Life around the scallop theorem

Eric Lauga

Department of Mechanical and Aerospace Engineering, University of California San Diego, 9500 Gilman Drive, La Jolla CA 92093-0411, USA.

(Dated: November 16, 2010)

Locomotion on small scales is dominated by the effects of viscous forces and, as a result, is subject to strong physical and mathematical constraints. Following Purcell’s statement of the scallop theorem which delimitates the types of swimmer designs which are not effective on small scales, we review the different ways the constraints of the theorem can be escaped for locomotion purposes.

I. INTRODUCTION

Swimming cells, such as bacteria (prokaryotes) or spermatozoa (eukaryotes), represent the prototypical example of active soft matter. They are active as they transform chemical energy (ATP for eukaryotes, ion flux for prokaryotes) into mechanical work [1] and, as a result, are able to continuously change shape and move in viscous environments [2]. As mechanical entities, cells belong to the world of soft matter, displaying complex rheological properties on a range of time and spatial scales and responding to external forcing in a time-dependent and nonlinear fashion [3].

In their micron-size environment, the fluid forces acting on swimming cells are dominated by the effect of viscous dissipation [4, 5]. Seminal papers in the 1950s laid the groundwork for detailed investigations on the hydrodynamics of cell locomotion [6–9], with the main goal of predicting cell kinematics, energetics, the interactions with their environment, and the general importance of fluid forces in biological form and function [10–15].

In 1977, Purcell’s influential paper “Life at low Reynolds number” put a somewhat different spin on a field which was already mature [16]. In it, Purcell brought to light the counter-intuitive physical and mathematical constraints arising from locomotion in an inertialess world. He demonstrated that for organisms moving in very viscous fluids, there exists a class of shape change that can never be used for locomotion, a result beautifully summarized under the name “scallop theorem”, borrowing the name of such an organism — a hypothetical microscopic scallop — which could not locomote in the absence of inertia.

In this short review, we look back at the scallop theorem, and pose the question: What are the basic ingredients necessary to design swimmers able to move on small scales? What are the different ways offered by physics to get around the constraints of the theorem? After stating the various assumptions for the theorem to be valid (§II), we show how non-reciprocal shape changes (§III), inertia (§IV), hydrodynamic interactions (§V), and coupling with the physical environment (§VI) can all be exploited to provide locomotion on small scales.

II. THE SCALLOP THEOREM

The scallop theorem has a relatively simple statement [16]. Consider a body changing shape in a time-periodic fashion. In the absence of inertia, the equations describing the motion of an incompressible Newtonian fluid are Stokes equations, which are linear and independent of time [4, 5]. In addition, in the absence of inertia, the swimmer remains perpetually force- and torque-free [15]. Purcell’s scallop theorem can then be stated as follows. If the sequence of shapes displayed by the swimmer is identical to the sequence of shapes displayed when seen in reverse — so-called reciprocal motion — then the average position of the body cannot change over one period. Another manner to describe reciprocal motion is stated in Purcell’s original paper as:

*Electronic address: elauga@ucsd.edu*
“... I change my body into a certain shape and then I go back to the original shape by going through the sequence in reverse... So, if the animal tries to swim by a reciprocal motion, it can’t go anywhere.”

Time is not explicitly mentioned in the theorem and in fact, because of the linearity and time-independence of the equations, the rate at which the sequence of shapes is being displayed is irrelevant [15]. In Purcell’s own words,

“Fast, or slow, it exactly retraces its trajectory, and it’s back where it started.”

Physically, the absence of time in the equations of motion means there is no intrinsic time scale to the swimming problem, which prevents distinguishing between forward and backward in a reciprocal motion.

Purcell’s statements may appear simple, but are in fact far-reaching. They form the basis of a purely geometrical approach to cell locomotion [17–20] and have sparked considerable attention in the area of biolocomotion from the physics and soft matter community — so much so that “Life at low Reynolds number” is now the most cited paper in the field.

The name for the theorem originates from the simplest kind of reciprocal swimmers, namely those deforming with a single degree of freedom, such as the hinge of a hypothetical micron-scale scallop. For any swimmer with a single geometrical degree of freedom, say $\theta(t)$, then by properties of Stokes equations, its swimming speed, $u$, necessarily scales as $u \sim \dot{\theta} F(\theta)$, which is always an exact derivative, and thus averages in time to zero $\langle u \rangle = 0$. Swimmers with only one degree of freedom can thus never swim on small scales.

Strictly speaking, the scallop theorem is valid only with the following assumptions: a single swimmer displaying reciprocal motion in an infinite quiescent Newtonian fluid and in the absence of inertia and external body forces.

Examining each of these assumptions in detail suggests a way around the theorem and a design for a swimmer, which we now review. As in Purcell’s original paper, we will focus only on swimming by shape change or motion and we will thus not consider chemical swimmers [21–24] or solid bodies powered by external fields [25].

III. NON-RECIPROCAL KINEMATICS

A. Biological swimmers: waves

The main message of Purcell’s paper is that swimmers should change their shapes in a non-reciprocal fashion. The manner in which motion occurs should thus indicate a clear direction of time, which leads naturally to the occurrence of waves. Indeed, most swimmings cells locomote by using traveling wave-like deformation of their bodies or appendages [10–15]. Swimming bacteria rotate one or more helical flagella using rotary motors embedded in the cell walls [38–40], leading to flagella kinematics akin to that of traveling helical waves, and thus propulsion [41, 43] (Fig. 1a). Other types of bacteria swim using whole-body wave deformation propelled by flagella beneath the cell’s outer membrane [44] or wave-like propagation of kinks in their shapes in the absence of flagella [45]. Spermatozoa and other singly flagellated eukaryotes swim using traveling waves [12] induced by molecular motors-driven internal sliding of polymeric filaments inside the flagellum [46–48] (Fig. 1b). The flagella kinematics can be planar [49], helical [43], or even doubly helical [50]. The many cilia covering some eukaryotes also deform as so-called metachronal waves [52, 54] (Fig. 1c).

B. Synthetic swimmers

Beyond the swimming methods displayed by biological swimmers, some simpler modes of non-reciprocal motion can be devised theoretically and in the lab.

---

1 The theorem is also valid in a confinement environment as long as the boundaries display no motion.
1. Imposing non-reciprocal kinematics

As shown by Purcell, swimmers with a single degree of freedom cannot move. One needs therefore at least two degrees of freedom and their prescribed variation in time should sweep a finite area in parameter space. In his original paper, Purcell proposed such a swimmer \[16\], namely an elongated body with two rotational hinges \[55–57\] (Fig. 1d).
Subsequently, non-reciprocal swimmers of very simple shapes have been devised theoretically, including ones composed of three spheres [30, 58–61] (Fig. 1e), two volume-changing spheres [62], and two-orientation changing spheres or ellipsoids [64]. Beyond geometry, the two degrees of freedom could also be physical parameters, for example the volume and spontaneous curvature of a lipid vesicle [31] (Fig. 1f). Alternatively, the swimmer’s shape and deformation change could be topologically equivalent to the inside-out rotation of a torus [65, 66] or tank-treading [67] for which periodicity is achieved by a continuous series of displacements tangent to the swimmer shape. Continuous normal flows in the form of fluid jets can also be used [68].

2. Flexible swimmers: Non-reciprocal kinematics from reciprocal forcing

A second class of simple swimmers can be designed for which a reciprocal actuation combined with flexibility or elasticity can lead to kinematics of shape change which are non-reciprocal, and thus to locomotion.

The prototypical example of this class of swimmers is a flexible filament actuated periodically up and down at one end where it is clamped, and free on the other [69]. If the filament is rigid, its motion is reciprocal and cannot be used for propulsion. In contrast, if the filament is flexible and is actuated near the typical frequency at which viscous drag and elastic forces balance, its shape as it is actuated up (respectively down) is concave (respectively convex), leading to non-reciprocal kinematics and propagation of an elasto-hydrodynamic wave. Mathematically, the scallop theorem breaks down because time enters the problem through the viscous drag term in the equation for the filament shape (via a partial time-derivative), and thus a relevant time scale can be defined.

The generation of propulsive force and locomotion using flexibility filaments has been the center of many theoretical and computational investigations [69–72]. A macro-scale experiment confirmed the physical picture outlined above [72]. Related phenomena include elastic buckling instabilities [74–76] and shape transitions [77–79] for rotated elastic filaments. At the micro-scale, an experimental realization of a flexible swimmer was achieved using elastic superparamagnetic filaments [80] actuated by external magnetic fields and attached to a red blood cell (Fig. 1g) [32], prompting subsequent modeling efforts [81–84]. A similar implementation was achieved using a nanometric silver filament attached to an externally-driven ferromagnetic nickel head (Fig. 1h) [33]. In all these cases however, it is the presence of external torques (via external magnetic fields) that allows locomotion, and thus they do not represent true self-propelled motion.

IV. INERTIA

For the scallop theorem to be valid, all inertial terms in the equation of motion of the swimmer should be set to zero. Naturally, they cannot exactly disappear unless no motion occurs, and thus a fundamental question arises, namely how much inertia is needed to escape the constraints of the theorem? Is the scallop theorem valid only asymptotically, or does it stand as long as inertia is below a certain limit? These questions were first posed by Dudley and Childress [85] who studied the behavior of a mollusk able to use both reciprocal and non-reciprocal modes of locomotion, and who postulated that a finite amount of inertia was necessary for locomotion to be able to occur.

Mathematically, three qualitatively different Reynolds numbers can be defined. Consider a swimmer of typical size $L$ and density $\rho_s$ undergoing reciprocal motion of amplitude $A$ and frequency $\omega$ in a Newtonian fluid of density $\rho$ and shear viscosity $\mu$. Using a typical velocity scale $U \sim A\omega$, the natural Reynolds number for the reciprocal motion is given by $Re = \rho L A \omega / \mu$, and is the one corresponding to the nonlinear advection term in the Navier Stokes equations (for example, in water, $Re \approx 10^{-4}$ for $E. coli$ while $Re \approx 10^{-2}$ for human spermatozoa). The oscillatory Reynolds number, corresponding to the linear unsteady Stokes term, is given by $Re_\omega = \rho L^2 \omega / \mu$. Finally, the Reynolds number based on the body inertia is $Re_s = \rho_s L^2 \omega / \mu$, sometimes called a Stokes number, which quantifies the typical ratio between the rate of change of the swimmer momentum and the magnitude of the viscous forces in the fluid.

For small amount of inertia, the breakdown of the scallop theorem occurs either continuously or discontinuously with these Reynolds numbers depending on the geometrical symmetries in the reciprocal actuation. In the case of
symmetric shapes — typically simple flappers — experiments and modeling demonstrated that a finite, order one, amount of inertia is necessary, indicating a discontinuous transition through an inertial hydrodynamic instability \[34, 85–89\] (Fig. 1i). As a difference, in the case of asymmetric shapes or actuation, the transition is continuous, with locomotion occurring either as some power of Re \[90\] or both Re and Re \[371x687\] (with Re/Re \[380x689\] constant) \[35, 91\] (Fig. 1j). Interestingly, for asymmetric shapes, a continuous transition with swimmer inertia was obtained in the absence of fluid inertia \(Re = Re_\omega = 0\), with locomotion occurring as powers of Re \[92\].

V. HYDRODYNAMIC INTERACTIONS

The inertialess scallop envisioned by Purcell as the prototypical non-swimmer is isolated in the fluid. It turns out however that hydrodynamic interactions with other such non-swimmers, or more generally flexible entities, can be exploited to swim. Physically, as cells or other synthetic swimming devices do work on the surrounding fluid, they act as hydrodynamic disturbances on the otherwise-quiescent environment, thereby setting up flow fields which are in general dipolar \[11\]. In biology these flow fields have important consequence on the generation of collective modes of locomotion \[93–97\] and rheology at the whole-population level \[98, 99\].

Although a body undergoing reciprocal motion cannot swim, two bodies undergoing reciprocal motion with nontrivial phase differences are able to take advantage of the unsteady hydrodynamic flows they create to undergo nonzero collective and relative dynamics; there is thus no many-scallop theorem \[100, 101\]. As each reciprocal swimmer behaves in general as an unsteady dipole, the collective effect arises from the time-rectification of such unsteadiness, and thus decays generically as \(1/d^3\), where \(d\) is the typical swimmer-swimmer distance (or even faster if additional geometrical symmetries are present \[100, 102\]). Naturally, two reciprocal non-swimmers taken as a whole are not unlike a single non-reciprocal swimmer, although the qualitative details of their locomotion do differ \[100\].

Experimentally, this effect was demonstrated for hydrodynamic interactions between a rigid flapper, beating in a reciprocal fashion, and a flexible boundary (free surface). The rectification of the reciprocal flow by the free surface motion leads to flow and forces scaling quadratically with the applied flapping frequency, and the creation of a reciprocal pump \[36\] (Fig. 1k). The experimental application of these ideas to a collection of free-swimming bodies remains however to be confirmed. To generate reciprocal motion with nontrivial phase-differences, one possibility would be to use elastic field-responsive particles under a uniform AC forcing; particles with different relaxation times would respond to fields with different phases, and thus would be able to move collectively \[100\]. In the case of purely identical non-swimmers, two of them cannot swim, but three or more are able to move \[102\]. In that case, the phase differences in body kinematics are induced by hydrodynamic flows, leading to a slow \(1/d^7\) effect \[102\].

VI. PHYSICAL ENVIRONMENT

In the scallop theorem, the assumption that locomotion takes place in a Newtonian environment is crucial, as it allows the inertialess equations of fluid motion to be linear and independent of time. A change of the mechanical and rheological properties of the fluid would however naturally lead to a different type of conclusion. Complex fluids are abundant in biology, and cell locomotion often takes place in strongly elastic polymeric fluids \[103–111\], which has been the focus of much recent work \[106, 111\].

As the fluid becomes non-Newtonian, three different physical effects can potentially be exploited to generate small-scale locomotion \[112, 113\]. First, complex fluids possess in general rheological properties which are rate dependent. In particular, viscosities often display shear-thinning behavior, meaning they decrease with shear rates. In this type of fluid, and in contrast with the Newtonian case, the rate at which the reciprocal sequence of shapes is being displayed would matter, a result which could be used to design a reciprocal swimmer. This was recently demonstrated theoretically for bodies swimming using a reciprocal helical actuation at different rates in model polymeric fluids \[109\].

The second physical effect to be exploited is that of normal stress differences, which arise from the stretching by the flow of the microstructure suspended in the complex fluids. Normal stress differences scale quadratically with the
applied shear [112] and remain thus identical under a reversal a time, allowing propulsion. Locomotion using normal stress differences was demonstrated theoretically for a three-dimensional body undergoing small-amplitude reciprocal motion at constant rate [110]. The generation of forces and flow by reciprocal flapping was also reported [37, 114] (Fig. 1l).

The last physical effect to be exploited is that of stress relaxation. Even for small-amplitude motion and linearized dynamics, the simplest evolution equation for the stress in a polymeric fluid contains a memory term in the form of a partial time derivative times a relaxation time. Whether, even in the linear regime, stress relaxation can be taken advantage of for locomotion purposes is an intriguing, but yet unexplored, possibility.

VII. CONCLUSION

In this short review, we have used Purcell’s scallop theorem as a framework to lay out the basic physical principles behind the design of small-scale swimming devices. We have shown how non-reciprocal kinematics, inertia, hydrodynamic interactions, and the nature of surrounding environment can all be physically exploited to achieve small-scale propulsion. With advances in micro- and nano-fabrication, the discussion on the theorem can now move from that akin to a mathematical exercise to a true engineering challenge.

As briefly mentioned in Ref. [13], there exists at least another class of body motion which always leads to zero locomotion in a Newtonian fluid, namely those for which the time-reversal of the motion is identical to its mirror-image (for example, the motion of a rod sweeping the envelope of a cone). The formal derivation of the complete class of non-swimming body kinematics would provide a new thrust in small-scale locomotion research by allowing novel opportunities to get around these mathematical constraints.

Acknowledgements

Discussions with Denis Bartolo are gratefully acknowledged. I thank B. Chan, A. E. Hosoi, A. Aubusson-Fleury and the CNRS Photothèque for providing me with the images reproduced in Fig. 1c and d, as well as the authors from Refs. [26, 27, 29–37] who gave us permission to reproduce their images. This work was supported in part by the US National Science Foundation through grant number CBET-0746285.

[1] D. Bray, Cell Movements (Garland Publishing, New York, NY, 2000).
[2] G. T. Yates, Am. Sci. 74, 358 (1986).
[3] B. D. Hoffman and J. C. Crocker, Annu. Rev. Biomed. Eng. 11, 250 (2009).
[4] J. Happel and H. Brenner, Low Reynolds Number Hydrodynamics (Prentice Hall, Englewood Cliffs, NJ, 1965).
[5] S. Kim and J. S. Karilla, Microhydrodynamics: Principles and Selected Applications. (Butterworth-Heinemann, Boston, MA, 1991).
[6] G. I. Taylor, Proc. Roy. Soc. A 209, 447 (1951).
[7] G. I. Taylor, Proc. Roy. Soc. A 211, 225 (1952).
[8] G. J. Hancock, Proc. Roy. Soc. Lond. A 217, 96 (1953).
[9] J. Gray and G. J. Hancock, J. Exp. Biol. 32, 802 (1955).
[10] J. Lighthill, Mathematical Biofluidodynamics (SIAM, Philadelphia, 1975).
[11] J. Lighthill, SIAM Rev. 18, 161 (1976).
[12] C. Brennen and H. Winet, Ann. Rev. Fluid Mech. 9, 339 (1977).
[13] S. Childress, Mechanics of Swimming and Flying (Cambridge University Press, Cambridge U.K., 1981).
[14] L. J. Fauci and R. Dillon, Ann. Rev. Fluid Mech. 38, 371 (2006).
[15] E. Lauga and T. R. Powers, Rep. Prog. Phys. 72, 096601 (2009).
[16] E. M. Purcell, Am. J. Phys. 45, 3 (1977).
[17] A. Shapere and F. Wilczek, Phys. Rev. Lett. 58, 2051 (1987).
[59] C. H. Wiggins and R. E. Goldstein, Phys. Rev. Lett. 80, 3879 (1998).
[60] C. P. Lowe, Philos. Trans. R. Soc. London, Ser. B 358, 1543 (2003).
[61] M. C. Lagomarsino, F. Capuani, and C. P. Lowe, J. Theor. Biol. 224, 215 (2003).
[62] E. Lauga, Phys. Rev. E 75, 041916 (2007).
[63] T. S. Yu, E. Lauga, and A. E. Hosoi, Phys. Fluids 18, 091701 (2006).
[64] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[65] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[66] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[67] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[68] E. Lauga, Phys. Rev. E 75, 041916 (2007).
[69] T. S. Yu, E. Lauga, and A. E. Hosoi, Phys. Fluids 18, 091701 (2006).
[70] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[71] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[72] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[73] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[74] E. Lauga, Phys. Rev. E 75, 041916 (2007).
[75] T. S. Yu, E. Lauga, and A. E. Hosoi, Phys. Fluids 18, 091701 (2006).
[76] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[77] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[78] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[79] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[80] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[81] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[82] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[83] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[84] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[85] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[86] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[87] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[88] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[89] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[90] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[91] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[92] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[93] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[94] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[95] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[96] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[97] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[98] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[99] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[100] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[101] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[102] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[103] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[104] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[105] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[106] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).
[107] M. Manghi, X. Schlagberger, and R. R. Netz, Phys. Rev. Lett. 96, 068101 (2006).
[108] C. W. Wolgemuth, T. R. Powers, and R. E. Goldstein, Phys. Rev. Lett. 84, 1623 (2000).
[109] S. Lim and C. S. Peskin, SIAM J. Sci. Comput. 25, 2066 (2004).
[110] H. Wada and R. R. Netz, Europhys. Lett. 75, 645 (2006).