

Review Article

Coordination of eukaryotic cilia and flagella

Kirsty Y. Wan^{1,2}

¹Living Systems Institute, University of Exeter, Exeter, U.K.; ²College of Engineering Mathematics and Physical Sciences, University of Exeter, Exeter, U.K.

Correspondence: Kirsty Y. Wan (K.Y.Wan2@exeter.ac.uk)



Propulsion by slender cellular appendages called cilia and flagella is an ancient means of locomotion. Unicellular organisms evolved myriad strategies to propel themselves in fluid environments, often involving significant differences in flagella number, localisation and modes of actuation. Remarkably, these appendages are highly conserved, occurring in many complex organisms such as humans, where they may be found generating physiological flows when attached to surfaces (e.g. airway epithelial cilia), or else conferring motility to male gametes (e.g. undulations of sperm flagella). Where multiple cilia arise, their movements are often observed to be highly coordinated. Here I review the two main mechanisms for motile cilia coordination, namely, *intracellular* and *hydrodynamic*, and discuss their relative importance in different ciliary systems.

Introduction

A major signature of living organisms is their ability to generate and coordinate movement. Even plants, which are often considered to have a static existence, exhibit purposeful, directed movement for growth optimisation (e.g. circumnutation) [1]. Throughout evolution, multiple divergent strategies for locomotion have arisen in land, air and sea, including swimming, crawling, galloping and flying. By far the most ancient of these is locomotion through a fluid environment (known as motility), which gave motile organisms the ability to navigate towards favourable conditions (light, nutrients), and consequently a significant selective advantage over their non-motile counterparts. Here, we focus on motility associated with surface-attached appendages known as *cilia*, or interchangeably *flagella*. This ubiquitous and evolutionarily successful organelle appears in virtually all extant branches of eukaryotes [2]. Figure 1 illustrates a number of different species spanning several orders of magnitude in size, which invariably use cilia to swim or to move fluid from one region to another. At the unicellular extreme, spermatozoa propagate bending waves from base to tip to push themselves through the fluid [3], the biflagellate green alga *Chlamydomonas* coordinates two flagella in a synchronous breaststroke [4–6], while larger ciliates including *Paramecium* and *Stentor* are covered in cilia which have become specialised either for efficient swimming or feeding [7]. In the ventricles of the brain, cilia generate a complex, directional transport network for precise control of substance redistribution [8] and even provides structural support [9], while in the human trachea, the coordinated sweeping of cilia clears debris and mucus up and out of the lungs over distances of tens of centimetres [10,11]. Do these diverse ciliary systems rely upon unified mechanistic principles to achieve coordinated patterns of activity?

The green algal flagellates, in particular, have emerged as a preferred model system not only for studying the structural biology of cilia and their relation to human ciliopathies [22–25] but also the fluid physics of cilia-driven flows [16,26–28]. Here, we focus on select species of microalgae that exhibit significant differences in size, number and spatial organisation of their flagella, and which, presumably through adaptation to various ecological niches, are able to produce a surprising diversity of swimming gaits. These species encompass single-celled organisms with a diameter of 10 μm or less, but also larger, multicellular species such as *Volvox* spp which are several hundreds of microns in size. As we shall see, their distinct ciliary coordination patterns could only have arisen from a delicate interplay between two very different physical mechanisms.

Received: 28 August 2018
Revised: 04 October 2018
Accepted: 08 October 2018

Version of Record published:
21 November 2018

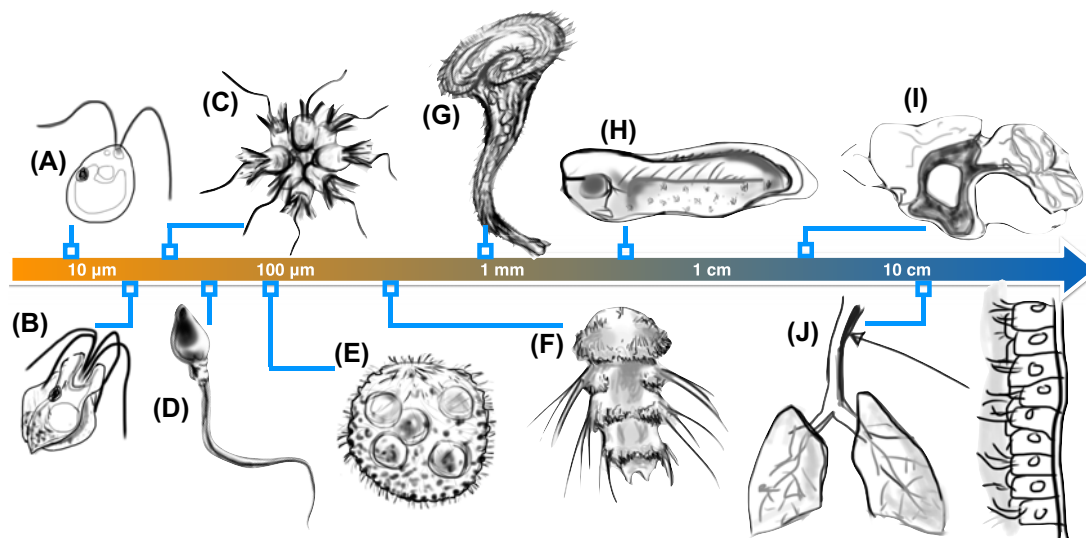


Figure 1. Unity and diversity of ciliary systems

The same fundamental structure occurs in the tiniest of microorganisms as well as ciliated tissues, but exhibits drastic differences in number and localisation. Examples include (A) the algal biflagellate *Chlamydomonas reinhardtii* [4], (B) a quadriflagellate Prasinophyte alga *Pyramimonas* sp. [12], (C) rosette-forming choanoflagellates [13,14], (D) human sperm [15], (E) the spherical alga *Volvox carteri* [16,17], (F) the ciliated larvae of the marine annelid *Platynereis dumerilii* which have segmental multiciliated cells, and long, stiff chaetae [18], (G) the trumpet-shaped ciliate *Stentor coeruleus* [19], (H) ciliated epithelia of *Xenopus laevis* embryos [20], (I) ependymal cilia in mouse brain ventricles which direct cerebrospinal fluid flows [8] (cilia are localised to shaded region) and (J) multiciliated columnar cells in the human trachea [10,21].

Swimming with cilia and flagella

The act of waving an appendage through a fluid creates a local disturbance of decaying magnitude away from the source [29]. Micron-sized organisms dwell in a regime dominated by viscous effects, in which there is zero inertial coasting, so that they must employ very different mechanisms of self-propulsion compared with larger organisms such as fish. Consequently, cilia have evolved to harness drag-based propulsion [30]. There is a surge of recent interest in mimicking the success of this design in the manufacture of artificial robotic microswimmers for biomedical applications [31].

Self-propelling bodies in the viscous (so-called low Reynolds number) regime experience no net forces or torques, so that motion is completely specified by their shape kinematics. There are two main considerations for generating net propulsion, first, a cilium's characteristic slender shape ensures drag-anisotropy when moving through the fluid, second, microorganisms actively prescribe a time-varying distribution of bending moments along the cilium to generate a cyclical, but importantly non-reciprocal sequence of shape changes – in other words a *stroke*. A typical ciliary beat consists of a power stroke, in which the long axis is perpendicular to the direction of motion, and a recovery stroke, in which it is much more curved, and aligned with the direction of motion. Eukaryotic cilia and flagella are distinctive in their capacity to propagate large-amplitude bending waves of activity. Indeed, the observation of slow amplitude decay first led Machin [32] to postulate the existence of active force generating components distributed along the entire length of the filament, before the first experiments were conducted showing these putative components to be dyneins (from the Greek for 'force') residing inside the axoneme. The precise mechanism by which distributed dynein activity leads to emergence of ciliary beating is still not fully understood, and remains a highly active field of research [33–40].

Individual as well as groups of cilia and flagella show a remarkable sensitivity and mutability to extracellular as well as intracellular perturbations. For multiple cilia, different propulsion modes are produced by careful modulation of the phase difference between the periodic strokes of neighbouring cilia: zero phase difference for the biflagellate breaststroke, or a fixed, non-zero phase difference for metachronal waves in ciliary arrays.

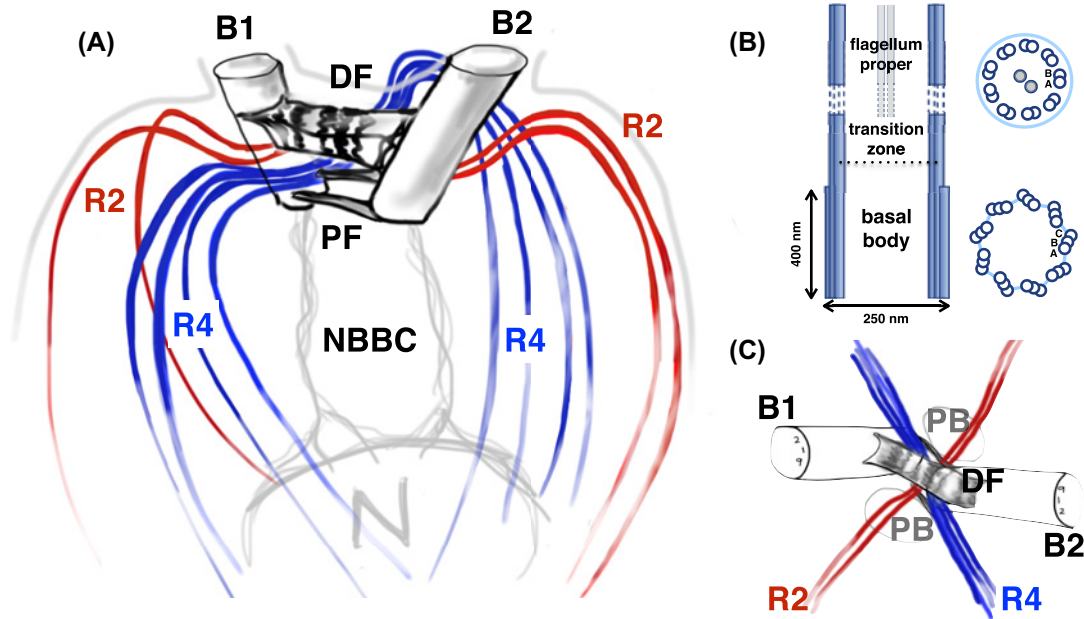


Figure 2. The *Chlamydomonas* flagellar apparatus

(A) Schematic showing the cytoskeletal architecture of basal bodies (B1,2), microtubular roots (two-membered rootlets R2 and four-membered rootlets R4), and fibrous/contractile connections (NBBCs to the nucleus, proximal and distal striated fibres PF and DF). (B) Longitudinal section of a flagellum showing a structural change from triplet microtubules to doublets, two characteristic cross-sections are shown: one through the basal body and the second through flagellum proper. (C) Top view, highlighting radial symmetries in the flagellar apparatus, the locations of the two PB, the cruciate arrangement of microtubule bundles and the DF connecting specific microtubule doublets in the mature basal bodies (B1,2),

Centrioles in the algal flagellar apparatus

Across diverse evolutionary phyla, a high degree of conservation pertains not only to the cilium itself but also to the centrioles to which these microtubule-based organelles are attached [41,42]. Centrioles are present in many unicellular eukaryotes but lost in most land plants [43], and have been studied extensively in several model species including *Chlamydomonas reinhardtii*, *Drosophila melanogaster*, *Caenorhabditis elegans* and human cell lines [44–46], and it is to *C. reinhardtii* in particular that we owe much of our knowledge of centriole assembly and the microtubular composition of cilia and flagella [25,47]. Centrioles also serve as the structural basis of centrosomes, which in turn organise spindles during cell division and regulate cytokinesis [48]. Centrosomal duplication is tightly coupled to the cell-cycle [49–51], and in mammalian cells are intimately involved in cell proliferation, migration and determining polarity [52,53]. Mature centrioles, in this context then known as *basal bodies*, can dock to the plasma membrane (Figure 2) where they are responsible for templating sensory or primary cilia for signal transduction, or motile flagella for cell motility.

Eukaryotic centrioles/basal bodies are cylindrical, have a chiral arrangement of triplet microtubules (termed A, B, C) which become doublets extending into the axoneme proper, giving rise to a distinctive nine-fold symmetry. In the flagellar apparatus of *C. reinhardtii*, individual triplets also exhibit a gradual longitudinal twist going from the basal to distal end [54]. *C. reinhardtii* flagella are approx. 12 μm long and 250 nm wide, and have a ubiquitous structure comprising nine doublet microtubules encircling a central pair, which consists of two microtubules (Figure 2B). Basal bodies associated with such ‘9 + 2’ axonemes are thought to be the ancestral form present in the common unicellular ancestor of eukaryotes [44]. The C tubules terminate in the transition zone, which gates protein entry into the axoneme, while A and B tubules terminate in the distal part of the flagellum [55]. The peripheral doublets are transiently linked by tens of thousands of dynein motors, regulatory components and other complexes in 96 nm repeating units [56], and beating occurs through the distributed activity of these various dynein isoforms [57]. Signalling and mechanical interactions via the central pair/radial spokes are also thought to be involved in axonemal beat modulation [40,58].

Algal basal bodies are particularly important for organising the cytoskeleton and for organellar placement [59,60]. In interphase, *C. reinhardtii* cells have two mature flagella-bearing basal bodies (~400 nm) and two nascent/pro-basal bodies (PB, ~86 nm), anchored in a fixed orientation (Figure 2C). Unlike metazoan centrosomes where mother and daughter centrioles are oriented perpendicularly [41], in *C. reinhardtii* they assume a V-shape [61,62], to facilitate breaststroke swimming. Two types of fibrous structures are present in the flagellar apparatus [63,64]: microtubular roots, and contractile, centrin-based fibres. Each mature basal body has a four-membered and a two-membered microtubule rootlet, containing acetylated α -tubulin, which are assembled to form a cruciate pattern characteristic of green algal flagellates. The four-membered rootlets demarcate the cleavage furrow, and determine the placement of a *de novo* assembled eyespot (photosensor) near the daughter basal body. Each cell has a unique eyespot, thereby breaking bilateral symmetry. A large distal striated fibre (DF) connects the two mature basal bodies just below the transition zone. Fibrous bundles called nuclear basal body connectors (NBBCs) connect the basal bodies to the nucleus [65]. The precise functions of many of these accessory structures remain to be further elucidated [66,67].

More than one cilium – a question of coordination

Given all of the above, how do cilia and flagella cooperate to optimise the movement of fluid given physical constraints such as placement of basal bodies within a precisely defined cytoskeletal architecture? In the human ciliopathy known as primary ciliary dyskinesia (PCD), mutations were first identified in ciliary ultrastructure, but variants of the syndrome have also been found that are due entirely to disorientation of the cilia [68]. Likewise, in unicellular protists such as *Tetrahymena* or *Paramecium* the main function of the ciliate cortex is to order basal bodies and nucleate cilia for motility. In all these cases, maintaining correct ciliary orientation and coordination is crucial. Neighbouring cilia which inhabit a shared fluid environment must interact hydrodynamically due to their physical proximity, but cilia may also be constrained intracellularly, which of these two (possibly antagonistic) contributions dominate can only be ascertained on a case-by-case basis.

Proof of hydrodynamic interactions

The colonial alga *Volvox* (Figure 1E) holds clues to the ancient evolutionary origins of multicellularity [69]; spheroids have two cell types, large germ cells in the interior, and thousands of flagellated somatic cells adorning the surface are oriented with their basal bodies directed away from the A-P axis to produce large-scale flows and directed swimming [70]. Individual flagellated somatic cells, when isolated from their parental colonies and placed in pairs with different relative beat orientations on separate micropipettes, can synchronize their beating purely as a result of interactions through the fluid [71]. Pairs of cells with parallel stroke orientations exhibited in-phase synchrony, but anti-phase synchrony when the power strokes faced away from each other (Figure 3A,B). Moreover, the decay of synchronization with increasing distance of separation was shown to be functionally consistent with hydrodynamic predictions [71]. Theoretical models can also account for the emergence of phase synchrony and even large-scale metachronal waves of ciliary activity, consistent with experimental observations [17,72,73]. A corollary of this is that flagellar beating must be compliant: mechanical forces and hydrodynamic loading can alter the engagement of axonemal dyneins, thereby changing the beating waveform and frequency. The physics of these non-local, feedback-driven phenomena is currently under further investigation [74–77]. More recently, hydrodynamic stresses have even been suggested as a mechanism for ultrafast and efficient cell–cell communication in the protist *Spirostomum ambiguum* [78].

Proof of intracellular control

Contrary to the case of *Volvox*, the in-phase synchronous breaststroke of *Chlamydomonas* (Figure 3D–F) could not be reconciled with the same hydrodynamic theory (see also [79]). Instead, can fibrous connections in the algal flagellar apparatus, which not only lie in the plane of flagellar beating, but are even found attached to each basal body at specific numbered microtubule doublets, provide additional intracellular coupling [61,65,80]? Recent experiments examining the motility phenotypes of *vfl-3* mutants – which have a variable number of full-length, fully motile flagella ($0 \leq N \leq 5$), suggest this is indeed the case [12]. In this mutant, flagella are also found in aberrant positions and orientations [81], but importantly are missing or defective in the distal striated fibre (recall Figure 2). No in-phase synchronous breaststrokes were observed, indicating that internal coupling must have been necessary to coordinate the biflagellate breaststroke of wildtype cells. Indeed, flagella in certain configurations (e.g. triplets) exhibited hydrodynamic synchronisation in the absence of these intracellular connections (Figure 4A).

Notably, it has been predicted in multiple theoretical models that changes in the amount of sliding between microtubule doublets at the flagellar base or other boundary conditions can effect global changes in the beating dynamics

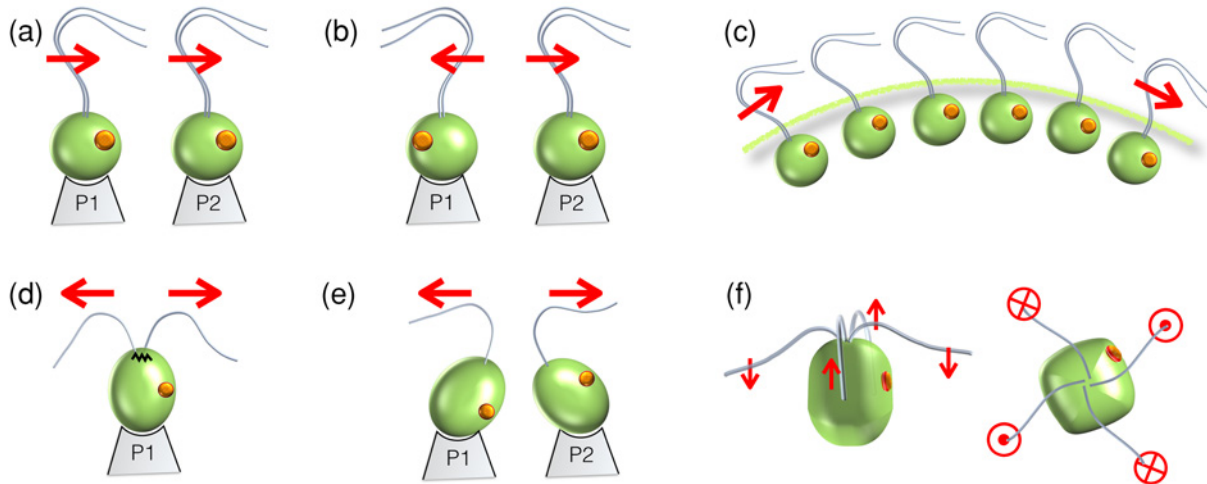


Figure 3. Experimental versus natural configurations of algal flagella

In all cases, the conspicuous orange dots represent algal eyespots (rudimentary photoreceptors), whose positioning is related to the developmental age of the flagella. A pair of *V. carteri* somatic cells held on nearby micropipettes, interacting hydrodynamically, exhibits either in-phase synchrony (A) or anti-phase synchrony (B) depending on their relative orientation. (C) *In vivo*, arrays of these cells coordinate metachronal waves in the *Volvox* colony. By contrast, the in-phase breaststroke of *C. reinhardtii* (D) cannot be reproduced in pairs of wildtype cells that have been rendered uniflagellate (E), implicating an internal (possibly spring-like) coupling provided by the distal striated fibre. Arrows indicate power stroke directions (A–E). (F) In a different species (see also Figure 1B), a quadriflagellate beat pattern (aka trot) is observed, comprising two pairs of breaststrokes displaced temporally by 1/2 beat cycle.

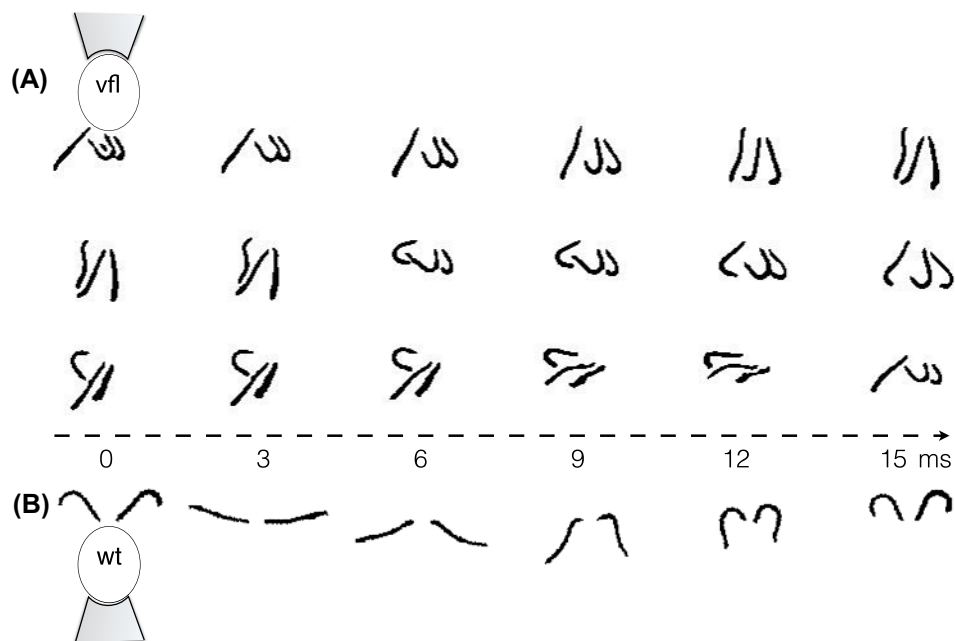


Figure 4. Aberrant flagellar coordination patterns in a basal-coupling mutant.

Flagellar waveforms and dynamics are tracked in a tri-flagellate *vfl-3* mutant (A) and wildtype *C. reinhardtii* (B). (Cell body/pipette not shown on subsequent image frames.) The distal striated fibre is missing or defective in *vfl-3* [81]. Consequently, coordination between the three flagella reverts to hydrodynamic interactions, in which the pair beating with power strokes in the same direction tends to synchronize in in-phase, but in anti-phase with respect to the third singlet flagellum. The wildtype cell on the other hand, maintains an in-phase synchronous breaststroke. (For further examples, see [12].)

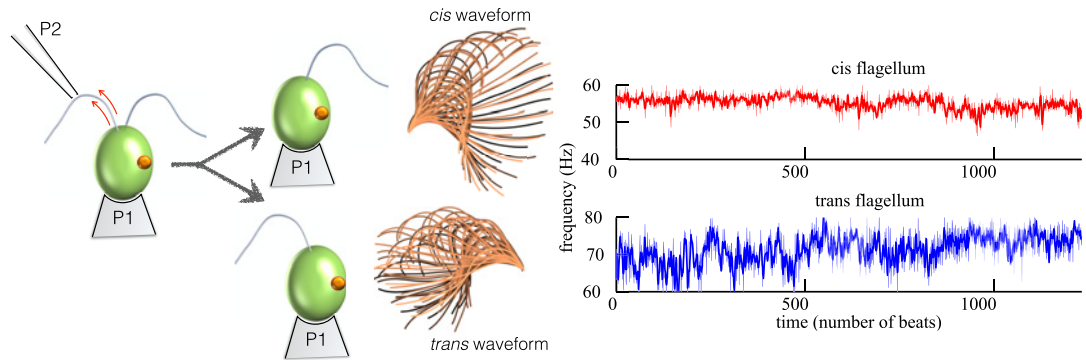


Figure 5. Revealing intrinsic differences between the two *C. reinhardtii* flagella

The two flagella are similar but not identical. *In vivo* micromanipulation and microsurgery revealed a differential sensitivity to deflagellation-induced intracellular calcium elevation – here inferred indirectly by measuring ciliary beat frequencies in the two flagella of the same cell, after successive rounds of deflagellation and regrowth. The stereotypical *trans*-flagellum beating mode is markedly faster, has an attenuated waveform, and exhibits greater frequency variations than the *cis*.

in a non-trivial manner [36,40,82,83]. Such considerations may yet help to explain observations of a novel (biflagellate) anti-phase gait in another *C. reinhardtii* mutant known as *ptx1* [83,84]. More generally, elaborate networks of intracellular structures are associated with the basal bodies of other species of algal flagellates including quadri- and octoflagellates, revealing a surprising diversity of gaits involving precise phase relationships between the flagella [12]. Further investigation into the morphology and function of these fibrous structures may yield fresh insights into the evolution of the algal flagellar apparatus and the origins of active, biochemical gait-control.

The generational age of basal bodies

For many simple eukaryotes, cilia and flagella are used simultaneously for sensing and motility, which leads to one further complication. Organisms must be able to produce asymmetric behavioural changes in response to environmental cues: in addition to directed swimming, they must be able to turn. This is of particular importance in phototaxis – motility with respect to light, where they adjust to photostimuli by altering the beating of cilia/flagella [85,86]. In *C. reinhardtii*, this is accomplished through asymmetric actuation of the two flagella [87], which are termed *cis* or *trans* depending on relative distances from the unique eyespot (Figure 5). Both flagella insert into the apical region of the cell and are of equal length, and have no visible ultrastructural or morphological differences. An inherent centriolar asymmetry exists however, rooted in the difference in generational age of the *cis* versus *trans* basal body; at each round of cell division, the *cis* basal body is always formed anew yet the *trans* is inherited from the parent cell. This sequence of semi-conservative basal body duplication forms the basis of a greater, cell-wide asymmetry [88,89].

At a behavioural level, evidence of differential Ca^{2+} sensitivities in the two *C. reinhardtii* flagella first emerged in demembrated cell models placed in buffers with varying concentrations of calcium [90,91], as well as in early micropipette experiments [4]. More recently, high-speed imaging was used in combination with micromanipulation and microsurgery in *live* cells to decouple the intrinsic beating dynamics of one flagellum from its partner, in which individual *cis*- or *trans*-flagella were carefully removed from each biflagellate cell by induced self-scission (Figure 5), allowing sufficient time for flagellar regeneration between the successive amputations [6]. Attributing these *in vivo* flagellar responses to deflagellation-induced calcium elevation [92], isolated *cis*-flagella were found to beat with a lower mean frequency as well as lower frequency noise [93] than isolated *trans*-flagella. Biochemical explanations for these differences remain elusive, but entry of signalling proteins into a basal body may well depend on its generational age [35]. In the normal wildtype breaststroke (a putative lower calcium state), intrinsic *cis*–*trans* frequency and amplitude differences are presumed sufficiently suppressed [91] to enable intracellular fibres to lock the two flagella into synchrony.

Conclusions

For many eukaryotes, the importance of the centriole/basal body complex is well established. Highly conserved basal body and axonemal architectures are thought to have originated in unicellular organisms that adopted the very first proto-cilium for motility [44,94]. The algal flagellates display a great diversity in form yet conservatism in function,

emphasising their value in helping us disentangle the ancient evolutionary and phylogenetic origins of the flagellar apparatus. By performing comparative studies across different species spanning extremes in size and complexity, causal relationships between ultrastructure and motile behaviour were deduced, uncovering two distinct mechanisms for achieving robust beat coordination that are exemplified by dominance of *hydrodynamic interactions* in the colonial *Volvox*, and of *intracellular coupling* in the unicellular *Chlamydomonas*.

In metazoa however, or even in the larger ciliates, causality is not usually so clear-cut and additional mechanisms must also be considered [95]. In *Platynereis* larvae for example (Figure 1F), ciliary coordination is closely gated by neural activity [18]. In *Tetrahymena*, striated kinetodesmal fibres help maintain basal body orientation and resist hydrodynamic stresses produced by ciliary beating [96], while in the mouse trachea, the apical cytoskeleton was found to be necessary for correct basal body alignment [97]. Conversely, ciliary flows are implicated in the establishment of order: ciliary patterning in the larval skin of *Xenopus* is a two-stage process in which basal bodies are first oriented relative to the tissue axis but subsequently refined through feedback by extracellular fluid flows [98], in mouse brain ventricles however, basal bodies begin randomly docked, before a coupling between hydrodynamic forces and intracellular Planar Cell Polarity signalling sets the correct orientation [99]. With these more confounding contexts in mind, further study of flagellar coordination in algae and other protists will help reveal with greater clarity, the physics behind this complex interplay between flows, mechanical coupling and intracellular signalling.

Summary

- Motile cilia and flagella occur in diverse biological systems.
- Multiple cilia can interact intracellularly as well as hydrodynamically.
- Passive hydrodynamic interactions can drive the beating of nearby filaments into synchrony or metachrony, e.g. in ciliary arrays.
- In unicellular algae with only a few front-mounted flagella, additional coupling is instead provided by contractile fibres, implicating an essential role of the flagellar apparatus for the coordination and control of appendages.

Acknowledgements

I thank Raymond E. Goldstein, Marco Polin, Douglas R. Brumley and Kyriacos C. Leptos for collaborations and support, and other former colleagues from the Goldstein Lab at DAMTP, University of Cambridge, whose work have been discussed here. I am grateful to Gáspár Jékely for a critical reading of the manuscript.

Funding

This work was supported by a startup grant from the Living Systems Institute, University of Exeter, and a Springboard Award from the Academy of Medical Sciences.

Competing Interests

The author declares that there are no competing interests associated with the manuscript.

Abbreviations

DF, distal fibre; NBBC, nuclear basal body connector; PF, proximal fibre; PB, pro-basal bodies.

References

- 1 Darwin, C.R. (1875) *The Movements and Habits of Climbing Plants*, John Murray, London
- 2 Mitchell, D.R. (2007) The evolution of eukaryotic cilia and flagella as motile and sensory organelles. *Adv. Exp. Med. Biol.* **607**, 130–140, https://doi.org/10.1007/978-0-387-74021-8_11
- 3 Ishimoto, K., Gadelha, H., Gaffney, E.A., Smith, D.J. and Kirkman-Brown, J. (2017) Coarse-graining the fluid flow around a human sperm. *Phys. Rev. Lett.* **118**, 124501, <https://doi.org/10.1103/PhysRevLett.118.124501>
- 4 Ruffer, U. and Nultsch, W. (1987) Comparison of the beating of *cis*-flagella and *trans*-flagella of *Chlamydomonas* cells held on micropipettes. *Cell Motil. Cytoskeleton* **7**, 87–93, <https://doi.org/10.1002/cm.970070111>

- 5 Goldstein, R.E., Polin, M. and Tuval, I. (2009) Noise and synchronization in pairs of beating eukaryotic flagella. *Phys. Rev. Lett.* **103**, 168103, <https://doi.org/10.1103/PhysRevLett.103.168103>
- 6 Wan, K.Y., Leptos, K.C. and Goldstein, R.E. (2014) Lag, lock, sync, slip: the many ‘phases’ of coupled flagella. *J. R. Soc. Interface* **11**, 20131160, <https://doi.org/10.1098/rsif.2013.1160>
- 7 Tang, S. K.Y. and Marshall, W.F. (2017) Self-repairing cells: how single cells heal membrane ruptures and restore lost structures. *Science* **356**, 1022–1025, <https://doi.org/10.1126/science.aam6496>
- 8 Faubel, R., Westendorf, C., Bodenschatz, E. and Eichele, G. (2016) Cilia-based flow network in the brain ventricles. *Science* **353**, 176–178, <https://doi.org/10.1126/science.aae0450>
- 9 Mahuzier, A., Shihavuddin, A., Fournier, C., Lansade, P., Faucourt, M., Menezes, N. et al. (2018) Ependymal cilia beating induces an actin network to protect centrioles against shear stress. *Nat. Commun.* **9**, 2279, <https://doi.org/10.1038/s41467-018-04676-w>
- 10 Smith, D.J., Gaffney, E.A. and Blake, J.R. (2008) Modelling mucociliary clearance. *Respir. Physiol. Neurobiol.* **163**, 178–188, <https://doi.org/10.1016/j.resp.2008.03.006>
- 11 Leopold, P.L., O’Mahony, M.J., Lian, X.J., Tilley, A.E., Harvey, B.G. and Crystal, R.G. (2009) Smoking is associated with shortened airway cilia. *PLoS One* **4**, e8157, <https://doi.org/10.1371/journal.pone.0008157>
- 12 Wan, K.Y. and Goldstein, R.E. (2016) Coordinated beating of algal flagella is mediated by basal coupling. *Proc. Natl. Acad. Sci. USA* **113**, E2784–E2793, <https://doi.org/10.1073/pnas.1518527113>
- 13 Alegado, R.A., Brown, L.W., Cao, S.G., Dermenjian, R.K., Zuzov, R., Fairclough, S.R. et al. (2012) A bacterial sulfonolipid triggers multicellular development in the closest living relatives of animals. *Elife* **1**, UNSP e00013, <https://doi.org/10.7554/eLife.00013>
- 14 Kirkegaard, J.B., Marron, A.O. and Goldstein, R.E. (2016) Motility of colonial choanoflagellates and the statistics of aggregate random walkers. *Phys. Rev. Lett.* **116**, 038102, <https://doi.org/10.1103/PhysRevLett.116.038102>
- 15 Gaffney, E.A., Gadelha, H., Smith, D.J., Blake, J.R. and Kirkman-Brown, J.C. (2011) Mammalian sperm motility: observation and theory. *Annu. Rev. Fluid Mech.* **43**, 501–528, <https://doi.org/10.1146/annurev-fluid-121108-145442>
- 16 Solari, C.A., Drescher, K., Ganguly, S., Kessler, J.O., Michod, R.E. and Goldstein, R.E. (2011) Flagellar phenotypic plasticity in volvocalean algae correlates with Péclet number. *J. R. Soc. Interface* **8**, 1409–1417, <https://doi.org/10.1098/rsif.2011.0023>
- 17 Brumley, D.R., Polin, M., Pedley, T.J. and Goldstein, R.E. (2015) Metachronal waves in the flagellar beating of *Volvox* and their hydrodynamic origin. *J. R. Soc. Interface* **12**, 20141358, <https://doi.org/10.1098/rsif.2014.1358>
- 18 Veraszto, C., Ueda, N., Bezares-Calderon, L.A., Panzera, A., Williams, E.A., Shahidi, R. et al. (2017) Ciliomotor circuitry underlying whole-body coordination of ciliary activity in the *Platynereis* larva. *Elife* **6**, e26000, <https://doi.org/10.7554/eLife.26000>
- 19 Slabodnick, M.M. and Marshall, W.F. (2014) *Stentor coeruleus*. *Curr. Biol.* **24**, R783–R784, <https://doi.org/10.1016/j.cub.2014.06.044>
- 20 Brooks, E.R. and Wallingford, J.B. (2014) Multiciliated cells. *Curr. Biol.* **24**, R973–R982, <https://doi.org/10.1016/j.cub.2014.08.047>
- 21 Sears, P.R., Thompson, K., Knowles, M.R. and Davis, C.W. (2013) Human airway ciliary dynamics. *Am. J. Physiol. Lung Cell. Mol. Physiol.* **304**, L170–L183, <https://doi.org/10.1152/ajplung.00105.2012>
- 22 Witman, G.B., Rosenbaum, J.L., Berliner, J. and Carlson, K. (1972) *Chlamydomonas flagella*. 1. Isolation and electrophoretic analysis of microtubules, matrix, membranes, and mastigonemes. *J. Cell Biol.* **54**, 507–539, <https://doi.org/10.1083/jcb.54.3.507>
- 23 Pazour, G.J., Dickert, B.L., Vucica, Y., Seeley, E.S., Rosenbaum, J.L., Witman, G.B. et al. (2000) *Chlamydomonas* IFT88 and its mouse homologue, polycystic kidney disease gene tg737, are required for assembly of cilia and flagella. *J. Cell Biol.* **151**, 709–718, <https://doi.org/10.1083/jcb.151.3.709>
- 24 Lehtreck, K.F., Johnson, E.C., Sakai, T., Cochran, D., Ballif, B.A., Rush, J. et al. (2009) The *Chlamydomonas reinhardtii* BBSome is an IFT cargo required for export of specific signaling proteins from flagella. *J. Cell Biol.* **187**, 1117–1132, <https://doi.org/10.1083/jcb.200909183>
- 25 Silflow, C.D. and Lefebvre, P.A. (2001) Assembly and motility of eukaryotic cilia and flagella. Lessons from *Chlamydomonas reinhardtii*. *Plant Physiol.* **127**, 1500–1507, <https://doi.org/10.1104/pp.010807>
- 26 Hill, N.A. and Bees, M.A. (2002) Taylor dispersion of gyrotactic swimming micro-organisms in a linear flow. *Phys. Fluids* **14**, 2598–2605, <https://doi.org/10.1063/1.1458003>
- 27 Pedley, T.J. and Kessler, J.O. Hydrodynamic phenomena in suspensions of swimming microorganisms. *Annu. Rev. Fluid Mech.* **24**, 313–358, <https://doi.org/10.1146/annurev.fl.24.010192.001525>
- 28 Goldstein, R.E. (2015) Green algae as model organisms for biological fluid dynamics. *Annu. Rev. Fluid Mech.* **47**, 343–375, <https://doi.org/10.1146/annurev-fluid-010313-141426>
- 29 Drescher, K., Goldstein, R.E., Michel, N., Polin, M. and Tuval, I. (2010) Direct measurement of the flow field around swimming microorganisms. *Phys. Rev. Lett.* **105**, 168101, <https://doi.org/10.1103/PhysRevLett.105.168101>
- 30 Lauga, E. and Powers, T.R. (2009) The hydrodynamics of swimming microorganisms. *Rep. Prog. Phys.* **72**, 096601, <https://doi.org/10.1088/0034-4885/72/9/096601>
- 31 Nelson, B.J., Kaliakatos, I.K. and Abbott, J.J. (2010) Microrobots for minimally invasive medicine. *Annu. Rev. Biomed. Eng.* **12**, 55–85, <https://doi.org/10.1146/annurev-bioeng-010510-103409>
- 32 Machin, K.E. (1958) Wave propagation along flagella. *J. Exp. Biol.* **35**, 796–806
- 33 Edwards, B. F.L., Wheeler, R.J., Barker, A.R., Moreira-Leite, F.F., Gull, K. and Sunter, J.D. (2018) Direction of flagellum beat propagation is controlled by proximal/distal outer dynein arm asymmetry. *Proc. Natl. Acad. Sci. USA* **115**, E7341–E7350, <https://doi.org/10.1073/pnas.1805827115>
- 34 Han, J.H. and Peskin, C.S. (2018) Spontaneous oscillation and fluid–structure interaction of cilia. *Proc. Natl. Acad. Sci. USA* **115**, 4417–4422, <https://doi.org/10.1073/pnas.1712042115>
- 35 Ishikawa, T. (2017) Axoneme structure from motile cilia. *Cold Spring Harb. Perspect. Biol.* **9**, a028076, <https://doi.org/10.1101/cshperspect.a028076>
- 36 Oriola, D., Gadelha, H. and Casademunt, J. (2017) Nonlinear amplitude dynamics in flagellar beating. *R. Soc. Open Sci.* **4**, 160698, <https://doi.org/10.1098/rsos.160698>

- 37 Sartori, P., Geyer, V.F., Scholich, A., Julicher, F. and Howard, J. (2016) Dynamic curvature regulation accounts for the symmetric and asymmetric beats of *Chlamydomonas flagella*. *Elife* **5**, e13258, <https://doi.org/10.7554/eLife.13258>
- 38 De Canio, G., Lauga, E. and Goldstein, R.E. (2017) Spontaneous oscillations of elastic filaments induced by molecular motors. *J. R. Soc. Interface* **14**, 20170491, <https://doi.org/10.1098/rsif.2017.0491>
- 39 Bayly, P.V. and Dutcher, S.K. (2016) Steady dynein forces induce flutter instability and propagating waves in mathematical models of flagella. *J. R. Soc. Interface* **13**, 20160523, <https://doi.org/10.1098/rsif.2016.0523>
- 40 Hu, T.C. and Bayly, P.V. (2018) Finite element models of flagella with sliding radial spokes and interdoublet links exhibit propagating waves under steady dynein loading. *Cytoskeleton* **75**, 185–200, <https://doi.org/10.1002/cm.21432>
- 41 Azimzadeh, J. and Marshall, W.F. (2010) Building the centriole. *Curr. Biol.* **20**, R816–R825, <https://doi.org/10.1016/j.cub.2010.08.010>
- 42 Carvalho-Santos, Z., Azimzadeh, J., Pereira-Leal, J.B. and Bettencourt-Dias, M. (2011) Tracing the origins of centrioles, cilia, and flagella. *J. Cell Biol.* **194**, 165–175, <https://doi.org/10.1083/jcb.201011152>
- 43 Hodges, M.E., Scheumann, N., Wickstead, B., Langdale, J.A. and Gull, K. (2010) Reconstructing the evolutionary history of the centriole from protein components. *J. Cell Sci.* **123**, 1407–1413, <https://doi.org/10.1242/jcs.064873>
- 44 Gupta, A. and Kitagawa, D. (2018) Ultrastructural diversity between centrioles of eukaryotes. *J. Biochem. (Tokyo)* **164**, 1–8, <https://doi.org/10.1093/jb/mvy031>
- 45 Mirvis, M., Stearns, T. and Nelson, W.J. (2018) Cilium structure, assembly, and disassembly regulated by the cytoskeleton. *Biochem. J.* **475**, 2329–2353, <https://doi.org/10.1042/BCJ20170453>
- 46 Nigg, E.A. and Raff, J.W. (2009) Centrioles, centrosomes, and cilia in health and disease. *Cell* **139**, 663–678, <https://doi.org/10.1016/j.cell.2009.10.036>
- 47 Satir, P. and Christensen, S.T. (2007) Overview of structure and function of mammalian cilia. *Annu. Rev. Physiol.* **69**, 377–400, <https://doi.org/10.1146/annurev.physiol.69.040705.141236>
- 48 Yubuki, N. and Leander, B.S. (2013) Evolution of microtubule organizing centers across the tree of eukaryotes. *Plant J.* **75**, 230–244, <https://doi.org/10.1111/tbj.12145>
- 49 Stearns, T. (2001) Centrosome duplication: a centriolar pas de deux. *Cell* **105**, 417–420, [https://doi.org/10.1016/S0092-8674\(01\)00366-X](https://doi.org/10.1016/S0092-8674(01)00366-X)
- 50 Bornens, M. (2012) The centrosome in cells and organisms. *Science* **335**, 422–426, <https://doi.org/10.1126/science.1209037>
- 51 Nabais, N., Gomes Pereira, S. and Bettencourt-Dias, M. (2018) Noncanonical biogenesis of centrioles and basal bodies. *Cold Spring Harb. Perspect. Biol.*
- 52 Dawe, H.R., Farr, H. and Gull, K. (2007) Centriole/basal body morphogenesis and migration during ciliogenesis in animal cells. *J. Cell Sci.* **120**, 7–15, <https://doi.org/10.1242/jcs.03305>
- 53 Fu, J.Y., Hagan, I.M. and Glover, D.M. (2015) The centrosome and its duplication cycle. *Cold Spring Harb. Perspect. Biol.* **7**, a015800, <https://doi.org/10.1101/cshperspect.a015800>
- 54 Li, S., Fernandez, J.J., Marshall, W.F. and Agard, D.A. (2012) Three-dimensional structure of basal body triplet revealed by electron cryo-tomography. *EMBO J.* **31**, 552–562, <https://doi.org/10.1038/emboj.2011.460>
- 55 O'Toole, E.T., Giddings, T.H., McIntosh, J.R. and Dutcher, S.K. (2003) Three-dimensional organization of basal bodies from wild-type and delta-tubulin deletion strains of *Chlamydomonas reinhardtii*. *Mol. Biol. Cell* **14**, 2999–3012, <https://doi.org/10.1091/mbc.e02-11-0755>
- 56 Oda, T., Yanagisawa, H., Kamiya, R. and Kikkawa, M. (2014) A molecular ruler determines the repeat length in eukaryotic cilia and flagella. *Science* **346**, 857–860, <https://doi.org/10.1126/science.1260214>
- 57 Lin, J.F. and Nicastro, D. (2018) Asymmetric distribution and spatial switching of dynein activity generates ciliary motility. *Science* **360**, 396, <https://doi.org/10.1126/science.aar1968>
- 58 Barber, C.F., Heuser, T., Carbajal-Gonzalez, B.I., Botchkarev, V.V. and Nicastro, D. (2012) Three-dimensional structure of the radial spokes reveals heterogeneity and interactions with dyneins in *Chlamydomonas flagella*. *Mol. Biol. Cell* **23**, 111–120, <https://doi.org/10.1091/mbc.e11-08-0692>
- 59 Dutcher, S.K. (2003) Elucidation of basal body and centriole functions in *Chlamydomonas reinhardtii*. *Traffic* **4**, 443–451, <https://doi.org/10.1034/j.1600-0854.2003.00104.x>
- 60 Feldman, J.L., Geimer, S. and Marshall, W.F. (2007) The mother centriole plays an instructive role in defining cell geometry. *PLoS Biol.* **5**, e149, <https://doi.org/10.1371/journal.pbio.0050149>
- 61 Ringo, D.L. (1967) Flagellar motion and fine structure of flagellar apparatus in *Chlamydomonas*. *J. Cell Biol.* **33**, 543–+, <https://doi.org/10.1083/jcb.33.3.543>
- 62 O'Toole, E.T. and Dutcher, S.K. (2014) Site-specific basal body duplication in *Chlamydomonas*. *Cytoskeleton* **71**, 108–118, <https://doi.org/10.1002/cm.21155>
- 63 Melkonian, M. (1980) Ultrastructural aspects of basal body associated fibrous structures in green-algae – a critical-review. *Biosystems* **12**, 85–104, [https://doi.org/10.1016/0303-2647\(80\)90040-4](https://doi.org/10.1016/0303-2647(80)90040-4)
- 64 O'Kelly, C.J. and Floyd, G.L. (1983) Flagellar apparatus absolute orientations and the phylogeny of the green algae. *Biosystems* **16**, 227–251, [https://doi.org/10.1016/0303-2647\(83\)90007-2](https://doi.org/10.1016/0303-2647(83)90007-2)
- 65 Geimer, S. and Melkonian, M. (2004) The ultrastructure of the *Chlamydomonas reinhardtii* basal apparatus: identification of an early marker of radial asymmetry inherent in the basal body. *J. Cell Sci.* **117**, 2663–2674, <https://doi.org/10.1242/jcs.01120>
- 66 Salisbury, J.L. (1989) Centrin and the algal flagellar apparatus. *J. Phycol.* **25**, 201–206, <https://doi.org/10.1111/j.1529-8817.1989.tb00114.x>
- 67 Wingfield, J.L. and Lechtreck, K.F. (2018) *Chlamydomonas* basal bodies as flagella organizing centers. *Cells* **7**, 79, <https://doi.org/10.3390/cells7070079>
- 68 Rayner, C. F.J., Rutman, A., Dewar, A., Greenstone, M.A., Cole, P.J. and Wilson, R. (1996) Ciliary disorientation alone as a cause of primary ciliary dyskinesia syndrome. *Am. J. Respir. Crit. Care Med.* **153**, 1123–1129, <https://doi.org/10.1164/ajrccm.153.3.8630555>

- 69 Herron, M.D. (2016) Origins of multicellular complexity: *Volvox* and the volvocine algae. *Mol. Ecol.* **25**, 1213–1223, <https://doi.org/10.1111/mec.13551>
- 70 Kirk, D.L. (1998) *Volvox*, Cambridge University Press
- 71 Brumley, D.R., Wan, K.Y., Polin, M. and Goldstein, R.E. (2014) Flagellar synchronization through direct hydrodynamic interactions. *Elife* **3**, e02750, <https://doi.org/10.7554/eLife.02750>
- 72 Feriani, L., Juenet, M., Fowler, C.J., Bruot, N., Chioccioli, M., Holland, S.M., Bryant, C.E. and Cicuta, P. (2017) Assessing the Collective Dynamics of Motile Cilia in Cultures of Human Airway Cells by Multiscale DDM. *Biophys. J.* **113**, 109–119, <https://doi.org/10.1016/j.bpj.2017.05.028>
- 73 Guo, H.L., Fauci, L., Shelley, M. and Kanso, E. (2018) Bistability in the synchronization of actuated microfilaments. *J. Fluid Mechanics* **836**, 304–323, <https://doi.org/10.1017/jfm.2017.816>
- 74 Elfring, G.J. and Lauga, E. (2009) Hydrodynamic phase locking of swimming microorganisms. *Phys. Rev. Lett.* **103**, 088101, <https://doi.org/10.1103/PhysRevLett.103.088101>
- 75 Klindt, G.S., Ruloff, C., Wagner, C. and Friedrich, B.M. (2016) Load response of the flagellar beat. *Phys. Rev. Lett.* **117**, 258101, <https://doi.org/10.1103/PhysRevLett.117.258101>
- 76 Man, Y., Koens, L. and Lauga, E. (2016) Hydrodynamic interactions between nearby slender filaments. *Epl* **116**, 24002, <https://doi.org/10.1209/0295-5075/116/24002>
- 77 Goldstein, R.E., Lauga, E., Pesci, A.I. and Proctor, M. R.E. (2016) Elastohydrodynamic synchronization of adjacent beating flagella. *Phys. Rev. Fluids* **1**, 073201, <https://doi.org/10.1103/PhysRevFluids.1.073201>
- 78 Mathijssen, A., Culver, J., Saad Bhamla, M. and Prakash, M. (2018) Collective intercellular communication through ultra-fast hydrodynamic trigger waves. *bioRxiv*, URL <https://www.biorxiv.org/content/biorxiv/early/2018/09/26/428573.full.pdf>
- 79 Quaranta, G., Aubin-Tam, M.E. and Tam, D. (2015) Hydrodynamics versus intracellular coupling in the synchronization of eukaryotic flagella. *Phys. Rev. Lett.* **115**, 238101, <https://doi.org/10.1103/PhysRevLett.115.238101>
- 80 Salisbury, J.L. (1988) The lost neuromotor apparatus of *Chlamydomonas* – rediscovered. *J. Protozool.* **35**, 574–577, <https://doi.org/10.1111/j.1550-7408.1988.tb06128.x>
- 81 Wright, R.L., Chojnacki, B. and Jarvik, J.W. (1983) Abnormal basal-body number, location, and orientation in a striated fiber-defective mutant of *Chlamydomonas reinhardtii*. *J. Cell Biol.* **96**, 1697–1707, <https://doi.org/10.1083/jcb.96.6.1697>
- 82 Riedel-Kruse, I.H., Hilfinger, A., Howard, J. and Julicher, F. (2007) How molecular motors shape the flagellar beat. *HFPJ* **1**, 192–208, <https://doi.org/10.2976/1.2773861>
- 83 Klindt, G.S., Ruloff, C., Wagner, C. and Friedrich, B.M. (2017) In-phase and anti-phase flagellar synchronization by waveform compliance and basal coupling. *New J. Phys.* **19**, 113052, <https://doi.org/10.1088/1367-2630/aa9031>
- 84 Leptos, K.C., Wan, K.Y., Polin, M., Tuval, I., Pesci, A.I. and Goldstein, R.E. (2013) Antiphase synchronization in a flagellar-dominance mutant of *Chlamydomonas*. *Phys. Rev. Lett.* **111**, 158101, <https://doi.org/10.1103/PhysRevLett.111.158101>
- 85 Harz, H. and Hegemann, P. (1991) Rhodopsin-regulated calcium currents in *Chlamydomonas*. *Nature* **351**, 489–491, <https://doi.org/10.1038/351489a0>
- 86 Jekely, G., Colombelli, J., Hausen, H., Guy, K., Stelzer, E., Nedelec, F. et al. (2008) Mechanism of phototaxis in marine zooplankton. *Nature* **456**, 395–399, <https://doi.org/10.1038/nature07590>
- 87 Bennett, R.R. and Golestanian, R. (2015) A steering mechanism for phototaxis in *Chlamydomonas*. *J. R. Soc. Interface* **12**, 20141164, <https://doi.org/10.1098/rsif.2014.1164>
- 88 Marshall, W.F. (2012) Centriole asymmetry determines algal cell geometry. *Curr. Opin. Plant Biol.* **15**, 632–637, <https://doi.org/10.1016/j.pbi.2012.09.011>
- 89 Pearson, C.G. (2014) Choosing sides – asymmetric centriole and basal body assembly. *J. Cell Sci.* **127**, 2803–2810, <https://doi.org/10.1242/jcs.151761>
- 90 Bessen, M., Fay, R.B. and Witman, G.B. (1980) Calcium control of waveform in isolated flagellar axonemes of *Chlamydomonas*. *J. Cell Biol.* **86**, 446–455, <https://doi.org/10.1083/jcb.86.2.446>
- 91 Kamiya, R. and Witman, G.B. (1984) Submicromolar levels of calcium control the balance of beating between the 2 flagella in demembrated models of *Chlamydomonas*. *J. Cell Biol.* **98**, 97–107, <https://doi.org/10.1083/jcb.98.1.97>
- 92 Wheeler, G.L. and Brownlee, C. (2008) Rapid spatiotemporal patterning of cytosolic Ca²⁺ underlies flagellar excision in *Chlamydomonas reinhardtii*. *Plant J.* **53**, 401–413, <https://doi.org/10.1111/j.1365-313X.2007.03349.x>
- 93 Wan, K.Y. and Goldstein, R.E. (2014) Rhythmicity, recurrence, and recovery of flagellar beating. *Phys. Rev. Lett.* **113**, 238103, <https://doi.org/10.1103/PhysRevLett.113.238103>
- 94 Satir, P., Mitchell, D.R. and Jekely, G. (2008) How did the cilium evolve? *Curr. Top. Dev. Biol.* **85**, 63–82, [https://doi.org/10.1016/S0070-2153\(08\)00803-X](https://doi.org/10.1016/S0070-2153(08)00803-X)
- 95 Wallingford, J.B. (2010) Planar cell polarity signaling, cilia and polarized ciliary beating. *Curr. Opin. Cell Biol.* **22**, 597–604, <https://doi.org/10.1016/j.ceb.2010.07.011>
- 96 Galati, D.F., Bonney, S., Kronenberg, Z., Clarissa, C., Yandell, M., Elde, N.C. et al. (2014) DisAp-dependent striated fiber elongation is required to organize ciliary arrays. *J. Cell Biol.* **207**, 705–715, <https://doi.org/10.1083/jcb.201409123>
- 97 Herawati, E., Taniguchi, D., Kanoh, H., Tateishi, K., Ishihara, S. and Tsukita, S. (2016) Multiciliated cell basal bodies align in stereotypical patterns coordinated by the apical cytoskeleton. *J. Cell Biol.* **214**, 571–586, <https://doi.org/10.1083/jcb.201601023>
- 98 Mitchell, B., Jacobs, R., Li, J., Chien, S. and Kintner, C. (2007) A positive feedback mechanism governs the polarity and motion of motile cilia. *Nature* **447**, 97–U8, <https://doi.org/10.1038/nature05771>
- 99 Guirao, B., Meunier, A., Mortaud, S., Aguilar, A., Corsi, J.M., Strehl, L. et al. (2010) Coupling between hydrodynamic forces and planar cell polarity orients mammalian motile cilia. *Nat. Cell Biol.* **12**, 341–U86