Microswimming in viscoelastic fluids
Gaojin Li a,b, Eric Lauga c, Arezoo M. Ardekani d,*

a State Key Laboratory of Ocean Engineering, Shanghai 200240, China
b School of Naval Architecture, Ocean and Civil Engineering, Shanghai Jiao Tong University, Shanghai 200240, China
c Department of Applied Mathematics and Theoretical Physics, University of Cambridge, Wilberforce Road, Cambridge CB3 0WA, UK
d School of Mechanical Engineering, Purdue University, West Lafayette, IN 47907, USA

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A B S T R A C T

The locomotion of microorganisms and spermatozoa in complex viscoelastic fluids is of critical importance in many biological processes such as fertilization, infection, and biofilm formation. Depending on their propulsion mechanisms, microswimmers display various responses to a complex fluid environment: increasing or decreasing their swimming speed and efficiency, modifying their propulsion kinematics and swimming gaits, and experiencing different hydrodynamic interactions with their surroundings. In this article, we review the fundamental physics of locomotion of biological and synthetic microswimmers in complex viscoelastic fluids. Starting from a continuum framework, we describe the main theoretical approaches developed to model microswimming in viscoelastic fluids, which typically rely on asymptotically-small dimensionless parameters. We then summarize recent progress on the mobility of single cells propelled by cilia, waving flagella and rotating helical flagella in unbounded viscoelastic fluids. We next briefly discuss the impact of other physical factors, including the micro-scale heterogeneity of complex biological fluids, the role of Brownian fluctuations of the microswimmers, the effect of polymer entanglement and the influence of shear-thinning viscosity. In particular, for solution of long polymer chains whose sizes are comparable to the radius of flagella, continuum models cannot be used and instead Brownian Dynamics for the polymers can predict the swimming dynamics. Finally, we discuss the effect of viscoelasticity on the dynamics of microswimmers in the presence of surfaces or external flows and its impact on collective cellular behavior.

1. Introduction

In their natural biological environment, microorganisms and spermatozoa often swim in complex fluids with non-Newtonian characteristics. Examples include mammalian spermatozoa swimming in viscoelastic mucus in the cervix or along the fallopian tubes [1–7], the bacterium *Helicobacter pylori* moving through the mucus layer covering the stomach and causing ulcers [8,9], the Lyme disease spirochete *Borrelia burgdorferi* penetrating the connective tissues in our skin [10,11], and bacteria producing extracellular polymeric substances (EPS) and forming biofilms (Fig. 1a) [12–19]. In marine environments, transparent exopolymer particles made of dissolved organic matter or polymer chains released by phytoplankton can initiate bacteria attachment to surfaces and serve as nutrient hot spots [20]. Another important example of a biological flow of complex fluids at the micron scale is the motion of cilia transporting layers of mucus that cover the epithelium of organs and glands. This motion plays a crucial role in the removal of foreign substances in the mammalian respiratory tracts, the transport of fallopian tube isthmic mucus and ovum, and the transcervical spermatozoa migration in reproductive tracts [6,21–23]. In addition,ependymal cilia found in specialized brain cells are involved in the transport of cerebrospinal fluid on small scales [24].

In fluid mechanics, the Reynolds number $Re$, which is ratio of inertial force to viscous force, is one of the most important dimensionless parameter. In the context of micro-scale locomotion at the heart of this paper, we define $Re \equiv \rho U c / \mu$, where $l_c$ and $U_c$ are the characteristic length and swimming speed of the microorganism, while $\rho$ and $\mu$ are the mass density and dynamic viscosity of the fluid, respectively. Since cells move in a low-Reynolds number world, $Re \ll 1$, when they swim in a Newtonian fluid they must adopt non-reversible strokes to escape the constraints of Purcell’s scallop theorem and achieve self-propulsion [27].

A variety of swimming strategies are employed by motile cells in nature [28]. Some protozoa, such as *Paramecium*, and algae, such as Volvox, rely on metachronal waves generated by the collective beating motion of their cilia. Many bacteria, such as the model organism *Escherichia coli* (*E. coli*), swim by rotating helical flagellar filaments...
attached to the cell body, while spirochete bacteria, such as *Spiroplasma*, swim by the propagation of rotational waves or shape kinks along their spiral cell bodies. In contrast, most spermatozoa and some nematodes, such as *Caenorhabditis elegans* (*C. elegans*), swim using a planar waving deformation of their flagella or slender body. In the lab, different types of artificial swimmers have been designed and fabricated to mimic the swimming behavior of microorganisms; this includes self-phoretic colloids [29], particles externally actuated by magnetic, acoustic, electric fields [30–32], vibrated granular matter [33] and magnetic torque-driven helical robots [34,35].

To understand the fundamental hydrodynamics at play in swimming cells, a number of theoretical and computational models have been developed historically. The most widely studied ones include the squirmer model [36,37], Taylor’s infinitely-long planar waving sheet [38] and waving filament [39], and the rotating helix model [40,41]. These simplified models usually prescribe the actuation kinematics of the microswimmers (so-called swimming gait) by assuming that they are not influenced by the surrounding fluid environment. Understanding the impact of complex fluids in these fundamental models has attracted significant research effort in the past decade, much of which will be the focus of this review. For a discussion of the fundamentals of microswimming and biophysical modeling in a Newtonian fluid, we refer the readers to review articles [19,21,42–48] and books [49–51].

The non-Newtonian rheology of complex fluids breaks in general the time-reversal symmetry of the Newtonian Stokes equations. This in turn enables self-propulsion under reciprocal motion, as experimentally demonstrated using artificial swimmers [52,53]. More importantly, complex fluids can significantly impact the swimming kinematics and motility of individual microswimmers and their collective behavior. The bacterium *E. coli* swims faster in a polymer solution than in a Newtonian environment and it follows straighter trajectories with suppressed wobbling of the cell body [54]. The visualization of fluorescently-labeled polymers reveals that the rotation of the flagellar filaments of *E. coli* strongly stretches the polymers in the fluid and generate local elastic stresses that suppress the unbundling of flagellar filaments, thereby impacting its run-and-tumble dynamics [54]. In the case of the algal cell *C. reinhardtii*, the viscoelasticity of the fluid changes the wave envelope of the beating flagellar strokes by restricting the displacement of the flagella close to the cell body. The swimming speed of the cell is then hindered despite the flagella beating at a higher frequency in a polymer solution [55]. Similar effects are also observed for swimming spermatozoa, whose flagella beat at a higher frequency and with a smaller amplitude near the cell body in a viscoelastic cervical mucus compared to a Newtonian fluid (see Fig. 1b) [1,25,56,57]; surprisingly, the swimming speed of the cells is less affected [1]. Collectively, bovine spermatozoa show disorganized individual swimming in Newtonian fluids, but they form dynamic clusters in viscoelastic fluids (Fig. 1c) with cluster sizes and cell–cell alignments increasing with the viscoelasticity of the fluid [26]. These examples demonstrate that the impact of complex fluids on microswimming depends strongly on the type of swimming cells as well as on the interplay between the swimming gait and the rheology of the fluid environment.

Many non-Newtonian biological fluids, including biofilm and mucus, are hydrated polymeric gels made of 90% water by weight; the remaining components, which are typically conglomeration of extracellular polysaccharides, proteins and lipids, give rise to the elasticity of the fluids and to its shear-thinning viscosity. The rheological properties of biofilms show high variability; their elastic modulus $G$ is in the range $10$–$10^3$ Pa for most biofilms and can reach $10^5$ Pa for biofilms from natural hot springs [58–62]; the viscosity of biofilms is typically in the range $\mu \sim 10^2$–$10^7$ Pa·s, while their relaxation time is $\lambda \sim 10$ s for *S. epidermidis* biofilm [63] and $\sim 10^1$ s for a wide range of environmental biofilms [60]. In the case of mucus, we have $\mu \sim 10^2$–$10^7$ Pas, $G \sim 0.1$–$10^2$ Pas, and $\lambda \sim 1$–$10$ s [64,65]. Since the relevant time scale for microswimming along the size of the swimmer itself is $l_i/U_i \sim 0.1$–$1$ s and the flagellum beating period is $T \sim 10^2$–$0.1$ s, we see that the relaxation time of these complex fluids is comparable to, or much larger than, the time scales of microswimming. It is thus expected that

Fig. 1. Examples of biological microswimming in complex fluids. (a) Polymicrobic biofilm showing heterogeneous micro-scale structure (adapted with permission from Ref. [13]). (b) Superimposed images of swimming bull spermatozoa in a Newtonian fluid (top) and a viscoelastic solution of long-chain polyacrylamide (bottom) (adapted with permission from Ref. [25]). (c) Bovine spermatozoa swim in a disordered fashion in a Newtonian fluid (left) but aggregate and swim parallel to each other, while forming clusters in a viscoelastic fluid (right) (adapted with permission from Ref. [26]).
microswimmers should be strongly influenced by elastic stresses in a complex fluid and thus be subject to the full range of non-Newtonian effects, including stress relaxation, strain retardation, and normal stress differences. A review of these classical topics can be found in textbooks on non-Newtonian fluids, including the different constitutive equations models that have been used by the community to model swimming in polymeric fluids (second-order fluids, Oldroyd-B, Giesekus, FENE models) [66,67].

While many studies model complex fluids as continuum media, it is important to realize that most biological examples are highly heterogeneous at the microscopic scale [68]. Many microorganisms experience a biological environment of entangled fibers of around 10–100 nm in diameter and pores ranging in size from 0.1 to 1 μm [69]. For example, an aerosol biofilm is a porous structure with polymer conglomations connected by voids and channels of about 10–100 μm in diameter that enhance the transport of oxygen and nutrients through the colony [70]. Such micro-scale heterogeneity impacts the motility of microswimmers.

In dermis and gelatin matrices, the Lyman disease spirochete *B. burgdorferi* may temporally adhere to the matrix and undergo wriggling and lunging movements that are not observed in a liquid medium [71]. In a semi-dilute polymer solution, the bacterium *E. coli* swims faster than in a Newtonian fluid because its rotating flagella strongly stretches the polymer molecules in its vicinity and thus carves out a low-viscosity tunnel free of polymers around the flagellum. The flagellum motor increases its rotation rate, which leads to an increase of the cell speed [72–74]. Microorganisms may also explicitly change the local environment to assist their motility in complex biological fluids. For example, in the acidic environment of the stomach, the pathogenic bacterium *H. pylori* increases the local pH by secreting an enzyme that transforms the viscoelastic mucin gel into a viscous liquid thereby facilitating the penetration of the cell through the epithelial surface of the stomach [9,75]. In cervical mucus, “following” spermatozoon swim slower than “vanguard” ones despite having the same flagellar beating frequencies and shapes. Such a difference in propulsive efficiency is due to the alteration of the local mucus properties resulting from the sustained permeation by spermatozoa [3]. To better understand the impact of micro-scale heterogeneity, recent studies have thus proposed to use coarse-grained polymer models [73,74].

Beyond microorganism swimming, the motion of small-scale bodies in complex fluids is encountered in other related settings. For example, in microrheology, the local mechanical properties of a soft, complex material can be extracted from the thermal or forced motion of colloidal probes [76–78]. Another example is that of the electrophoresis of charged colloidal particles or macromolecules in gel matrices, which is a common biochemical technique to separate biomolecules based on their size and charge [79], and for which some of the models developed to address microswimming are applicable.

In this article, we review recent theoretical, computational and experimental progress on biological and synthetic microswimming in complex fluids. Our work follows naturally from previous reviews devoted to active colloids in complex fluids [80] and to the role of complex fluids in biological systems [81,82]. Our paper, written in celebration of the legacy of J. G. Oldroyd, focuses on the impact of fluid viscoelasticity on microswimming, and includes both individual motion as well as hydrodynamic interactions in viscoelastic fluids (typically less covered in previous reviews). We discuss as well the influences of non-continuum effects and thermal noise, both of which may be important at the micron scale. Polymer entanglement and the role of shear-thinning viscosity are also briefly mentioned.

This review is organized as follows. We first summarize the key dimensionless parameters and the relevant theoretical continuum framework in Section 2. We next discuss in Section 3 the impact of viscoelasticity on the motility of individual microswimmers. We then focus in Section 4 on the role of hydrodynamic interactions in complex fluids: swimmer/surface, swimmer/flow and swimmer/swimmer. Finally we conclude in Section 5 with our perspective on the direction of research in this area.

Fig. 2. The space of dimensionless parameters relevant to microswimming in viscoelastic fluids: Reynolds number *Re*, Deborah number *De*, Weissenberg number *Wi*, elasticity number *El* and Knudsen number *Kn*.

### 2. Theoretical framework

#### 2.1. Parameters and dimensionless numbers

We start by introducing the classical dimensionless numbers governing fluid dynamics in a viscoelastic (polymeric) fluid and the governing equations relevant for microswimming. We denote by *λ* the polymer relaxation time, while *c* and *l* are the characteristic time, shear rate and length scales for the flow around the microswimmer. The mass density of the fluid is *ρ*, and *ρ* = *ρ* + *ρ* is the total fluid viscosity, while *ρ* and *ρ* are the viscosity of the solvent and polymer, respectively.

Beyond the Reynolds number *Re*, the key dimensionless parameters for the fluid flow are the Deborah number *De*, Weissenberg number *Wi*, elasticity number *El*, the dimensionless polymer and solvent viscosities *μ* and *μ* = *μ* + *μ*, the suspension Knudsen number *Kn*, and the translational and rotational Péclet number, *Pe*, respectively. They are classically defined as

\[
De = \frac{\lambda}{t_c}, \quad (1a)
\]

\[
Wi = \frac{\lambda}{\gamma_c}, \quad (1b)
\]

\[
El = \frac{\lambda_{\mu}}{Re} = \frac{\lambda_{\mu}}{\rho \gamma_c^2}, \quad (1c)
\]

\[
\mu_p = \frac{\mu_p}{\mu}, \quad \mu_s = 1 - \mu_p, \quad (1d)
\]

\[
Kn = \frac{l_p}{l_c}, \quad (1e)
\]

\[
Pe_t = \frac{l_p U_s}{D_s}, \quad Pe_r = \frac{U_s}{D_s l_c}. \quad (1f)
\]

The Deborah number *De* represents the ratio between the polymer relaxation time and observation time of the fluid flow. For a swimming microorganism, the flow time scale is typically defined as either *t* = 1/ω or *t* = 1/ω, where *U* is the swimming speed and *ω* is the frequency for a periodic motion of the swimmer, for example the waving motion of the flagella.

The Weissenberg number *Wi* is the product of the polymer relaxation time *λ* with the characteristic flow shear rate *γ*. It also measures the ratio between the elastic force (first normal stress difference in a steady simple shear flow) ∼ *λμ*γ² [83] and the viscous force ∼ *ργ*γ. In some flows, e.g. steady squeezing motion, we have *Wi* ∼ *De*, while in general these two dimensionless numbers can be quite different. For example, the unsteady viscoelastic flow induced by an oscillatory shear on a squirmer surface can have different magnitudes for *De* and *Wi*.
In the classical oscillatory shear flow of viscoelastic fluids, the Pipkin diagram classifies various regimes as 'viscometric' for $De \ll 1$, 'linear viscoelastic' for $Wi \ll 1$, and 'nonlinear viscoelastic' for non-small $De$ and $Wi$ [84].

The Weissenberg and Deborah numbers are often related to one another via a length scale relevant to the microswimmer. For example, in the case of a self-propelled photoretic particle, the shear rate inside the thin interaction layer is much stronger than that in the flow in the bulk region and we have $Wi = De/\delta$ with $\delta \ll 1$ being the dimensionless boundary layer thickness [55]. In the case of a waving flagellum, the ratio between the Deborah and Weissenberg numbers depends on the dimensionless beating amplitude, $Wi = Ae/\delta$, with $A = \lambda^* l_c$, where $\lambda^*$ is the dimensional amplitude of the wave. However, even at small-amplitude, the nonlinear impact of viscoelastic stress is important [86].

The third dimensionless number, termed the elasticity number and defined as $El = Wi/Re$, is independent of the swimming kinematics and is only a function of the fluid properties and the size of the swimmer. It corresponds to the ratio of the polymer relaxation time $\lambda$, to the time scale for momentum diffusion, $\sqrt{L^2/(\mu/\rho)}$. For typical microswimmers in biofilms and mucus, we have $El \gg 1$.

The dimensionless polymer viscosity, $\bar{\mu}_p$, characterizes the ratio of the polymeric viscosity to the solution viscosity $\mu$ (which includes both polymer and solvent), while $\bar{\mu}_s$ is the solvent viscosity normalized by the solution viscosity. Consequently, $\bar{\mu}_p + \bar{\mu}_s = 1$. For many classical constitutive models, such as Oldroyd-B, Giesekus and the FENE family of models, $\bar{\mu}_p \approx 1/\lambda$ is the ratio between the retardation time and the relaxation time of the polymer. Note that the viscosity in an Oldroyd-B model is constant in a shear flow, whereas it displays a shear-thinning behavior in a Giesekus fluid or FENE-P fluid.

In a dilute polymer solution, i.e. one for which the polymer concentration $c$ is lower than the overlap concentration, $c < c^*$, the polymer viscosity is the product of the concentration with the intrinsic viscosity of the polymer, $\bar{\mu}_p \sim c [\eta]$ with $[\eta] \sim 1/\epsilon^* \sim M^{1+\epsilon}$ follow a power-law dependence on the polymer molecular weight $M$; $\nu = 3/5$ and $1/2$ are the scaling exponents of polymer solutions of good and athermal solvents, respectively. In contrast, for semi-dilute or entangled polymer solutions, the polymer viscosity scales as $\bar{\mu}_p \sim M^\alpha$, and it grows quickly with increasing polymer molecular weight since $\alpha = 3.4-4$ [87]. For relevant biological fluids, the dimensionless polymer viscosity $\bar{\mu}_p$ can span a wide range of values, from 0.01 to around 1.

Next, the dimensionless suspension Knudsen number $Kn_s$ quantifies non-continuum aspects and measures the ratio between the size of the dispersed phase $l_d$ and the characteristic swimming flow length scale $l_p$. It is sometimes called the gradient number to reflect the fact that the two ends of a long bead–spring dumbbell experience non-local flow [73]. In a viscoelastic polymer solution, the relevant value $l_g$ is the radius of gyration of the polymer, while in a solution of entangled polymers, $l_p$ is the typical mesh size of the network. In a suspension of rigid colloids, $l_g$ is the typical size of the colloids. When $Kn_s \ll 1$, the complex fluid can be modeled as a continuum around the microswimmer, otherwise it is not a continuum medium and the details of its microstructure need to be taken into account. See Fig. 2 for the space of dimensionless parameters relevant to microswimming in viscoelastic fluids.

Finally, the Péclet numbers, $Pe_i$ and $Pe_{\psi}$, characterize the ratio between the swimming motion and the translational $(Pe_i)$ or rotational Brownian motion $(Pe_{\psi})$ of the swimmer due to the thermal fluctuations or swimmer’s own random reorientation (such as the tumbling motion of flagellated bacteria, e.g. E. coli), where in Eq. (1f)$D_i$ and $D_{\psi}$ are the translational and rotational diffusivities of the swimmer. These two parameters, which are used in mean-field theories, play an important role in the collective motion of microswimmers [88,89]. Depending on the constitutive models, other dimensionless parameters also affect microswimming in viscoelastic fluids. This includes the mobility factor $a_{\psi}$ in the Giesekus model, which measures the anisotropic hydrodynamic drag on the polymer molecules, and the maximum dimensionless length of the polymers in the FENE-type models.

2.2. Governing equations

The motion of a microswimmer in a non-Newtonian fluid is controlled by the interplay between the low-Re hydrodynamics governed by the Cauchy momentum equation, the swimming kinematics or the internal actuation mechanism of the swimmer (which provides the boundary conditions for the fluid), and the fluid rheological characteristics (i.e. the constitutive equation for the complex fluid). Classical governing equations for continuum viscoelastic fluids can be written as [90,91]

$$\nabla \cdot \tau = \nabla \cdot \mathbf{r} \quad (2a)$$

and

$$\mu = \eta (1 + A) \tau + M(\tau, \mathbf{u}) = \mu (1 + B) \tau + \mathcal{N}(\tau, \mathbf{u}) \quad (2b)$$

where $\mathbf{r}$ is the pressure, $\tau$ is the deviatoric part of the stress tensor, which includes both Newtonian and non-Newtonian contributions to the stress. The Cauchy momentum in the absence of inertia, Eq. (2a), states therefore simply that the divergence of the total stress tensor, $\sigma = -p \mathbf{1} + \tau$, is zero. In Eq. (2b), $\mu$ is the total zero-shear rate viscosity of the fluid, $\mathcal{N}$ is the shear rate tensor, while $A$ and $B$ are two linear differential operators in time that capture the polymer relaxation and retardation. In contrast, the symmetric nonlinear operators $\mathcal{M}$ and $\mathcal{N}$ represent convection, stretching and nonlinear relaxation; they include in particular all terms arising from the objective derivatives of the deviatoric stress and of the shear rate tensor. The constitutive relationship stated in Eq. (2b) describes all the classical differential constitutive relationship for polymeric fluids, including all Oldroyd models. The equations above are then to be solved with kinematic and/or stress boundary conditions provided by the swimmer; in parallel, the dynamics of a microswimmer is subject to instantaneous force-free and torque-free conditions, allowing to determine in a quasi-steady fashion its linear and angular swimming velocities.

2.3. Reciprocal theorem

In cases where the swimmer velocity is of interest and where the swimmer moves by imposing velocity boundary conditions, one can use the Lorentz' reciprocal theorem (a form of the principle of virtual work [92,93]) to determine the velocity in some asymptotic limits. Rather than directly solving for the flow field, the reciprocal theorem uses a dual problem for the swimmer with the same instantaneous shape undergoing rigid-body motion. Such theoretical analysis using the reciprocal theorem typically requires an asymptotic expansion in a small parameter [86,90,94], although it can be shown to also work for a strictly tangential squirming motion in a linear viscoelastic fluid [91]. In these cases, Eq. (2b) takes a simpler form and the deviatoric stress can always be written as

$$\tau = \mu \gamma + c \mathcal{S}(\mathbf{u}) \quad (3)$$

where $c$ is a small dimensionless parameter and the symmetric tensor $\mathcal{S}$ is an explicit nonlinear function of the velocity field $\mathbf{u}$. The small parameter $c$ in Eq. (3) could be the Deborah number $De$ in a weakly nonlinear polymeric fluid, the viscosity ratio $\bar{\mu}_p$ in a dilute polymer solution, or the small amplitude $A$ of the swimming kinematics. For a generalized linear viscoelastic fluid, the above constitutive relationship is expressed in Fourier domain and $\mu$ is the frequency-dependent complex viscosity [91]. Note that the results obtained with the generalized linear viscoelastic model should be used with caution since that model is not objective (i.e. frame invariant) [66].

One can then apply the reciprocal theorem at each order in $c$ in order to determine the velocity of the swimmer asymptotically. We assume that the swimmer imposes a known swimming gait velocity $\mathbf{u}_{\psi}$ on its surface $S$, which is taken to be either the fixed reference surface for the small-amplitude perturbations $\mathbf{u}_{\psi}$ is the leading order
3. Motility of individual swimmers in viscoelastic fluids

In this section, we discuss the dynamics of a single swimmer in an unbounded viscoelastic fluid. The two most fundamental questions are: (i) for Newtonian non-swimmers, can viscoelasticity lead to locomotion? and (ii) for Newtonian swimmers, does the viscoelasticity of the fluid enhance or impede microswimming? As we see below, while the answer to (i) is broadly yes due to the generation of nonlinear stresses, the answer to (ii) is not obvious but depends on various factors, including the geometry of the swimmer, its swimming gait, the dimensionless parameters of the flow, and the values of $De$ and $Wi$.

3.1. Reciprocal swimmer

In a Newtonian fluid, microswimming is constrained by Purcell’s scallop theorem [27], stating that reciprocal deformation of a deforming body (i.e one for which the sequence of shapes is identical under a time-reversal symmetry) leads to zero net propulsion on average due to the linearity and time reversibility of the Stokes equations. Indeed, as seen in Eq. (4), time only enters the problem as a parameter in the boundary conditions so that the locomotion of any swimmer in a Newtonian fluid is instantaneously and linearly determined by the velocity boundary condition $u_{s1}$. A consequence is that the total distance traveled by a swimmer over a period is independent of the rate of actuation of the swimmer body but depends only on the sequence of body shapes. In contrast, complex fluids display nonlinear rheological properties (including shear-dependent viscosities and normal-stress differences) and therefore the stresses exerted on any swimmer are expected to break the constraints of the scallop theorem and to enable net propulsion in general.

Two types of reciprocal swimming in complex fluids have been demonstrated. The first type consists of swimming enabled by normal-stress differences arising in shear flows. For a tethered semi-infinite rigid plate undergoing small-amplitude sinusoidal motion with flapping amplitude $A$ (Fig. 3a), an asymptotic analysis showed that the non-Newtonian stresses generate a net flow under time-periodic actuation, and the flapping motion leads to non-Newtonian propulsive forces exerted on the plate at $O(A^2)$ [99]. In contrast, a net pumping flow is induced at $O(A^4)$ [96], with a strength that has a non-monotonic dependence on the Deborah number [96].

Experimentally, the viscoelastic propulsion of reciprocal flapping swimmers was demonstrated using a magnetically polarized dumbbell-swimmer in an oscillating magnetic field, with net motion occurring for swimmers with asymmetric shapes or for symmetric swimmers near a wall in viscoelastic fluids, with no motion in a Newtonian fluid [52].

A related example of a swimmer driven by normal stress differences in viscoelastic fluids is the dumbbell (‘snowman’) swimmer rotating around its symmetry axis [35,97]. The rotation of the spheres in the dumbbell induces elastic hoop stresses along the curved streamlines in the azimuthal direction, leading to an inward contraction in the radial direction and thus secondary streaming flows which push the two spheres away from each other. The strengths of these flows are unbalanced if the two spheres have different sizes and hence for an asymmetric dumbbell they lead to a net motion (towards the side of the small sphere, see Fig. 3b). A similar mechanism leads to the propulsion of an axisymmetric swimmer performing a harmonic torsional oscillation in a viscoelastic fluid [100]. Due to elasticity-induced streaming flows, a reciprocal swimmer composed of two spheres of unequal sizes which oscillate periodically along its axis of symmetry can self-propel in a viscoelastic fluid [101]. For both in-phase and anti-phase oscillations, the net motion is always in the direction of the smaller sphere. A swimmer composed of two spheres that can deform elastically is seen to swim in the same direction in a Newtonian fluid [101]. Recently, the motion of a torque-free swimmer composed of two spheres rotating around their symmetry axis in opposite directions was studied [102]. The swimmer is seen to always translate towards the larger sphere (the one with the slower rotation), presumably because the fast rotation of the smaller sphere creates a stronger streaming flow directed away from the larger sphere. Other shapes of revolution and arrangements of spheres were also studied, leading to translation possible in both directions.
A second type of reciprocal microswimming in viscoelastic fluids does not exploit steady-state normal stresses but instead relies on the rate-dependence of the flows and forces generated in the fluid. Indeed, unlike the Newtonian limit, the instantaneous swimming speed in a viscoelastic fluid is expected to depend nonlinearly on the rate at which the swimmers deforms its body, and therefore in general an asymmetry in a periodic actuation should result in an asymmetry in swimming kinematics. This strategy was proposed in Ref. [98] using the example of a helically waving filament propagating waves at different speeds in forward and backward strokes (specifically the wave moved forward by one wavelength in the first third of the time period and then backward by one wavelength in the final two thirds of the period, see Fig. 3c). In both periods, the helical filament moved in the direction opposite to the wave propagation but with a smaller total displacement in the first (forward) cycle due to a larger value of $D_e$. Since no locomotion occurs in the Newtonian limit $D_e = 0$ and propulsion is expected to be poor in the high-$D_e$ limit, net swimming is maximal for $D_e \sim O(1)$ [98]. Note that in this case, the velocity of the flagellum is not simply obtained as the sum of the velocities of each stroke because the time-history of the polymer plays an important role.

### 3.2. Squirmer model

A popular model in the theoretical treatment of swimming at low Reynolds number is the squirmer model [36,37]. A squirmer takes the form of a spherical or spheroidal cell that acts on the surrounding fluid with a tangential slip velocity prescribed along its surface. The squirmer model has been widely used to study the motion of ciliates such as the green algae Volvox and the protozoan Paramecium driven by the synchronized metachronal beating of short cilia covering the cell body. A squirmer is also the appropriate mathematical model to describe the motion of a self-propelled droplet driven by marangoni stresses [103,104] and autophoretic colloidal particles driven by electrophoresis, diffusophoresis or thermophoresis, be they externally driven or self-generated [29,105,106].

In the squirmer model, the tangential velocity is typically taken to be axisymmetric (and therefore with the same symmetry axis as the cell body) and is given by an expansion in Legendre polynomials (and associated Legendre polynomials) with either steady or time-periodic coefficients. In a Newtonian fluid, the flow field around the squirmer is then computed using spherical and spheroidal harmonics [36,37,107]. In most applications only the first few modes of the squirmer motion, which decay spatially the slowest, are considered; the influence of higher-order modes on swimming in a viscoelastic fluid was considered in [94].

In the case of a spherical squirmer, the slip velocity is often written as the sum of two modes, $u_s' = B_1 \sin \theta + B_2 \sin 2\theta/2$; the first mode determines the swimming speed ($U_0 = 2B_1/3$ in an unbounded Newtonian fluid) and represents a source dipole singularity decaying as $u \sim r^{-3}$ in the far-field; the second mode $B_2$ corresponds to a Stokeslet dipole (or stresslet) decaying as $u \sim r^{-2}$ and a source quadrupole decaying as $u \sim r^{-4}$. When $\beta \equiv B_2/B_1 > 0$, the squirmer is a puller with propulsion generated in front of the body, as in cells of the green alga genus Chlamydomonas; in contrast, when $\beta < 0$ the squirmer is a pusher with the propulsion generated from its rear side, as in the flagellated bacterium E. coli. The particular case $\beta = 0$ corresponds to a neutral squirmer. Note that squirmers with time-periodic modes have also been studied [108], as have squirmers with swirls and azimuthal components to their boundary conditions [109,110].

In the case $D_e \ll 1$ or $\tilde{\nu}_p \ll 1$, the motion of a squirmer in an unbounded viscoelastic fluid can be tackled using asymptotic expansions and the reciprocal theorem. In both cases, the leading-order solution is the squirmer motion in a Newtonian fluid, which induces the polymer deformation. The difference is that for $D_e \ll 1$, components of the polymer stress tensor are decoupled and are determined solely by the local strain rate, while in the limit $\tilde{\nu}_p \ll 1$, the components are coupled to one another. De Corato et al. [111] considered spherical squirmers in a second-order fluid in the limit $D_e \ll 1$ and obtained their swimming velocity and power as

$$\frac{U}{U_0} = 1 - \frac{3}{10} \left(1 + \frac{2\nu_s}{\nu_f}\right) \beta D_e,$$

$$\frac{P}{P_0} = 1 - \frac{9}{10} \left(1 + \frac{2\nu_s}{\nu_f}\right) \left(\frac{3 + \beta^2}{2 + \beta^2}\right) D_e,$$

where $U_0 = 2B_1/3$ and $P_0 = 6\alpha(2 + \beta^2)$ are the velocity and power in a Newtonian fluid of the same total viscosity $\mu = \mu_s + \mu_p$ as the viscoelastic fluid, $\nu_s$ and $\nu_f$ are the first and second normal stress difference coefficients for the fluid ($\nu_s > 0$ and $-\nu_f/2 \leq \nu_s \leq 0$ for a realistic polymeric fluid), and the Deborah number is defined as $D_e = \lambda U_0/a$ with $a$ the radius of the squirmer and $\lambda = \nu_f/(2\mu_p)$.

This second-order fluid model is valid for a slow and slowly varying flow of a polymeric fluid and most nonlinear constitutive equations reduce to the second-order fluid in the limit $D_e \ll 1$. We have in particular $\nu_s = 2\lambda \mu_p$, $\nu_f = 0$ for the Oldroyd-B and FENE models, and $\nu_s = 2\lambda \mu_p$, $\nu_f = -3\lambda \mu_p a_n$ for the Giesekus model (here $a_n$ is the mobility factor). The result in Eq. (6) shows that pusher cells ($\beta < 0$) swim faster but consume more power in a weakly viscoelastic fluid than in a Newtonian fluid; the opposite is true for pullers ($\beta > 0$), that swim slower but expend less power to do so. At leading-order in $D_e$, neutral squirmers are not affected by the viscoelastic fluid. Numerical simulation with a Giesekus model show that the above asymptotic solution remains valid as long as $D_e < 0.02$ and the squirmer speed is always reduced for $D_e \geq 0.1$ [111]. Expansion at higher orders in $D_e$ shows that the neutral squirmer actually swims slower in a Giesekus fluid [117].

Similar conclusions appear to apply qualitatively to autophoretic Janus particle [118,119]. In that case, the type of swimmer (pusher vs. puller) is determined by the sign of the local phoretic mobility and the ratio between active and inert regions. By expanding in the viscosity ratio $\tilde{\nu}_p$, Li and Koch [85] considered the motion of an electrophoretic particle in a dilute polymer solution at finite $D_e$. While
in a Newtonian fluid, a uniformly-charged particle with a thin ionic Debye layer disturbs the fluid as a neutral squirmer, in a polymer solution the particle becomes a puller at small $D_e$. The particle velocity is reduced for all $D_e$, with a minimum occurring at $D_e = O(1)$. At high $D_e$, the polymer is strongly stretched by the local extensional flow at the rear side of the particle in both Debye layer and bulk region forming a birefringent strand. These results indicate that the swimming kinematics of a swimmer is strongly influenced by the polymer.

At high Deborah numbers, numerical simulations are needed to capture the full impact of viscoelasticity for large deformations. Zhu et al. [112,120] studied the motion of a squirmer with both spherical and prolate spheroidal shapes in a polymeric fluid using finite elements. For all types of squirmers, the viscoelasticity in the fluid always reduces both the swimming speed and the power consumption. The squirmer reaches a minimum velocity at $D_e = O(1)$, at which point the swimming efficiency $\eta \sim U^2/P$ peaks. As expected, the polymers are strongly stretched in regions of local extensional flows close to the squirmer surface, at the front and the rear in the case of pushers and on the sides in the case of pullers (Fig. 4a).

### 3.3. Waving slender bodies and flexible flagella

For undulatory swimmers, both enhancement and decrease of locomotion speeds have been observed in viscoelastic fluids depending on the details of the swimming gaits and the fluid rheology. The nematode *C. elegans* always swims slower in a semi-dilute solution of high-molecular-weight polymer (carboxymethyl cellulose) than in a Newtonian fluid of the same viscosity [121]. In contrast, in a polymer solution of xanthan gum, *C. elegans* swims slower in a semi-dilute solution but faster in a concentrated solution of entangled polymer network [122]. Theoretically, in what was historically the first paper in the field of low-Reynolds number swimming, Taylor proposed to use a two-dimensional (2D) waving sheet as a model for the propagation of sinusoidal wave on flexible spermatozoa flagella [38] and later extended it to the case of three-dimensional (3D) filaments [39]. The sheet model also allows to capture the waving motion of the envelope of cilia tips for large ciliated organisms [50,51]. Both the sheet and filament models predict, at small amplitude, the same swimming speed in a Newtonian fluid, $U_0 = \omega k A^2/2$, where $\omega$, $k$ and $A$ are the frequency, wavenumber and the amplitude of the waving motion (the small amplitude limit corresponds to the assumption $Ak \ll 1$).

The results for both 2D and 3D undulatory swimmers in a viscoelastic fluid are in broad agreement. Early studies showed that the speed of a small-amplitude waving slender swimmer is unchanged in a linear viscoelastic fluid [123] and in a second-order fluid at $Re = 0$ [124]; adding the inertia from the fluid increases the speed if $Re$ is below a critical value, while the speed decreases at large $Re$ [124]. However as pointed out in several studies [86,98,125], geometrically nonlinear viscoelastic effects always need to be included even for a small-amplitude swimmer: a constitutive relationship needs to be objective, and the transport of viscoelastic stresses naturally leads to geometrically nonlinear terms in the constitutive modeling [66].

In that case, Sturges [125] showed that the time derivative of the Rivlin–Ericksen tensor at each degree is of the same order as the swimming amplitude, therefore the polymeric stress cannot be simplified using a linear model. Based on an integral constitutive equation, the analysis shows that, at a fixed $E1$, the speed of a waving sheet in a nonlinear viscoelastic fluid decreases monotonically with increasing $Re$. For a fixed value of $Re$, their result shows that the speed decreases monotonically with $De$, consistent with the result in Eq. (7) for a flagellum in the absence of inertia.
In a classical Oldroyd-B polymeric fluid, the swimming speed of both the small-amplitude waving sheet and waving filament models is \[ \frac{U}{U_0} = \frac{1 + \hat{\mu}P}{1 + \hat{D}e^{-c}}, \tag{7} \]
where \( \hat{D} = \lambda \omega \) is defined using the undulatory frequency \( \omega \) of the sheet and where \( U_0 \) is the swimming speed in a Newtonian fluid. Importantly, this result arises from the (geometrically) nonlinear response of the viscoelastic fluid to the small amplitude waving motion since in a linear viscoelastic fluid the speed of the flagellum is unchanged [123]. The power \( P \) and the swimming efficiency \( \sim U^2/P \) follow the same dependence on \( \hat{D} \) as in Eq. (7) if the Newtonian fluid chosen for comparison is taken to have the zero-shear rate viscosity of the non-Newtonian fluid.

The result in Eq. (7) turns out to be valid for waves of either tangential and normal motion [86], for filaments undergoing small-amplitude helical waves [127], and for pumping using small-amplitude peristaltic waves (for which the pumping rate is analogous to the swimming speed) [128,129]. A squirmer driven by small-amplitude oscillating slip velocities also shows a similar speed dependence on \( \hat{D} \) in a viscoelastic fluid [90,91]. Interestingly, the speed ratio in Eq. (7) is exactly the same as the dimensionless dynamic viscosity \( \mu'/\omega/\mu \) (i.e. the real part of the complex viscosity) in a small-amplitude oscillatory shear (SAOS) flow [66,67]. The result in Eq. (7) was also found to be independent of the particular constitutive model chosen for the polymeric fluid as along as they have the same complex viscosity as those of Oldroyd-B, Johnson–Segalman, Giesekus and FENE-P [86].

Extensions of the result in Eq. (7) to higher orders in the waving amplitude shows that the swimming speed of a large-amplitude waving sheet also monotonically decreases with \( \hat{D} \) as in Eq. (7). For a waving flagellum when the feedback from the fluid stresses on the flagellum, the waving amplitude is seen to increase in a viscoelastic fluid, so much so that it may overcome the viscoelastic hindrance from Eq. (7) and gain an overall speed enhancement.

which agrees with the Eq. (7) in the limit \( Sp < 1 \) and becomes \( U/U_0 = [1 + \hat{\mu}P]/[1 + \hat{D}e^{-c}] > 1 \) for \( Sp \gg 1 \). For a stiff swimmer with \( Sp \ll 1 \), the active stress scaling as \( \sim f k^2 \) (\( f \) is the imposed active bending moment per unit length) is balanced by the bending resistance of the body (\( \sim kA^2 \)), so the waving amplitude scales as \( A \sim f/k^2 \) and it is not affected by the fluid. In that limit, the velocity decreases in a viscoelastic fluid as described in Eq. (7). For \( De \gg 1 \), the speed is \( \hat{\mu} \), the speed in a Newtonian fluid (\( \hat{\mu} \), is the dimensionless solvent viscosity). In contrast, for a soft swimmer with \( Sp \gg 1 \), the active stress is balanced by the viscous resistance from the fluid. In a Newtonian fluid, that fluid resistance scales as \( \sim \mu U_0/A \sim \mu \alpha kA \), where we have used Taylor’s result for swimming speed \( U_0 = A^2/\alpha k \). In a viscoelastic fluid with \( De \gg 1 \), the fluid resistance \( \mu U_0/A \) becomes \( \mu \alpha kA \), due to \( U_0 \sim \hat{\mu}U_0 \) (Eq. (7)) and \( U_0 \sim A^2/\alpha k \omega \) where \( \alpha \) is the amplitude in a viscoelastic fluid, \( \mu \) and \( \mu_1 \) are the total and solvent viscosity, respectively. The two fluid resistances are balanced by the same active stress \( \sim f k^2 \), therefore \( A \sim A/\hat{\mu} \) and \( U_0 \sim U_0/\hat{\mu} \). For a soft flagellum, the waving amplitude is seen to increase in a viscoelastic fluid, so much so that it may overcome the viscoelastic hindrance from Eq. (7) and gain an overall speed enhancement.

For a swimmer of finite length, the waveform may no longer be a simple sinusoidal wave. In addition to the effects above, the analysis of a finite filament undergoing small-amplitude waving showed that the interaction between an elastic body and a viscoelastic fluid can modify the swimming speed by qualitatively affecting the beating pattern of the filament [113,126]. An increase in \( De \) tends to reduce the swimming speed monotonically for a filament of large wavelength; for a filament of medium wavelength the speed first slightly increases and then it decreases; finally an increase in \( De \) can reverse the swimming direction of a filament with small wavelength by dramatically changing its beating pattern.

Physically, the elasticity of the fluid increases the bending length of the filament and tends to suppress its undulations at its proximal end (i.e. the junction to the cell body) and in the middle portion (Fig. 4b). The results in Ref. [113] qualitatively reproduce the beating patterns of spermatozoa flagella observed in Newtonian and viscoelastic fluids [1,25,56]. A similar effect can be observed for the flagella on the algae C. reinhardtii whose lateral displacements near the cell body decrease in a viscoelastic fluid [55].

Two-dimensional numerical simulation demonstrated that a waving swimmer of finite length and large amplitude can swim faster in a viscoelastic fluid [134]. It has a maximum speed at \( De \sim 1 \) and generates strong polymer extension behind it. This result turns out to be very different from the polymer stress distribution around an infinitely long flagellum, for which the polymer is periodically stretched on the windward side of the traveling wave and relaxes on the leeward side (Fig. 4c). Thomases and Guy [115] further showed that, beyond the fluid, the swimming speed in a viscoelastic fluid depends critically on the elasticity of the body and the swimming stroke. Specifically, a swimmer with a decreasing amplitude from head to tail (e.g., C. elegans) slows down in a viscoelastic fluid, and so does a stiff swimmer whose waving amplitude increases (e.g. flagellated spermatozoa). In contrast, if that swimmer is softer, the swimming speed can go up due to the combined effects of the increase in the stroke amplitude [135] and the favorable asymmetric distribution of the polymer stress around the swimmer (Fig. 4d). This result, which is consistent with theoretical results for an infinitely-long elastic swimming sheet internally driven by active stresses [133], can help explain the experiments in Refs. [121,122]. In a concentrated polymer solution, the viscosity of the fluid substantially increases, which leads to an increase in the Sperm number from Eq. (8), or equivalently, a reduction of the effective stiffness of the swimmer. In three dimensions, the polymer stresses become more concentrated near the end of the swimmer and lead to a weaker speed reduction for a model C. elegans with prescribed kinematics (i.e. in the infinitely stiff limit) compared to the 2D case [136].
Experimentally, an increase in the swimming speed was demonstrated in a viscoelastic fluid for a flexible swimmer composed of a magnetic head and a soft tail actuated in a time-periodic magnetic field [137]. Within the range of parameters tested in the experiments ($De \leq 5$), the ratio of swimming speed for swimmers in Boger fluids (i.e. viscoelastic fluids that have a constant viscosity) and Newtonian fluids of the same viscosity was seen to continuously increase with the Deborah number. Related work showed that the speed of a waving cylindrical sheet increases in a Boger fluid, while it decreases in a shear-thinning viscoelastic fluid [138], showing that the shear-dependence of the viscosity can also have a strong impact on microswimming.

### 3.4. Helical locomotion

Microswimming driven by rotating slender helical shapes is the locomotion method used by bacteria with slender flagellar filaments and by sprochetes. Early experiments showed that many bacteria cannot only swim in polymer solutions [139–143] but that in fact they swim faster than in the solvent alone, despite the fact that the polymer solution is much more viscous [140]. For a wide range of bacteria species, the average speed varies non-monotonically with the solution viscosity and reaches a peak value at a critical viscosity [139,141,142].

Berg and Turner [143] showed that the angular velocity of the body of a tethered *E. coli* cell is inversely proportional to the fluid viscosity in a branched polymer solution, consistent with Newtonian dynamics; in contrast, it is less affected in a solution of unbranched polymer chains, suggesting strong non-Newtonian effects in that case. Berg and Turner suggested that in a solution of unbranched polymers, bacteria swim in a loose quasi-rigid network of microsized pores and experience as a result two different resistances when moving through the network (low friction inside the pores and high friction against the network), a strong anisotropy at the origin of the unusual swimming behavior. In comparison, a solution of branched polymers is more homogeneous and has a behavior closer to that of a Newtonian fluid. Based on this physical idea, Magariyama and Kudo [144] modified the classical resistive-force theory empirically by introducing different apparent viscosities for motions tangent and normal to a body surface and successfully reproduced experimental results.

A recent study [72] reexamined the Berg and Turner argument and demonstrated convincingly that the rotating rate of the cell body seen in a solution of unbranched polymers originates from a viscosity contrast between that experienced by the flagellum and that near the body. Indeed, around the fast-rotating flagellum, the transition of polymers from a coiled to an elongated state creates locally a low viscosity zone and results in an enhancement of the swimming speed [72]. This study highlighted therefore that the swimming dynamics is strongly influenced by the microstructure of the non-Newtonian fluid rather than the properties at macroscales assumed in a classical continuum fluid model. We will revisit this idea in the next subsection and focus first below on studies that modeled the viscoelastic fluid as a continuum material.

For an infinitely-long filament passing a helical traveling wave of small amplitude, the swimming speed in a continuum viscoelastic fluid monotonically decreases with increasing $De$ and follows exactly the same expression (7) as a waving sheet and filament [98]. By solving the Stokes equations in a helical coordinate system, Li and Spagnolie [127] studied the locomotion of helical bodies of arbitrary cross-section using helical waves. They showed that, in a viscoelastic fluid and for helices of arbitrary cross-section, the speed of each mode of the small-amplitude helical wave is proportional to the real part of the complex viscosity, a result that remains true for swimming confined inside a cylindrical tube.

In contrast, a helical filament with a large pitch angle was shown to swim with a speed that depends non-monotonically on the Deborah number [116,145]. Experiments demonstrated that macro-scale force-free rigid helices rotating along their axes in viscous and viscoelastic Boger fluids experience a maximum speed enhancement for $De \sim O(1)$, which was confirmed by numerical simulations [116]. At the same Deborah number, the speed enhancement was found to be stronger for a helix of high pitch angle and small thickness than for low pitch angle and large radius. Using simulations to solve for the distribution of polymer stresses, the speed increase was suggested to be caused by the interaction of the rotating helical shapes with their own viscoelastic wakes, which is strongest when the time scale of the helical motion matches the polymer relaxation time (Fig. 4e) [116].

#### 3.5. Non-continuum effects

As seen in this review so far, most studies on viscoelastic microswimming have modeled the fluid as single-phase continuum medium. However, complex fluids are usually chemically and mechanically heterogeneous on the length scales relevant to biological swimmers, so non-continuum effects are therefore expected to be important.

In dilute and semi-dilute polymer solutions, microswimmers see individual and overlapping polymer coils of radius of gyration $\sim 10^{-8}$ m. The rotating flagellar filaments of bacteria induce a local shear flow which strongly stretches polymers [54,73], reducing the fluid viscosity around the flagellum. This in turn increases the cell swimming speed by increasing the flagellum rotation and decreasing the cell body rotation [72,73].

Brownian dynamics simulations for a helix rotating in a suspension of polymer dumbbells show that, at the same Deborah number and solvent and polymer viscosities, long polymers are less deformed than short polymers since they experience a weaker nonlocal shear (Fig. 5a) [73]. The magnitude of first normal stress difference exerted on the helix, and the modification of the swimmer speed, are also found to be more pronounced when the polymer is smaller than the swimmer [73].

By tracking the motion of a single polymer chain near a rotating helix, Balin et al. [148] showed that polymers are pumped along the helical flagellum, while migrating radially towards the rotating helix and being stretched, eventually depositing at the rear end of the flagellum and relaxing to their equilibrium configuration. When attached to a cell body, the rotating helix creates a depletion zone of long polymer chains leading to an apparent slip velocity between the swimmer and the fluid, which increases the swimming speed [74] (Fig. 5b). Note that apparent slip was proposed in earlier theoretical models as a mechanism to increase swimming speeds [149]. The study in Ref. [74] then predicts that the flagellum rotates slower in a polymer solution than in a Newtonian fluid, which is the opposite of the argument based on the viscosity difference for cell body and flagellum from Ref. [72], and future experiments will be necessary to measure the angular speed of the flagellum to test these hypotheses.

Beyond flagellated cells, active Brownian colloidal particles are significantly influenced by polymers in suspension. Experiments show that, by increasing the Deborah number, the rotational diffusivities of the particles increase monotonically by more than two orders of magnitude and saturates at $De \sim 0.1$, and the translational diffusivities increase by one order of magnitude [150]. Above a critical Deborah number $De > De_c \sim \dot{\gamma}$, active particles undergo a transition from enhanced diffusion to a persistent rotational motion [146] (Fig. 5c).

Unlike in a Newtonian environment, the relaxation of polymers means that the flow induced by an active particle in a viscoelastic fluid is influenced by its time history. Such a fluctuating flow field exerts in turn long-memory random stresses on the particle and interacts with the Brownian fluctuations. Depending on how the persistence time compares to the time scale of the particle motion, this can enhance the particle diffusion or even lead to its circular motion [146]. Coarse-grained simulations further showed that the rotational enhancement of active particles is affected by two other effects, namely the reduced absorption of polymers on the particle surface and a front–back
asymmetric encounter of polymers at the front of the swimmer [151].

The second effect is more pronounced when the polymer concentration is near the overlap concentration, beyond which the rotational diffusivity is reduced because of strong and homogeneous polymer absorption [151]. In the entangled polymer regime, recent work showed that active particles can escape entanglements on time scales much shorter than the polymer relaxation time [152]. The competition between caging in polymer solutions and the persistent motion of active particles leads to a non-monotonic dependence of the translational diffusivity on the particle radius, or the inverse of $D_e$, in a viscoelastic fluid [153].

3.6. Effects of polymer entanglement and shear-thinning viscosity

At high concentrations, because of crosslinking, polymer molecules might overlap and entangle with one another, thereby forming gel-like networks. Leshansky [154] showed that, in a fluid consisting of fixed microstructures, the strong anisotropy in hydrodynamic resistance enhances the speed and efficiency of microswimming with prescribed propulsion gaits. Studies on two-fluid models that treat the polymer network and solvent as two coupled elastic and viscous continuum phases showed that microswimming speed is influenced by the deformability of the network and the nature of the interactions between the swimmer and the network [155,156]. The strongest speed enhancement is observed for a swimmer in a sparse network of stiff polymers when the swimmer/polymer interaction is mediated by the solvent [155]. A similar conclusion was obtained by simulations of a discrete model of a viscoelastic network immersed in a viscous fluid [147] (Fig. 5d).

Besides elasticity, the shear-thinning behavior of the viscosity is another important property for complex fluids. The impact of shear-dependent rheology on microswimming has been studied in previous studies and we refer to them for further details [114,157–161]. Typically, wherever high shear is induced, the fluid viscosity decreases locally (Fig. 5e) and causes two effects: a local confinement of the swimmer, which enhances the swimming speed [114,161], and reduction of fluid forces which tends to hinder locomotion [158,160,162]. The power consumption is however always reduced due to the reduction in viscosity [159].

4. Hydrodynamic interactions in viscoelastic fluids

In the previous section, we discussed the motion of a single swimmer in an unbounded viscoelastic fluid, focusing on the change in the swimming speed and energetics. Beyond individual locomotion, other physical effects are impacted by viscoelasticity, including the response to external flows [95,163–166], the interactions with surfaces and interfaces [17,19,167–173], swimmer-swimmer interactions [167,174,175] and collective motion [44,176–182]. These can all significantly influence the behavior of microorganisms in nature. For example, biofilm formation is usually initiated by the accumulation and the irreversible attachment of bacteria on surfaces [183]. External flows can enhance bacteria aggregation, trigger biofilm formation in vortical regions [17] and lead to the formation of biofilm streamers in confined environments [184]. Interaction with clean and contaminated interfaces can affect the spatial distribution of bacteria [185–188]. In contrast to the extensive literature on individual swimming in a
bulk viscoelastic fluid, fewer studies have been devoted to aspects of interactions. In this section, we summarize work on the three types of relevant hydrodynamic interactions, namely swimmers with surfaces or interfaces, swimmers with external flows and between neighboring swimmers.

4.1. Interactions with surfaces

Swimming in a viscoelastic fluid near a boundary has been addressed using Taylor’s waving sheet model [167,169,170]. In the small-amplitude limit, the swimming speed is simply given by the speed obtained in an unbounded Newtonian fluid, \( U_0 = \omega k A^2/2 \), times two ratios, which are due to two separate effects: a term identical to that for near-wall swimming in a Newtonian fluid (which is always above one and thus enhances swimming) [190], and the term from Eq. (7) for swimming in an unbounded viscoelastic fluid (which is always below one and thus always hinders swimming) [130,170]. At a finite amplitude, numerical simulations show that the combination of the wall and viscoelasticity effects can lead to a non-monotonic speed dependence on \( De \) and produce a stronger speed enhancement than the wall effect alone when both the flagellum-wall distance and the normalized polymer viscosity are small (\( \beta_2 = 0.1 \)) [170]. It was conjectured that this speed enhancement is caused by the combination of a wall-induced increase of polymer stretching in regions of extensional flow between the alternating vortices near the waving sheet and the modification of the fluid velocity due to the advection and relaxation of the polymer stress. Chrisepp et al. [167] further studied numerically the hydrodynamic interactions between an actuated swimming sheet and a rigid wall or with a passive elastic membrane, observing a deterioration of the swimming performance in an Oldroyd-B fluid compared to a Newtonian one, resulting from the relatively large flagellum-surface distance and high polymer viscosity ratio (\( \beta_2 = 0.5 \)).

An important impact of viscoelasticity on near-wall microswimming is that it can strongly modify a swimmer’s trajectory and change its near-wall residence time. Of course this does not occur for an infinitely-long sheet model since the fluid in the gap is incompressible, and therefore the volume of fluid between the swimmer and the wall cannot change. In a Newtonian fluid, it has long been observed experimentally that swimming bacteria are attracted by surfaces [186,191–193]. This phenomenon results from a combination of multiple effects, including far-field hydrodynamic interactions [192], bacteria reorientation due to collisions [194] or Brownian fluctuations [193,195], and surface-induced suppression of flagellar unbinding [196].

When only hydrodynamic interactions are included, numerical simulations of squirmer models in a Newtonian fluid show that the squirmers of relatively weak dipoles (\( |\beta| \leq 2 \)) tend to swim parallel to a no-slip surface during a finite time that depends on many factors (including the incoming angle, the distance between the swimmer and the wall and the squirmer parameter \( \beta \)) before escaping from the surface, while squirmers with larger values of \( |\beta| \) are weakly attracted to the surface, following periodic bouncing trajectories [197,198]. In comparison, a pusher squirmer in a viscoelastic fluid can be permanently trapped near the surface at high Deborah numbers due to the large forces generated by polymer extension behind the swimmer [168] (Fig. 6a), with pullers less affected. Similar results are also observed for finite-length undulatory swimmers [169]. A theoretical analysis of a squirmer in a weakly viscoelastic fluid (\( De < 1 \)) shows that, depending on the initial orientation and distance of the swimmer from the wall, its near-wall residence time may increase or decrease, and the near-wall periodic trajectories turn into unstable spirals [172]. No permanent wall attraction of the swimmer was however observed in this analysis. For a squirmer actuating the fluid with a time-reversible small-amplitude slip velocity, a perturbation analysis predicts a permanent wall attraction for both pushers and pullers initially either swimming close to the wall [171] or being oriented in a specific range of angles relative to the wall [199].

Many of these results may differ when non-hydrodynamic effects are included. For a self-propelled particle influenced by Brownian noise, adding polymers into the fluid actually reduces its wall-trapping time [173] because the presence of polymers increases its rotational motion (Fig. 6b) [150,151].

4.2. Interactions with external flows

The reciprocal theorem was used to compute the motion and hydrodynamic forces of a microswimmer in an arbitrary background flow of a weakly non-Newtonian fluid [95]. The final expression resembles the results in Eqs. (4) and (5) with additional contributions from the background flow. A similar approach was applied to a spherical squirmer in a viscoelastic shear flow at low \( De \) and showed that it induced a secondary rotation of the swimmer with angular velocity of magnitude \( O(De\beta) \) [165]. At high \( De \), further numerical simulations showed that a neutral squirmer always realigns its swimming direction with the vorticity axis. Pushers and pullers have more complex dynamics; pushers align with the vorticity axis if they have a small swimming speed and relatively low \( De \), otherwise they reorient into the shear plane and tumble around the vorticity axis; pullers display the opposite behavior [165]. This feature might be exploited to separate swimming microorganisms based on their propulsion mechanisms.

The dynamics of microorganisms suspended in an external flow of viscoelastic fluids was also addressed using simplified point-swimmer models [166,189]. Assuming weak viscoelasticity, these studies describe the orientation of a discrete swimmer using (Newtonian) Jeffery’s orbits for an inert particle [200] and incorporating a viscoelasticity-induced migration into the swimmer velocity [201]. In a vortical flow, viscoelasticity leads to an aggregation of swimming microorganisms and dynamics along limit cycles, whose sizes and shapes depend on both swimmer motility and fluid viscoelasticity [189] (Fig. 6c). In a Poiseuille flow, the oscillating upstream-swimming trajectory of swimmers in Newtonian fluids turns into damped oscillatory motion approaching the centerline of the channel in a weakly viscoelastic fluid with shear-dependent viscosity [166]. Other phenomena reported for spermatozoa in Newtonian fluids, such as cross-stream migration in Poiseuille flows [163] or buckling in an extensional flow [164] may also be affected by viscoelasticity fluid.

4.3. Interactions between microswimmers

Biological organisms and spermatozoa often swim together, forming clusters and influencing one another through hydrodynamic interactions. For example, adjacent spermatozoa are observed to swim cooperatively with their flagella beating in synchrony, leading to higher waving frequency and swimming speed [174]. In a Newtonian fluid, a front–back asymmetry in the flagellar waveform is required for two infinitely-long flagella to synchronize [202]; the final synchronized state (in phase or opposite phase) is then just a function of the flagellar geometry. In contrast, in a viscoelastic fluid pairwise interaction between two symmetric flagella always leads to in-phase synchronization, and is associated with the lowest energy dissipation in the surrounding fluid [175]. Asymptotic analysis in the limit of a small dimensionless amplitude \( A \) shows that synchronization in a viscoelastic fluid occurs on a time scale \( t \sim A^{−2} \), which is much faster than the typical phase-locking time in a Newtonian fluid \( t \sim A^{−4} \) [175]. Numerical simulations further showed that the speed and power efficiency for two synchronized flagella are lower than in the case of a single flagellum, a conclusion valid for both Newtonian and viscoelastic fluids [167].

In the case of multiple microswimmers, viscoelasticity in the fluid can also impact collective behavior. Bozorgi and Underhill analyzed the linear stability of a homogeneous and isotropic distribution of microswimmers in a viscoelastic fluid using a continuum mean-field theory [176,177]. The average velocity of an infinitesimal volume of swimmer suspension is the sum of the swimming velocity for an
isolated swimmer, the velocity in the fluid and a diffusive contribution for the center of mass of the swimmer cluster, while the average angular velocity is given by the sum of Jeffery's reorientation and rotational diffusion. The presence of a viscoelastic stress in the fluid is seen to have two influences: the extra polymer viscosity $\mu^p$ stabilizes the suspension, while the linear relaxation of the polymeric stress destabilizes it. Similar to the Newtonian cases, a puller suspension is always stable, while a pusher suspension is unstable. In a Newtonian fluid, the most unstable mode occurs at zero wavenumber and is of zero wavespeed. Increasing the value of $De$ increases its growth rate and generates a new peak of maximum growth rate at a non-zero wavenumber associated with non-zero wavespeed (see Fig. 6d) [179]. This result indicates that viscoelasticity reduces the size of the coherent vortex structures induced by the collective motion of pusher swimmers, as was later confirmed in simulations [178,179].

Numerical simulations of rod-like pusher swimmers showed that viscoelasticity enhances aggregation by generating strong polymer stresses inside the gap between two swimmers parallel to each other (Fig. 6e) [179]. A similar effect is observed for bovine sperm cells in viscoelastic fluids [26] (Fig. 1c). Pullers are less affected by fluid viscoelasticity, and instead the swimmers stick together at their ends like an asterisk.

For an active nematic suspension [180,181], adding polymers enhances the stability of a system where polymer and nematic fields are coupled only by fluid velocity, mainly due to the increase of the viscosity. A direct coupling between polymers and nematics destabilizes the system and generates rich dynamics and flow states. At high $De$, viscoelasticity reduces drag in the transient active turbulence state by generating strong polymer stresses acting against the local extensional flow and freezing the pattern of defects in the flow.

Fig. 6. Interactions for swimmers in viscoelastic fluids. (a) A pusher squirmer is permanently trapped near a surface due to large polymer stresses at the tail of the swimmer (adapted with permission from Ref. [168]). Inset: polymer extension around the squirmer represented by ellipsoids oriented and scaled by the eigenvectors of the polymer conformation tensor. (b) Trajectories of a self-propelled spherical particle (diameter: 7.82 μm) in a circular domain of a viscoelastic fluid. The particle escapes from the wall more frequently at higher $De$ [173]. (c) Microswimmers in a vortical flow are randomly distributed if the fluid is Newtonian but they concentrate along limit cycles if the fluid is viscoelastic, with swimmers of higher motility forming a larger cycle (adapted with permission from Ref. [189]). (d) Growth rate of the linear perturbation for a homogeneous and isotropic distribution of pushers in a viscoelastic fluid (adapted with permission from Ref. [176]). (e) Flow field (velocity $u$ and elastic energy $\tau^p/2$) and distribution of swimmers in Newtonian (top) and viscoelastic fluids (bottom). Viscoelasticity in the fluid reduces the size and intensity of the coherent flow structure in a pusher suspension and enhances the pusher aggregation, while it has less effects on the pullers (adapted with permission from Ref. [179]).
More recently, cell division and motility in a viscoelastic fluid environment were considered theoretically using a two-phase fluid model and polymers were found to suppress the flow at the interface and damp the interfacial instabilities [182].

5. Conclusion and perspective

In this review, we summarized our knowledge on microswimming in complex viscoelastic fluids, with an emphasis on mathematical theory, engineering applications and biological relevance. Complex fluids, which typically have multi-scale structures, are involved in many biological processes so the interactions between microswimmers and complex fluids might have played an important role in the evolution of organisms. Past research in this field, within the continuum framework of low-Reynolds number hydrodynamics and non-Newtonian rheology, provides fundamental physical insight in the complex behaviors of motile microorganisms in nature. By introducing multiple nonlinearities, viscoelasticity in the fluid breaks the time-reversal symmetry of Newtonian Stokes flows, thereby changing the dynamics of microswimmers and their hydrodynamic interactions with their environment.

We summarized in detail the impact of viscoelasticity on the motility of a single microswimmer and many of these results can be related to classical knowledge in the dynamics of suspensions in complex fluids. For a spherical squirmer, the velocity change at small De is due to the asymmetric distribution of polymer stresses. At large De, strong polymer stresses in the local extensional regions reduce the speed, an effect that is reminiscent of the motion of passive particle and droplets in viscoelastic fluids [203]. A small-amplitude flagellum undergoing waving motion swims slower in a viscoelastic fluid because it creates a small-amplitude oscillatory shear flow and thus experiences a reduced dynamic viscosity [66,67]. For models of flagella with more realistic features (flexibility, finite-length, asymmetric waving form, etc.), the speed change is more intricate and depends on the details of both swimmer and fluid [169].

The viscoelasticity of the fluid also affects hydrodynamic interactions, similarly to many other viscoelastic flows. Viscoelasticity-induced surface/swimmer and swimmer/swimmer attractions for pushers resembles the surface attraction of a particle settling near a wall and the mutual attraction between settling particles in viscoelastic fluids [204–206]. Aggregation and alignment occur for both swimmers and passive particles in a vortical viscoelastic flow [204,207]. The role of polymer in destabilizing swimmer suspensions is reminiscent of the occurrence of viscoelastic turbulence at negligible Reynolds numbers [208,209]. Future studies will be necessary to reveal the extent to which these phenomena are fundamentally identical and how the swimmer motility affect these results. For example, numerical methods used to study suspensions of interacting particles and cells in viscoelastic fluids [210–214] could be extended to investigate the role of fluid elasticity on the transport of microswimmers.

As this review makes clear, there remain many important questions to answer in future studies of microswimming in complex fluids. One key challenge will be to distinguish the universal effects of viscoelasticity on microswimming from those specific to particular swimming modes. An interdisciplinary bridge between fluid mechanics, microbiology, and rheology will be necessary to build the full physical picture. One promising avenue is the extension to microswimmers of our current understanding of the hydrodynamics and rheology of suspensions of passive particles in simple flows [215–217]: we refer readers to a recent review of the rheology of suspensions in viscoelastic fluids [218]. Due to the nonlinearities in the case of complex fluids, the classical framework of Stokesian Dynamics cannot be directly applied for viscoelastic fluids beyond the weakly nonlinear regime so numerical simulations able to fully resolve interactions between microstructure and microswimmer will therefore be needed. Another topic of potential interest is the comparison between microswimming in polymer solutions and liquid crystals, which display non-isotropic fluid elasticity due to the director field [219,220]. Inspired by active and passive micro rheology [221,222], we could also envision a situation where synthetic microswimmers become useful as micro rheological probes for complex fluids. Once we fully unravel the dynamics of microswimmers in complex fluids, we can consider practical applications where synthetic microswimmers can be used in targeted drug delivery and in vivo biochemical detection studies.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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