

Bistability, Oscillations, and Bidirectional Motion of Ensemble of Hydrodynamically Coupled Molecular Motors

P. Magaretti,^{1,2,*} I. Pagonabarraga,^{3,4,5} and J.-F. Joanny^{6,7}

¹Max-Planck-Institut für Intelligente Systeme, Heisenbergstr. 3, D-70569 Stuttgart, Germany

²IV. Institut für Theoretische Physik, Universität Stuttgart, Pfaffenwaldring 57, D-70569 Stuttgart, Germany

³Departament de Física de la Matèria Condensada, Facultat de Física, Universitat de Barcelona, Carre Martí i Franques 1, Barcelona 08028, Spain

⁴UBICS, Institute of Complex Systems, Universitat de Barcelona, Barcelona 08028, Spain

⁵CECAM, Centre Européen de Calcul Atomique et Moléculaire, École Polytechnique Fédérale de Lausanne, Batochime, Avenue Forel 2, 1015 Lausanne, Switzerland

⁶Physicochimie Curie (Institut Curie/CNRS-UMR168/UPMC), Institut Curie, Centre de Recherche, PSL Research University, 26 rue d'Ulm 75248 Paris Cedex 05, France

⁷ESPCI 10 rue Vauquelin 75005 Paris, France

(Received 11 May 2016; published 19 October 2017)

We analyze the collective behavior of hydrodynamically coupled molecular motors. We show that the local fluxes induced by motor displacement can induce the experimentally observed bidirectional motion of cargoes and vesicles. By means of a mean-field approach we show that sustained oscillations as well as bistable collective motor motion arise even for very large collection of motors, when thermal noise is irrelevant. The analysis clarifies the physical mechanisms responsible for such dynamics by identifying the relevant coupling parameter and its dependence on the geometry of the hydrodynamic coupling as well as on system size. We quantify the phase diagram for the different phases that characterize the collective motion of hydrodynamically coupled motors and show that sustained oscillations can be reached for biologically relevant parameters, hence, demonstrating the relevance of hydrodynamic interactions in intracellular transport.

DOI: 10.1103/PhysRevLett.119.168101

The transport of macromolecules, vesicles, and organelles inside cells relies on the active motion of molecular motors along biofilaments [1]. When motors pull on organelles or vesicles, the fluid flow that they induce provides motor-motor hydrodynamic interactions (HI), hence, providing an additional mechanism for molecular motors coupling [2] that, possibly, is responsible of the experimentally observed enhanced velocity of motors pulling on fluid vesicles [3]. In particular, the motion of these cargoes has been observed to be mono- and bidirectional, the latter relying on the presence of motors pulling the cargoes in opposite directions [4–10]. While previous studies have focused preferentially on rigidly coupled motors [11–14], in this Letter we show that the bidirectional motion observed in experiments can be induced and controlled via the HI induced by motors active displacement. Exploiting a mean-field approach we identify the key parameters controlling the onset of the bidirectional motion and we show the relevance of HI for biologically relevant scenarios.

While a few motors at the cargo tips can pull against the cargo drag, the rest of the motors can move along the cargo. Their net motion is responsible for the onset of HI. We model the molecular motor dynamics exploiting the two-state-model [11] that regards molecular motors as particles with two internal states. In order to account for motors

pulling in opposite directions, we model the two families of motors as a single family of *effective* motors that in the “bound” state experience a symmetric periodic force $f(x) = -\partial_x V = f_0 \cos(2\pi x/L) = f_0 \tilde{f}(x)$ of period L [14]. With rate $\omega_{\text{off}}(x)$ motors jump to a “weakly bound” state, in which they diffuse freely. Motors bind with rate $\omega_{\text{on}}(x)$. We describe the system in terms of densities of bound $[\rho(x)]$ and weakly bound $[\sigma(x)]$ motors. The total density, $\rho(x) + \sigma(x)$, cannot exceed the maximal filament occupation prescribed by excluded volume, and in order to keep analytical insight, we assume that the motors remain in a dilute regime. The motion of molecular motors occurs in the low Reynolds regime and it generates a fluid flow whose magnitude reads

$$\gamma v(x) = f(x) + \int f(y)\rho(y)W(x,y)dy, \quad (1)$$

where $W(x,y)$ is the dimensionless Oseen tensor accounting for HI [15] and γ is the single motor drag coefficient. According to Eq. (1), the induced fluid velocity is characterized by the local flow due to the force that the filament exerts on the molecular motor [first term in the right-hand side of Eq. (1)] and the collective hydrodynamic flow induced by the rest of the molecular motors at the position of the reference motor. This separation between local and

collective induced flows is characteristic of softly interacting motors, and it is absent for rigidly coupled motors [14]. Since the HI extends over distances large compared to the spatial variation of the motor-filament interaction, for a large system size Λ , $\Lambda \gg L$, we assume a mean field HI,

$$\int_{\Lambda} f(y)\rho(y)W(x,y)dy \approx k\frac{L}{\Lambda}\int_{\Lambda} f(y)\rho(y)dy + O(1),$$

$$k = \frac{1}{L}\int_{\Lambda} W(x,y)dy, \quad (2)$$

which holds when $f(x) = f_0$ and $\rho(x) = \rho_0$, and becomes exact for an infinite system, $\Lambda \rightarrow \infty$, if the motor density follows the spatial dependence imposed by the motor or filament periodic force. In this regime, reasonable when boundary effects can be disregarded, the spatial dependence of the hydrodynamic long-range coupling $W(x,y)$ experienced by the motors becomes negligible and it can be described by an effective dimensionless parameter k . For example, for motors of linear size R moving in a three-dimensional environment $W(x,y) = (3R/2)[1/(|x-y|)]$, leading to

$$k_{3D} = \int_{2R}^{\Lambda/2} \frac{3R}{2} \frac{dr}{r} = \frac{3R}{2L} \ln(\Lambda/4R), \quad (3)$$

which grows logarithmically with system size. Alternatively, when motors are pulling on a membrane-coated cargo, such as organelles or vesicles, their tails are linked to molecules embedded in the fluidlike membrane (characterized by a 2D viscosity $\eta_{2D,mem}$) quite more viscous than the cytoplasm (characterized by a 3D viscosity $\eta_{3D,cyt}$). When the intrinsic dynamics of motor tails does not affect motors dynamics, we can identify the dynamics of the motors with that of the tracers and therefore we can calculate the 2D HI. In this regime, we expect the 2D HI to dominate the three-dimensional flows induced in the cytoplasm for motor-motor separations smaller than $l = \eta_{2D,mem}/\eta_{3D,cyt}$. It is then reasonable to assume $k = k_{3D} + k_{2D}$ with

$$\frac{k_{2D}}{2} = \int_{2R}^M \ln \frac{l}{r} dr = \frac{M}{L} \left(1 + \ln \frac{l}{M}\right) - 2\frac{R}{L} \left(1 + \ln \frac{l}{2R}\right), \quad (4)$$

where $2 \ln(l/r)$ is the dimensionless Oseen tensor in two dimensions and $M = \min[l/e, (\Lambda/2)]$. Equations (3), (4) capture the diverging nature of HI for increasing system sizes Λ . Therefore, for large system sizes, $\Lambda/L \gg 1$, we can neglect the $O(1)$ terms in Eq. (3), and Eq. (1) reduces to

$$v(x) = \frac{f_0}{\gamma} (\tilde{f}(x) + k\langle \tilde{f}(x)\tilde{\rho}(x) \rangle_x), \quad (5)$$

where we have introduced the dimensionless density $\tilde{\rho}(x) = L\rho(x)$ [and similarly we introduce $\tilde{\sigma}(x) = L\sigma(x)$].

Accordingly, we can write, in dimensionless units, the evolution of the mean bound $\hat{\rho}(x) = \langle \tilde{\rho}(x) \rangle_{\tilde{\rho}, \tilde{\sigma}}$ and weakly bound $\hat{\sigma}(x) = \langle \tilde{\sigma}(x) \rangle_{\tilde{\rho}, \tilde{\sigma}}$ motor densities [16]

$$\begin{aligned} \dot{\hat{\rho}}(x) &= -\partial_x \lambda \hat{\rho}(x) [\tilde{f}(x) + k\langle \tilde{f}(x)\hat{\rho}(x) \rangle_x] + \\ &\quad - \tilde{\omega}_{\text{off}}(x)\hat{\rho}(x) + \tilde{\omega}_{\text{on}}(x)\hat{\sigma}(x), \\ \dot{\hat{\sigma}}(x) &= -\partial_x \lambda \hat{\sigma}(x) k\langle \tilde{f}(x)\hat{\rho}(x) \rangle_x + \tilde{\omega}_{\text{off}}(x)\hat{\rho}(x) - \tilde{\omega}_{\text{on}}(x)\hat{\sigma}(x) \\ &\quad + k_{\text{on}}[c_{\infty} - \hat{\rho}(x) - \hat{\sigma}(x)] - k_{\text{off}}\hat{\sigma}(x), \end{aligned} \quad (6)$$

where $\langle \psi(x) \rangle_{\rho} = \int \psi P[\rho(x)] dx$ stands for the average of ψ over the probability distribution of density profiles $\rho(x)$ while $\langle \psi(x) \rangle_x = (1/L) \int_0^L \psi(x) dx$ corresponds to the spatial average of ψ over the filament period (see Supplemental Material [17]). The first term in the right-hand side of Eqs. (6) describes motor advection according to the local velocity the motors are exposed to, the second term describes the motor binding kinetics, while the third term in the evolution equation for weakly bound motors describes the fact that motors can detach from the filament with rate k_{off} and bind with rate k_{on} , proportional to the available space $c_{\infty} - \hat{\rho}(x) - \hat{\sigma}(x)$, where c_{∞} is the motor concentration in bulk [18].

The motor flux in Eqs. (6) is proportional to $\lambda = f_0/\bar{\omega}L\gamma$, the ratio of the typical time a motor needs to slide down the potential $L\gamma/f_0$, and the characteristic inverse hopping time $\bar{\omega} = \langle \omega_{\text{on}}(x) + \omega_{\text{off}}(x) \rangle_x/2$. The bound and weakly bound rate densities depend on the filament structure. We consider the simple, periodic form

$$\tilde{\omega}_{\text{on/off}} = \max(\Delta\omega_{\text{on/off}} \mp \delta\omega_{\text{on/off}} \sin(2\pi x/L), 0), \quad (7)$$

which interpolates between regimes where the hopping rates are essentially homogeneous along the filament, $\Delta\omega_{\text{on/off}} \gg \delta\omega_{\text{on/off}}$, or only take place at localized regions on the filament, $\Delta\omega_{\text{on/off}} \ll \delta\omega_{\text{on/off}}$. The latter, together with an opposite sign between the bound and weakly bound rates accounts for the fact that binding and unbinding processes are localized at different regions along the filament. Normally, unbinding rates are more spatially localized than bounding rates, implying $\delta\omega_{\text{off}}/\Delta\omega_{\text{off}} > \delta\omega_{\text{on}}/\Delta\omega_{\text{on}}$.

For fast bulk kinetics $k_{\text{on,off}} \rightarrow \infty$, the concentration of weakly bound motors, $\hat{\sigma}$ is homogeneous along the filament. In this regime, Eqs. (6) decouple and the collective behavior of the motors is controlled by the evolution of $\hat{\rho}(x)$,

$$\begin{aligned} \dot{\hat{\rho}}(x) &= -\lambda\partial_x \hat{\rho}(x) [\tilde{f}(x) + k\langle \tilde{f}(x)\hat{\rho}(x) \rangle_x] + \\ &\quad - \tilde{\omega}_{\text{off}}(x)\hat{\rho}(x) + \frac{k_{\text{on}}}{k_{\text{off}} + k_{\text{on}}} \tilde{\omega}_{\text{on}}(x)[c_{\infty} - \hat{\rho}(x)], \end{aligned} \quad (8)$$

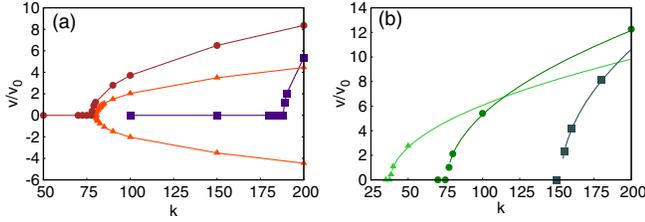


FIG. 1. Average molecular motor velocity normalized by the single bound motor velocity $v_0 = f_0/\gamma$ as a function of the dimensionless coupling k , as obtained from Eq. (8). Hydrodynamically (panel a) and rigidly (panel b) motor coupling, both characterized by $\Delta\omega_{\text{on}} = \Delta\omega_{\text{off}} = 1$ and $\delta\omega_{\text{on}} = \delta\omega_{\text{off}} = 1/2$ and $c_\infty = 1$. $\lambda = 0.015, 0.05, 0.1$ (square, circle, triangle) and $\lambda = 0.017, 0.033, 0.066$ (square, circle, triangle) for panels (a) and (b), respectively.

in terms of rescaled binding rates that keep the same functional dependence as in Eq. (7). We have numerically solved Eq. (8) using a Lax-Wendroff scheme with periodic boundary conditions applied at the ends of a period of the ratchet potential. Figure 1(a) shows that the configuration where motors do not have a net velocity, stable for weak HI, becomes unstable at a critical coupling, k_0 , above which a net motor current, which breaks left or right symmetry, develops; a similar scenario has been described for rigidly coupled motors [14]. Despite the apparent similarity between the emergence of net motion for soft and rigidly coupled motors, the different underlying physical mechanisms responsible for symmetry breaking lead to significant differences in collective motor dynamics. While hydrodynamically coupled motors are characterized by a non-monotonic dependence of k_0 on λ , see Fig. 1(a), the opposite holds true for rigidly coupled motors, Fig. 1(b) [19]. A linear stability analysis around the quiescent state shows

$$k > \frac{4}{\pi\lambda\left(\frac{\delta\omega}{\Delta\omega} - \pi\lambda\right)c_\infty} \quad (9)$$

as the sufficient condition [20] for symmetry breaking and onset of net motor currents. Since $k \geq 0$, Eq. (9) identifies an interval $\lambda \in [0; \lambda_{\text{max}}]$ for which symmetry breaking occurs, with $\lambda_{\text{max}} = (1/\pi)(\delta\omega/\Delta\omega)$. According to Eq. (9), k diverges for both $\lambda = 0, \lambda = \lambda_{\text{max}}$ leading to the existence of a minimum value $k = k_c$ at $\lambda = \lambda_c = \frac{1}{2}\lambda_{\text{max}}$. For the parameters used in Fig. 1(a) the stability analysis predicts $\lambda_c \sim 8 \times 10^{-2}$ and $k_c \lesssim 70$ in good agreement with numerical results. For rigidly bound motors, the second term in the denominator of Eq. (9) disappears, leading to an inverse proportionality between the coupling constant k and the dimensionless forcing λ , consistent with the results shown in Fig. 1(b). This difference has significant implications. For example, while for $\lambda > \lambda_c$ a reduction in λ will favor the onset of net fluxes, when $\lambda < \lambda_c$ decreasing λ will hinder, or even prevent, the development of a net motor flux.

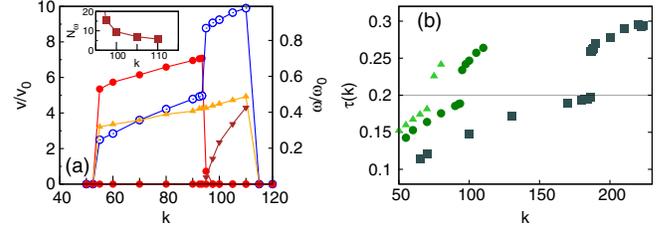


FIG. 2. (a) Average velocity (solid circles), velocity variance (open circles), (both normalized by the bound single motor velocity, $v_0 = f_0/\gamma$), and sustained oscillations frequency, ω_v (upward triangles) and frequency of inversion in the bistable regime Ω_v (downward triangles) (both normalized by the hopping rate peak value ω_0) upon variation of k . Ω_v has been magnified by a factor of 10 for sake of clearness. Hopping rates are characterized by $\Delta\omega_{\text{on,off}} = -1, \delta\omega_{\text{on,off}} = 2$. Inset: number of oscillation between subsequent velocity switches $N_\omega = 2\omega_v/\Omega_v$ in the bistable regime. (b) dimensionless time $\tau(k) = 2v(k)/\omega_0\epsilon$ governing the stability of the sustained oscillation (see text) as a function of the dimensionless coupling k for $\lambda = f_0/\bar{\omega}L\gamma = 2 \times 10^{-2}, 3.3 \times 10^{-2}, 4 \times 10^{-2}$ (squares, circles, triangles). Points below the dashed lines are for motors undergoing sustained oscillation, whereas above the dashed line for motors in the bistable regime.

In the opposite regime, when the exchange of molecular motors with the bulk is negligible, $k_{\text{on,off}} = 0$, the total number of motors moving along a filament is conserved. When the evolution of bound $\hat{\rho}(x)$ and weakly bound $\hat{\sigma}(x)$ motors are coupled, Eqs. (6) must be solved consistently. In this regime motors tend to accumulate spatially, leading, in some cases, to large local motor densities. It is known that conservation of the overall number of motors moving along a filament promotes cluster [2] and shock wave formation [21]. The morphological details of these structures are sensitive to excluded volume and short range interactions. However, for binding and unbinding rates sharply peaked at the potential extrema, $\Delta\omega_{\text{on,off}} < \delta\omega_{\text{on,off}}$, the development of regions of high molecular motor density only occurs for large coupling parameters. Hence, we can address the instability of the homogeneous, quiescent molecular motor profile avoiding the development of shock waves. We observe that this quiescent configuration destabilizes above a threshold coupling parameter k_1 , characterized by a Hopf bifurcation, as shown in Fig 2(a) (for $k_1 = 55$). Above k_1 the stable state is characterized by a nonzero mean velocity and an oscillation of frequency ω_v , as shown in Fig. 2(a) [22]. Motor velocity oscillations emerge as a result of the periodic change in the density of the bound motors. While moving under the action of the driving potential, the fluid flow generated by bound motors advects weakly bound motors along. After reaching the bottom of the potential, bound motors cease to move and jump to the diffusive state with rate ω_{off} . This leads to an increase of diffusive motors, hence inducing an overall decrease in the average motor velocity, which relies in the small fraction of bound motors still displacing. Once diffusing motors reach the hopping

region, they bind strongly to the filament at a rate ω_{on} , starting a new cycle. As a result of this alternate correlated motor exchange, both strongly $\hat{\rho}(x)$ and weakly bound $\hat{\delta}(x)$ motor densities develop traveling waves.

As we increase k/λ a second bifurcation is observed, k_2 , above which molecular motors exhibit bistability. In this regime the motor density increases gradually, where the motor states can be switched, leading eventually to an overall motor density that exceeds the maximum occupancy. In order to explore this regime we then redistribute uniformly the motor excess [23].

For the system shown in Fig. 2, above the threshold value $k_2 = 95$ the motor velocity still oscillates with frequency ω_v around a nonvanishing mean velocity, v . However, in contrast to the regime $k_1 < k < k_2$, where the sign of v is fixed, for $k > k_2$ the sign of v changes with frequency Ω_v . For time scales larger than Ω_v^{-1} , the average motor velocity vanishes and a bistable behavior emerges (see Fig. 1 in the Supplemental Material [17]), analogous to the one experimentally observed [4–10,24]. This second transition is captured by the dimensionless time, $\tau(k) = 2v(k)/\omega_0\epsilon$, defined as the ratio between the characteristic hopping time $\frac{1}{2}\omega_0^{-1}$ [25], and the time a particle spends in the region in which the hopping rate is nonvanishing, ϵ , being pushed at a speed $v(k)$ [26]. As shown in Fig. 2(b), when $\tau(k) \lesssim 0.2$ the time needed to jump between the two states is much smaller than the time that a particle spends in crossing the hopping region; therefore, the majority of the motors rebind and the system undergoes sustained oscillations. On the contrary, when $\tau(k) \gtrsim 0.2$ part of the motors in the weakly bound state cannot jump back to the bound state and do not contribute to the next cycle. The loss of active motors affects subsequent oscillations. These effects sum up until the system switches the direction of the average velocity on time scales of the order of Ω_v . The number of oscillations between two subsequent switching events, $N_\omega = 2\omega_v/\Omega_v$, decreases for increasing k , as shown in the inset of Fig. 2(a). For increasing k the two time scales approach, $N_\omega \rightarrow 1$, and the bistability disappears; subsequently, motors remain in a quiescent state.

The bistable state observed is typical of soft, hydrodynamically coupled motors for which the local density of both bound and weakly bound motors can adjust dynamically. This feature is absent both if we disregard weakly bound motor dynamics (see Fig. 1), or for rigidly coupled motors [14], when motor density rearrangements are suppressed. Bistability can be recovered for rigidly coupled motors [14], also when weakly bound motors are in contact with a reservoir (data not shown), only if the hopping dynamics is noisy. However, this mechanism vanishes for large system sizes, for which the noise becomes negligible. On the contrary, for hydrodynamically coupled motors bistability arises by increasing the system size, encoded in k , and persists for a finite range of values of k as shown in Fig. 2(a), regardless of system size.

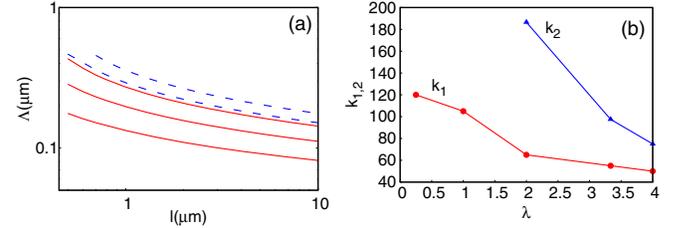


FIG. 3. (a) Minimum system size, Λ , calculated from Eq. (4) for symmetry breaking (solid lines) and for bistable onset (dashed lines) as a function of $l = \eta_{2\text{D},\text{mem}}/\eta_{3\text{D},\text{cyt}}$ for motor pulling on membranes. The membrane-embedded tracer size R is of the order of the membrane thickness ~ 4 nm, leading to $R \sim 1/2L$. Thicker lines stand for larger values of k : $k = 50, 70, 90$ (solid lines) and $k = 95, 110$ (dashed lines), respectively. (b) Values of the coupling parameters k_1, k_2 at which the two bifurcations occur as a function of the motor properties encoded in $\lambda = f_0/(\bar{\omega}L\gamma)$.

The collective phases identified for hydrodynamically coupled motors are controlled by the coupling parameter k that, according to Eqs. (3), (4), depends on the system size Λ . Therefore, the 2D or 3D nature of the HI determines the relation between k and Λ . Assuming a cytoplasm viscosity $\eta_{3\text{D},\text{cyt}} \in [10^{-1}; 10^{-2}]$ Pa s [27] and membrane viscosity $\eta_{2\text{D},\text{mem}} \in [5 \times 10^{-7}, 10^{-8}]$ Pa s m [28], we obtain $l = \eta_{2\text{D},\text{mem}}/\eta_{3\text{D},\text{cyt}} \in [10^{-1}, 10]$ μm that lies within the typical range of biological situations for which the typical velocity is $v \sim 0.1$ μm [29]. The hopping rate can be assumed $\omega_0 \approx 10^2 \alpha \text{ s}^{-1}$, α being the inverse of the efficiency [30]. For these values of parameters we get $\tau \approx 0.2$ that fits in the range of values identified by Fig. 2(b). By inverting Eq. (4) we can calculate the dependence of Λ upon k when the 2D contribution dominates over the 3D. Figure 3(a) shows that for $l \sim 1$ μm , systems as small as $\Lambda \sim 0.5$ μm can undergo hydrodynamically induced symmetry breaking, while slightly larger systems $\Lambda \sim 0.7$ μm develop bistability. For larger values of l the 3D contribution dominates and we expect hydrodynamically induced symmetry breaking for systems of order of 10–100 μm emphasizing the relevance of the 3D HI for larger systems such as neurons, or in technological applications as in microfluidic devices. The onset of symmetry breaking and bistability depends on the parameters governing motors dynamics, namely, the force motor can provide f_0 and their hopping rate $\bar{\omega}$. The ratio of f_0 and $\bar{\omega}$ is encoded in the dimensionless parameter λ . As shown in Fig. 3(b) the values of both k_1 and k_2 decrease upon increasing the strength of the motors or decreasing their hopping rate—the latter being easily controlled in experiments by tuning the ATP concentration.

In conclusion, the HI between bidirectional molecular motors strongly affects their dynamics. For intermediate system sizes, HI triggers the onset of net motor currents [31]. For these regimes the motor velocity has been observed to be oscillatory about a nonvanishing average, or bistable, when motors switch their direction over larger

time scales, leading to an overall vanishing currents, as observed experimentally [5,8–10]. Such features rely on the local variation of the motor density, typical of HI, absent when this degree of freedom is neglected (rigid coupling) or reduced, as happens when the molecular motors are not conserved, e.g., through bath exchange. The typical system sizes over which the soft HI leads to symmetry breaking, or to bistability, are compatible with typical biologically relevant sizes, $\Lambda \in [0.1, 10] \mu\text{m}$, typical for Golgi apparatus displacement [8] or bistable cargo transport [5,9,10].

P. M. acknowledges MINECO for financial support under Project No. FIS 2015-67837-P and Institute Curie for financial support. I. P. acknowledges MINECO and DURSI for financial support under Projects No. FIS 2015-67837-P and No. 2014SGR-922, respectively, and from *Generalitat de Catalunya* under Program *Icrea Acadèmia*.

*Corresponding author.
malgaretti@is.mpg.de

- [1] J. Howard, *Mechanics of Motor Proteins and Cytoskeleton* (Sinauer, Sunderland, 2001).
- [2] P. Magaretti, I. Pagonabarraga, and D. Frenkel, *Phys. Rev. Lett.* **109**, 168101 (2012).
- [3] S. Nelson, K. Trybus, and D. Warshaw, *Proc. Natl. Acad. Sci. U.S.A.* **111**, E3986 (2014).
- [4] S. Ma and R. L. Chisholm, *J. Cell Sci.* **115**, 1453 (2002).
- [5] M. Yusuf Ali, G. G. Kennedy, D. Safer, M. Trybus, H. Lee Sweeney, and D. M. Warshaw, *Proc. Natl. Acad. Sci. U.S.A.* **108**, E535 (2011).
- [6] A. Kunwar, S. K. Tripathy, J. Xu, M. K. Mattson, P. Anand, R. Sigua, M. Vershinin, R. J. McKenney, C. C. Yu, A. Mogliner *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* **108**, 18960 (2011).
- [7] M. Badoual, F. Jülicher, and J. Prost, *Proc. Natl. Acad. Sci. U.S.A.* **99**, 6696 (2002).
- [8] A. Nebenführ, L. A. Gallagher, T. G. Dunahay, J. A. Frohlick, A. M. Mazurkiewicz, J. B. Meehl, and L. A. Staehelin, *Plant Physiol.* **121**, 1127 (1999).
- [9] A. Hendricks, E. Perlson, J. L. Ross, H. W. Schroeder III, M. Tokito, and E. Holzbaur, *Curr. Biol.* **20**, 697 (2010).
- [10] Y. Goldman and E. Holzbaur, *Curr. Opin. Cell Biol.* **22**, 4 (2010).
- [11] F. Jülicher and J. Prost, *Phys. Rev. Lett.* **75**, 2618 (1995).
- [12] F. Jülicher and J. Prost, *Phys. Rev. Lett.* **78**, 4510 (1997).
- [13] S. Klumpp, R. Lipowsky, and M. J. I. Mu, *Biophys. J.* **98**, 2610 (2010).
- [14] T. Guérin, J. Prost, and J.-F. Joanny, *Phys. Rev. Lett.* **106**, 068101 (2011).
- [15] The Oseen approximation is valid for dilute systems and constitutes a lower limit for denser systems (see Supplemental Material [17]).
- [16] Equations (6) are obtained via a mean-field approximation (See Supplemental Material [17]).
- [17] See Supplemental Material at <http://link.aps.org/supplemental/10.1103/PhysRevLett.119.168101> for detailed discussion concerning the use of the Oseen tensor, we provide the derivation of Eqs. (6), (8), and (9) of the main text and we provide an additional figure showing the bidirectional motion.
- [18] Since bound and weakly bound species correspond to different states of the same motor, their sum cannot exceed a maximum density, ζ_{\max} , corresponding to complete local occupancy. Accordingly, for nondilute regimes, a global factor on the right-hand side of Eq. (6) must be included to ensure such a constraint.
- [19] The rigid coupling regime is obtained by solving Eq. (8) disregarding the local contribution, $\tilde{f}(x)$.
- [20] Eq. (9) constitutes only a sufficient and not necessary condition since it has been obtained imposing that the dynamic system of linearized perturbations around the quiescent state has at least three real solutions without constraining their signs (see Supplemental Material [17]).
- [21] Y. Aghababaie, G. I. Menon, and M. Plischke, *Phys. Rev. E* **59**, 2578 (1999).
- [22] For peaked hopping rates, the relevant time scale determining the oscillations is controlled by the hopping rates ω_0 , rather than by their average values $\bar{\omega}$.
- [23] When exceeding the maximum value, $\hat{\rho}_{\max}$, the excess of motors is equally redistributed along the period.
- [24] T. Guérin, J. Prost, P. Martin, and J.-F. Joanny, *Curr. Opin. Cell Biol.* **22**, 14 (2010).
- [25] The prefactor $\frac{1}{2}$ accounts for the hopping rate variable shape.
- [26] $v(k)$ is defined as the average velocity, $\langle v(k) \rangle_{x,\hat{\rho},\bar{\omega}}$, for $k < k_2$, whereas $v(k) = \sqrt{\langle v^2(k) \rangle_{x,\hat{\rho},\bar{\omega}}}$ for $k > k_2$, i.e., when $\langle v(k) \rangle_{x,\hat{\rho},\bar{\omega}} = 0$.
- [27] C. Margraves, K. Kihm, S. Y. Yoon, S. Choi, C. K. Lee, J. Liggett, and S. J. Baek, *Biotechnol. Bioeng.* **108**, 2504 (2011).
- [28] P. Cicuta, S. L. Keller, and S. L. Veatch, *J. Phys. Chem. B* **111**, 3328 (2007).
- [29] O. Campas, C. Leduc, P. Bassereau, J. Casademunt, J.-F. Joanny, and J. Prost, *Biophys. J.* **94**, 5009 (2008).
- [30] ω_0 has been calculated assuming instantaneous hopping and motors velocity $v = 1 \mu\text{m/s}$. The time required to move one period $L \sim 10 \text{ nm}$ is $\tau = 10^{-2} \text{ s}$ for 100% efficiency [11]. For smaller efficiencies we estimate $\tau = 1/\alpha 10^{-2} \text{ s}$.
- [31] Such a coupling is amplified for nonprocessive motors that are more prone to show collective effects due to their low duty ratio [32].
- [32] M. Bloemink and M. Geeves, *Semin. Cell Dev. Biol.* **22**, 961 (2011).