speed \hat{U} . Force balance then yields the singlecell swimming speed $U_s \equiv (F/\zeta_s)[1 - 3/(2\ell) + 1/(2\ell^3)]$. Thus, the closer the point force is to the cell (i.e., the smaller is ℓ), the more drag the cell experiences and the slower is U_s . Setting $\ell = 3$ yields $U_s \sim 118 \,\mu$ m/s, consistent with observations.

This intuitive picture extends to a raft of cells. As the raft moves at velocity $U\hat{\mathbf{e}}_x$, it experiences a Stokes drag $-\zeta U\hat{\mathbf{e}}_x$. The disturbance flow created by the point forces $F\hat{\mathbf{n}}_i$ acting at $\mathbf{r}_i + L\hat{\mathbf{n}}_i$, produces a disturbance drag $D = F\Sigma_i\hat{\mathbf{n}}_i \cdot \mathbf{u}_d(\mathbf{r}_i + L\hat{\mathbf{n}}_i)$, where \mathbf{u}_d is the (dimensionless) disturbance flow from the raft when it is dragged along $\hat{\mathbf{e}}_x$ with unit speed. Force balance then yields

$$U = -\frac{F}{\zeta} \sum_{i=1}^{N} \hat{\mathbf{n}}_{i} \cdot \left[\hat{\mathbf{e}}_{x} - \mathbf{u}_{d} \left(\mathbf{r}_{i} + L \hat{\mathbf{n}}_{i} \right) \right], \qquad (1)$$

where $-F\Sigma_i \hat{\mathbf{n}}_i$ is the sum of reaction forces propelling the raft along $\hat{\mathbf{e}}_x$, and where \mathbf{u}_d has been rendered dimensionless by the unit speed. In practice, we compute \mathbf{u}_d and ζ using a Boundary Element Method [10]. Because of the curved geometry, point forces are closer to neighboring cells in the *flag-in* state than in the *flag-out* state. Thus, as in Fig. 2(d), for a geometry with a given |R|, the *flag-in* state has a larger disturbance drag than the *flaq-out* state, and a smaller speed U.

The difference in swimming speed between the two states can also be explained in terms of $\mathbf{u}_d(\mathbf{r}_i + L\hat{\mathbf{n}}_i)$ in (1). Figure 2(f) shows that \mathbf{u}_d inside the raft is close to $\hat{\mathbf{e}}_x$ because of the curved geometry and screening effects. Hence, U is small when the point forces are inside. Meanwhile, \mathbf{u}_d outside decays with the distance from the raft, so U is large when $\mathbf{r}_i + L\hat{\mathbf{n}}_i$ is outside.

Previous work on filter-feeding in choanoflagellates focused first on the far-field limit based on a stresslet description [18], but later work showed near-field effects can significantly affect capture rates [19]. To estimate the filter-feeding flux Q passing through a colony of C. flexa, we measure, in the body frame, the flux passing through the surface S_f projected a distance of 1.2*a* from the cell center along $\hat{\mathbf{n}}$, as in Fig. 2(g). By the reciprocal theorem, Q can be written in terms of the disturbance flow \mathbf{u}_f around a stationary raft and the hydrodynamic forces \mathbf{F}_f on the raft when the surface S_f applies a unit normal pressure $\hat{\mathbf{p}}$ on the fluid,

$$Q = F \sum_{i} \hat{\mathbf{n}}_{i} \cdot \mathbf{u}_{f}(\mathbf{r}_{i} + L\hat{\mathbf{n}}_{i}) + U\hat{\mathbf{e}}_{x} \cdot \left(\mathbf{F}_{f} - \int_{S_{f}} \hat{\mathbf{p}} \, dA\right), \quad (2)$$

where \mathbf{u}_f and \mathbf{F}_f acquire the units of velocity/pressure and area, respectively, by scaling with $|\hat{\mathbf{p}}|$. Numerical results in Fig. 2(d) show that the flux due to point forces $\sum_i F \hat{\mathbf{n}}_i \cdot \mathbf{u}_f / \hat{p}$ strongly dominates Q. Therefore, the difference in Q between the two states can be explained by $\mathbf{u}_f(\mathbf{r}_i + L \hat{\mathbf{n}}_i)$ (Fig. 2(g)). To maintain incompressibility under pressure $\hat{\mathbf{p}}$, the disturbance flow \mathbf{u}_f is much stronger inside the raft than outside. Hence, point forces placed inside the raft pump more flow through the raft than when placed outside.

Figure 2(d) shows the effect of changes in the raft curvature and packing fraction. There is one R that maximizes swimming speed in the *flaq-in* state and another one that maximizes feeding flux in the *flag-out* state. This arises from a balance between the screening effect mentioned above and the alignment of forcing. In the flaq-out state, an initial decrease in curvature aligns the forcing direction with the swimming direction, increasing swimming speed, but a further reduction in curvature reduces the screening effect as cells are now more spread out in the plane orthogonal to the swimming direction. A similar argument applies to the flow rate maximum in the *flag-in* state. Comparing these maxima, Fig. 2(d)shows that a spread-out colony results in more flux, while a closely-packed colony results in faster motility. Thus, through the interconversion between the two states, C. flexa takes advantage of the hydrodynamics effect of the curved geometry for efficient filter-feeding and swimming.

Mechanics of inversion – Detailed studies suggest that inversion requires an active process within each cell, likely driven by contraction of an F-actin ring at the apical pole through the action of myosin [8]. Thus, a full treatment would address the complex problem of elastic filaments responding to the apical actomyosin system and adhering to each other. We simplify this description by considering as in Fig. 2(e) that each cell *i*, located at \mathbf{r}_i and surrounded by m_i neighbors, has m_i rigid, straight filaments emanating from it. Two filaments from neighboring cells i and j meet at vertex ρ located at \mathbf{r}_{ρ} , with $\phi_{i\rho}$ the angle between $\mathbf{r}_{\rho} - \mathbf{r}_{i}$ and the cell normal vector $\hat{\mathbf{n}}_{i}$. Any two adjacent filaments emanating from cell *i*, and which meet neighboring filaments at vertices ρ and σ , define a plane whose normal $\hat{\mathbf{n}}_{i\rho\sigma}$ points toward the apicobasal axis $\hat{\mathbf{n}}_i$. That normal and its counterpart $\hat{\mathbf{n}}_{j\sigma\rho}$ on cell j determine the angle $2\psi_{ij\rho\sigma}$ between the two planes.

As above, we use the geodesic icosahedron to define the cell positions and thus determine the filament net-



FIG. 3. Inversion dynamics from numerical studies. (a)-(d) A colony, initially at a hemispherical minimum with $(\phi_0, \psi_0) = (0.55, 0.65)$, inverts after a change to (0.71, 0.41), with $\ell_0 = 0.5$, $K_{\psi} = 2$ and $K_{\ell} = 5$. Connections between collar vertices are shown in blue, apicobasal axes as red arrows at cell body positions. (e) Minimum energy E in the $\psi_0 - \phi_0$ plane, where C < 0 (*flag-in*) above the black line $\psi_0 = \phi_0$, and C > 0 (*flag-out*) below. (f) Evolution of E vs C as the colony relax towards a minimum energy state after instantaneous changes in (ϕ_0, ψ_0) shown by the dotted and solid red and blue lines in (e).

work connecting neighboring cells. The two sets of angles $\{\phi\}$ and $\{\psi\}$ are used to define a Hookean elastic energy that mimics the elasticity of the microvilli, allowing for preferred intrinsic angles ϕ_0 and ψ_0 that encode the effects of the apical actomyosin system on the microvilli and the geometry of microvilli adhesion. Allowing also for stretching away from a rest length ℓ_0 , the energy is

$$E = \frac{1}{2}k_{\phi}\sum_{i,\rho}\delta\phi_{i\rho}^{2} + \frac{1}{2}k_{\psi}\sum_{i,j,\rho,\sigma}\delta\psi_{ij\rho\sigma}^{2} + \frac{1}{2}k_{\ell}\sum_{i,\rho}\delta\ell_{i\rho}^{2}, \quad (3)$$

where $\delta \phi_{i\rho} = \phi_{i\rho} - \phi_0$, $\delta \psi_{ij\rho\sigma} = \psi_{ij\rho\sigma} - \psi_0$, and $\ell_{i\rho} = |\mathbf{r}_i - \mathbf{r}_{\rho}| - \ell_0$. The moduli k_{ϕ} , k_{ψ} and k_{ℓ} and quantities ϕ_0, ψ_0 and ℓ_0 are assumed constant for all cells.

The energy (3) is intimately tied to the lattice geometry of the raft. If the cells are arranged in a hexagonal lattice $(m_i = 6)$ the system of filaments can achieve E = 0by setting all cell-collar angles to ϕ_0 , all collar-collar interface angles to ψ_0 , and $\phi_0 = \psi_0$. This corresponds to a flat sheet. Increasing $\phi_0 = \psi_0$ leads to uniform, isotropic sheet expansion. In a non-planar raft, curvature is introduced through topological defects $(m_i \neq 6)$, such as pentagons, and mismatch between the local values of ψ and ϕ . While pentagonal defects are known to cause outof-plane buckling in crystal lattices [14], they do not by themselves select a particular *sign* of the induced curvature. Thus, there is inherent bistability in the cellular raft that can be biased by changes in the geometry of the out-of-plane filaments, somewhat akin to the role of "apical constriction" in the shapes of epithelia [20].

For the case of two cells lying in a plane, each with two filaments, and with one vertex between them, if $\phi = \phi_0$, $\psi = \psi_0$, and $r = \ell_0$, then the filament tips lie on a circle

of radius $R_0 = 1/C_0$, where $C_0 = \sin(\psi_0 - \phi_0)/\ell_0 \sin \phi_0$. While, in general, the equilibrium state of a curved raft will not have $\phi_{i\rho} = \phi_0$, $\psi_{ij\rho\sigma} = \psi_0$ and $r_{i\rho} = \ell_0$ everywhere, we may nevertheless use this relationship to define a proxy for the average curvature of the raft. Recognizing that in numerical studies stretching effects are small, we ignore variations in $r_{i\rho}$ and define $C = \sin(\langle \psi \rangle - \langle \phi \rangle)/\ell_0 \sin(\langle \phi \rangle)$, where $\langle \cdot \rangle$ is an average over cells and vertices. The colony is in the *flag-in* (*flag-out*) state when C > 0 (C < 0).

The simplest model of raft dynamics localizes the viscous drag to the individual cell and vertex positions \mathbf{r}_{γ} according to a gradient flow $\zeta \partial_t \mathbf{r}_{\gamma} = -\partial E/\partial \mathbf{r}_{\gamma}$ driven by the force derived from (3). We solve this dynamics numerically with forward integration. Since the $\hat{\mathbf{n}}_i$ are constrained to have unit length, they are normalized after each step in the direction of the negative gradient, making the dynamical algorithm follow a projected gradient descent [21]. Via a rescaling of time we may set one of the elastic constants to unity (say, k_{ϕ}) and need only consider the ratios $K_{\psi} = k_{\psi}/k_{\phi}$ and $K_{\ell} = k_{\ell}/k_{\phi}$.

Interconversion between the *flag-in* and *flag-out* states is shown in Figs. 3(a-d) following an abrupt change in the preferred angle pair (ϕ_0, ψ_0) that crossing the line of equality $\psi_0 = \phi_0$ that divides the states (Fig. 3(e)), as during raction/relaxation of the F-actin ring in response to a stimulus. The intermediate shapes exhibit a ring of inflection points similar to those seen in experiments on *C. flexa* and also in the inversion the algae *Pleodorina* [3] and larger species [22, 23]. Tracking the energy as each of the two equilibria is achieved, the picture that emerges in Fig. 3(f) is evolution on a double-well potential energy landscape as a biasing field is switched in sign.

We have shown that simple models can explain the swimming, feeding, and inversion of the recently discovered multicellular choanoflagellate C. flexa [8]. These results suggest further exploration on a possible continuum description of the sheets, fluid-structure interactions during locomotion, dynamics of photokinesis, and developmental processes of these remarkable organisms.

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