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Hydrodynamics of diatom chains and semiflexible fibres

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Diatoms are non-motile, unicellular phytoplankton that have the ability to form colonies in the form of chains. Depending upon the species of diatoms and the linking structures that hold the cells together, these chains can be quite stiff or very flexible. Recently, the bending rigidities of some species of diatom chains have been quantified. In an effort to understand the role of flexibility in nutrient uptake and aggregate formation, we begin by developing a three-dimensional model of the coupled elastic–hydrodynamic system of a diatom chain moving in an incompressible fluid. We find that simple beam theory does a good job of describing diatom chain deformation in a parabolic flow when its ends are tethered, but does not tell the whole story of chain deformations when they are subjected to compressive stresses in shear. While motivated by the fluid dynamics of diatom chains, our computational model of semiflexible fibres illustrates features that apply widely to other systems. The use of an adaptive immersed boundary framework allows us to capture complicated buckling and recovery dynamics of long, semiflexible fibres in shear.

1. Introduction

Flexible, elastic fibres that either move around in a fluid or move a fluid around are ubiquitous in nature and biology. In mammals, cilia are excellent examples of actuated elastic fibres that are responsible for mucus clearance in the respiratory tract and promote ovum transport in the oviduct [1]. Biological polymers such as DNA and actin filaments or synthetic polymers such as nylon are examples of passive elastic fibres that may bend or buckle in a moving fluid [2–5]. In order to predict the dynamics of a semiflexible fibre moving in a viscous, incompressible fluid, its mechanical properties must be known.

Here, we focus on the intriguing flexible fibres that are formed by some types of diatoms, non-motile phytoplankton. Phytoplankton account for approximately 40% of the primary marine production as well as one-fourth of the oxygen production on the Earth. The local flow environment around phytoplankton affects processes such as nutrient uptake and predator–prey interactions [6,7]. While diatoms are unicellular, some types have evolved the ability to form colonies in the form of chains using a variety of linking structures that hold the cells together (figure 1) [8]. Figure 1*a,b* shows micrographs of *Lithodesmium undulatum* and *Guinardia delicatula*. While these chains appear to be fairly stiff, figure 1*c* shows images of *Thalassiosira* sp., which appear to be much more flexible. Does flexibility enhance or hinder nutrient transport and acquisition by diatom chains in a dynamic flow environment? This question was addressed by Musielak *et al.* [9] using a computational fluid dynamic model in two dimensions. It was found that stiffer chains enjoy enhanced nutrient uptake in a patchy environment, perhaps because they bend less and sweep out a larger volume of fluid than more flexible chains. The correspondence between the mechanical properties of the model diatom chains in [9] and real diatom chains could not be directly assessed because, at that time, no laboratory measurements of bending rigidity or flexural stiffness of diatom chains were available. However, recently, Young *et al.* [8] performed experiments that quantified the flexural stiffness of different types of diatom chains by measuring their deflections under flow when held across

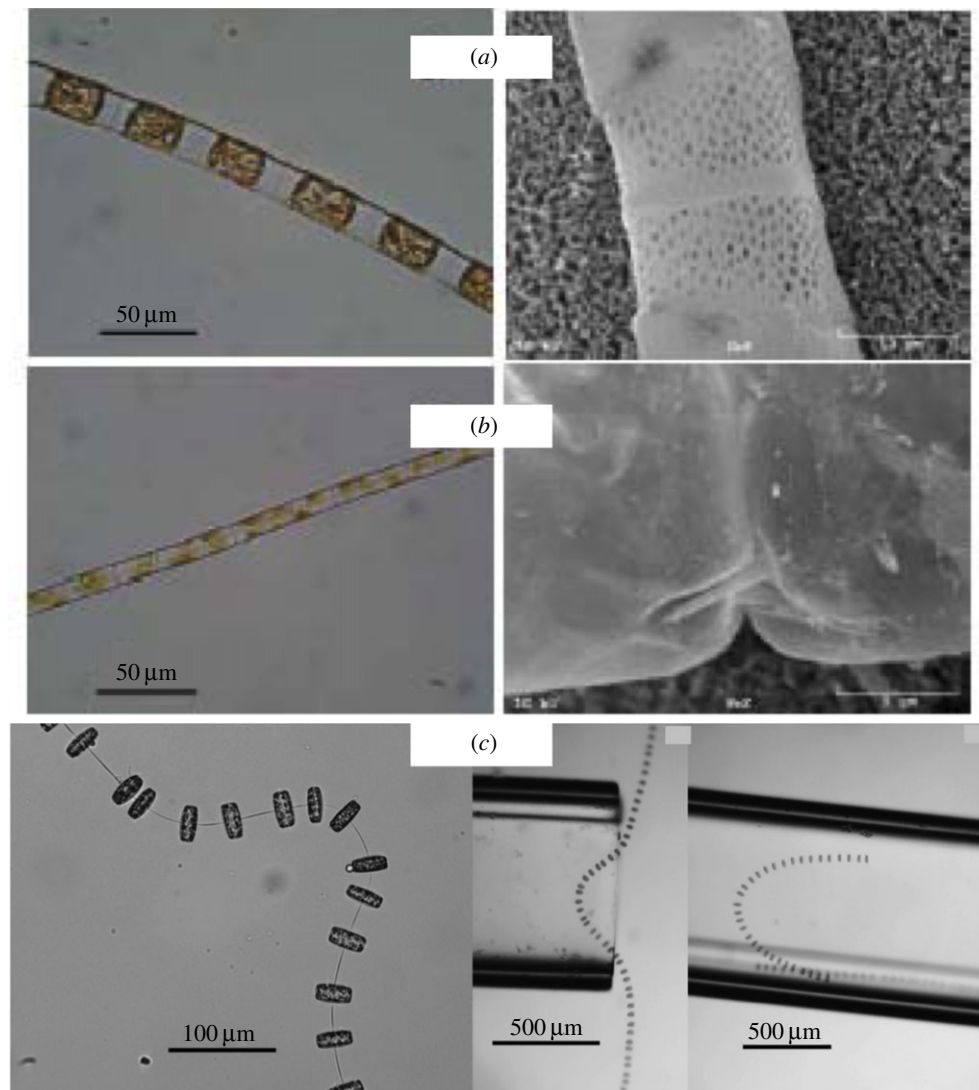


Figure 1. Light (left) and scanning electron (right) micrographs of (a) *L. undulatum* and (b) *G. delicatula*, courtesy of A. Young and L. Karp-Boss, University of Maine. (c) Images of *Thalassiosira* sp. adapted from [8]. (Online version in colour.)

a capillary tip. A simple beam theory was applied to approximate the flexural stiffness given the measured deflections and imposed volumetric flow rate. Here, we describe the calibration of these laboratory experiments [8] with the development of a full three-dimensional computational model of an inhomogeneous elastic chain. Having verified that the computational elastic structure has captured elastic properties of true diatom chains, we examine the dynamics of these chains in shear flow.

While motivated by the fluid dynamics of diatom chains, our three-dimensional computational model of semiflexible fibres illustrates features that apply widely to other systems. Namely, we show that using a three-dimensional adaptive immersed boundary framework [10] allows us to capture complicated buckling and recovery of long, flexible fibres in shear flow. Moreover, we argue that the dynamics of complex elastic fibres cannot always be predicted by assuming that they are homogeneous, linear beams. In the following sections, we will describe the structure of the computational diatom chain and the computational experiments that were performed to mirror the laboratory experiments in [8]. We will then examine how the orbits of these semiflexible fibres in shear flow depend upon their elastic properties and the connectivity properties of the network of points and springs that build them.

2. Calibration of experiments and computations

2.1. Construction of elastic diatom chain

Motivated by the structure of *L. undulatum* (figure 1a), we construct an elastic model of a diatom chain by creating a network of nodes and springs. The nodes are placed regularly on the surface of the cylindrical fibre, equally spaced around cross sections that are orthogonal to the centreline of the cylinder. We will consider three types of structures based upon the specified connectivity of the nodes. In the *surface-spring type*, we specify that linear springs connect closest points on the surface to each other and to next-to-closest points on the surface. The *full-spring type* not only has linear springs connecting nodes on the surface as in the *surface-spring type* but also has additional nodes placed along the centreline that are connected by internal springs to the surface nodes on the corresponding cross section (spokes). The *diatom-chain type* comprises an alternating sequence of each of the previous types, with the *full-spring* structure modelling the valve or cellular part of a single diatom and the *surface-spring* structure modelling the cell-connection between two diatoms in the chain. Figure 2 shows a schematic of the node-connectivity that builds the diatom chain. Figure 3 shows the long fibre that is the computational diatom

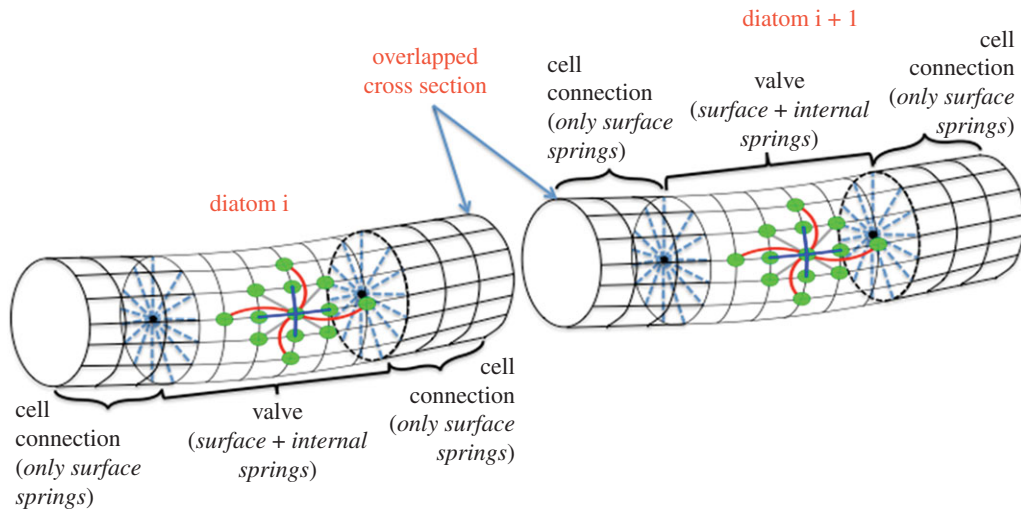


Figure 2. A schematic that shows the node–spring connectivity structure that is used to build a model diatom chain. (Online version in colour.)

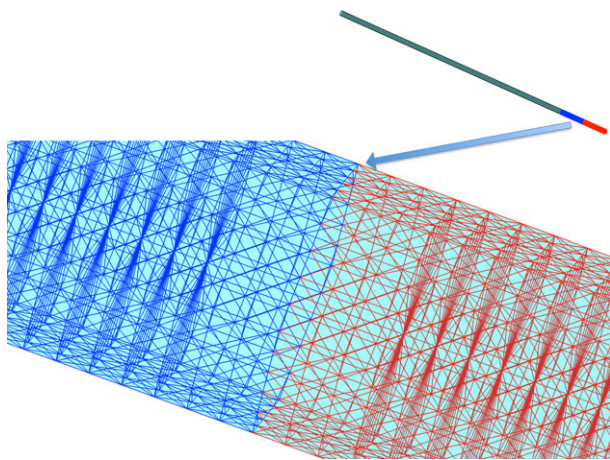


Figure 3. The computational diatom chain structure. The zoomed-in portion shows the alternating valve and cell structures. The valve structures that connect the diatom cells do not have internal spokes. (Online version in colour.)

chain, with the ‘zoomed-in’ portion showing the connectivity structure of two adjacent diatoms.

Assume that \mathbf{x}_k and \mathbf{x}_l are two nodes of the diatom chain that are connected by a linkage or spring of rest length Δs_{kl} . The force at \mathbf{x}_k due to this spring is

$$\mathbf{f}(\mathbf{x}_k) = \tau_{kl} \left(\frac{\|\mathbf{x}_l - \mathbf{x}_k\| - \Delta s_{kl}}{\|\mathbf{x}_l - \mathbf{x}_k\|} \right) \frac{(\mathbf{x}_l - \mathbf{x}_k)}{\|\mathbf{x}_l - \mathbf{x}_k\|}. \quad (2.1)$$

There is an equal and opposite force due to this spring at the node \mathbf{x}_l . Here, τ_{kl} (kg s^{-2}) is the stiffness constant of the linear spring and Δs_{kl} (m) is its resting length. This resting length is taken to be the distance between the nodes \mathbf{x}_l and \mathbf{x}_k when the fibre is initialized as the straight cylinder shown in figure 3. The total elastic energy E ($\text{kg m}^2 \text{s}^{-2}$) in this node–spring system is given by

$$E(\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_N) = \sum_{\text{springs}} \frac{\tau_{kl}}{2} (\|\mathbf{x}_l - \mathbf{x}_k\| - \Delta s_{kl})^2, \quad (2.2)$$

where N is the total number of nodes. Note that $E = 0$ when the model diatom chain is in its straight, equilibrium configuration.

For prescribed stiffness constants τ_{kl} , we can estimate the macroscopic bending modulus of the model diatom chain using the non-hydrodynamic approach outlined by Lim &

Peskin [11]. The cylindrical structure is bent at a constant curvature β and the total potential energy E stored in all of the springs is calculated using equation (2.2). Assuming that the node–spring structure approximates a homogeneous elastic beam, the stored energy in the bent structure is related to curvature by

$$E = \frac{1}{2} A \beta^2 L, \quad (2.3)$$

where L is the length of the chain and $A = EI$ is its flexural stiffness or bending rigidity. Here, E is the Young’s modulus and I is the chain’s second moment of area. Note that A has units of N m^2 ; and a quadratic relationship between the total energy E and β is noted. A least-squares fit is used to determine the macroscopic bending rigidity A .

Alternatively, one may use the above procedure to determine spring stiffness constants τ_{kl} that will result in a prescribed macroscopic bending rigidity A . This is the approach that we will take in building the computational fibres to match the bending rigidities measured for diatom chains in [8]. For simplicity, in the model diatom chains discussed herein, we choose the same spring stiffness $\tau_{kl} = S_0$ for each of the linkages in the structure.

In order to further characterize the material properties of these computational fibres, we pinned one end of the fibre down and imposed a series of fixed strains to the fibre by extending the other end and fixing it so that the extended fibre was up to twice its equilibrium length. We then found the minimum energy configuration of the spring–node structure in the strained configuration. We found that the energy in the strained systems was a quadratic function of the strain. Therefore, the overall structures are neither strain-hardening nor strain-softening, but Hookean, like the individual spring elements that comprise them.

2.2. Estimating flexural rigidity of diatom chains

Young *et al.* [8] discuss their procedure to calculate flexural rigidities of four distinct types of chain-forming diatoms. The overall strategy was to measure the bending of the chains in response to an applied force. Individual diatom chains were suspended across the rim of a hollow glass capillary tube submerged in a seawater-filled Petri dish. Chains were then exposed to a constant hydrodynamic force that bent the chain into the tube (figure 4a). The volumetric flow rate of

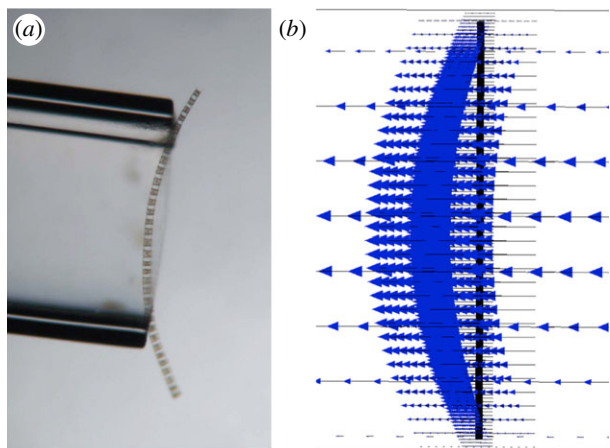


Figure 4. (a) A chain of *G. delicatula* placed across the rim of the capillary tube (courtesy of A. Young and L. Karp-Boss, University of Maine). (b) A computational chain placed in a channel subject to an imposed flow. (Online version in colour.)

the imposed flow was measurable, along with the chain dimensions and the resulting chain deflection. Assuming that the diatom chains in these experiments are governed by small-deflection beam theory, the flexural stiffness (bending rigidity) may be calculated as [12]

$$A = EI = \frac{5fL^4}{384y}, \quad (2.4)$$

where f is the force per unit length at the point of maximum deflection of the chain, y is the maximum deflection of the chain and L is the length of the straight chain. In order to directly apply equation (2.4), an estimate of the force f was needed. Young *et al.* [8] used the relationship,

$$f_{\text{drag}} = C_d \rho u^2 r, \quad (2.5)$$

that relates the drag force on a cylindrical fibre to its radius r , the velocity at the point of maximum deflection u , the density of the seawater ρ and a drag coefficient C_d . This drag coefficient C_d was estimated using an empirical formula based upon the Reynolds number of the flow. We refer the reader to [8] for details. In order to validate this procedure, the flexural stiffness for nylon was also estimated and it compared favourably with previously reported values. Table 1 shows the estimated flexural stiffnesses calculated in a subset of the experiments reported in [8] and provided to us by Young and Karp-Boss during the course of these experiments. Note that the range of measured flexural stiffnesses for different chain types varies over four orders of magnitude. The stiffest diatom chain, *L. undulatum*, has tightly fused, silica-based connections, whereas *G. delicatula*, the most flexible diatom chain measured has a single silica spine connection. Moreover, some types such as *Thalassiosira* sp. were too flexible for the experimental procedure to yield an estimate of the bending rigidity (figure 1c).

Table 1 also shows spring stiffnesses chosen for the model diatom chain so that its bending rigidity based upon computing the total energy in a bent configuration (equation (2.3)) matches the measured bending rigidity. For instance, consider a model of a *L. undulatum* chain that comprised 12 diatoms, with each valve of length 60 μm , and a cell-connection portion of 7.2 μm at each end, for a total length of 0.9 mm. This is discretized using 504 cross sections, with 30 nodes around each hexagonal cross section, and one node along the

Table 1. Chain diameters and average flexural stiffnesses of different diatom chain types measured by A. Young and L. Karp-Boss at the University of Maine (2011, personal communication). These data come from a subset of those experiments reported in [8]. The third column shows the associated stiffness constants used for the springs of the computational diatom chain so that the bend modulus of the model computed using equation (2.3) matches column 2.

diatom chain type	chain diam. (m)	flex stiff (N m^2)	stiff constant (N m^{-1})
<i>L. undulatum</i>	3.24×10^{-5}	1.93×10^{-13}	6.00
<i>S. turris</i>	2.37×10^{-5}	1.23×10^{-13}	7.00
<i>L. annulata</i>	3.08×10^{-5}	6.30×10^{-15}	2.44×10^{-1}
<i>G. delicatula</i>	1.547×10^{-5}	5.73×10^{-17}	7.44×10^{-3}

centreline. The portion of this diatom chain that corresponds to the valve has springs connecting the centreline to the surface. By choosing the spring constant $S_0 = 6.00 \text{ N m}^{-1}$, we assert that the overall bending rigidity of the model diatom chain comprised these 12 diatoms and discretized as stated is $A = 1.93 \times 10^{-13} \text{ N m}^2$.

2.3. Coupling of elastic diatom chain with fluid

We choose an immersed boundary framework to model the interaction of the diatom chain with a viscous, incompressible fluid [13,14]. The system is governed by the three-dimensional Navier–Stokes equations

$$\rho \left[\frac{\partial \mathbf{u}}{\partial t} + \mathbf{u} \cdot \nabla \mathbf{u} \right] = -\nabla p + \mu \Delta \mathbf{u} + \mathbf{F}(x, t), \quad (2.6)$$

$$\nabla \cdot \mathbf{u} = 0, \quad (2.7)$$

$$\mathbf{F}(x, t) = \sum_j \int_{\Gamma_j} \mathbf{f}_j(s, t) \delta(\mathbf{x} - \mathbf{X}_j(s, t)) ds \quad (2.8)$$

$$\text{and} \quad \frac{\partial \mathbf{X}_j(s, t)}{\partial t} = \mathbf{u}(\mathbf{X}_j(s, t), t) \\ = \int_{\Omega} \mathbf{u}(\mathbf{x}, t) \delta(\mathbf{x} - \mathbf{X}_j(s, t)) d\mathbf{x}, \quad (2.9)$$

where ρ is the fluid density, \mathbf{u} is the fluid velocity, \mathbf{X}_j denotes the Lagrangian coordinates of the j th longitudinal filament Γ_j comprising the cylindrical diatom chain (either along its surface or its centreline), p denotes pressure, μ is the dynamic viscosity and Ω denotes the fluid domain. Note that the forces \mathbf{f}_j defined on the filaments Γ_j are precisely sums of the spring forces in equation (2.1). The Eulerian elastic force density \mathbf{F} on the fluid is a Dirac delta-function layer of force that is supported at the material points of the neutrally buoyant diatom chain. Away from these points, the force of the fibre on the fluid is zero.

Traditionally, the immersed boundary method uses a Cartesian background grid to keep track of Eulerian fluid quantities such as velocity and pressure, whereas the material points of the immersed structure are treated in a Lagrangian manner. Grid-based, discretized versions of the Dirac delta function are used to communicate force on the immersed structure to the background grid and to interpolate fluid velocities on the grid to the immersed structure [14]. Here, we use an adaptive and parallel implementation of the immersed boundary method, IBAMR (Griffith *et al.* [10]), that dynamically

Table 2. Maximum deflection and drag measured by the laboratory experiments of A. Young and L. Karp-Boss at the University of Maine (2011, personal communication). These data come from a subset of those experiments reported in [8]. Also shown are corresponding maximum deflection and drag force resulting from our coupled three-dimensional fluid/fibre model.

species	(lab) max. defl. (m)	(comp) max. defl. (m)	(lab) drag (N m ⁻¹)	(comp) drag (N m ⁻¹)
<i>Lithodesmium</i>	4.0×10^{-5}	1.3×10^{-5}	8.8×10^{-5}	3.8×10^{-5}
<i>S. turris</i>	4.4×10^{-5}	2.8×10^{-5}	1.8×10^{-4}	7.4×10^{-5}
<i>L. annulata</i>	6.4×10^{-5}	5.4×10^{-5}	4.2×10^{-5}	1.9×10^{-5}
<i>Guinardia</i>	6.2×10^{-5}	2.9×10^{-5}	6.4×10^{-8}	8.5×10^{-8}

refines the Cartesian mesh to its finest level around the immersed structure as it moves throughout the fluid domain. Adaptivity allows the use of a large computational domain Ω , because a coarse mesh may be used away from the immersed boundary, while details of the fluid–structure interaction near the immersed boundary are captured by a fine mesh. In particular, a cell-centred, second-order projection method on a hierarchical structured grid is used to solve the incompressible Navier–Stokes equations. The viscous terms are treated implicitly, and a second-order Godunov method for the explicit treatment of the advective term is employed. A second-order Runge–Kutta method is used to update the system in time (see [10] for implementation details).

To simulate the experiments done in the laboratory, we immerse the diatom chain models in a background parabolic flow (figure 4b). We set $\rho = 10^3 \text{ kg m}^{-3}$ and $\mu = 10^{-3} \text{ Pa s}$. Note that the spring stiffnesses were chosen as in table 1 to give an overall bending rigidity that was measured for each diatom chain type. Cross-sectional diameters of the chains were calibrated for each of the types, but the same overall connectivity structures as shown in figure 3 were used in each case. Tethering forces to keep the cross sections at the ends of the chains fixed near the no-slip boundary were used. In addition, the parameters for the parabolic flow were chosen to match the volumetric flow rates as in the laboratory set-up. The diatom chain, whose equilibrium configuration is a straight cylinder, is initially placed vertically, but is bent by the imposed parabolic flow. Soon the coupled fluid-chain system reaches a steady state. As in the laboratory experiments, we are able to measure the maximum displacement of the chain. However, unlike the laboratory experiments, we can directly measure (rather than estimate) the drag force at the point of maximum displacement. The maximum displacement and the drag force based upon laboratory experiments are shown in table 2 along with the corresponding computational measurements of both.

Here, we see that the computational measurements of maximum displacement and drag force agree with those estimated using laboratory experiments to well within an order of magnitude for all diatom chain types. While neither the true diatom chains nor the spring–node computational models of the diatom chain are homogeneous beams, the use of small-deflection beam theory allows us to quantify an effective macroscopic bending rigidity of the chain.

Similar computational experiments were performed for model chains that were made up of only surface-spring connections and full-spring connections. When compared to the more heterogeneous diatom chain model, one would guess that the surface-spring chain model obtained by removing all of the spoke connections in the valve portions would result in a structure with a smaller macroscopic bending rigidity. Similarly, a

full-spring chain model obtained by adding spoke connections at every cross section would presumably have a larger macroscopic bending rigidity. In fact, this is the case. For instance, for a model of *G. delicatula* of length 0.9 mm and bending rigidity $A = 5.73 \times 10^{-17} \text{ N m}^2$, the corresponding surface-spring structure has a bending rigidity of $A = 5.65 \times 10^{-17} \text{ N m}^2$, and the corresponding full-spring structure has a bending rigidity of $A = 5.89 \times 10^{-17} \text{ N m}^2$. When these three structures were subjected to the imposed parabolic flow, their maximum deflections and drag forces at the point of maximum deflection were basically identical. By contrast, we will see below that these three structures do not behave similarly when subjected to a shear flow, where compressive stresses give rise to buckling behaviour.

3. Semiflexible fibres in shear flow

The seminal assessment of Munk & Riley [15] indicates that when the size of phytoplankton is more than 10 or so micrometres, ambient fluid motion can enhance fluxes of solutes to or from cells. Lazier & Mann [16] have also suggested that because phytoplankton are typically smaller than the Kolmogorov length scale in the ocean, phytoplankton experience turbulence as a linear shear. For this reason, the fluid dynamics of diatom chains driven by linear shear flows is important to the study of nutrient delivery to phytoplankton. While recent work suggests that more complex background flows around phytoplankton, such as dissipative vortices, should be considered [17]; here, we focus on the behaviour of diatom chains in shear flow. In a steady shear flow, chains tend to align with their long axis parallel to the flow and tumble periodically [18]. These orbits are reminiscent of the rotational orbits of an ellipsoid in shear, whose analysis appears in the classic work of Jeffery [19]. Jeffery showed that the rotational period T of a spheroid in Stokes flow with an imposed shear rate κ was a function of the spheroid's aspect ratio a_r (major axis/minor axis)

$$T = \frac{2\pi}{\kappa} \left(a_r + \frac{1}{a_r} \right). \quad (3.1)$$

Karp-Boss & Jumars [18] studied the motion of two types of chain-forming diatoms, *Skeletonema costatum* and *Thalassiosira*, and compared the observed rotational periods to that predicted by Jeffery's theory for a spheroid of the same aspect ratio. They observed that rotational periods of the elongated chains fell well below those predicted by the theory for the corresponding spheroid and that the more flexible chains typically rotated faster than more rigid ones of the same aspect ratio.

The study of semiflexible fibres in shear flows certainly pre-dates the above-mentioned investigation of diatom chains in

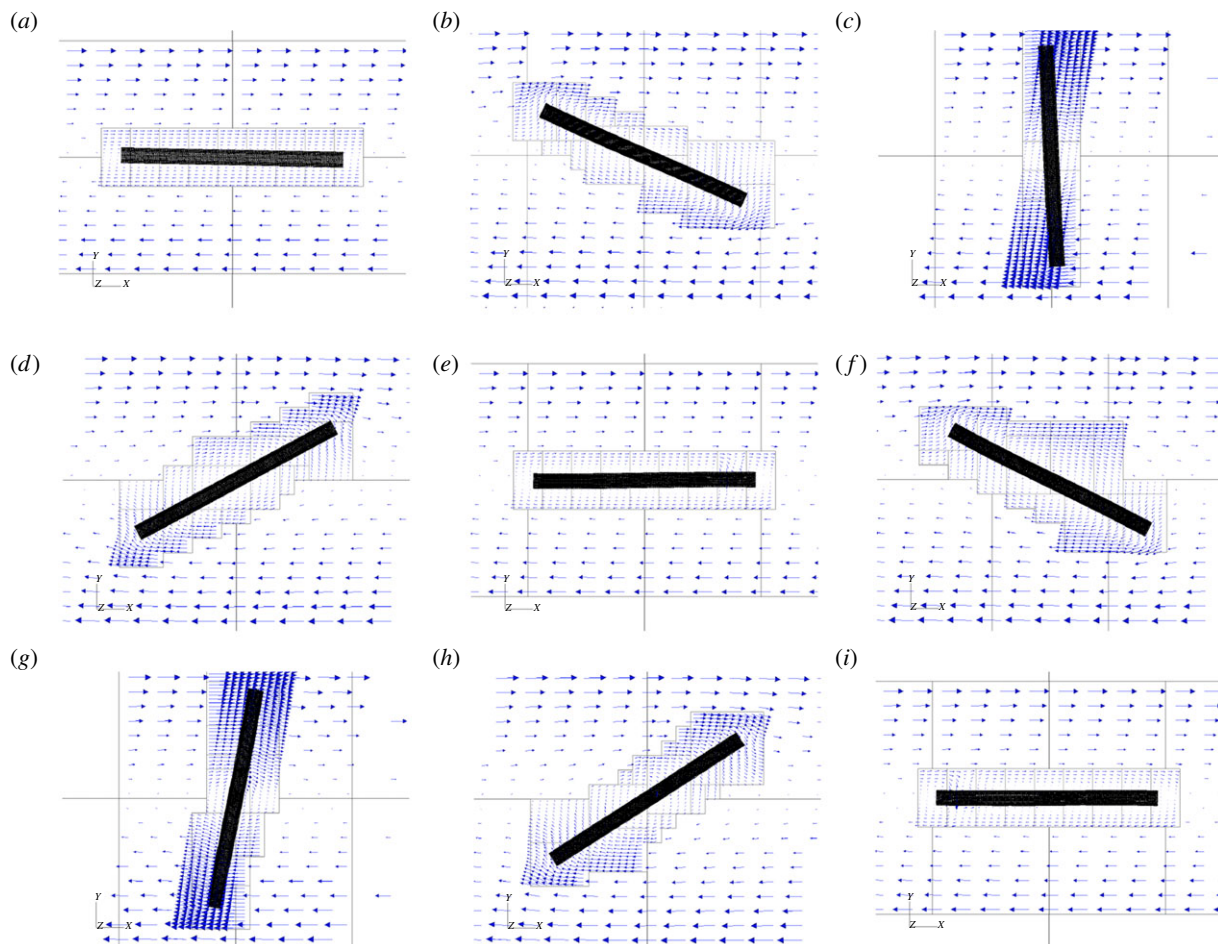


Figure 5. One rotational period of a diatom-chain structure of non-dimensional length $L_c = 1$ in an imposed shear flow. The fibre length is 14.43 times its diameter. The Reynolds number $Re = 0.06$ and the effective flow forcing $\bar{\mu} = 1.15 \times 10^3$. (Online version in colour.)

shear. For instance, the classical work of Forgacs & Mason [4] characterized the qualitative shape deformations of flexible fibres in shear by tracking the orbits of pulp, Dacron and elastomer fibres. Springy rotations, snake rotations and coil formations were observed for fibres of different lengths. As in the diatom chain experiments [18], more flexible fibres were found to have shorter rotational periods than stiffer fibres with the same aspect ratio. Forgacs and Mason also noted that for a given fluid viscosity and shear rate, a critical fibre length existed beyond which the threadlike particle bent. The deformation increased with the fibre length. In the past decades, this fibre buckling has been analysed using slender body hydrodynamics [20–22], bead models of the fibre that neglect hydrodynamic interaction of fibre segments [23] and computational models that couple the fluid equations with the fibre forces [24–26]. Below we will examine buckling in our fibre models, but first we will introduce the non-dimensional parameters that govern this elastic-hydrodynamic system.

3.1. Reynolds number and effective flow forcing

The Reynolds number that governs the dynamics of the cylindrical fibre in a background shear flow measures the relative importance of inertial forces to viscous forces. To define the Reynolds number, we use the fibre length L and the characteristic velocity κL , where κ is the shear rate (s^{-1})

$$Re = \frac{\rho \kappa L^2}{\mu}. \quad (3.2)$$

The other governing dimensionless parameter measures the relative importance of fluid forces to elastic forces

$$\bar{\mu} = \frac{8\pi\mu\kappa L^4}{A}. \quad (3.3)$$

This dimensionless parameter is referred to in [22] as the effective viscosity, but here we refer to $\bar{\mu}$ as the effective flow forcing. Here, A is the flexural stiffness or bending rigidity of the fibre. We can see the appearance of the non-dimensional bending rigidity when we consider the local slender body model of a filament in flow that gives a local relation between the velocity of the filament centreline and the force the filament exerts on the fluid

$$8\pi\mu \left(\frac{dx}{dt} \right) = c(I + x_s x_s) f. \quad (3.4)$$

Here, $x(s, t)$ is the position of the fibre's centreline, x_s is the tangent vector to the centreline and $c \approx \ln \epsilon^{-1}$, where ϵ is the ratio of the filament's radius to its length. If equation (3.4) is non-dimensionalized using the filament's length L as a characteristic length, the velocity κL as a characteristic velocity and A/L^3 as a characteristic force per unit length, the effective flow forcing emerges. Note that the effective flow forcing $\bar{\mu}$ that is felt by a fibre increases as the fluid viscosity increases, increases as the length of the fibre increases, increases as the shear rate increases, but decreases as it becomes stiffer (as its bending rigidity increases).

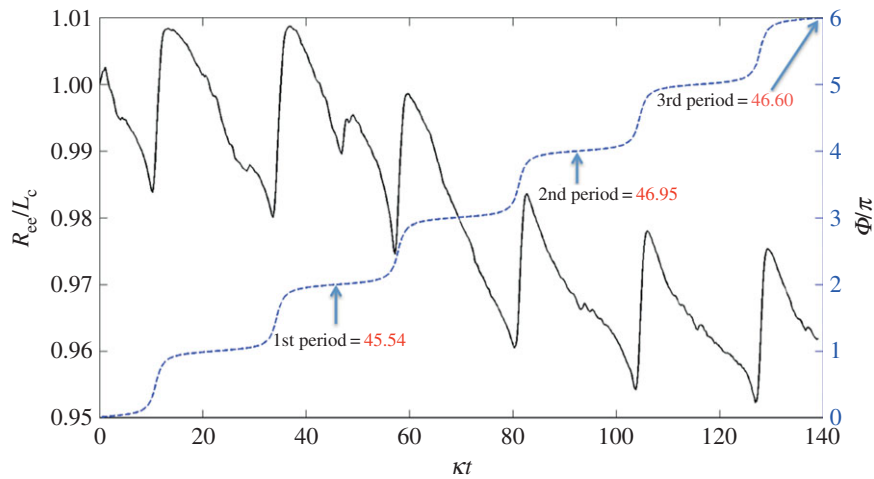


Figure 6. Evolution of the ratio of end-to-end distance R_{ee} to equilibrium length L_c (solid line) and orientational angle of the diatom-chain orbit (dashed line) shown in figure 5. (Online version in colour.)

Typical shear rates experienced by phytoplankton in the ocean are estimated to be $0.01 \text{ s}^{-1} \leq \kappa \leq 1 \text{ s}^{-1}$ and typical lengths of diatom chains are estimated to be $0.25 \text{ mm} \leq L \leq 4 \text{ mm}$ [7]. For a fluid with the density and viscosity of seawater, and the types of diatom chains whose bending rigidities are shown in table 1, the resulting Reynolds numbers and effective flow forcing experienced are in the range

$$6.25 \times 10^{-4} \leq Re \leq 16 \quad \text{and} \\ 5.00 \times 10^{-4} \leq \bar{\mu} \leq 1.1 \times 10^5. \quad (3.5)$$

The upperbound for effective flow forcing, for example, would be achieved by a *G. delicatula* that is 4 mm long, exposed to a shear rate of 1 s^{-1} . It is possible that more flexible diatom chain types such as *Thalassiosira* sp., whose bending rigidity could not be measured using the aspiration technique in [8], experience effective flow forcings that are orders of magnitude more.

3.2. Diatoms in shear

We will now examine the results of numerical experiments where we subject our computational chains to a background shear flow. The connectivity of these chains will be either full-spring, surface-spring or that of the alternating diatom chain structure.

First, we will consider a diatom chain of non-dimensional length $L_c = 1$ whose structure is shown in figure 3. Here, L_c is the length of the entire chain, and, in this case, the chain comprised three connected diatom cells. The aspect ratio of the chain (length/diameter) is 14.43. The three-dimensional computational domain is a square box of dimensions $ML_c \times ML_c \times ML_c$, where $M = 67$ in the following simulations. The background shear flow is achieved by specifying velocity boundary conditions at the upper and lower planes bounding the domain. Note that the use of an adaptive mesh allows us to use this large computational domain, while still resolving the fluid grid around the diatom chain. Here, we chose Reynolds number $Re = 0.06$ and an effective flow forcing of $\bar{\mu} = 1.15 \times 10^3$. This effective flow forcing would correspond to a *Lauderia annulata* whose length is approximately 2.4 mm in a shear flow of $\kappa = 1 \text{ s}^{-1}$, or a *G. delicatula* whose length is approximately 1.27 mm in the same flow. We note that here

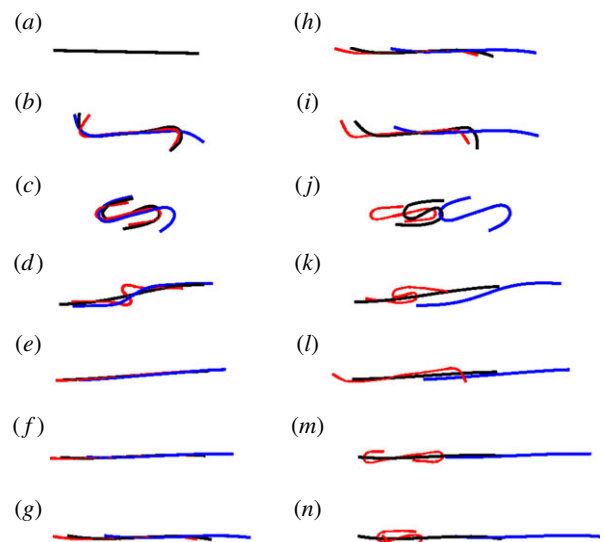


Figure 7. Three chain types subjected to the same shear. Diatom-chain type (black, $A = 5.73 \times 10^{-17} \text{ N m}^2$), surface-spring (red, $A = 5.65 \times 10^{-17} \text{ N m}^2$) and full-spring type (blue, $A = 5.89 \times 10^{-17} \text{ N m}^2$). Although the macroscopic bending rigidity of these chains differs by less than 5%, their orbits are very different.

we are computing at a Reynolds number somewhat smaller than that experienced by chains of these lengths, at the given shear, in seawater.

Figure 5 shows successive snapshots of this diatom chain as it goes through a complete orbit. Note that at time $t^* = 0$ (upper left frame), the initial position is slightly perturbed from the horizontal. The imposed shear flow is evident by the velocity vectors depicted on the dynamic, adaptive mesh. The centreline of this mostly rigid fibre remains in a plane due to symmetry. Figure 6 tracks the evolution of the angle that the fibre makes with the horizontal as a function of the non-dimensional time $t^* = \kappa t$. As is typical of Jeffery orbits, the diatom chain spends most of its time aligned with the flow in a horizontal position, interspersed with quick flipping events. Using Jeffery's theory (equation (3.1)) for an ellipse of aspect ratio $a_r = 14.43$, the approximate rotational period would be $t^* = 91.1$. Using a slender body

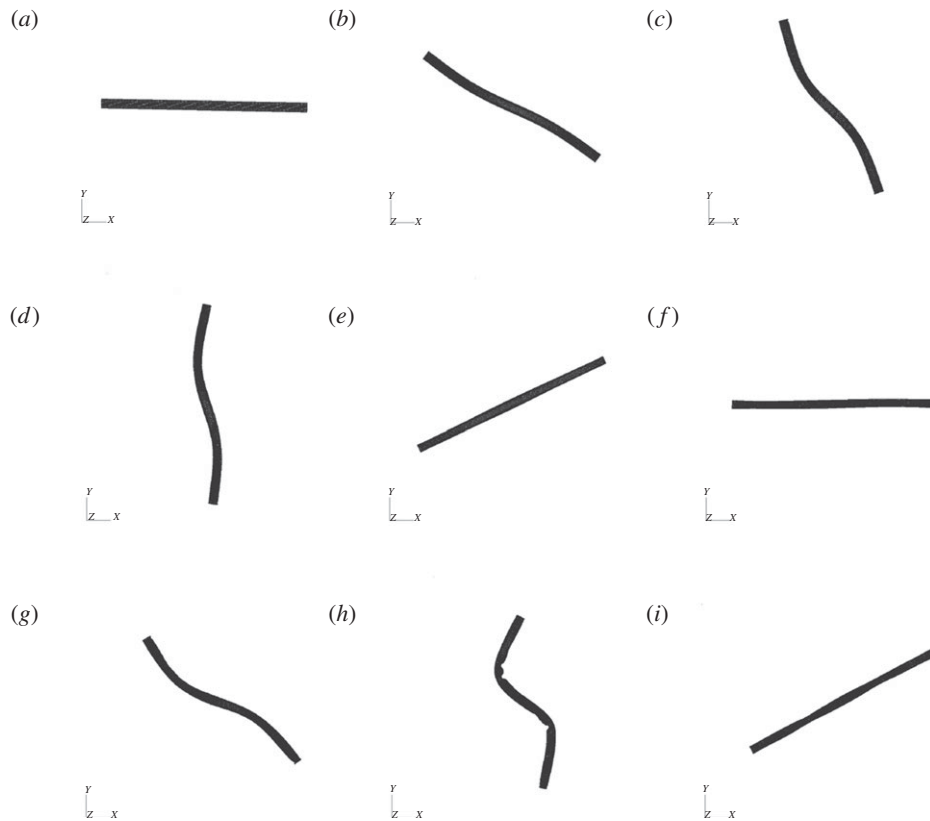


Figure 8. (a–e) Orbit of a surface-spring structure of non-dimensional length $L_c = 1.33$ in an imposed shear flow. The fibre length is 19.24 times its diameter. The Reynolds number $Re = 0.10$ and the effective flow forcing $\bar{\mu} = 3.64 \times 10^3$.

approximation, Cox [27] determined an effective aspect ratio $\hat{a}_r = 1.24a_r/\sqrt{\ln a_r}$ to be used in Jeffery's formula for a rigid slender cylinder of aspect ratio a_r . This effective aspect ratio ($\hat{a}_r = 10.95$) would result in an approximate rotational period of $t^* = 69.38$. Figure 6 shows that the rotational period of our computational fibre is approximately $t^* = 47$. We note two main reasons for the difference between the computed rotational period and that predicted by the theory. Firstly, the aspect ratio of this diatom chain does not fall well within the range valid for slender body approximation. Recently, Zhang *et al.* [28] discussed the applicability of the slender body approximation [27] to fibres with smaller aspect ratios. Secondly, this fibre is not perfectly rigid. For thin, rod-like structures, the rotational period is dominated by the time that the rod stays in the horizontal position, aligned with the flow. For even a modest amount of flexibility, that horizontal position is disturbed, causing an early onset flipping event. Figure 6 also shows the evolution of the ratio of the end-to-end distance of the fibre to its equilibrium length. This ratio would be identically one for a rigid cylinder. Note that the computational fibre underwent both compression and extension from its equilibrium length, and, although no significant deformation is evident in figure 5, the end-to-end length of the fibre does vary by as much as 5% of its equilibrium. Flexibility was noted by Forgacs & Mason to give rise to rotational periods that were shorter than those of a rigid fibre with the same aspect ratio [4]. We also remark that we have validated the computational method used here for an ellipsoid rotating in shear flow by comparing the resulting rotational dynamics with that predicted by Jeffery in [29].

The related surface-spring and full-spring structures of non-dimensional length $L_c = 1$, as discussed above, have

macroscopic bending rigidities that fall within 5% of this diatom chain structure. Although not shown here, we subjected these other structures to the same shear and observed dynamics that were basically indistinguishable from the diatom chain dynamics shown in figure 5. There was little evidence of deformation, even for the structure with links only between surface nodes (surface-spring).

Although similar orbits are observed for these relatively short, rigid fibres with different spring connectivities (diatom-chain, surface-spring and full-spring), this is no longer the case when we increase the fibre length fourfold ($L_c = 4$). Increasing the length results in a Reynolds number of $Re = 0.94$ and an effective flow forcing of $\bar{\mu} = 2.95 \times 10^5$. The aspect ratio of these longer chains is $a_r = 57.71$. Figure 7 shows the positions and deformations of the three chain types as they undergo a single orbit. As expected, the full-spring (blue) and diatom-chain (black) deform less than the chain with surface-springs (red). The dynamics of these three chains in shear, when they are experiencing compressive stresses, are very different. Such a difference did not appear when the chains were subjected to the parabolic flow, as in the aspiration experiments described above.

For even the full-spring fibre of length $L_c = 4$ in figure 7 above, we note the onset of buckling. This did not happen at the length $L_c = 1$. Keeping all other parameters fixed, is there a critical fibre length that will result in buckling? Consider the dynamics of a surface-spring structure in shear that we know did not buckle at $L_c = 1$, but did at $L_c = 4$. Changing its length to $L_c = 1.33$, and its aspect ratio to $a_r = 19.24$ with all other parameters fixed results in a Reynolds number of $Re = 0.104$ and an effective flow forcing of $\bar{\mu} = 3.64 \times 10^3$. Figure 8 shows a series of snapshots of the fibre as it undergoes a single orbit. At this larger value of effective flow forcing, the

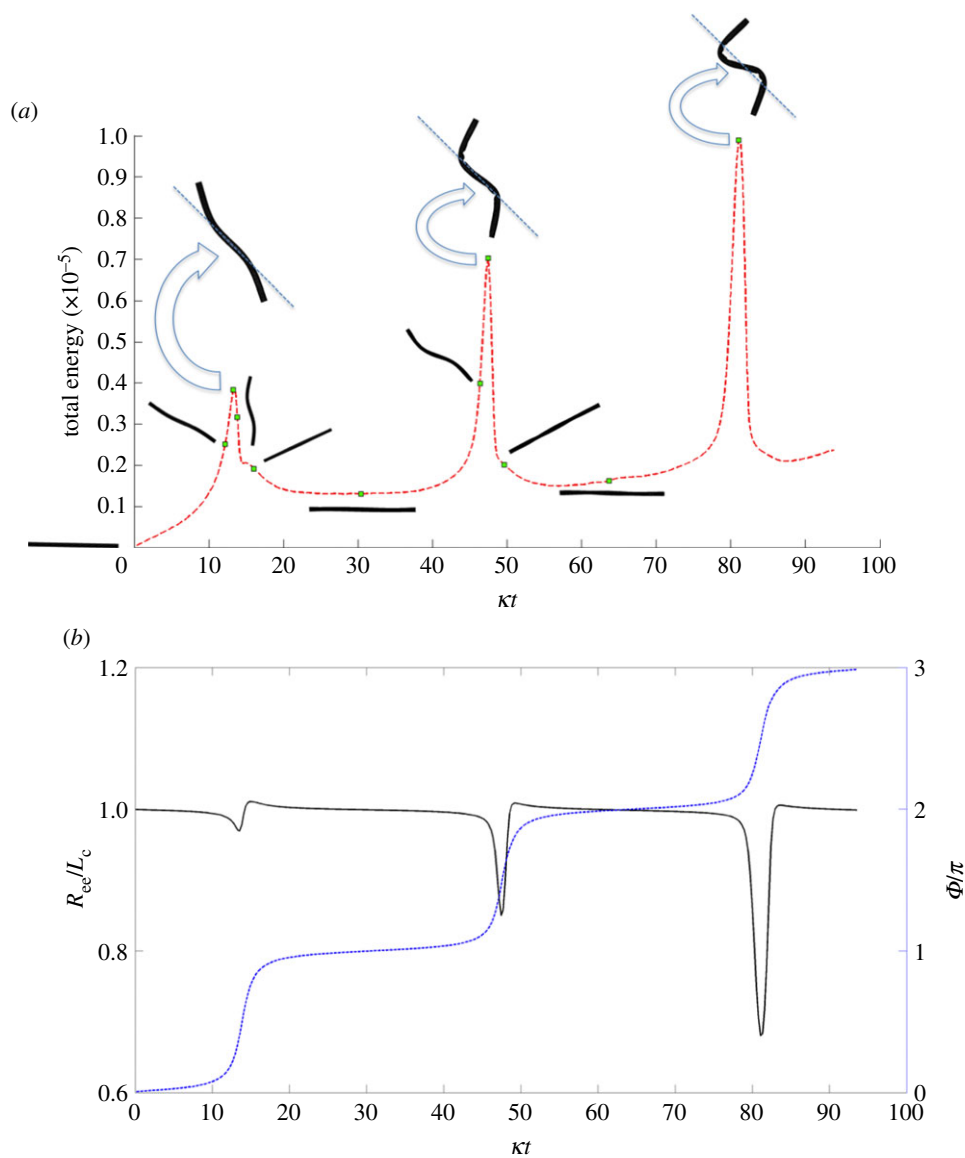


Figure 9. (a) Evolution of the total elastic energy of the surface-spring fibre as it rotates in shear. The square dots represent the total energy associated with the shapes of the fibre at the different times shown in figure 8. (b) Evolution of ratio of end-to-end distance R_{ee} to equilibrium length L_c (solid line) and orientational angle of the surface-spring fibre's orbit (dashed line) shown in figure 8. (Online version in colour.)

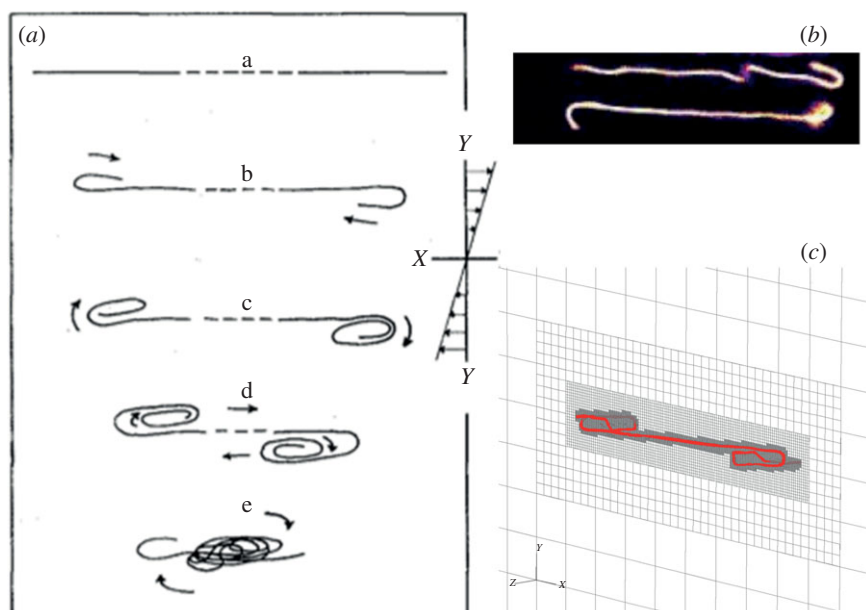


Figure 10. (a) Schematic sketch from Forgacs & Mason [4] showing coiling of a very long fibre in shear; (b) multiple bending events occurring for a long actin filament in shear (images from Harasim *et al.* [2]); (c) a snapshot from our simulation for $L_c = 16.67$. (Online version in colour.)

buckling due to compressive stresses is evident. Because this fibre is built out of a network of nodes and springs, we can measure the total amount of stored energy in the springs as a function of non-dimensional time (figure 9a). We see local peaks in stored energy at the instances where the fibre is 45° from the horizontal, when it is undergoing the maximal compressive stress. While the energy in the springs does decrease as the fibre rotates from this position, we see that the fibre springs do not have enough time to get back to equilibrium, and the magnitude of these peaks in energy increase during successive periods. Figure 9b shows the evolution of the ratio of the end-to-end distance of the fibre to its equilibrium length, along with its rotational angle. Here, we see that the end-to-end distance during buckling gets to as low as 70% of the equilibrium length.

Qualitatively, our results compare well with those of Tornberg & Shelley [22]. They demonstrated that buckling occurs at a threshold value of effective flow forcing (effective viscosity) about $\bar{\mu} = 2 \times 10^5$, while our calculations show this buckling occurs at about $\bar{\mu} = 3.64 \times 10^3$. However, the aspect ratio of their slender fibres was assumed to be $a_r = 10^3$, while our fibre that first exhibited buckling had an aspect ratio of $a_r = 19.24$. Therefore, the appropriate non-dimensional quantity to compare would be

$$\bar{\bar{\mu}} = \frac{\bar{\mu}}{\ln(a_r^2/e)}, \quad (3.6)$$

which takes into account the aspect ratio [3,21]. The buckling observed in [22] occurred at $\bar{\bar{\mu}} = 1.56 \times 10^4$ and our simulations show buckling at $\bar{\bar{\mu}} = 7.4 \times 10^2$. While these differ by a factor of about 20, we do remark that the fibres of [22] are perfectly inextensible, which is certainly not the case for the computational fibres presented here.

3.3. Complex fibre orbits in shear

While our motivation for this research is to understand the local flow environment around diatom chains and the resulting implications on nutrient uptake, the computational model developed here is applicable to fibre dynamics in flow where the Reynolds numbers and effective flow forcings fall outside the range relevant to diatom geometries and flow parameters. Following the classic paper of Forgacs & Mason [4], we sought to capture the coil dynamics of long semiflexible fibres, which were only sketched by the authors (figure 10a). It is noted that the deformation of the fibre, while it is aligned with the flow direction, begins first with the bending of the ends. More recently, experiments by Harasim *et al.* [2] on long actin filaments reveal similar dynamics (figure 10b).

Using the same surface-spring structure and parameters as in the simulation of figure 8, but with an increased non-dimensional length of $L_c = 16.67$ and aspect ratio of $a_r = 240.46$, we subject the fibre to a shear flow with Reynolds number $Re = 1.67 \times 10^1$ and the effective flow forcing of $\bar{\mu} = 8.9 \times 10^7$. Figure 11 shows a sequence of snapshots of the coil dynamics of this long, flexible fibre in shear as it undergoes one rotation. Also shown are the adaptive meshes that capture this complex coiling and recovery. We first note the correspondence between the dynamics of this fibre and the sketch of [4] shown in figure 10a. We also see, in the second and third frame, that the bending of the ends is accompanied by complex buckling in the centre of the fibre, as also seen in the actin polymer micrographs of Harasim *et al.* [2] in figure 10b. In the course

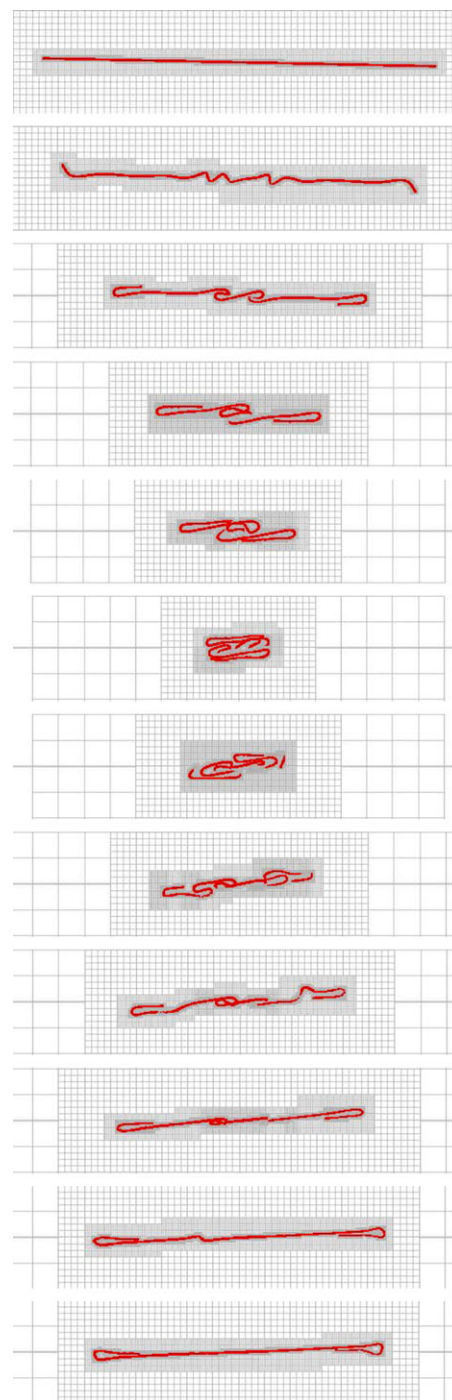


Figure 11. Orbit of a surface-spring structure of non-dimensional length $L_c = 16.67$ in an imposed shear flow. The fibre length is 240.47 times its diameter. The Reynolds number $Re = 1.67 \times 10$ and the effective flow forcing $\bar{\mu} = 8.9 \times 10^7$. (Online version in colour.)

of one period, the fibre is able to recover from its entangled coil, but the ends remain bent (electronic supplementary material, movie S1). Although only one period is shown here, this long fibre continued to undergo a few more orbits going from a coiled configuration back to an elongated configuration, but the fibre did not recover from its coiled state. During this complex motion, because of the slight asymmetry caused by adaptive mesh generation, the centreline of the fibre does not remain in the plane, and the bent ends have a three-dimensional component to them (figure 10c).

Lindstrom & Uesaka [24] presented a fluid dynamic model of long, flexible fibres in shear that could exhibit coiled dynamics. However, it was proposed that such

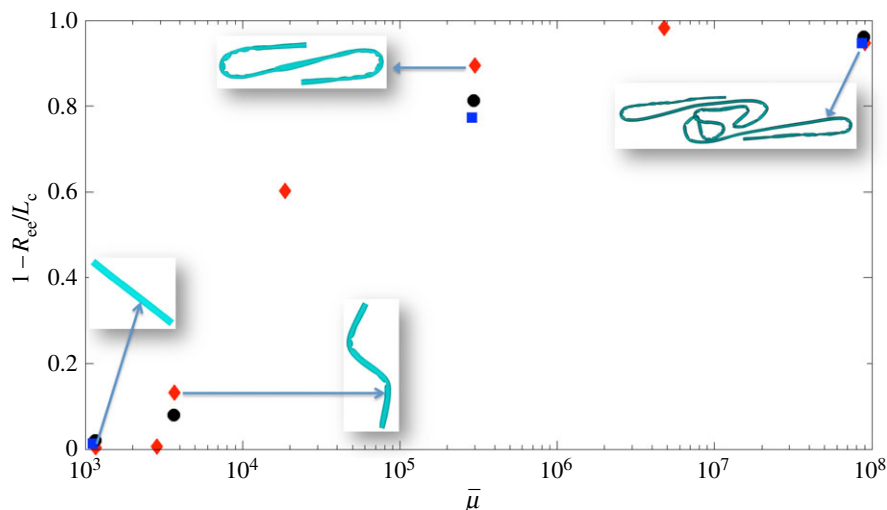


Figure 12. The order parameter $1 - R_{ee}/L_c$ plotted versus $\bar{\mu}$ for each of the simulations described above. Here, the minimum value of end-to-end distance R_{ee} achieved within the first rotational period was used. The diamonds indicate surface-spring fibres, the dots indicate diatom-chain structures and the squares indicate full-spring structures. (Online version in colour.)

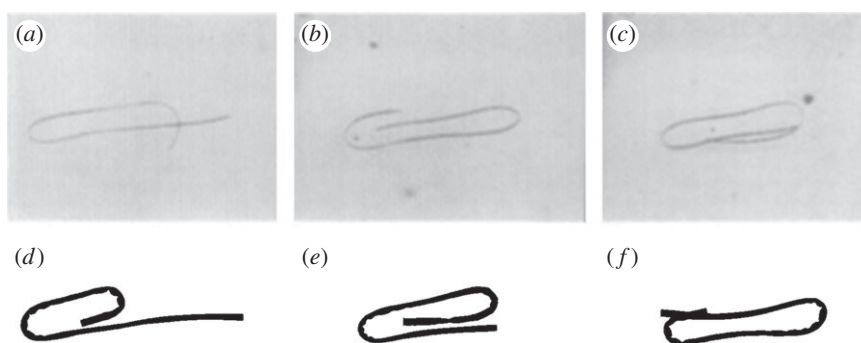


Figure 13. (a–c) Snapshots of an elastomer fibre's motion from Forgaçs & Mason [4]; and (d–f) equivalent snapshots from our simulation.

coiled motion could only be achieved when the fibre was endowed with an intrinsic, non-zero curvature. However, the complex coil dynamics we present above assumed an equilibrium configuration of a straight cylinder.

A convenient order parameter that measures the extent of buckling of a fibre is $1 - R_{ee}/L_c$ [3], where L_c is the equilibrium length of the straight fibre, and R_{ee} is the actual end-to-end distance in its evolving configuration. We summarize the results of the previous simulations in figure 12 by choosing the minimum value of end-to-end distance achieved within the first period of rotation of the fibre to compute the order parameter and plot that versus the effective flow forcing $\bar{\mu}$ corresponding to the simulation. The diamonds indicate surface-spring fibres, the dots indicate diatom-chain structures and the squares indicate full-spring structures.

Figure 13 demonstrates that this computational model can also capture the closed loop dynamics that are also described for an elastomer filament by Forgaçs & Mason [4] (electronic supplementary material, movie S2). We note that the dynamics depicted here are fully three-dimensional.

We also note the interesting dynamics of a non-uniform chain where the first third of its length has full-spring connections while the remaining two-thirds just has surface-spring connections. Figure 14 shows a few snapshots of this fibre as it undergoes rotation and deformation (electronic supplementary material, movie S3).

4. Discussion

Here, we have studied the elasto-hydrodynamics of semiflexible fibres in a moving fluid. In particular, our motivation is to understand the local flow environment of phytoplankton and its role in nutrient uptake and aggregate formation. Of particular interest are the diverse species of chain-forming diatoms [18]. Only recently has there been success in quantifying the flexural rigidities of some types of these diatom chains [8]. While some species were too flexible for the experimental method to yield a rigidity estimate, the bending rigidities of those species measured varied over four orders of magnitude. We have developed a three-dimensional computational model, based upon an immersed boundary framework [14], that couples an inhomogeneous elastic chain to an incompressible fluid. The elastic properties of the node-spring network comprising the computational fibre were chosen to match the macroscopic bending rigidities of the diatom chains measured in [8].

The computational model presented here does not treat the diatom chain as a simple beam, nor does it make the assumptions of slender body theory. Nevertheless, we have found that simple beam theory does a good job of describing the deformation of the inhomogeneous fibre when it is subjected to a parabolic flow. However, we found that although computational fibres with different connectivity (surface-spring and full-spring) have nearly equal macroscopic bending rigidities,

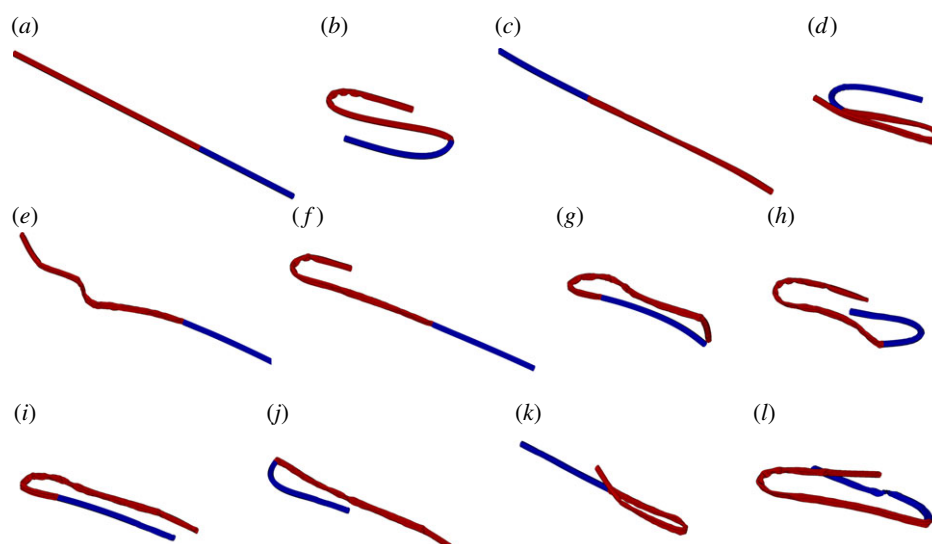


Figure 14. Snapshots of rotational orbits of an inhomogeneous chain, where one-third of the length has full-spring connections (blue) while the remaining two-thirds just has surface-spring connections (red). Note that these frames are not equally spaced in time.

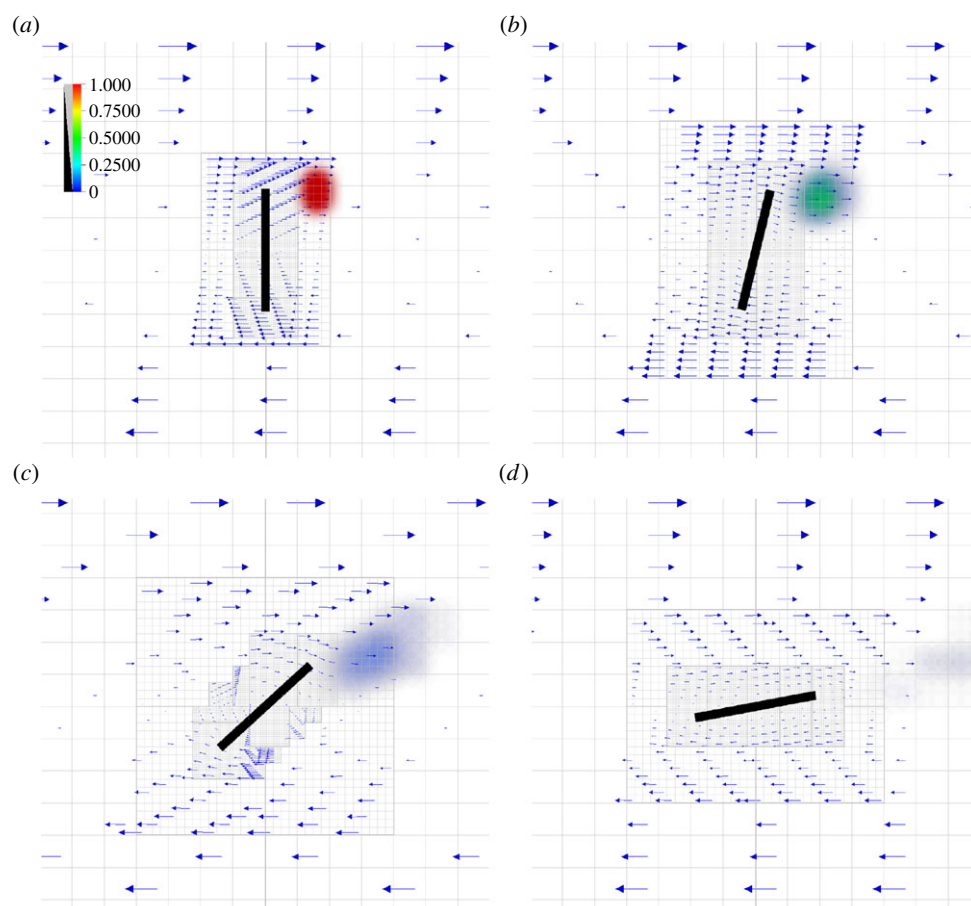


Figure 15. From (a–d), evolution of a bolus of nutrient concentration in the diatom-shear system. The Peclet number that measures ratio of the diffusion timescale to the advection timescale, LU/D , here is about 76, where L is the length of the diatom, $U = L\kappa$, where κ is shear rate, and D is the diffusivity of the nutrient.

when subjected to a shear flow with large enough effective flow forcing, their shape evolutions vary dramatically.

Our computational model also suggests that for the species of diatoms whose flexural rigidities were quantified in [8], *L. undulatum* and *Stephanopyxis turris* would not exhibit buckling in shear flows typically encountered in the ocean, but *L. annulata* and *G. delicatula* chains of more than 1–2 mm could. While the bending rigidity of more flexible species

such as *Thalassiosira* was not quantified, their deformation in shear was reported in [18]. There it was conjectured that the flexibility of these chains may allow them to resist breakage when subjected to shear.

In order to gain more insight into the role of flexibility in diatom chains, we plan to extend this computational model in several directions. Firstly, as in the earlier two-dimensional studies of nutrient acquisition by diatom chains [9], the

advection, diffusion and consumption of a chemical species around the diatom chain will be incorporated. This may readily be done since the velocity field is available at grid nodes in the entire domain. To demonstrate the future development of this model, we show the evolution of a bolus of nutrient that is diffusing and being advected by the diatom-shear system in figure 15. Just as this adaptive method [10] allows the computational model to resolve complicated fluid–structure interactions around the diatom chain, the resolution of nutrient dynamics will be achieved by adding grid-adaptivity that tags spatial regions with large nutrient gradients for refinement. (This has not been done in figure 15.) Secondly, assessing the effects of turbulence on phytoplankton solely by analysing simple shear flow has recently been questioned [17]. There it was suggested that dissipative vortices, such as a Burgers

vortex, may better represent the local turbulent environment felt by diatom chains. Although here we focused upon diatom chain orbits when subjected to simple, steady shear flow, a richer set of fully three-dimensional background flows will be examined.

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