Turning Bacteria Suspensions into Superfluids

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The rheological response under simple shear of an active suspension of Escherichia coli is determined in a large range of shear rates and concentrations. The effective viscosity and the time scales characterizing the bacterial organization under shear are obtained. In the dilute regime, we bring evidence for a low-shear Newtonian plateau characterized by a viscosity decreasing with concentration. In the semidilute regime, for particularly active bacteria, the suspension displays a “superfluidlike” transition where the viscous resistance to shear vanishes, thus showing that, macroscopically, the activity of pusher swimmers organized by shear is able to fully overcome the dissipative effects due to viscous loss.

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Owing to its relevance in medicine and ecology and its importance for technological applications, the hydrodynamics of active suspensions is at the center of many recent fundamental studies [1,2]. In nature, wide classes of living microorganisms move autonomously in fluids at very low Reynolds numbers [3]. Their motility stems from a variety of propulsive flagellar systems powered by nanomotors. For bacteria such as Bacillus subtilis or E. coli, the propulsion comes from the rotation of helix-shaped flagella creating a propulsive force at the rear of the cell body [4]. Consequently, many original fluid properties stem from the swimming activity [5–11]. Because of hydrodynamic interactions, bacteria may produce mesoscopic patterns of collective motion sometimes called “bioturbulence” [12–17]. In a flow, these bacteria may organize spatially [18], and under shear, for pusher swimmers, the swimming activity yields the possibility to decrease the macroscopic viscosity to values below the suspending fluid viscosity [5]. In the dilute regime, kinetic theories via a simple account of the dominant long range hydrodynamic field [19–22] provide closed forms for shear viscosity as a function of shear rate. Remarkably, at a low shear rate, these theories predict a Newtonian plateau with a viscosity decreasing linearly with concentration [19–21]. On the other hand, phenomenological theories were also proposed to describe macroscopically active suspensions via a coupling of hydrodynamic equations with polar and/or nematic order parameters [2,5,6,23–25]. A striking outcome of these theories is that, for a set of coupling parameters rendering essentially a high swimming activity, a self-organized motive macroscopic flow may show up in response to shear [23–25]. This onset of a dissipationless current is described in analogy with the superfluidity transition [23,24] of liquids. Experimental evidence for viscosity reduction to values below the suspending fluid viscosity was brought for B. subtilis [8] and E. coli [26] suspensions. However, no viscosity vs shear rate and vs time under steady and uniform shear exists. Moreover, these pioneering experiments did not provide evidence for the low-shear viscous plateau which is at the core of all theoretical predictions in the dilute regime. Finally, the phenomenological predictions for the nonlinear regime have remained so far unobserved. Noticeably, for unicellular algae, viewed as “puller” swimmers, the predicted low-shear rate increase of viscosity was measured experimentally [9]. In this Letter, in addition to a viscosity vs shear rate and vs time characterization of an E. coli suspension, we provide, in the dilute regime and at a low shear rate, experimental evidence for a linear decrease of the apparent viscosity with bacteria concentration. We also explore regimes of higher concentration and describe the conditions where we observe a transition to a dissipation-free macroscopic flow.

The active fluids considered here are prepared out of two strains of wild type E. coli (ATCC9637 and RP437) suspended into a minimal medium where the bacteria are still motile but do not divide. ATCC9637 is cultured overnight at 25°C in LB medium shaken at 240 rpm. RP437 is cultured overnight at 30°C and shaken at 240 rpm in M9 minimal medium supplemented with 1 mg/ml casamino acids and 4 mg/ml glucose. Next, the culture is washed twice by centrifugation (2300 g for 10 min), and the cells are resuspended into a motility medium containing 10 mM potassium phosphate pH 7.0, 0.1 mM K-EDTA, 34 mM K-acetate, 20 mM sodium lactate, and 0.005% polyvinylpyrrolidone (PVP-40). To avoid bacterial sedimentation, the suspension is mixed with Percoll (1 vol/1 vol). The bacteria concentration n is represented by its volume fraction φ = n/Vb, where Vb is the bacteria body volume chosen as the classical value Vb = 1 μm³.
Shear stress is measured in a low-shear Couette rheometer (Contraves 30) designed especially for probing low-viscosity fluids. The inner bob (radius $R_i = 5.5$ mm, length 8 mm, and underside cone angle 20°) is suspended by a torsion wire into a cup (inner radius $R_i = 6$ mm). The cup rotates at an angular rate $\Omega$ controlled by a computer. The corresponding shear rate is $\dot{\gamma} = (\Omega R_0/R_0 - R_i)$ and is given with a precision of 0.5%. The central feature, making this instrument very precise for low stress measurements, is that the central bob is kept fixed by a feedback counter-rotation of the suspending wire. The instrument measures the compensating torque required to keep the torsion wire at its null position. The torque is then converted into shear stress every 0.7 s with a sensitivity of $10^{-3}$ mPa.

Importantly, due to the small surface area between the fluid and the air, the flux of $O_2$ is insufficient to compensate the amount of $O_2$ consumed by the bacterial activity. To avoid bacteria suffocation and, consequently, a severe drop of activity, we supplement the suspension with L-serine, an amino acid allowing the bacteria to keep a significant swimming activity in the absence of oxygen [27,28]. Therefore, in the early instants of the measurements, the bacteria are still in oxygenated conditions, but, since they consume the oxygen, their mean velocity and diffusion coefficient decrease and stabilize within about 10 min. Consequently, we observe a continuous increase in the suspension viscosity until a constant value is reached.

TO obtain a full rheogram as displayed in Fig. 1(b), the following protocol is used. A volume (1.25 ml) of the suspension is poured into the rheometer’s cup, and then the bob is set into place. After 30 s of rest, the cup is rotated for 30 s at a steady state shear rate. The rotation is then stopped for 30 s. These steps are repeated with increasing shear rate values. Once the highest $\dot{\gamma}$ is reached, the procedure is

![FIG. 1 (color online). Shear stress response for an E. coli suspension (ATCC9637 strain, $T = 25 \degree C$).](image)

(a) Shear stress $\Sigma$ rescaled by the applied shear rate $\dot{\gamma}$ during the rotation to display an effective viscosity $\Sigma/\dot{\gamma}$ in the sheared regime. Gray and black lines: Fluid without bacteria ($\phi = 0$). Colored solid lines: Fluid with bacteria ($\phi = 0.67\%$). Various $\dot{\gamma}$ are applied ranging from 64 (dark blue line) down to 0.022 s$^{-1}$ (dark red line).

(b) Relative viscosity $\eta/\eta_0$ averaged over three realizations as a function of $\dot{\gamma}$ (empty square, $\phi = 0.11\%$; empty circle, $\phi = 0.21\%$; empty triangle, $\phi = 0.44\%$; empty diamond, $\phi = 0.67\%$). The solid line is an adjustment by the Carreau law: $\eta/\eta_0 = 1.08 - 0.795/[1 + (\dot{\gamma}/0.6)^2]$. The vertical dashed line shows $\dot{\gamma}_c$; below this shear rate $\eta(\dot{\gamma})/\eta_0$ is less than 1.

(c) Values of the plateau viscosity $\eta_p/\eta_0$ as functions of the bacteria volume fraction $\phi$ for very low shear rates (empty circle, $\dot{\gamma} = 0.022$ s$^{-1}$; empty square, $\dot{\gamma} = 0.04$ s$^{-1}$; empty triangle, $\dot{\gamma} = 0.075$ s$^{-1}$).
Rheology measurements were first performed to obtain viscosity at volume fractions between $\phi = 0.1\%$ and 0.67%. In Fig. 1(a), we display the stress responses obtained for a suspension at a given $\phi$, for various shear rates. For the suspending fluid alone (a Newtonian fluid, $\eta_0 = 1.4$ mPa · s), stress-time responses, at the start or at the stop of the applied shear, are fast and correspond to the device compliance (gray and black lines). A similar behavior is observed for the suspensions probed at high shear rates which moreover display a viscosity higher than $\eta_0$ as observed classically for suspensions of passive particles. However, at low shear rates, a strikingly different behavior is observed. When shear starts, the stress jumps to the value measured in the absence of bacteria, and then, after an exponential decrease, lasting for a few seconds, a steady effective viscosity—$\eta$—is reached. When shear stops, the stress decreases abruptly and eventually changes sign. Finally, the stress relaxes exponentially to 0 with a characteristic time $\tau_r$ not very different from the time scale $\tau^+$, needed to reach a steady viscous response under shear [data in Fig. 2(c)]. In this last stage, the bacterial motion induces a motive stress on the inner bob.

Figure 1(b) shows the suspension viscosity $\eta$ as a function of $\dot{\gamma}$ for different volume fractions ranging from $\phi = 0.11\%$ ($1.1 \times 10^9$ bact/mL) up to $\phi = 0.67\%$ ($6.7 \times 10^9$ bact/mL). We observe the three regimes predicted by the theories [19,20], (i) at high shear rates ($\dot{\gamma} > 1$ s$^{-1}$), the active contribution to viscosity is negligible and a Newtonian plateau appears akin to suspensions of passive particles of the same shape; (ii) below a critical
shear rate value \( \gamma_\ell \leq 1.5 \text{ s}^{-1} \), the suspension viscosity is lower than the suspending fluid viscosity; (iii) at low shear rates \( \gamma \leq 0.1 \text{ s}^{-1} \), an “active viscous plateau” \( \eta_p(\phi) \) appears. Furthermore, the theories also predict a linear dependence of \( \eta_p \) with \( \phi \) given by

\[
\eta_p = \eta_0 + K \left( \frac{r}{t_c} \right) \phi,
\]

where \( K \propto [A - B(\tau/t_c)] \); \( t_c \) is the time taken by a bacterium to drag the fluid over its size, and \( \tau \) characterizes the directional persistence of a swimming trajectory \[30\]; \( A \) and \( B \) depend solely on the bacterium shape \[19–22\].

As shown in Fig. 1(c), \( \eta_p(\phi) \) decreases linearly with the concentration as \( \eta_p/\eta_0 = 1 + K\phi \) with \( K \approx -120 \pm 10 \), as long as the shear rate is sufficiently low (i.e., in the range 0.022–0.075 \text{ s}^{-1}). The experimental results are very consistent with active suspension theories for pusher swimmers in the dilute regime. Noticeably, within the framework of closed form theories established in the dilute regime, such a complete determination of the viscous response can be used to give an estimation of the microscopic bacterial activity (see \[29\] for an explicit derivation of the dipolar strength and other microscopic parameters via the Saintillan \[20\] kinetic model).

We next compare the viscous response of the bacteria for two different activity levels. When both \( \text{O}_2 \) and \( \text{L-serine} \) are present in the suspension, one obtains an "hyperactivated" regime characterized by high values of bacteria diffusivity and swimming velocity \( D = 7 \times 10^{-11} \text{ m}^2/\text{s} \) and \( V_0 = 28 \mu\text{m/s} \) for ATCC9637). After 10 min, when \( \text{O}_2 \) is consumed, the motility is maintained by the metabolization of \( \text{L-serine} \). In this case, one has a lower activity with \( D = 1.2 \times 10^{-11} \text{ m}^2/\text{s} \) and \( V_0 = 20 \mu\text{m/s} \) (see \[29\], Fig. 2).

The first regime only lasts a few minutes, and the measurements were thus restricted to a single \( \tilde{\gamma} \) sufficiently low (here 0.04 \text{ s}^{-1}) to estimate \( \eta_p \), the active plateau viscosity. In the “hyperactive” domain, \( \eta_p \) is again found for \( \phi < 0.6\% \) to decrease linearly with \( \phi \) but with a larger slope \( K' = -200 \pm 3 \) [see Fig. 2(b)]. This result demonstrates that the slope \( K \) is strongly related to the bacterial activity. Next, these experiments were repeated with the second \( \text{E. coli} \) strain (RP437). These bacteria have a lower swimming velocity \( V_0 = 20 \mu\text{m/s} \) and also a slightly longer body length 2.2 \( \mu\text{m} \) instead of 1.7 \( \mu\text{m} \). A linear \( \eta_p \) vs \( \phi \) relation is also found but with a larger negative slope \( K'' \approx -259 \pm 13 \) [red squares in Fig. 2(b)], demonstrating a correlation between the bacterial characteristics and the rheological response at a low shear rate.

We finally increase the number of bacteria in the solution to values above 0.6\%. For the ATCC9637 strain in a medium that does not contain oxygen [empty symbols in Fig. 2(b)], we observe that the viscosity becomes constant and independent of \( \phi \) for \( \phi > 0.7\% \) with \( \eta_p/\eta_0 \sim 0.2 \). In oxygenated conditions and for highly motile batches, we measure that the viscosity \( \eta_p/\eta_0 \) also reaches a constant value beyond \( \phi \sim 0.5\% \) and \( \phi \sim 0.4\% \) for ATCC9637 and RP437, respectively. For the three cases, the active plateau viscosity becomes thus independent of the concentration over a significant domain (from \( \phi = 0.6\% \) up to 2.4\%).

In addition under hyperactivated conditions, one observes a viscous response reaching zero [see Fig. 2(a)], meaning that the local viscous dissipation is macroscopically entirely compensated by the swimming activity. Moreover, with the very active RP strain, negative values for \( \eta_p \) [see Figs. 2(a) and 2(b)] could be obtained at the edge of the transition. To validate the existence and robustness of this low viscosity regime, other experiments at lower shear rate and higher temperature were conducted: They all display a \( \eta \sim 0 \) regime. These results point towards the idea of an organization process triggered by the shear flow which may last even when the shear ceases. In Fig. 2(c), we plot the relaxation times \( r^+ \) and \( r^- \) as functions of the mean distance \( \bar{x} = (\phi^{1/3}) \) between bacteria. The transition between the dilute and the semidilute regimes occurs here for a distance \( \bar{x} \approx 6 \mu\text{m} \). In fact, below this length, the relaxation time is found to decrease linearly with \( \bar{x} \) (while the viscosity \( \eta_p \) remains constant). The result seems to be consistent with the phenomenological picture proposed for an active polar swimmer and with the clustering interaction recently observed in the absence of flow \[16\] and suggests that the collective effect becomes important for distances shorter than 6 \( \mu\text{m} \).

In conclusion, our experiments show that the bacterial activity has a measurable influence on shear viscosity. We brought direct experimental evidence, at low shear rates, for an active viscous plateau whose value decreases linearly with concentration (for \( \phi < 0.3\% \)). This confirms a central prediction for active pusher suspensions in the linear kinetic regime and validates the assumptions of the model, the main ones being the possibility to use the Bretherton Jeffery equations as a reliable swimming kinetic model and, most importantly, the hypothesis of the existence of an effective Gaussian disorientation noise, that eases the resolution of the Fokker-Planck equation. The most striking feature of the rheological response is indeed the emergence of a viscousless superfluidity regime (\( \eta \sim 0 \) or even lower). Presently, there is no ab initio microhydrodynamic calculation describing the impact of bacteria interactions and, possibly, the influence of collective organization on the macroscopic rheology. Therefore, one has to rely on phenomenological arguments to identify the possibly relevant macroscopic hydrodynamic contributions \[2,23\]. In this framework, several authors \[23–25\] investigated the responses to shear of active polar particles, and, remarkably, they predicted the possibility of a transition to a “zero-viscosity” regime when the activity is increased. This result can also be cast in the framework of recent experimental works pointing out the possible use of bacterial motion to
drive mechanical devices [31–33]. Actually, this Letter goes in this direction, as we show, at least in principle, that rotational macroscopic power could be extracted from the swimming activity as for a rotatory motor [25]. Finally, such a strong viscosity reduction may be a crucial element when considering macroscopic transport and particle dispersion in porous systems or in capillary networks, a central question to many applications involving bacterial fluids.

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[29] See Supplemental Material at http://link.aps.org/supplemental/10.1103/PhysRevLett.115.028301 for (1) the variation of the viscosity \( \eta \), of the mean bacteria velocity and diffusion coefficient with the oxygen concentration and (2) the details of the calculation to estimate the strength of the force dipole based on the rheological data.