

This is a repository copy of *The counterbend dynamics of cross-linked filament bundles* and flagella.

White Rose Research Online URL for this paper: <a href="https://eprints.whiterose.ac.uk/id/eprint/117164/">https://eprints.whiterose.ac.uk/id/eprint/117164/</a>

Version: Accepted Version

#### Article:

Coy, Rachel and Gadelha, Hermes Augusto Buarque orcid.org/0000-0001-8053-9249 (2017) The counterbend dynamics of cross-linked filament bundles and flagella. Interface. 20170065. ISSN: 1742-5662

https://doi.org/10.1098/rsif.2017.0065

#### Reuse

Items deposited in White Rose Research Online are protected by copyright, with all rights reserved unless indicated otherwise. They may be downloaded and/or printed for private study, or other acts as permitted by national copyright laws. The publisher or other rights holders may allow further reproduction and re-use of the full text version. This is indicated by the licence information on the White Rose Research Online record for the item.

#### **Takedown**

If you consider content in White Rose Research Online to be in breach of UK law, please notify us by emailing eprints@whiterose.ac.uk including the URL of the record and the reason for the withdrawal request.



# ROYAL SOCIETY OPEN SCIENCE

rsos.royalsocietypublishing.org





Article submitted to journal

#### **Subject Areas:**

XXXXX, XXXXX, XXXX

#### Keywords:

XXXX, XXXX, XXXX

#### Author for correspondence:

Insert corresponding author name e-mail: hermes.gadelha@york.ac.uk

# The counterbend dynamics of cross-linked filament bundles and flagella

Rachel Coy<sup>1</sup>, Hermes Gadêlha<sup>2</sup>

<sup>1</sup>CoMPLEX, University College London, London WC1E 6BT, UK.

<sup>2</sup>Department of Mathematics, University of York, York YO10 SDD, UK.

Cross-linked filament bundles, such as in cilia and flagella, are ubiquitous in biology. They are considered in textbooks as simple filaments with larger stiffness. Recent observations of flagellar counterbend, however, show that induction of curvature in one section of a passive flagellum instigates a compensatory counter-curvature elsewhere, exposing the intricate role of the diminutive cross-linking proteins at large-scales. We show that this effect, a material property of the cross-linking mechanics, modifies the bundle dynamics, and induces a bimodal  $L^2 - L^3$  length-dependent material response that departs from the Euler-Bernoulli theory. Hence, the use of simpler theories to analyse experiments can result in paradoxical interpretations. Remarkably, the counterbend dynamics instigates counter-waves in opposition to driven oscillations in distant parts of the bundle, with potential impact on the regulation of flagellar bending waves. These results have a range of physical and biological applications, including the empirical disentanglement of material quantities via counterbend dynamics.

#### 1. Introduction

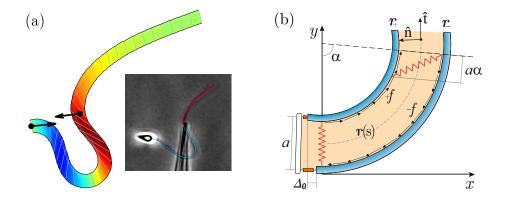
The spontaneous generation of harmonic bending waves along a sperm flagellum has been a source of fascination since it was reported on in the late 17th century [1]. It was not until 1968, however, that the fundamental mechanism behind the flagellar wave propagation was unveiled [2–4]. ATP-induced inter-microtubule tangential motion is converted into transversal forces that are capable of bending the flagellar assembly altogether, laying the empirical basis for the sliding filament theory for eukaryotic flagellum.

© 2014 The Authors. Published by the Royal Society under the terms of the Creative Commons Attribution License http://creativecommons.org/licenses/by/4.0/, which permits unrestricted use, provided the original author and source are credited.

Notably, almost one decade before the discovery of the interfilament sliding [2,3,5], the existence of such active elements along the sperm flagellum was theorized via a simple fluid-structure interaction model [6]. Machin demonstrated that the combined action of viscous and elastic dissipation experienced by a slender filament rapidly damps any driven oscillation along its length, thus requiring the action of contractile elements in order to sustain large waving amplitude [7]. Later, Goldstein and co-workers [8] elegantly demonstrated that the elastohydrodynamics of any simple Euler-Bernoulli filament moving in a viscous fluid leads to a hyperdiffusive dissipation of bending, characterized by a bending penetration length  $\ell_{\omega}$ , which can be further exploited to extract material parameters from biological filaments in a wide range of length-scales [9]. Hitherto the elastohydrodynamics of active and passive filaments have generated a vast literature of analytical, computational and empirical studies across disciplines [10–19].

Despite the inherent complexity of filament bundles [20–31], as exemplified by the axonemal flagellum [23,24,24], with its 9+2 cross-linked microtubule doublets arranged in a cylindrical fashion [23], the textbook elastic bending stiffness has been estimated using a simplistic linear relation between bending moment and curvature [27-31], as derived from Euler-Bernoulli rod theory [32]. Incidentally, the inadequacy of classical rod theories, from Euler-Bernoulli to Timoshenko and Cosserat [32], emerged via paradoxical counterbend empirical responses, Fig. 1(a), first observed by Lindemann and co-workers [33,34], and later captured via a geometrically exact mechanical model by Gadêlha et al [35] (Fig. 1(a)). These studies revealed how the induction of curvature in one section of a passive sperm flagellum instigates compensatory countercurvature elsewhere, namely the counterbend phenomenon [33,34]. They established the critical role of the diminutive elastic linking-proteins while instigating large-amplitude deformations, inherently coupling distant parts along the bundle assembly, despite their small slenderness ratio [33-35]. More recently, the counterbend phenomenon was also exploited in order to extract material quantities from Chlamydomonas flagella [36], despite the relatively short length flagella. The dynamical response of the counterbend phenomenon in passive cross-linked bundles still remains unexplored in the literature.

The discovery of the counterbend phenomenon highlighted the current need to reassess both the established material measurements [20-22,25-31,37], and the resulting mechanical response, from statics to dynamics, of cross-linked filament bundles [27-31]. The former is crucial in a broad range of biological structures, from the cytoskeleton of eukaryotic cells to cellular division, cross-bridge muscle contraction, and locomotion, via structures like the axoneme. A fundamental challenge, both experimentally and theoretically, is therefore to understand how this complex structure yields bulk material properties and overall cellular mechanical responses and, ultimately, function. In active bundles, the consequences of using inadequate material parameters, which have been used for the past 30 years to investigate flagellar waves [7,11,38-49], are still unknown. This is further confronted with an increasing number of, repeatedly contradicting, active control models for the flagellar wave coordination [7,11,39,40,43,44,46–52]. Paradoxically, in order to induce bending waves, flagellar control models rely on the implementation of filament bundle deformations, in distinct material directions, that are yet to be scrutinized in isolation, from curvature [38,40,45,49] to interfilament sliding [7,11,44,46,52,53], and axial distortions [43,47,50]. This is aggravated by the strong coupling between the unknown activity, and the passive and dissipative components, leading to the non-identifiability of parameters when contrasted against experiments [54,55]. Without the disentanglement between the passive and active elements, and without the rationalization of the resultant mechanical response of cross-linked filament bundles, it is unclear, for example, which competing flagellar control hypothesis [7,11,39,40,43,44,46–51], if any, is able to provide a quantitative understanding of the flagellar regulation and, crucially, function of the internal mechanics and structure. Indeed, any comprehensive model of flagellar bending self-organization depends on reliable measurements of mechanical and material properties of the system in absence of activity [35,36].



**Figure 1.** The counterbend phenomenon and geometry of deformation: (a) Micrograph showing the static configuration of a sea urchin sperm rendered passive with its head attached to the coverslip while forced externally by a micro-probe [34], together with the geometrically exact filament-bundle model prediction. Red curve show the model curve fitting result from [35]. (b) 2D representation of the axoneme and the sliding filament mechanism with basal compliance [7,35,37,46]. Micrograph adapted from Gadêlha et al. [35].

Here, we complement the seminal work by Machin [6] and Goldstein [8] on the dynamics of passive filaments, and demonstrate how the nanometric cross-linking proteins that are present in passive cross-linked filament bundles instigate novel dynamical counterbend phenomena. This is in contrast with previous models on flagellar wave coordination [7,11,39,40,43,44,46–52], which incorporate the cross-linking interaction in conjunction with molecular motor dynamics. We consider the dynamical situation in which only the structural passive elements are present. For axonemal filament-bundles, this corresponds to the empirical situation in which molecular motors are rendered passive [33,34,36], Fig. 1(a). The filament-bundle elastohydrodynamical model unveils the occurrence of counter-travelling waves in distant parts of bundle, reducing the propulsive potential of driven oscillations, and even reversing the propulsive direction, from pushing to pulling hydrodynamics. We show that the interplay between the interfilament sliding at the base, and cross-linking dissipation elsewhere, give rise to a bimodal  $L^2 - L^3$ length-dependent material response that departs from canonical Euler-Bernoulli theory. Hence the use of simpler rod theories to analyse experiments can result in paradoxical interpretations. Furthermore, the counterbend dynamics offers a robust way to measure material quantities empirically, bypassing cumbersome force-displacement experiments at the microscale [30,31,33, 34,36], Fig. 1(a). These results further suggest that the dynamical counter-wave phenomena is likely to play a critical role on the waveform organization, and the subsequent wave direction, of long flagella [48,49,56].

## 2. Cross-linked filament bundle elastohydrodynamics

We consider a planar representation of cross-linked filament bundles and flagellar axonemes, as depicted in Fig. 1(b), used interchangeably hereafter, composed of two elastic, inextensible filaments that resist deformation with an elastic bending modulus E [7,35,37,46]. Each constituent filament  $\mathbf{r}_{\pm}(s,t) = \mathbf{r}(s,t) \pm a/2$   $\hat{\mathbf{n}}(s,t)$  is separated by a distance a, much smaller than the filament length  $a \ll \mathbf{L}$ , normal to the to the centreline  $\mathbf{r}(s,t)$  at every point in arclength s and time t. Geometry constrains the normal vector  $\hat{\mathbf{n}}(s,t) = -\sin\alpha \mathbf{e}_x + \cos\alpha \mathbf{e}_y$  to the plane, where  $\alpha$  is the angle between the fixed frame s-axis and the tangent to the centreline  $\hat{\mathbf{t}} = \mathbf{r}_s$ . Like a rail-track [37], the constituent filaments travel distinct contour lengths forcing a geometrical arclength

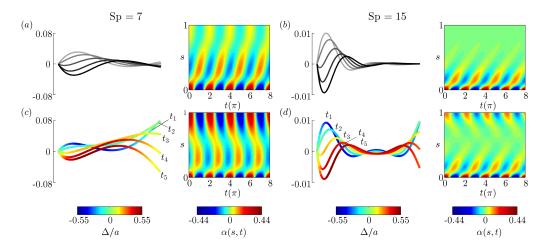


Figure 2. Counter travelling wave formation in filament bundles. (a,b) Euler-Bernoulli hyperdiffusive waveforms and (c,d) filament bundle waveforms for  $\gamma=0, \mu=100$ . The rescaled sliding displacement  $\Delta/a$  is overlaid in the waveform in (c,d). The time progression of the waveforms runs from  $t_1$  to  $t_5$ , covering one half of the periodic solution. For the  $\alpha$  plots, the time is labelled in multiples of  $\pi$ .

mismatch  $\Delta(s,t) = \Delta_0(t) + a(\alpha(s,t) - \alpha_0(t))$ , where  $\Delta_0$  and  $\alpha_0$  are the length mismatch and tangent angle at s=0 (Fig. 1)(b). Points of equal contour length along the filament-bundle are connected by elastic links which generate a shearing force, and thus an internal moment, proportional to the sliding displacement  $f(s,t) = k\Delta(s,t)$  with an elastic sliding resistance k. At the basal end, the additional connecting compliance across the filaments, commonly found in spermatozoa and inhomogeneous bundles, is Hookean  $\kappa_e \Delta_0(t) = -\int_0^L f(s',t) \mathrm{d}s'$  with a spring constant  $\kappa_e$  [35,46] (Fig. 1(b)).

For asymptotically slender filament-bundles, the hydrodynamic forces experienced by an infinitesimal element is anisotropic and linearly related to the local velocity  $\mathbf{f}_{\mathrm{vis}} = -\zeta_{\perp}(\hat{\mathbf{n}}.\mathbf{r}_t)\hat{\mathbf{n}} - \zeta_{\parallel}(\hat{\mathbf{t}}.\mathbf{r}_t)\hat{\mathbf{t}}$ , where  $\zeta_{\perp},\zeta_{\parallel}$  are the lowest order resistive coefficients derived from inertialess hydrodynamics [27,28]. Contact forces are not defined constitutively due to the inextensibility constraint [32]. The filament-bundle elastohydrodynamics is governed by the balance of contact forces and contact moments

$$-E\alpha_{ssss} + a^2k\alpha_{ss} = \zeta_{\perp}\alpha_t, \tag{2.1}$$

simplified here for small curvatures [8,9,11,46]. The filament-bundle shape is given by the initial value problem  $r(s,t)=r(0,t)+\int_0^s(\cos\alpha(s',t),\sin\alpha(s',t))\,\mathrm{d}s'$  for an arclength s and time t. Boundary conditions ensure the total balance of forces,  $F(s)=-E\alpha_{ss}+af(s,t)$ , and torques,  $M(s)=-E\alpha_s+a\int_s^L f(s',t)\mathrm{d}s'$ , acting on the bundle [32,35], as detailed in the Supporting Information (SI). The resulting cross-linking mechanics couples distant parts along the bundle via the total momentum balance, now modified non-locally by  $f(s,t)=-\gamma ka\int_0^L (\alpha(s')-\alpha_0)\,\mathrm{d}s'+ka(\alpha(s)-\alpha_0)$  (SI). Dissipation from different material directions are mediated by the hydrodynamic drag. The filament-bundle dynamics is dictated by the interplay between the elastohydrodynamic hyperdiffusion [8,9] and the cross-linking diffusion [7,37,46]. These boundary moments alter the hyperdiffusion balance in Eq. 2.1, and instigate novel long-range phenomena as we explore below.

#### (a) The counterbend dynamics: angular actuation

The post-transient behaviour of the shape dynamics is captured by single frequency solutions of the form  $\alpha(s,t)=\mathrm{Re}\{\tilde{\alpha}(s)e^{-i\omega t}\}$ . After convenient rescaling, the eigenvalue problem reduces to  $r^4-\mu r^2-i\mathrm{Sp}^4=0$ , with eigenfunctions  $\alpha(s,t)=\mathrm{Re}\{\sum_{j=1}^4 C_j e^{r_j s-it}\}$ , where the coefficients

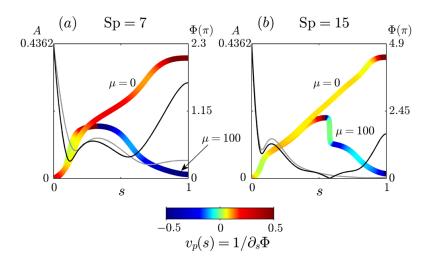


Figure 3. The amplitude A of the solutions in the  $\mu=0$  and  $\mu=100$  cases are represented by the grey and black lines respectively; the amplitude of the coloured lines indicates the values of  $\Phi$  at each point along the flagella as a multiple of  $\pi$ , with the colour denoting the velocity. Note that all velocities below -0.5 and above 0.5 are coloured the same as each of these values respectively. For all of the cases featured in the plot,  $\gamma=0$ .

 $C_j$  are determined by non-local boundary moments (SI). The dimensionless filament-bundle compliance parameter, also referred as sperm number,  $\mathrm{Sp} = \mathrm{L}(\zeta_\perp \omega/E)^{1/4}$ , captures the battle between elastic and viscous forces [8], whilst the sliding resistance parameter,  $\mu = a^2 \mathrm{L}^2 k/E$ , compares bending stiffness with the cross-linking resistance [35,46]. The basal compliance is given by  $\gamma = k\mathrm{L}/(k\mathrm{L} + \kappa_e)$ , and varies from  $\gamma = 0$ , corresponding to no interfilament sliding at the base, to  $\gamma = 1$ , for free basal sliding [35,46].

The emergence of the non-local, counterbend dynamics is depicted in Fig. 2 for a sinusoidal angular actuation of the proximal end (Supplementary Movie 1) with  ${\bf r}_t(0,t)=0$ , and zero force and torque condition at the distal end (SI). An angular amplitude of 0.4362 rad is used to limit the maximum radius of curvature to 0.1. Fig. 2 contrasts Machin's original solutions, Fig. 2(a,b), with the filament-bundle post-transient dynamics, Fig. 2(c,d). Travelling waves originating from the distal end, indicative of the non-local counterbend effect [35], can be clearly seen in Fig. 2(c,d). A relatively small sliding displacement  $\Delta$  (overlaid colour in Fig. 2(c,d)), equivalent to only half of the bundle diameter, is capable of deforming the bundle non-locally with the same magnitude of the imposed actuation.

The amplitude modulation A(s), phase  $\Phi(s)$  and velocity of propagation  $v_p(s)=1/\partial_s\Phi$  are depicted in Fig. 3, following suitable transformation to solutions of the form  $\alpha(s,t)=A(s)\cos{(t-\Phi(s))}$ . The abrupt change in wave direction is triggered by the loss in monotonicity of the phase. Non-local counter-waves propagate in opposite direction from the distal end with a non-uniform decaying magnitude along the arclength due to the high-order dissipation. This is in contrast to the one-directional wave propagation of Euler-Bernoulli filaments,  $\mu=0$  in Fig 3. The sharp change in wave direction coincides with reduced wave amplitudes at the point in arclength Fig. 3). Interestingly, the non-local actuation of cross-linking moments at distal parts of the bundle is delayed by the overdamped dynamics, as indicated by the proximal-distal phase difference. Higher Sp causes larger proximal-distal phase mismatch, faster decay of the counterbend wave speed and amplitude, indicative of a destructive interference between proximal and distal waves in Figs. 2(d) and 3 (b), demonstrated by the sharp jump in phase in Fig. 3.

The counterbend dynamics impacts significantly the resulting hydrodynamic propulsion. The time-averaged propulsive force  $\bar{F}_x$  exerted by fluid on the filament bundle is obtained by integrating the x component of the local drag force,  $\mathbf{f}_{\text{vis}} = -\zeta_{\perp}(\hat{\mathbf{n}}.\mathbf{r}_t)\hat{\mathbf{n}} - \zeta_{\parallel}(\hat{\mathbf{t}}.\mathbf{r}_t)\hat{\mathbf{t}}$ , along the

length of the bundle and averaging over one period of oscillation [8,9,14,15,57], as detailed in SI text. Due to the symmetry of the oscillations around the x axis, the y component of the propulsive force averages to zero. Following Refs. [8,15], we conveniently rescale length by  $(E/\omega\zeta_{\perp})^{1/4}$  and time by  $2\pi/\omega$ , so that the averaged propulsive force in the x direction may be written as

$$\bar{F}_x = (\zeta_{\perp} - \zeta_{\parallel})\omega/2\pi\sqrt{E/\omega\zeta_{\perp}} \Upsilon_x(\operatorname{Sp}, \mu, \gamma),$$

with a common elastohydrodynamic factor multiplying the force scaling function  $\Upsilon_x(\operatorname{Sp}, \mu, \gamma)$ , which is now modified by the cross-linking mechanics. The force scaling function  $\Upsilon_x(\operatorname{Sp}, \mu, \gamma)$ depicted in Fig. 4 captures the comparative propulsive potential between different points in the parameter space  $(Sp, \mu, \gamma)$ , where the convention is that the force exerted by the filament bundle on the fluid is  $-\bar{F}_x$  [8,15,57]. For a simple Euler-Bernoulli filament ( $\mu = 0$ ), dashed curve in Fig. 4, the propulsive force is always positive, except for effectively stiff filaments (Sp = 0), in which no propulsion can be generated  $\Upsilon_x = 0$  [8,15,58]. This is in accordance with Purcell's scallop theorem where no net propulsion can be achieved via time reversible motion in inertialess fluids [ibid]. As Sp increases, the filament becomes effectively more flexible, breaking Purcell's scallop theorem, thus inducing an increasingly large hydrodynamic propulsion, characterized by the emergence of a maximum value (dashed curve in Fig. 4). Increased filament floppiness, large Sp, causes the hydrodynamic force to decay and plateau. In this limit, the main contribution of the propulsive force arises from the tangential component of the hydrodynamic drag. Floppy filaments are unable to induce sizeable deformations, causing the filament to move effectively back and forth tangentially, along its length in the x direction. The sweet-spot of the hydrodynamic propulsion depicted in Fig. 4 (dashed curve) is the signature of the competition between viscous drag and bending stiffness, captured by the so-called penetration length  $\ell_{\omega} = (E/\omega \zeta_{\perp})^{1/4}$  [8,9,15]; first predicted by Wiggins et al [8] and empirically observed by Yu et al. [15].

A very distinct scenario is observed for filament bundles. As the basal compliance becomes stiffer, by reducing  $\gamma$ , the cross-linking mechanics switch from a mostly local contribution with small counterbend deformations ( $\gamma=1$  in Fig. 4), to non-local effects, with increasingly large amplitudes at the distal end ( $\gamma=0$  in Fig. 4 inset). Ultimately, this causes the propulsive force to vanish (circles in Fig. 4), and even switch the propulsive direction (negative values of  $\Upsilon_x$ ), thus equivalent to a backward net motion. Imposed oscillations are now counteracted by waves travelling in opposition at distal parts of the bundle (compare top and bottom net plots of each inset indicated by \* in Fig. 4). Small values of  $\gamma$  increases the non-local counterbending amplitude and causes the propulsive force to switch direction (negative values) as Sp increases ( $\gamma=0,0.4$  and 0.6 in Fig. 4). However, similarly to the Euler-Bernoulli case, increased floppiness of the filament bundle, i.e. large Sp, causes the amplitude of deformation to decay (both amplitude of imposed oscillation and counter-waves), reducing in this way the magnitude of the propulsive force which is now characterized by the appearance of a minimum value (see  $\gamma=0$  in Fig. 4).

The zero values of  $\Upsilon_x$  for  $\gamma=0,0.4$  and 0.6, circles in Fig. 4, are therefore a genuine manifestation of the perfect hydrodynamic balance between the imposed oscillation at the proximal end and the counter-waves at the distal end, with amplitudes modulated by the effective bundle stiffness Sp. Increased bundle floppiness causes the counter-wave amplitude to vanish, thus the change in sign to positive values of the propulsive force  $\bar{F}_x$  before it plateaus to a constant positive level, similarly to the Euler-Bernoulli filaments. The lower plateau value for filament bundles is due to the added effective stiffness arising from the cross-linking mechanics. The separatrix in Fig. 5 captures the region in parameter space where the local extrema of  $\Upsilon_x(\operatorname{Sp}_m)$  changes sign, the non-trivial roots of  $\Upsilon_x$  (circles) in Fig. 4. This indicates the region where a significant influence of non-local counterbend effect is predicted, and illustrates how the triad (Sp,  $\mu$ ,  $\gamma$ ) may be conveniently tuned to achieve zero, forward or backward propulsion (Fig. 5). Reversal in swimming direction may be achieved by simply increasing the frequency of oscillation for instance, recalling that  $\operatorname{Sp} = \operatorname{L}(\zeta_\perp \omega/E)^{1/4}$ . It is worth noting that the cross-linking dissipation does not affect the main elastohydrodynamic characteristics, governed by the penetration length

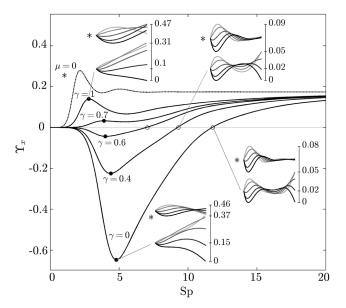


Figure 4. Scaling function  $\Upsilon_x$ , representative of the propulsive force. All cases plotted use the sliding resistance parameter value  $\mu=100$ , except for the dashed line which plots the case  $\mu=0$ . Waveforms corresponding to this  $\mu=0$  case are indicated by \*. The local maxima/minima  $\Upsilon_x(\operatorname{Sp}_m)$  in the interval  $\operatorname{Sp}_m \in [0,5]$  are labelled by black dots. Positions of the values where  $\Upsilon_x=0$  are labelled by circles.

 $\ell_{\omega} = L/\mathrm{Sp}$  [8], as indicated by the positive asymptote for both Euler-Bernoulli filaments and filament bundles in Fig. 4.

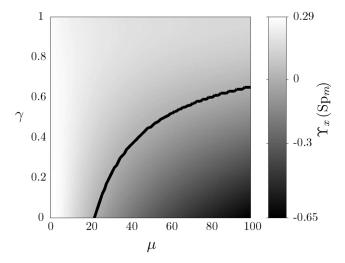


Figure 5. Local maxima/minima  $\Upsilon_x(\operatorname{Sp}_m)$ , with  $\operatorname{Sp}_m \in [0,5]$ , across the parameter space  $(\mu,\gamma)$ . The black line separates the values for which  $\Upsilon_x(\operatorname{Sp}_m)$  is negative from those for which it is positive. The region above (below) the black line indicates the positive maxima (negative minima) of  $\Upsilon_x(\operatorname{Sp}_m)$  in  $(\mu,\gamma)$ .

### (b) The bimodal length-dependent relaxation dynamics

The cross-linking mechanics introduces a diffusion-like time-scale,  $L^2\zeta_\perp/a^2k$ , with a somewhat weaker  $L^2$  geometrical dependence, in addition to the high-order, hyperdiffusive scaling  $L^4\zeta_\perp/E$ . Indeed, the cross-linking resistance  $\mu=a^2L^2k/E$  contrasts the elastohydrodynamic and cross-linking time-scales. The cross-linking resistance depends on geometrical aspects of the bundle, as  $\mu$  measures the ratio between the natural cross-linking elastic length  $\ell=\sqrt{E/a^2k}$  relative to total axial length via  $\mu=(L/\ell)^2$ . The cross-linking elastic length  $\ell$  has an important biophysical interpretation: it is the dimensional length by which cross-linking effects become prevalent. The cross-linking mechanics become increasingly important when  $L>\ell$ , while the opposite is found for  $L\sim\mathcal{O}(\ell)$  or smaller. The latter entails the possibility of studying relaxation counterbend phenomena by only varying the length of filament bundle, and motivates rescaling  $\mu$  relative to the reference length  $L_0$ , so that  $\mu(L)=L^2\mu_0$ ,  $\mu_0=a^2L_0^2k/E$  and now  $L=L/L_0$ , with smallest dimensionless length L=1.

The non-local, counterbend dynamics decaying from initial data, with an amplitude  $a_n$ , is depicted in Fig. 6 for filament-bundles that are clamped at the proximal end (SI). This is characterized by an effective relaxation constant  $\lambda_n^{-4}$  via  $\alpha(s,t) = \sum_n a_n S_n(s) \, e^{-\lambda_n^4 t}$ , with  $S_n = C_1 \sin(q_{1_n}s) + C_2 \cos(q_{1_n}s) + C_3 \sinh(q_{2_n}s) + C_4 \cosh(q_{2_n}s)$  for a given participating mode n. The triad of dissipative contributions acts as an effective dispersion medium for both bending and cross-linking deformations, dictated by the same dispersion relation. However, the mode shape is captured by the wavenumber-eigenvalue coupling  $q_{l_n} = \sqrt{(\sqrt{\mu^2 + 4\lambda_n^4} + (-1)^l \mu)/2}$  for l=1,2, which is not only influenced by the effective relaxation constant, reminiscent of pure high-order elastohydrodynamic dissipation, but also by the cross-linking diffusion. Boundary conditions define the transcendental solvability condition for  $\lambda_n$ , which depend implicitly on both  $\mu$  and  $\gamma$ , with an infinite number of mode solutions for each parameter set. Curve-fitting expressions for  $\lambda_1$  obtained from numerical solutions are presented in SI.

The non-local cross-linking diffusion introduces a bimodal length-dependent material response, as illustrated in Fig. 6 for the relaxation time of the fundamental mode

$$\tau_1(L) = \left(\frac{L}{\lambda_1(L)}\right)^4.$$

The relaxation dynamics of Euler-Bernoulli filaments decay with a characteristic  $L^4$ -dependence associated with the filament elastohydrodynamic hyperdiffusion [8]. When  $\mu=0$ ,  $\lambda_1$  is a constant solely defined by boundary conditions, thus  $\tau_1 \propto L^4$  as discussed above. This canonical case is shown by the black straight curve in Fig. 6, also indicated by the " $\propto L^4$ " at the top right corner of Fig. 6. The relaxation dynamics of filament bundles, on the other hand (depicted by the light blue and dark blue curves), deviate from the canonical  $L^4$ -dependence (straight black curve) as L increases, or equivalently  $\mu$  (the top horizontal axis in Fig. 6 shows the associated  $\mu$  values via the relationship  $\mu(L)=L^2\mu_0$ ). This is due to the fact that for filament bundles  $\lambda_1=\lambda_1(L)$ , thus inducing an additional length-dependence on the denominator of  $\tau_1$ . This causes the relaxation time to gradually deviate from the classical  $L^4$  dependence, with greater magnitude as L increases, depending on the details of the basal compliance. For  $\gamma=0$ , the deviation from the Euler-Bernoulli case starts approximately from  $\mu>1$ , while for  $\gamma=1$  occurs for approximately  $\mu>1$ 0. An  $L^2$  asymptote arises for long filament bundles with  $\gamma=0$ , while for  $\gamma=1$  the transition is to an  $L^3$  behaviour instead. These long-length asymptotes are indicated by the dashed grey curves labelled by " $\propto L^3$ " and " $\propto L^2$ " at the bottom left corner of Fig. 6.

The length-dependent transition between  $L^2$  and  $L^3$  asymptotes for long filament bundles is governed by the basal compliance. This  $\gamma$ -dependence is depicted in the inset (a) of Fig. 6, which plots the exponent  $\zeta$  of the relationship  $\tau_1 \propto L^{\zeta}$  in the limit of large L. For asymptotically long filament bundles, the exponent of  $\tau_1 \propto L^{\zeta}$  is quadratic ( $\zeta = 2$ ), and remains nearly quadratic until  $\gamma$  approaches 1, at which  $\zeta = 3$ . Such bijection of the material response entails that simultaneous measurement of the bundle mechanical properties, in different material direction, can be extracted

from simple relaxation experiments. In particular, increased inter-filament sliding, concentrated towards the clamped end, induces curvature-reversal for long filament bundles (Fig. 6 inset (b)), reminiscent of the counterbend phenomenon [35]. Boundary conditions require zero contact forces and torques at the free end, while clamped constraint facilitates the accumulation of cross-linking sliding towards the proximal end. Both bending and cross-linking deformations relax towards the reference configuration with the same effective rate  $\lambda_1$ . Fig. 6 inset (b) shows the simultaneous hyperdiffusive relaxation of bundle curvature (bundle shape) and sliding displacement (overlaid colour) towards the equilibrium straight configuration. Fig. 6 thus summarises the impact of the cross-linking mechanics on the relaxation time across the parameter space  $(\gamma, \mu)$ .

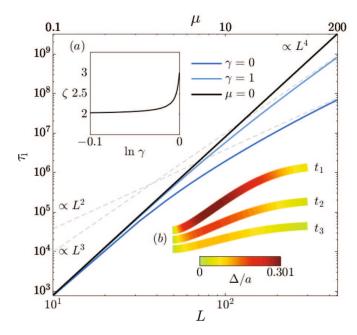


Figure 6. Rescaled relaxation time  $au_1$  for the first mode as a function of L for the Euler-Bernoulli filaments ( $\mu=0$ , black curve) and filament bundles (blue curves). The grey dashed lines indicate the long-length asymptote  $L^2$  and  $L^3$  for filament bundles (labelled by " $\propto L^3$ " and " $\propto L^2$ " at the bottom left corner). The top horizontal axis shows the associated  $\mu$  values via the relationship  $\mu(L)=L^2\mu_0$ . The inset Fig. 6(a) plots the exponent  $\zeta$  of the relationship  $au_1 \propto L^\zeta$ , as calculated for long filament bundles, against  $\ln \gamma$ ; Fig. 6(b) shows the time evolution of the the relaxation dynamics of the first mode, with the overlaid colours indicating the rescaled sliding displacement  $\Delta/a$ , for the case  $\mu=100$  and  $\gamma=0$ .

#### 3. Discussion

We studied the transient and post-transient dynamics of overdamped filament bundles that are interconnected by linking elastic proteins. Deformations in distinct material directions, arising from the cross-linking interfilament sliding and pure bending deformation, are coupled with local slender-body hydrodynamics. This leads to an effective dispersion mechanism governed by the superposition of short and long-range dissipation mechanisms. Cross-linking stresses are transmitted to distant parts of the bundle via boundary balance of moments. The cumulative moments are able to surpass the high-order elastohydrodynamic dissipation, and shape the bundle structure non-locally, with increased influence for long filament-bundles, or equivalently, large  $\mu$ .

The delicate interplay between the interfilament sliding at the base and the rest of the bundle results in a bimodal dynamic response, which departs from the classical Euler-Bernoulli theory [6,8,32]. When the basal sliding is permitted, cross-linking diffusion is mostly local, and acts to effectively reinforce the bundle structure. Long-range curvature-reversal events, however, are magnified when the basal sliding is constrained [35]. The counterbend dynamics generate spontaneous travelling waves in opposition to driven oscillations, which are capable of suppressing the propulsive potential, and even reverting the direction of propulsion (Fig. 4). Curvature perturbations diffuse more rapidly, a hundred times faster than Euler-Bernoulli hyperdiffusion with an equivalently higher bending rigidity (Fig. 6). Relatively small cross-linking deformations, up to only 30% of the bundle diameter, are capable of exciting large counterbend modes (Fig. 6 inset (b)), and induce a bimodal  $L^2 - L^3$  length-dependent deviation from the  $L^4$ -dependence of canonical filaments. Paradoxical measurements may arise if standard Euler-Bernoulli theory is used to interpret experiments [29,31,33,34], as exemplified by the paradoxical length-dependent bending stiffness in microtubules [59]. Indeed, the lengthdeviation predicted here may be mistakenly interpreted as an effective length-dependent bending rigidity via  $L^4\zeta_{\perp}/E(L)$  [8], if rather the Euler-Bernoulli theory is used; de facto, the Euler-Bernoulli theory is traditionally used since the first measurements of flagellar bundles [27,29, 31].

Static, force-displacement experiments that are often used to probe flagellar material quantities [29,31,34,36] are cumbersome, see Fig. 1(a). They require high-precision force calibrated probes and micro-manipulators, and often rely on the rare attachment of the filament's tip to the coverslip to micro-probe actuation [29,31,34]. This proximity of the filament bundle to the cover-slip can interfere with the interfilament sliding due to surface adhesion, biasing in this way force and shape measurements [35]. The counterbend dynamics provides a simpler and robust empirical route for the disentanglement of material parameters. This includes measurements of the basal interfilament elasticity, despite being deeply embedded at the connecting piece of the bundle (Fig. 1). As a result, standard microfluidic designs may be explored to induce shape changes dynamically [18]. Likewise, the dynamical counter-wave phenomenon may also inspire the design of artificial swimmers [60] that are able to reverse the swimming direction by simply increasing, for instance, the frequency of oscillation (see Fig. 4).

The counter-wave phenomena becomes increasingly important for bundles longer than  $\ell=\sqrt{E/a^2k}$ , typically  $5\mu m$  for flagella [2,7,46,49,61,62]. Interestingly, the majority of eukaryotic flagella exceed  $\ell$  by few orders of magnitude, from approximately  $30\mu m$  for *Chlamydomonas* and sea urchin sperm to almost  $200\mu m$  for quail sperm [61,62]. Cross-linking effects may also become increasingly important during flagellar growth, and influencing in this way the wave coordination during flagellar reconstitution in *Chlamydomonas* [63]. Molecular motors organization thus may operate differently for L <  $\ell$  and L >  $\ell$ . Indeed, local flagellar control models [7,49] recently gained empirical support when tested against short flagella experiments [49], a regime where counterbend phenomenon may be negligible (L <  $\ell$ ). This is despite of the well-known negative support of curvature control models [40,46], tested instead against long flagella (L >  $\ell$ ). The recurrent contradictions between sliding and curvature control models [7,11,40,46,49,52] may suggest the occurrence of distinct length-dependent flagellar regimes.

Linear models coupling the molecular motor reaction kinetics with interfilament sliding spontaneously propagate waves along the axoneme via a Hopf bifurcation [11,41,44,46,48]. The resulting wave train is observed to move from tip to base, i.e. in the direction opposite to what one would expect from a local dissipation theory (when the interfilament sliding is prevented at the base) [11,49,56]. Incidentally, the basal compliance was observed to influence the direction of wave propagation in flagella self-organizations models [46,56], demonstrating the sensitivity of the direction of the travelling wave to details of the connecting piece (basal part) and boundary conditions. This is in agreement with the bimodal response predicted here (see Fig. 2), in which counterbend is maximized when  $\gamma = 0$  [35]. The wave direction is influenced non-locally by cross-linking effects whose magnitude is regulated by the basal interfilament sliding (see Figs. 2 and 6).

This might explain the surprising significance of the basal compliance during the flagellar wave coordination observed recently in empirical studies [61], and even flagellar synchronization that may arise without recurring to hydrodynamical coupling [64]. Nevertheless, the mechanisms by which the flagellar wave direction is selected is poorly understood. Previous studies were reduced to the linear level, and at the nonlinear regime, the dynamical instability generates unstable traveling waves that can propagate in both directions, with potencial for multi-frequency modes [54,65]. The boundary conditions and basal mechanics assist the mode selection nonlinearly, and thus the direction of propagation, emphasizing how the flagellar dynamics is critically dependent on the underlying structural mechanics of the axoneme. Non-local hydrodynamic interactions [62], transversal axonemal deformations [43,47], and geometrical non-linearities [10,38] are also likely to affect the emergence of self-organization in flagellar systems.

The high-order diffusive interaction, intrinsic to elastohydrodynamic systems in Eq. 2.1, is observed throughout nature. In non-dilute systems, particles are affected by density variations beyond the nearest neighbours via, for example, biharmonic interactions  $u_t = D_1 \nabla^2 u - D_2 \nabla^4 u$  [66,67], thus closely related to Eq. 2.1. Despite the relatively short-range influence, such higher-order diffusion instigates complex spatio-temporal dynamics and self-organization in all fields of science [68,69]. They drive instabilities and even mediate the coexistence of spatial patterns and temporal chaos, as observed in Kuramoto-Sivashinsky systems [69]. Other exemplars of local, higher-order diffusion are found in Ginzburg-Landau superconductors, spatial patterning in Cahn-Hilliard and biochemical systems, plus generalized Fisher-Kolmogorov models, water waves and continuum mechanics systems, among others [66,68,69].

In contrast with canonical high-order diffusion systems [66,68,69], the flagellar scaffold, or equivalently, any cross-linked filament bundle immersed in a viscous fluid is governed by high-order dispersion medium that is inherently *non-local*. Here, the biharmonic diffusion arises instead via a *local* elastohydrodynamic dissipation [6,8], while the Fickian-like interaction arises through the *long-range* coupling reflecting the bundle mechanics [7,11,35,37], effectively connecting distant parts of the system via boundary bending moments. This unveils the potential for rich long-range phenomena via reaction-diffusion interactions [11,54,56], from non-local pattern formation to long-range synchronization of auto-oscillators, that are yet to be fully explored in the realm of mathematical biology.

We hope that these results will inspire theoreticians and experimentalists to study the dynamical effects of the conterbend phenomenon in filament-bundle as found throughout nature, including prospects for counterbend reaction-diffusion systems in flagellar dynamics, effectively bridging, non-locally, dynamical systems and PDE's.

# Acknowledgment

R.C. thanks Cambridge Bridgwater Summer Research Programme. H.G. acknowledges support by the Hooke Fellowship, University of Oxford, and WYNG Fellowship, Trinity Hall, Cambridge. The authors also thank Dr E.A. Gaffney for enlightening discussions. We dedicate this work in memory of Prof. John R. Blake, whose work and devotion will continue to inspire future generations of scientists.

#### References

- 1. Stuart S. Howards.
  - Antoine van Leeuwenhoek and the discovery of sperm. *Fertility and Sterility*, 67(1):16–17, January 1997.
- 2. Peter Satir.
  - STUDIES ON CILIA III. Further Studies on the Cilium Tip and a "Sliding Filament" Model of Ciliary Motility.
  - The Journal of Cell Biology, 39(1):77–94, October 1968.
- 3. Keith E Summers and IR Gibbons.

Adenosine triphosphate-induced sliding of tubules in trypsin-treated flagella of sea-urchin sperm.

*Proceedings of the National Academy of Sciences*, 68(12):3092–3096, 1971.

4. Fred D Warner and Peter Satir.

The structural basis of ciliary bend formation: radial spoke positional changes accompanying microtubule sliding.

The Journal of cell biology, 63(1):35, 1974.

5. Peter Satir.

Studies on cilia: Ii. examination of the distal region of the ciliary shaft and the role of the filaments in motility.

The Journal of cell biology, 26(3):805, 1965.

6. K. E Machin.

Wave Propagation Along Flagella.

Journal of Experimental Biology, 35(4):796–806, December 1958.

7. Charles J. Brokaw.

Flagellar Movement: A Sliding Filament Model.

Science, 178(4060):455-462, November 1972.

8. Chris H. Wiggins and Raymond E. Goldstein.

Flexive and propulsive dynamics of elastica at low reynolds number.

Phys. Rev. Lett., 80:3879, 1998.

9. C. H. Wiggins, D. Riveline, A. Ott, and R. E. Goldstein.

Trapping and wiggling: elastohydrodynamics of driven microfilaments.

Biophys J, 74(2 Pt 1):1043-1060, Feb 1998.

10. H. Gadêlha, E. A. Gaffney, D. J. Smith, and J. C. Kirkman-Brown.

Nonlinear instability in flagellar dynamics: a novel modulation mechanism in sperm migration?

*Journal of The Royal Society Interface*, 7:1689–, 2010.

11. SÃl'bastien Camalet, Frank Jülicher, and Jacques Prost.

Self-organized beating and swimming of internally driven filaments.

Phys. Rev. Lett., 82:1590, 1999.

12. L. Bourdieu, T. Duke, M. B. Elowitz, D. A. Winkelmann, S. Leibler, and A. Libchaber.

Spiral defects in motility assays: A measure of motor protein force.

Phys. Rev. Lett., 75:176–179, 1995.

13. R. E. Goldstein and S. A. Langer.

Nonlinear dynamics of stiff polymers.

Phys. Rev. Lett., 75:1094-1097, 1995.

14. H. C. Fu, C. W. Wolgemuth, and T. R. Powers.

Beating patterns of filaments in viscoelastic fluids.

Phys. Rev. E, 78:041913-041925, 2008.

15. T. S. Yu, , E. Lauga, and A. E. Hosoi.

Experimental investigations of elastic tail propulsion at low reynolds number.

Phys. Fluids, 18:0917011-0917014, 2006.

16. Sarah D. Olson, Sookkyung Lim, and Ricardo Cortez.

Modeling the dynamics of an elastic rod with intrinsic curvature and twist using a regularized Stokes formulation.

Journal of Computational Physics, 238:169–187, April 2013.

17. A. K. Tornberg and M. J. Shelley.

Simulating the dynamics and interactions of flexible fibers in stokes flows.

J. Comput. Phys., 196:8-40, 2004.

18. Vasily Kantsler and Raymond E. Goldstein.

Fluctuations, Dynamics, and the Stretch-Coil Transition of Single Actin Filaments in Extensional Flows.

Physical Review Letters, 108(3):038103, January 2012.

19. Tim Sanchez, Daniel T. N. Chen, Stephen J. DeCamp, Michael Heymann, and Zvonimir Dogic. Spontaneous motion in hierarchically assembled active matter.

Nature, 491(7424):431-434, November 2012.

20. Claus Heussinger, Felix Schüller, and Erwin Frey.

Statics and dynamics of the wormlike bundle model.

Phys. Rev. E, 81(2):021904, Feb 2010.

21. M. M. A. E. Claessens, C. Semmrich, L. Ramos, and A. R. Bausch.

Helical twist controls the thickness of f-actin bundles.

Proceedings of the National Academy of Sciences, 105(26):8819–8822, 2008.

22. Mireille MAE Claessens, Mark Bathe, Erwin Frey, and Andreas R Bausch.

Actin-binding proteins sensitively mediate f-actin bundle stiffness.

Nature materials, 5(9):748-753, 2006.

23. Don W. Fawcett, William Bloom, and Elio Raviola.

A Textbook of Histology.

Chapman & Hall, June 1994.

24. B. Afzelius.

Electron microscopy of the sperm tail.

Biophys. Cytol., 5:269, 1959.

25. JA Tolomeo and MC Hollev.

Mechanics of microtubule bundles in pillar cells from the inner ear.

Biophysical journal, 73(4):2241, 1997.

26. Itsushi Minoura, Toshiki Yagi, and Ritsu Kamiya.

Direct measurement of inter-doublet elasticity in flagellar axonemes.

*Cell structure and function*, 24(1):27–33, 1999.

27. B. Alberts.

Molecular Biology of the Cell.

Garland Science, New York, 2002.

28. J. Howard.

Mechanics of motor proteins and the cytoskeleton.

Sinauer Associates Sunderland, MA, 2001.

29. Charles B. Lindemann, Walter G. Rudd, and Robert Rikmenspoel.

The stiffness of the flagella of impaled bull sperm.

Biophysical Journal, 13(5):437 - 448, 1973.

30. M. Okuno.

Inhibition and relaxation of sea urchin sperm flagella by vanadate.

The Journal of Cell Biology, 85(3):712, 1980.

31. M. Okuno and Y. Hiramoto.

Direct measurements of the stiffness of echinoderm sperm flagella.

Journal of Experimental Biology, 79(1):235, 1979.

32. SS Antman.

Nonlinear Problems of Elasticity, volume 107 of Applied Mathematical Sciences.

Springer, 2005.

33. Charles B. Lindemann, Lisa J. Macauley, and Kathleen A. Lesich.

The counterbend phenomenon in dynein-disabled rat sperm flagella and what it reveals about the interdoublet elasticity.

Biophysical Journal, 89(2):1165 -1174, 2005.

34. Dominic W Pelle, Charles J Brokaw, Kathleen A Lesich, and Charles B Lindemann.

Mechanical properties of the passive sea urchin sperm flagellum.

Cell Motil Cytoskeleton, 66(9):721-735, Sep 2009.

35. H. Gadêlha, E. A. Gaffney, and A. Goriely.

The counterbend phenomenon in flagellar axonemes and cross-linked filament bundles.

Proceedings of the National Academy of Sciences, July 2013.

36. Gang Xu, Kate S Wilson, Ruth J Okamoto, Jin-Yu Shao, Susan K Dutcher, and Philip V Bayly. Flexural rigidity and shear stiffness of flagella estimated from induced bends and counterbends.

Biophysical Journal, 110(12):2759-2768, 2016.

37. R. Everaers, R. Bundschuh, and K. Kremer.

Fluctuations and stiffness of double-stranded polymers: railway-track model.

EPL (Europhysics Letters), 29:263, 1995.

38. M. Hines and JJ. Blum.

Bend propagation in flagella. i. derivation of equations of motion and their simulation. J. Biophys., 23:41, 1978.

39. Charles J. Brokaw.

Molecular mechanism for oscillation in flagella and muscle.

Proc. Natl Acad. Sci., 72:3102, 1975.

40. Charles J. Brokaw.

Computer simulation of flagellar movement. vi. simple curvature-controlled models are incompletely specified.

I. Biophys., 48:633, 1985.

41. C.J. Brokaw and D.R. Rintala.

Computer simulation of flagellar movement. iii. models incorporating cross-bridge kinetics. *J. Mechanochem. Cell. Motil.*, 3(2):77, 1975.

42. K. E. Machin.

The control and synchronization of flagellar movement.

Proc. R. Soc. London, Ser. B, 158:88, 1963.

43. Charles B Lindemann.

A" geometric clutch" hypothesis to explain oscillations of the axoneme of cilia and flagella. *Journal of theoretical biology*, 168(2):175–189, 1994.

44. M Hines and JJ Blum.

Bend propagation in flagella. ii. incorporation of dynein cross-bridge kinetics into the equations of motion.

Biophysical journal, 25(3):421, 1979.

45. Robert Rikmenspoel.

Contractile mechanisms in flagella.

Biophysical journal, 11(5):446, 1971.

46. Ingmar H. Riedel-Kruse, Andreas Hilfinger, Jonathon Howard, and Frank Jülicher.

How molecular motors shape the flagellar beat.

HFSP Journal, 1(3):192-208, 2007.

47. Philip V. Bayly and Kate S. Wilson.

Equations of Interdoublet Separation during Flagella Motion Reveal Mechanisms of Wave Propagation and Instability.

Biophysical Journal, 107(7):1756–1772, October 2014.

48. P. V. Bayly and K. S. Wilson.

Analysis of unstable modes distinguishes mathematical models of flagellar motion. *Journal of The Royal Society Interface*, 12(106):20150124, May 2015.

49. Pablo Sartori, Veikko F. Geyer, Andre Scholich, Frank Jülicher, and Jonathon Howard.

Dynamic curvature regulation accounts for the symmetric and asymmetric beats of Chlamydomonas flagella.

eLife, 5:e13258, May 2016.

50. Charles J Brokaw.

Computer simulation of flagellar movement x: doublet pair splitting and bend propagation modeled using stochastic dynein kinetics.

Cytoskeleton, 71(4):273-284, 2014.

51. Charles B Lindemann and Kathleen A Lesich.

Flagellar and ciliary beating: the proven and the possible.

Journal of Cell Science, 123(4):519-528, 2010.

52. Charles J Brokaw.

Computer simulation of flagellar movement ix. oscillation and symmetry breaking in a model for short flagella and nodal cilia.

Cell motility and the cytoskeleton, 60(1):35–47, 2005.

53. C.J. Brokaw and D.R. Rintala.

Computer simulation of flagellar movement. iii. models incorporating cross-bridge kinetics. *J. Mechanochem. Cell. Motil.*, 3(2):77, 1975.

54. David Oriola, Hermes Gadêlha, and Jaume Casademunt.

Nonlinear amplitude dynamics in flagellar beating.

Royal Society Open Science.

55. Franck Plouraboue, Ibrahima Thiam, Blaise Delmotte, Eric Climent, PSC Collaboration, et al. Identification of internal properties of fibers and micro-swimmers.

In APS Meeting Abstracts, 2016.

56. A. Hilfinger, A. K. Chattopadhyay, and F. Jülicher.

Nonlinear dynamics of cilia and flagella.

Phys. Rev. E, 79:051918-051925, 2009.

57. Eric Lauga and Thomas R Powers.

The hydrodynamics of swimming microorganisms.

Reports on Progress in Physics, 72(9):096601, 2009.

58. E.M. Purcell.

Life at low reynolds number.

Am. J. Phys., 45:3, 1977.

59. Francesco Pampaloni, Gianluca Lattanzi, Alexandr Jonáš, Thomas Surrey, Erwin Frey, and Ernst-Ludwig Florin.

Thermal fluctuations of grafted microtubules provide evidence of a length-dependent persistence length.

Proceedings of the National Academy of Sciences, 103(27):10248–10253, 2006.

60. Hermes Gadêlha.

On the optimal shape of magnetic swimmers.

Regular and Chaotic Dynamics, 18(1-2):75-84, 2013.

61. Kirsty Y Wan and Raymond E Goldstein.

Coordinated beating of algal flagella is mediated by basal coupling.

Proceedings of the National Academy of Sciences, 113(20):E2784–E2793, 2016.

62. E.A. Gaffney, H. Gadêlha, D.J. Smith, J.R. Blake, and J.C. Kirkman-Brown.

Mammalian sperm motility: Observation and theory.

Annual Review of Fluid Mechanics, 43(1):501-528, 2011.

63. Raymond E. Goldstein, Marco Polin, and Idan Tuval.

Emergence of synchronized beating during the regrowth of eukaryotic flagella.

Phys. Rev. Lett., 107:148103, Sep 2011.

64. Greta Quaranta, Marie-Eve Aubin-Tam, and Daniel Tam.

Hydrodynamics versus intracellular coupling in the synchronization of eukaryotic flagella. *Phys. Rev. Lett.*, 115:238101, Nov 2015.

65. David Oriola, Hermes Gadêlha, C Blanch-Mercader, and Jaume Casademunt.

Subharmonic oscillations of collective molecular motors.

EPL (Europhysics Letters), 107(1):18002, 2014.

66. James D Murray.

Mathematical Biology. II Spatial Models and Biomedical Applications {Interdisciplinary Applied Mathematics V. 18}.

Springer-Verlag New York Incorporated, 2001.

67. Donald S Cohen and James D Murray.

A generalized diffusion model for growth and dispersal in a population.

Journal of Mathematical Biology, 12(2):237-249, 1981.

68. Lambertus A Peletier and William C Troy.

*Spatial patterns: higher order models in physics and mechanics*, volume 45.

Springer Science & Business Media, 2012.

69. Arkady Pikovsky, Michael Rosenblum, Jürgen Kurths, and Robert C Hilborn.

Synchronization: a universal concept in nonlinear science.

American Journal of Physics, 70(6):655–655, 2002.