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Why Do Red Blood Cells Have Asymmetric Shapes Even in a Symmetric Flow?

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Understanding why red blood cells (RBCs) move with an asymmetric shape (slipperlike shape) in small blood vessels is a long-standing puzzle in blood circulatory research. By considering a vesicle (a model system for RBCs), we discovered that the slipper shape results from a loss in stability of the symmetric shape. It is shown that the adoption of a slipper shape causes a significant decrease in the velocity difference between the cell and the imposed flow, thus providing higher flow efficiency for RBCs. Higher membrane rigidity leads to a dramatic change in the slipper morphology, thus offering a potential diagnostic tool for cell pathologies.

Introduction.—Blood is a complex fluid that is primarily composed of red blood cells (RBCs), which occupy (in a healthy human body) about 45% of the blood volume. The rest consists of plasma, while the other blood elements (white blood cells, platelets, etc.) take up less than 1% of the total blood volume.

Because blood is a complex fluid, descriptions of its flow properties escape the traditional laws for simple fluids. The complex character results from an intimate coupling between the shape of RBCs and the ambient plasma, which leads to a rich set of RBC morphologies in the blood circulatory system. Understanding the selection of shapes and dynamics among a large manifold of possibilities is a challenging problem. This type of complexity is a characteristic property of nonequilibrium dissipative systems for which general thermodynamic principles, such as minimization of energy, maximization of entropy, etc., cannot be applied.

At equilibrium (i.e., in a quiescent fluid), healthy RBCs have a biconcave shape that corresponds to a minimal membrane bending energy. Under nonequilibrium conditions, as experienced in a simple shear flow, RBCs reveal a number of interesting shapes and dynamics. The most classical ones are tank-treading ellipsoids [1] (the cell elongates as an ellipsoid and orients itself with a fixed direction, while the fluid membrane shows a tank-tread-like motion) and tumbling motions [2]. These features, among others, are shared with phospholipid vesicles.

In a parabolic distribution of the velocity field (e.g., Poiseuille flow), a situation that is relevant for the blood circulatory system, RBCs assume a parachutelike shape. A clear distinction must be made between a symmetric shape, hereafter simply called a parachute, and an asymmetric one, referred to as a slipper. The reasons of why RBCs often show asymmetric shapes in a symmetric tube are a puzzle in blood microcirculatory research [3]. Earlier numerical studies reported on a manifestation of these slipper shapes [4,5]. The existence of the slipper has been linked with (i) confinement, and (ii) RBC elasticity (associated with the cytoskeleton a cross-linked network of proteins lying beneath the RBC membrane).

Elucidating the basic relevant mechanisms that are responsible for manifesting the slipper shape is highly desirable. The following questions represent the motivations for this study: (i) Is confinement relevant? (ii) Does the cytoskeleton play a necessary role in the development of the slipper? (iii) How does the slipper occur? And (iv) why does the slipper occur? These constitute questions of primary importance that we would like to address in this Letter.

We focus on the simplest configuration with the aim of identifying the minimal ingredients that are needed for the slipper shape. We consider a purely phospholipid membrane, known as a giant vesicle [6]. This allows us to shed light on whether or not the cytoskeleton is essential. Vesicles and RBCs both have inextensible membranes and exhibit bending modes of the same order of magnitude. The main difference is that RBCs are endowed with an in-plane shear elasticity by virtue of their cytoskeletons. We also disregard the presence of lateral walls, as we neglect confinement. In reality, a Poiseuille flow is bounded by walls (for a real system we have in mind a weak confinement).

We consider a 2D simulation, motivated by the fact that in several circumstances a 2D study [7] captures the 3D results [8,9].

Model.—We consider the limit of the vanishing Reynolds number (the Stokes limit) and focus on the situation where the internal fluid has the same viscosity as the ambient fluid. The imposed flow reads

\[ \nu = \nu_\infty \left[ 1 - \left( \frac{r}{W} \right)^2 \right] e_x, \]

(1)
where \( C = 2v_{\text{max}}/W^2 \) is the curvature of the flow and must be compared to the typical vesicle radius \( R_0 \). Since here we disregard confinement only \( C \) has a physical meaning.

Vesicle dynamics is formulated using a boundary integral formulation [10,11], in which we express the velocity at a point on the membrane as

\[
v(x_0) = v^* (x_0) + \frac{1}{\eta} \int G(x - x_0) \cdot f(x) ds,
\]

where the integral is performed along the membrane,

\[
G_{ij}(x - x_0) = -\delta_{ij} \ln|x - x_0| + \frac{(x - x_0)_i(x - x_0)_j}{|x - x_0|^2},
\]

is the free-space two dimensional Greens function (or Oseen tensor) for the Stokes operator [12], and \( \eta \) is the dynamic fluid viscosity. \( x \) and \( x_0 \) are 2D vector positions of a membrane point, and \( f \) is the membrane force per unit area, given by

\[
f = \kappa \left( \frac{d^2 c}{ds^2} + \frac{1}{2} c^3 \right) n - \zeta c n + \frac{d \zeta}{ds} t.
\]

This force balances the hydrodynamic force jump across the vesicle membrane. This force (derived in 2D in Refs. [7,13]) is obtained from the functional derivative of the Helfrich [14] bending energy \( E = \frac{\zeta}{2} \oint c^2 ds + \oint \zeta ds \), which includes the local arclength conservation constraint (expressing inextensibility) represented by the Lagrange multiplier \( \zeta \), the membrane curvature \( c \), the unit normal and tangent vectors \( n \) and \( t \), the arclength coordinate \( s \), and the membrane bending rigidity \( \kappa \). Equation (2) is solved numerically (see [15,16]).

Results.—Length is measured in units of the vesicle radius \( R_0 \) (defined as the radius of a circle having the same enclosed area). The first dimensionless number that can be formed is \( C_a = \eta v_{\text{max}} R^2/\kappa \) (or \( \eta CR^4/\kappa \) if we use \( C \), the only parameter characterizing the flow in an unbounded geometry). The vesicle deflation is defined as the ratio of the actual enclosed area over the area of a circle having the same perimeter \( p \):

\[
\nu = \frac{A}{\pi[p/2\pi]^2}.
\]

where \( \nu = 1 \) for a circle, and \( 0 < \nu < 1 \) otherwise. Thus, we have two dimensionless parameters \( \nu \) and \( C_a \). We also use physical units in our discussion in order to highlight the feasibility of experimental investigations and the connection to blood flow data. We postpone a detailed discussion of the results in a dimensionless form for future work. We have fixed the parameters associated with the imposed Poiseuille flow (\( v_{\text{max}} \) and \( W \)) and have varied \( \nu \). For definiteness, we first set \( v_{\text{max}} = 800 \mu\text{m/s} \) (a typical value in human venules [2]) and \( W = 10R_0 \), while \( \eta \) is fixed to the water viscosity (close to that of the blood plasma), and \( \kappa = 10^{-19} \text{J} \) (a typical value for vesicles and RBCs [17]). For large enough \( \nu (\nu > 0.7) \), we have found that vesicles that were initially placed off-center in the Poiseuille flow migrate towards the center, in accord with prior results [7,8]. At the center line, the vesicle assumes a parachute (symmetric) shape. This shape is stable against various perturbations. An interesting outcome is that below a critical value of \( \nu = \nu_c \approx 0.7 \), the parachute shape develops an instability that is characterized by the loss of up-down symmetry. The resulting shape is called a slipper (see movies in [18]).

We have analyzed, in some detail, the occurrence of this instability. We characterize this symmetry-breaking behavior using the \( y \) position of the center of mass, denoted as \( Y_G \). For \( \nu > \nu_c \), the shape is symmetric, and hence \( Y_G = 0 \). For \( \nu < \nu_c \), \( Y_G \) acquires a nonzero value. The results are reported in Fig. 1. Because of the symmetry, the \( \pm Y_G \) solutions are equivalent, as checked numerically for sev-

![Equilibrium Lateral Position](image1)

**Equilibrium Lateral Position**

![Slip Velocity](image2)

**Slip Velocity**

![Steady Tank-treading Velocity](image3)

**Steady Tank-treading Velocity**

![Graph1](image4)

**Graph1** (color online). Top panel: The behavior of the equilibrium lateral position of the center of mass \( Y_G \) as a function of \( \nu \) for different values of the flow parameter \( v_{\text{max}} \). The full lines are fits with \( (\nu_c - \nu)^{1/2} \). Middle panel: The slip velocity (normalized by \( v_{\text{max}} \)) as a function of \( \nu \). The slip velocity (or lag) is defined as the difference between the vesicle (when its vertical position has reached a plateau with time) and the corresponding unperturbed velocity at the position of the vesicle center of mass (\( Y_G = 0 \) for the parachute shape, and \( Y_G \neq 0 \) for the slipper). Bottom panel: The tank-treading velocity. The link with physical units is explained in the discussion.
eral cases that we do not report here. We can conclude that
the parachute-slipper transition corresponds to a supercriti-
cal bifurcation (the dynamical analogue of a second order
phase transition). A fit is shown in Fig. 1 (solid line in the
upper panel), and we find that in the vicinity of the bifur-
cation point \( Y_G \sim 0 \) and for \( \nu < \nu_c \), \( Y_G \sim (\nu_c - \nu)^{1/2} \)
constitutes a very good approximation. This is the signa-
ture of a supercritical (or pitchfork) bifurcation, which
belongs to the cusp catastrophe family. For the set of
parameters explored so far, we find that the supercritical
nature of the bifurcation persists. We cannot, however,
exclude the possibility of a subcritical bifurcation (the
dynamical analogue of a first order transition, where \( Y_G \)
would jump from zero to a finite value at a critical \( \nu \))
belonging to the butterfly catastrophe family. Our results
suggest that \( Y_G \) should obey, in the vicinity of the bifurca-
tion point, an equation of the form
\[
\frac{dY_G}{dt} = (\nu_c - \nu)Y_G - Y_G^3.
\] (5)

This equation has a trivial fixed point \( Y_G = 0 \) for all \( \nu \)
and a nontrivial pair of solutions \( Y_G = \pm \sqrt{\nu_c - \nu} \) for \( \nu < \nu_c \). It is a simple matter to show that \( Y_G = 0 \) is stable for
\( \nu > \nu_c \) and loses its stability for \( \nu < \nu_c \) in favor of two
stable branches \( Y_G = \pm \sqrt{\nu_c - \nu} \).

Once the question of how the slipper occurs is settled,
we are in a position to address the next natural question of
why a slipper forms at all. The shear rate is minimal
(actually it vanishes) at the center of the Poiseuille flow,
and it is expected that any deviation of the vesicle from the
center line would be penalized by higher dissipation as a
result of the higher shear rate. Furthermore, for a symmet-
ric (parachute) shape, membrane tank-treading is absent
because of an obvious central symmetry. As a conse-
quence, the relative velocity of the fluid inside the para-
chute vanishes. A slipper shape is accompanied by a tank-
treading membrane, and thus by flow circulation inside the
cell. Because of the presence of this additional flow, one
would expect higher dissipation, and from this point of
view a slipper should not be favored.

In fact, we find that a key property that dictates the
establishment of a slipper is associated with the notion of
lag between the vesicle and the imposed flow (i.e., the
difference between the vesicle velocity and that of the
bare imposed flow). One outcome of our study (Fig. 1,
middle panel) is that the lag increases with decreasing \( \nu \)
and attains a maximum (within numerical uncertainties) at
the bifurcation point, where the manifestation of a slipper
is accompanied by a decrease in the lag. This feature has
been found for every parameter explored so far. By assum-
ing an asymmetric shape, the vesicle reduces its lag, albeit
at the price of inducing membrane tank-treading and in-
ternal flow. The reduction of lag occurs as a subtle inter-
play between the shape adaptation to the flow and a
compromise between additional membrane and vesicle
internal dissipation. As a consequence, the slipper becomes
a favorable shape.

We have performed a systematic study to determine the
boundary in parameter space that separates the parachute
shape from the slipper one. In this brief exploration, we
keep the membrane bending rigidity fixed to a typical value
for vesicles used above and vary both the flow strength,
measured by \( v_{\text{max}} \), and the degree of deflation \( \nu \). The
results are shown in Fig. 2. There, we also show a panel
of shapes that are exhibited as a function of the flow and
deflation conditions. The slipper takes place in the yellow
region, while the parachute shape prevails in the violet one.
It was also useful to distinguish between a parachute with
a negative curvature at the rear and one with a positive
curvature (renamed a bullet).

Finally, we studied the evolution of the slipper shape as
a function of the membrane rigidity, for fixed \( v_{\text{max}}, W \), and \( \nu \).
We found a dramatic change in the morphology, even when
the rigidity increases by a factor of 2 only. Figure 3 shows
the results. Several RBC pathologies, such as malaria, are
accompanied by an increase in the elastic membrane
modulus. Exported proteins from parasites cause a signifi-
cant increase in membrane rigidity, which may attain
values that are 2.6 times higher than those of uninfected
RBCs [19]. This result, after adequate refinement that
includes shear elasticity and lateral walls, may be used as
a potential diagnostic for detecting RBC pathologies.
Multilayer vesicles, as well as polymer-decorated mem-
branes, exhibit higher rigidity, thus offering a model sys-
tem on which to directly test the present findings in a
weakly confined channel.

Discussion.—Vesicles and RBCs exhibit both membrane
inextensibility and bending modes. Membrane bending

![FIG. 2 (color online). Phase diagram in the plane of reduced
volume and maximum imposed velocity. Evolution of the shapes
is shown. Filled squares represent the boundary of the symmetry-
breaking bifurcation, and the solid line is a guide for the eyes.
The horizontal dashed line represents the boundary below which
the parachute shape has a negative curvature at the rear. Here
\( W/R_0 = 10 \).](188101-3)
rigidity for vesicles is of the order of $10^{-19}$ J and is close to that of RBCs [17]. Bending under a shear flow with shear rate $\gamma$ is characterized by the dimensionless number $C_s = \eta R_0^3 \gamma / \kappa$. Unlike vesicles, RBCs have an in-plane shear elasticity (as a result of the cytoskeleton). It is characterized by the shear modulus [17] $\mu_s \sim 2 - 6 \times 10^{-6}$ N/m [17], and by the dimensionless number $C_s = \eta R_0 \gamma / \mu_s$. The ratio between $C_s$ and $C_w$ shows that bending and shearing are of the same order. We have disregarded here the shear elasticity. This has allowed us to show that the slipper manifestation is not dictated by the cytoskeleton.

One possibility might be a necessity for the membrane to slip under a shear flow with shear rate $\gamma$ is characterized by the dimensionless number $C_s = \eta R_0^3 \gamma / \kappa$. Unlike vesicles, RBCs have an in-plane shear elasticity (as a result of the cytoskeleton). It is characterized by the shear modulus [17] $\mu_s \sim 2 - 6 \times 10^{-6}$ N/m [17], and by the dimensionless number $C_s = \eta R_0 \gamma / \mu_s$. The ratio between $C_s$ and $C_w$ shows that bending and shearing are of the same order. We have disregarded here the shear elasticity. This has allowed us to show that the slipper manifestation is not dictated by the cytoskeleton. Future studies should include shear elasticity and confinement for a quantitative application to RBCs.

For small arteries (with a diameter of about 100 $\mu$m), and venules (with diameters in the 20 $\mu$m range) a typical $\nu_{\text{max}}$ value falls in the interval between $10^5$ $\mu$m/s and $10^4$ $\mu$m/s [2]. Consequently, given the fact that $\nu \sim 0.65$ for RBCs, slippers (see Fig. 2) are more favorable in venules and capillaries than they are in arteries, a result in agreement with reported experiments [3]. We are not aware of a phase diagram like the one presented in Fig. 2 for vesi ces. Thus, this Letter should serve as a guide for future systematic experimental studies on vesicles, which, unlike RBCs, lend themselves to ample variation with respect to their reduced volume. This is an essential step in testing the present findings. It will also be interesting to study the role of confinement on RBCs as reported experimentally [20].

Conclusion.—We have shown that a major advantage of the slipper shape is a reduction in the lag (i.e., the slipper shape enhances transport efficiency). Neither confinement nor membrane shear elasticity seems to be a necessary ingredient [4,5,21]. It is not yet clear whether or not, beyond the transport efficiency found here, the slipper shape is dictated by other physiological determinants. One possibility might be a necessity for the membrane to tank-tread (recall that tank-treading is absent for a symmetric shape) in order to efficiently mix the hemoglobin, and thus, enhance oxygen distribution in tissues.

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FIG. 3 (color online). Evolution of the morphology of a slipper as a function of membrane rigidity. $\kappa = 1$ refers to the typical value $10^{-19}$ J for vesicles and RBCs. Here we have taken reference values (typical in venules), namely $V_{\text{max}} = 800$ $\mu$m/s and $W/R_0 = 10$.