Filament-motor protein system under loading: instability and limit cycle oscillations

Amir Shee, 1, 2, * Subhadip Ghosh, 3, † and Debasish Chaudhuri 1, 2, ‡

¹Institute of Physics, Sachivalaya Marg, Bhubaneswar 751005, India ²Homi Bhaba National Institute, Anushaktigar, Mumbai 400094, India ³Department of Physics, Faculty of Science, University of Zagreb, Bijeniĉka cesta 32, 10000 Zagreb, Croatia (Dated: August 18, 2021)

We consider the dynamics of a rigid filament in a motor protein assay under external loading. The motor proteins are modeled as active harmonic linkers with tail ends immobilized on a substrate. Their heads attach to the filament stochastically to extend along it, resulting in a force on the filament, before detaching. The rate of extension and detachment are load dependent. Here we formulate and characterize the governing dynamics in the mean field approximation using linear stability analysis, and direct numerical simulations of the motor proteins and filament. Under constant loading, the system shows transition from a stable configuration to instability towards detachment of the filament from motor proteins. Under elastic loading, we find emergence of stable limit cycle oscillations via a supercritical Hopf bifurcation with change in activity and the number of motor proteins. Numerical simulations of the system for large number of motor proteins show good agreement with the mean field predictions.

I. INTRODUCTION

Cytoskeletal filaments and associated motor proteins (MP) stabilize structure of the cell and determine its dynamics [1, 2]. The cross-linking MPs, while extending towards one end of the polar filament, can shear filament pairs against each other, in an active non-equilibrium process hydrolyzing ATP. Within a living cell, the filaments form a meshwork, in which each filament encounters forces due to its surrounding [3–5]. While a major contribution to this force comes from active processes [6], recent studies showed that entropic effects like that of depletion, and diffusible passive cross-linkers can lead to significant sliding forces on overlapping filaments [7–10].

The gliding motion of filaments on a motor protein assay has been used extensively to study dynamics of cytoskeletal filaments outside the living cell. The competition between opposing groups of MPs can lead to spontaneous oscillations in gliding assays [11]. Filament motion under cooperative MPs and position dependent load that could arise from passive cross-linkers or harmonic trap showed emergence of stable limit cycle oscillations [10, 12–14]. Similar spontaneous oscillations have been observed in many contexts in cell biology [15, 16], e.g., sarcomere oscillations, mitotic spindle oscillations, and chromosome oscillations [17–20].

Using mean field theory and stochastic simulations, we consider the motion of a filament in a gliding assay of MPs, in the presence of an external force. As has been shown recently, the depletion potential in filament bundles can change from a linear to harmonic form with increase in filament number [7]. We consider an external force that could be constant or be a function of filament

*Electronic address: amir@iopb.res.in †Electronic address: sghosh@phy.hr ‡Electronic address: debc@iopb.res.in position. Under a constant load, the filament on MP assay shows a dynamical crossover from stable to unstable phase. Whereas, in the presence of an elastic loading, the filament shows stable limit cycle oscillations when the number of MPs is larger than a critical value [12, 18]. We show how the onset of spontaneous oscillations depends on the MP activity in terms of its extension rate and detachment force, which can be tuned, e.g., by changing ATP concentration [21–23].

We present a linear stability analysis of mean field equations to find phase diagrams showing linearly stable and unstable phases, separated from stable and unstable spirals. This is compared them with numerical solutions of the non-linear equations. The boundary between the unstable spiral and linear instability disappears once nonlinearities are considered, and the whole region shows stable limit cycle oscillations. We show how the critical number of MPs required for the onset of spontaneous oscillations depend on the stiffness of the elastic load acting on the filament, a property that might be utilized by cells to sense stiffness of extra-cellular matrix. The mean field phase diagrams constitute our first main result.

We present a derivation of the mean field equation for bound MPs using a Fokker-Planck approach, to identify the limitations in the approximation. However, a full numerical simulation of the model, distinguishing the individual MPs and incorporating the stochastic nature of the dynamics, shows good agreement with the mean field prediction of the supercritical Hopf bifurcation boundary identifying the onset of spontaneous oscillations. This is our second main result. Deep inside the oscillatory phase, the dynamics shows a behavior typical of relaxation oscillators. The predictions of stable oscillations obtained from the mean field equations show good agreement with the stochastic simulations. In our numerical calculations, we use parameter values corresponding to microtubule and kinesin motor proteins, allowing our predictions amenable to direct experimental verification.

In Sec. II we present our model. The linear stabil-

ity analysis and its comparison with solutions of nonlinear mean field equations are presented in Sec. III A. In Sec. III B we present the derivation of mean number of attached MPs using the Fokker-Planck approach. We also present the evolution of probability density of attached and detached fraction of MPs in this section. Results of numerical simulations of the detailed stochastic model and their comparison against the Fokker-Planck mean field approach is presented in Sec. III C. Finally we conclude summarizing the main results in Sec. IV.

II. MODEL

We consider a gliding assay set up (Fig.1) in which the tail end of the MPs are attached irreversibly to a cover slip. The MPs are assumed to be active harmonic linkers having stiffness k_m . The head end of MPs can attach to a segment of rigid filament floating on the assay within a cutoff range r_c with a rate ω_a in a diffusion limited manner. The maximum number of MPs that can attach to the filament of length L is $N = L \phi_{MP}$, where ϕ_{MP} is the linear density of MPs attached to the substrate. The attached head of each MP extends along the filament in a directed fashion, from negative to positive end of the filament. This active extension requires energy consumption from ATP hydrolysis that brings the system out of equilibrium. The rate of extension in i-th MP is denoted by an active velocity \boldsymbol{v}_m^i that depends on the load force $f_l^i = k_m y^i$ exerted on the MP due to the extension y^i itself. We consider a piece-wise linear relation [11, 24]

$$v_m^i(f_l^i) = \begin{cases} v_0 & \text{for } f_l^i \le 0\\ v_0 \left(1 - \frac{f_l^i}{f_s}\right) & \text{for } 0 < f_l^i \le f_b, f_b > f_s \\ -v_{back} & \text{for } f_l^i > f_b \end{cases}$$
(1)

where f_s denotes the stall force and v_0 stands for the intrinsic MP velocity. For a load force beyond stall, $f_l \geq f_b > f_s$, the velocity saturates to an extremely small negative value v_{back} [11, 24], while supportive loads do not affect the intrinsic MP motion. Assuming the MPs to be forming slip bonds, the load dependent detachment rate is expressed as $\omega_{\text{off}} = \omega_d \exp(|f_l^i|/f_d)$. The attachment detachment ratio breaks detailed balance.

All the parameters v_0 , ω_a , ω_d , f_s , and f_d characterizing MPs are potentially functions of the ATP concentration in the ambient fluid. An assumption of Michaelis-Menten kinetics of ATP hydrolysis has been used to describe the ATP dependence of v_0 for kinesin, where v_0 increases linearly for small ATP concentrations to eventually saturate [22, 25]. Previous analysis of kinesin runlengths demonstrated the ATP dependence of f_d [10, 22]. A change in v_0 leads to various interesting dynamical regimes. We return to this point later in the paper.

The over-damped dynamics of the filament position x is determined by the mechanical force balance,

$$\gamma_f \dot{x} = F_m + F_e \tag{2}$$

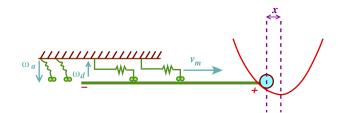


FIG. 1: (color online) Schematic diagram of the model where a motile MT filament is attached with a harmonic trap of stiffness constant K_T . When attached, kinesin walks along the filament towards right with a velocity v_m , pulling the filament towards left.

where the left hand side corresponds to the friction force characterized by γ_f and associated to the relative motion of the filament $\dot{x} := dx/dt$ with respect to the substrate. The n_m number of attached motor proteins exert a total force $F_m = -\sum_{i=1}^{n_m} f_i^i$, and F_e denotes the external loading that acts against the drive of the MPs. The filament motion can in turn drag the attached MPs along with it, such that the extension of *i*-th MP is given by

$$\dot{y}^i = v_m^i(f_l^i) + \dot{x}. \tag{3}$$

III. RESULTS

In this section we first present a mean field description of the model presented above. We utilize it to obtain linear stability predictions for dynamical phases and phase transitions in the presence of external loading on the filament. Numerical solutions of the non-linear mean field equations are used to compare with the linear stability results. In the second part, we present a Fokker-Planck description and derive the mean field equations to discuss its limitations. Finally, we perform detailed numerical simulations of the full stochastic model, and compare the results with mean field predictions.

A. Mean field theory

We assume all the MPs to be equivalent within the mean field approximation, and describe them using the same average extension $y = (1/n_m) \sum_{i=1}^{n_m} y^i$, where n_m denotes the number of attached MPs. To express the equations in a dimensionless form we use the energy scale set by $k_B T$, the time scale ω_d^{-1} , and the length scale $l_0 = \sqrt{k_B T/\gamma_f \omega_d}$. The unit of force is set by $f = \sqrt{k_B T \gamma_f \omega_d}$. Within mean field approximation, the dynamics is described by three coupled non-linear differential equations for dimensionless forms of filament position $\tilde{x} = x/l_0$, mean extension of MPs $\tilde{y} = y/l_0$, and the

attached fraction of MPs $\tilde{n}_m = n_m/N$,

$$\frac{d\tilde{x}}{d\tau} = \tilde{F}_e - N\tilde{n}_m \tilde{k}_m \tilde{y},$$

$$\frac{d\tilde{y}}{d\tau} = \tilde{v}_0 \left(1 - \frac{\tilde{k}_m \tilde{y}}{\tilde{f}_s} \right) + \frac{d\tilde{x}}{d\tau},$$

$$\frac{d\tilde{n}_m}{d\tau} = (1 - \tilde{n}_m)\tilde{\omega} - \tilde{n}_m \exp\left[\frac{\tilde{k}_m \tilde{y}}{\tilde{f}_d} \right].$$
(4)

In the above equations we used the dimensionless time $\tau = t\omega_d$, spring constant of MPs $\tilde{k}_m = k_m l_0/f$, attachment ratio $\tilde{\omega} = \omega_a/\omega_d$, stall force $\tilde{f}_s = f_s/f$, and detachment force $\tilde{f}_d = f_d/f$. In the presence of an external load acting against directed MPs, the mean MP extension remains positive. This allows us to express the mean detachment rate as $\omega_{\rm off} = \omega_d \exp(k_m y/f_d)$. We return to this point in Sec. III B.

In numerical estimates throughout this paper, we use parameter values typical of microtubule-kinesin assays shown in Table-I. These values set the unit of length $l_0 = \sqrt{k_B T/\gamma_f \omega_d} = 33$ nm, force $f = k_B T/l_0 = 0.125$ pN, and velocity $v = l_0 \omega_d = 33$ nm/s. In the following, we first perform a linear stability analysis of Eq.(4) using a constant loading \tilde{F}_e .

TABLE I: Parameters: Two values of v_0 and f_d correspond to ATP concentrations of $5\,\mu\mathrm{M}$ and $2\,\mathrm{mM}$ respectively.

active velocity	v_0	$0.006, 0.8 \mu \text{m/s} [22, 26]$
stall force	f_s	7.5 pN [24, 26]
back velocity	v_{back}	$0.02 \mu \text{m/s} [24]$
detachment force	f_d	1.8, 2.4 pN [10]
attachment rate	ω_a	5, 20/s [22, 27, 28]
detachment rate	ω_d	1/s [26]
motor stiffness	k_m	1.7 pN/nm [10] ^a , 0.3 pN/nm [30]
MT viscous friction	γ_f	$893 \ k_B \text{T-s}/\mu \text{m}^2 \ [9]$

^asame order of magnitude as in Ref. [18, 29]

1. Constant loading

The fixed points of the system of equations is obtained by setting all the time derivatives of Eq.(4) to zero to obtain.

$$\tilde{k}_m \tilde{y}_0 = \tilde{f}_s, \ \tilde{n}_m^0 = \tilde{\omega}/[\tilde{\omega} + \exp(\tilde{f}_s/\tilde{f}_d)] = \tilde{F}_e/N\tilde{f}_s, \quad (5)$$

The last relation determines the loading \tilde{F}_e corresponding to the fixed point of a system of N MPs characterized by attachment ratio $\tilde{\omega}$, stall force \tilde{f}_s and detachment force \tilde{f}_d . In the absence of position dependence of the external force, the fixed point is \tilde{x} - independent. Its stability can be analyzed considering the evolution $d|\psi\rangle/d\tau = a |\psi\rangle$ of a small perturbation $|\psi\rangle = (\delta \tilde{x}, \delta \tilde{y}, \delta \tilde{n}_m)$, where

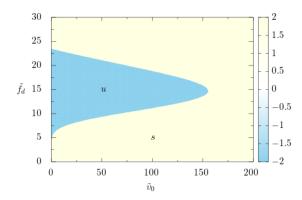


FIG. 2: (color online) Phase diagram under constant loading in $\tilde{v}_0 - \tilde{f}_d$ plane shown using the heat map of the quantity q defined in Eq.(6). We use parameter values typical of a microtubule- kinesin system, $\tilde{k}_m = 450 \, (k_m = 1.7 \, \mathrm{pN/nm})$, $\tilde{f}_s = 60$, $\tilde{\omega} = 20$ and use N = 5 number of MPs. The color-box shows the mapping of q-values to the color code. The light blue (yellow) region denotes unstable (stable) phase under perturbation.

a denotes the stability matrix with elements

$$a_{11} = 0, a_{12} = -\tilde{k}_m N \tilde{n}_m^0, a_{13} = -\tilde{f}_s N, a_{21} = 0,$$

$$a_{22} = -\left(\tilde{\mu} + \tilde{k}_m N \tilde{n}_m^0\right), a_{23} = a_{13}, a_{31} = 0,$$

$$a_{32} = -(\tilde{k}_m / \tilde{f}_d) \tilde{\omega} (1 - \tilde{n}_m^0), a_{33} = -\tilde{\omega} / \tilde{n}_m^0,$$

where $\tilde{\mu} := \tilde{v}_0/\tilde{y}_0$. Diagonalizing the stability matrix \boldsymbol{a} gives the characteristic equation $\lambda(\lambda^2 + p\lambda + q) = 0$ with solutions $\lambda_1 = 0$, and the other two eigenvalues given by $\lambda_{\pm} = (1/2)[-p \pm \sqrt{p^2 - 4q}]$. Here $p = -(a_{22} + a_{33}) > 0$ always, and

$$q = (a_{22}a_{33} - a_{13}a_{32}) (6)$$

may change sign, thereby controlling the stability of the fixed point. In this case, the smaller eigenvalue λ_{-} remains negative, however, λ_{+} can become positive when q changes sign from positive to negative. Thus q = 0line denotes the boundary between stable and unstable fixed points. This phase boundary is shown in Fig. 2. In the stable phase, the force balance is maintained as the filament moves in a direction opposite to the extension of MPs. In the unstable phase, the extension of MPs cannot stabilize the filament position, which slides in the direction of extension of the attached MPs. However, since the discriminant $p^2 - 4q = (a_{22} - a_{33})^2 + 4a_{13}a_{32} > 0$, the quadratic equation does not support any imaginary part in the eigenvalues. Thus oscillatory behavior, stable or unstable, is ruled out. Under constant external loading the filament driven by MPs cannot sustain oscillations.

2. Elastic loading

For elastic loading $\tilde{F}_e = -\tilde{K}_T \tilde{x}$. This might be generated on the filament by trapping one of its ends by a laser tweezer or atomic force microscope tip. As in the previous case, we perform a linear stability analysis. The fixed points of the mean field dynamics is given by

$$\tilde{k}_m \tilde{y}_0 = \tilde{f}_s, \ \tilde{n}_m^0 = \tilde{\omega}/[\tilde{\omega} + \exp(\tilde{f}_s/\tilde{f}_d)], \ \tilde{x}_0 = -\frac{\tilde{n}_m^0 N \tilde{f}_s}{\tilde{K}_T}.$$

The small perturbations around the fixed point $|\psi\rangle = (\delta \tilde{x}, \delta \tilde{y}, \delta \tilde{n}_m)$ evolves with $d|\psi\rangle/d\tau = a' |\psi\rangle$ where the elements of the stability matrix a' are given by

$$\begin{aligned} a'_{11} &= -\tilde{K}_T, a'_{12} = -\tilde{k}_m \tilde{n}_m^0 N, \\ a'_{13} &= -\tilde{f}_s N, a'_{21} = a'_{11}, \\ a'_{22} &= -(\tilde{\mu} + \tilde{k}_m N \tilde{n}_m^0), a'_{23} = a'_{13}, a'_{31} = 0, \\ a'_{32} &= -(\tilde{k}_m / \tilde{f}_d) \tilde{\omega} (1 - \tilde{n}_m^0), a'_{33} = -\tilde{\omega} / \tilde{n}_m^0, \end{aligned}$$
(7)

where, as before, $\tilde{\mu} := \tilde{v}_0/\tilde{y}_0$. In the case of constant loading, the matrix element a_{11} was zero, reducing one eigenvalue to zero. The other two eigenvalues were determined by a quadratic equation. However, for elastic loading, $a'_{11} \neq 0$ and the eigenvalues are given by the full cubic equation

$$\lambda^3 + A\lambda^2 + B\lambda + C = 0 \tag{8}$$

In terms of different matrix elements $A = -\text{Tr}(\boldsymbol{a'})$, $B = \frac{1}{2} (a_{ii}a_{jj} - a_{ij}a_{ji})$ where we implied summation over repeated indices, and $C = -\text{det}(\boldsymbol{a'})$. They can be expressed as

$$A = \tilde{\mu} + \tilde{K}_T + \tilde{\omega}/\tilde{n}_m^0 + \tilde{n}_m^0 \tilde{k}_m N$$

$$B = \tilde{\mu}\tilde{K}_T + \frac{\tilde{\omega}}{\tilde{n}_m^0} (\tilde{\mu} + \tilde{K}_T) + \tilde{k}_m \tilde{\omega} \left[1 - \frac{\tilde{f}_s}{\tilde{f}_d} (1 - \tilde{n}_m^0) \right] N$$

$$C = \tilde{\mu}\tilde{K}_T \tilde{\omega}/\tilde{n}_m^0. \tag{9}$$

Properties of these coefficients determine the existence of different phases and the dynamical behavior of the system. A cubic polynomial has eight possible combinations of real and complex roots. Eq.(9) shows that A is positive definite, C is positive semi definite, while B can change its sign. These strong restrictions eliminate four combinations for roots to the cubic polynomial $\lambda_{1,2,3}$. The remaining four combinations characterize the four different phases in the system. The possible combinations are as follows: (i) All three eigenvalues $\lambda_{1,2,3}$ are real negative, characterizing a linearly stable (s) phase. (ii) λ_1 is real negative, but $\lambda_{2,3}$ are real positive characterizing a linearly unstable (u) phase. (iii) λ_1 is real negative. On the other hand $\lambda_{2,3}$ are complex conjugate pairs with negative real part, $\lambda_{2,3} = -\alpha \pm i\beta$, characterizing a decaying oscillation of perturbations in stable spiral (ss) phase. (iv) λ_1 is real negative though $\lambda_{2,3}$ are complex conjugates with positive real part. $\lambda_{2,3} = \alpha \pm i\beta$

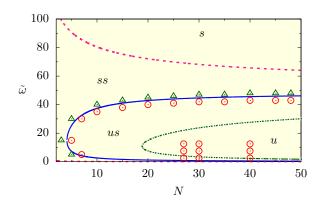


FIG. 3: Phase diagram for filament in MP assay under elastic loading in $N-\tilde{\omega}$ plane, with $\tilde{k}_m=450,\ \tilde{v}_0=24.24,\ \tilde{f}_s=60,\ f_d=19.2,\ \tilde{K}_T=83$ kept fixed. The lines denote the linear stability phase boundaries between linearly stable (s), stable spiral (ss), unstable spiral (us), and linearly unstable (u) phases. The points denoted by Δ and \bigcirc indicate decaying oscillations and limit cycle oscillations, respectively, corresponding to the full non-linear dynamics in Eq.(4).

with $\alpha > 0$ denotes oscillations with growing amplitude in unstable spiral (us) phase.

Phase transitions: (a) Phase boundary between linearly (un) stable and (un) stable spiral phase: The complex conjugate roots disappear as the minimum of the polynomial $p(\lambda) = \lambda^3 + A\lambda^2 + B\lambda$ touches the line $p(\lambda) = -C$, corresponding to two degenerate eigenvalues. The minimum of $p(\lambda)$ is at $\lambda_m = -\frac{A}{3} + \frac{1}{3}\sqrt{A^2 - 3B}$. This condition lead to the phase boundary,

$$C = \left[\frac{A}{3} + \frac{2}{3}\sqrt{A^2 - 3B}\right] \left[-\frac{A}{3} + \frac{1}{3}\sqrt{A^2 - 3B} \right]^2 \tag{10}$$

For $B \geq 0$, the boundary is between linear stable (s) and stable spiral (ss) phases. On the the hand, for B < 0, it denotes the boundary between unstable spiral (us) and linearly unstable (u) phases.

(b) Phase boundary between stable spiral and unstable spiral phases: As the sign of the real part α of complex conjugate roots $\lambda_{2,3} = \alpha \pm i\beta$ changes from negative to positive the system becomes unstable and start to oscillate. This transition is captured by setting $\alpha = 0$. This leads to the condition

$$C - AB = 0 \tag{11}$$

The growing amplitudes of oscillations in us phase predicted by linear stability analysis gets stabilized by nonlinearities into stable limit cycle oscillations, as we show later in Sec. IIIB using full stochastic simulations and further analysis. This denotes a supercritical Hopf bifurcation to a stable limit cycle, e.g., at a critical number of MPs N^* (Fig.3). The expression of N^* can be obtained by solving Eq.(11), a quadratic equation in terms of N^* . Note that in the absence of elastic loading $K_T = 0$ the

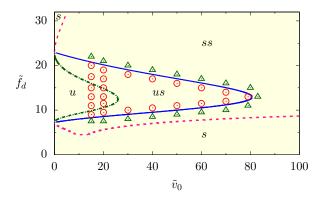


FIG. 4: Phase diagram for filament in MP assay under harmonic trap in $\tilde{v}_0 - \tilde{f}_d$ plane, keeping $\tilde{k}_m = 450$, $\tilde{\omega} = 20$, $\tilde{f}_s = 60$, N = 5 and $\tilde{K}_T = 83$ fixed. The phases and phase boundaries are indicated in the same manner as in Fig.3.

polynomial coefficient C=0. As a result Eq.(11) cannot be satisfied, and the Hopf bifurcation disappears.

At the supercritical Hopf bifurcation, the imaginary part of the eigenvalue $\beta = \sqrt{B}$, so that the MP- filament system shows oscillations with a frequency $f_{\omega} = \sqrt{B}/2\pi$. Clearly, the frequency of oscillations f_{ω} depends on the number of MPs, ATP-dependent activity of MPs determined by active velocity, attachment detachment rates, stall force, and detachment force, apart from the effective elastic constant of the external loading force.

Phase diagrams: Using Eq.(10) and Eq.(11) we present two phase diagrams showing the transitions between the above mentioned phases in Fig. 3, and 4. We use parameter values corresponding to kinesin-microtubule assays (Table-I).

In Fig. 3, we show the phase diagram in number of MPs N- and attachment ratio $\tilde{\omega}$ plane. This indicates the requirement of a threshold number of MPs N^* to get sustained oscillations in the us phase. In addition, sustained oscillations depend on the activity in the system, parametrized in terms of attachment-detachment ratio $\tilde{\omega}$, active velocity \tilde{v}_0 and the detachment force f_d . The lines in the plot show phase boundaries obtained from linear stability analysis, using Eq.s (10) and (11). The boundary between ss and us phase appears via a supercritical Hopf bifurcation. The full non-linear dynamics corresponding to Eq.(4) show decaying oscillations in ss phase (\triangle) , and limit cycle oscillations (\bigcirc) in the region denoted by us and u. Clearly, once the non-linearities are considered, the boundary between us and u phase become irrelevant, the whole region inside the us boundary shows limit cycle oscillations.

In Fig. 4 we further characterize these dynamical phase transitions in terms the of detachment force \tilde{f}_d and active velocity \tilde{v}_0 . As before, the linear stability analysis shows boundaries between s, ss, us and u phases. The consideration of non-linearities show that the whole region of us and u display limit cycle oscillations. The stable limit

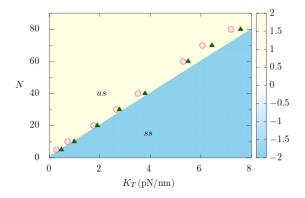


FIG. 5: The linear stability phase boundary between the stable spiral (blue: ss) and unstable spiral (yellow: us) phase in the plane of elastic loading stiffness K_T and MP number N is shown using the heat map of C-AB in Eq.(11). The color box shows the mapping for the values of the function. Parameters used correspond to kinesin-microtubule assay, keeping $\tilde{k}_m = 450$, $\tilde{\omega} = 20$, $\tilde{f}_s = 60$, $\tilde{f}_d = 19.2$, $\tilde{v}_0 = 24.24$ fixed. The points denoted by Δ and \odot indicate decaying oscillations (ss) and stable limit cycle oscillations (us), respectively, corresponding to the full non-linear dynamics shown in Eq.(4). Here we express K_T in units of pN/nm.

cycle phase appears from stable spiral via a supercritical Hopf bifurcation. This transition will be explored in further detail in Sec. III C using numerical simulations of the stochastic dynamics governing the MPs and filament. Note that the phase boundaries between ss and s in Fig.s 3 and 4 are inconsequential, as both the phases are stable in long time limit.

In Fig. 5, we show how the onset of stable limit cycle oscillations (us) depends on the number of MPs N recruited for a given rigidity K_T of the elastic loading. The plot uses parameter values corresponding to microtubulekinesin MP assay, at an ATP concentration of 2 mM. The minimum number of MPs required for the onset of spontaneous oscillations increases with the stiffness K_T of the substrate. While the particular calculations are performed for microtubule-kinesin system, the physical mechanism is equally applicable for acto-myosin systems. Our simple setup has a parallel in the rigidity sensing by cells, where contractile acto-myosin system couples to the extra-cellular matrix (ECM) via an adhesion complex consisting of alpha-actinin and integrin [31, 32]. The range of K_T values used in Fig. 5 belongs to the range of rigidities of sub-micron elastomeric pillars used in cell spreading experiments [32]. The cell may utilize an increase of processive myosin bundles, required for the onset of oscillations (tugging), as a strategy to sense the ECM stiffness [32, 33]. In fact, larger multifilament assemblies of myosin is noted near more rigid substrate [32].

The parameter values used in the above phase diagrams correspond to a gliding assay of microtubule on kinesin MPs (Table-I). The elastic loading on the filament can be applied by optical tweezers or atomic force

microscopes [12, 30, 34]. In Fig. 3 and 4, we used a value of dimensionless stiffness \tilde{K}_T that corresponds to 0.3 pN/nm. While the location of phase boundaries depends on \tilde{K}_T , the qualitative features remain unaltered. In the limit of extremely small \tilde{K}_T , however, the harmonic trap can act like a constant loading, as was shown in Ref. [35]. Within the cell, our study has relevance for the relative sliding motion of filaments where the loading might be provided by other cellular components, or the filament bending [18], and may have implications for rigidity sensing by cells [31, 32]. We considered length stabilized filaments, which are typically used in gliding assay experiments, thus disregarding the possible effects of active polymerization depolymerization of filaments in living cells [2].

B. Fokker-Planck approach to mean field

Having established the phase diagrams using mean field equations and linear stability analysis, in this section we use a Fokker-Planck approach [18] involving the probability distributions $P_{a,d}(y,t)$ of attached and detached fractions of MPs to derive and analyze the mean field equations. The distribution functions obey the normalization $\int_{-\infty}^{\infty} (P_a + P_d) dy = 1$, and evolve as

$$\partial_t P_a + \partial_y J_a = \omega_a P_d - \omega_d P_a$$

$$\partial_t P_d + \partial_y J_d = -\omega_a P_d + \omega_d P_a,$$
 (12)

where, the probability currents

$$J_a = \dot{y} P_a(y,t) - D_a \,\partial_y P_a(y,t)$$

$$J_d = -\nu y P_d(y,t) - D_d \,\partial_y P_d(y,t). \tag{13}$$

Here D_a , D_d are the diffusion coefficient of attached and detached MPs, respectively, and ν is the relaxation rate of the extension for the detached MPs.

As the typical relaxation rate is much faster than the attachment rate, $\nu \gg \omega_a$, one can assume the detached MPs relaxes immediately to equilibrium distribution,

$$P_d(y,t) = \tilde{n}_d(t) A \exp\left(-\frac{k_m y^2}{2k_B T}\right)$$
 (14)

with normalization $A=(k_m/2\pi k_BT)^{1/2}$. As a result, the fraction of detached MPs $\tilde{n}_d(t)=\int_{-\infty}^{\infty}dy P_d(y,t)$. This is related to the fraction of attached MPs $\tilde{n}_m(t)=\int_{-\infty}^{\infty}dy P_a(y,t)$ via the conservation of total probability $\tilde{n}_m(t)=1-\tilde{n}_d(t)$. Integrating the first equation of Eq.(12) we get

$$\frac{d\tilde{n}_m}{dt} = (1 - \tilde{n}_m)\omega_a - \langle \omega_d(y) \rangle \tilde{n}_m \tag{15}$$

where $\langle \omega_d(y) \rangle := \int_{-\infty}^{\infty} dy P_a(y,t) \omega_d(y) / \int_{-\infty}^{\infty} dy P_a(y,t)$. Using the expression for slip bond $\omega_d(y) = \omega_d \exp(k_m |y|/f_d)$ leads to

$$\frac{d\tilde{n}_m}{dt} = (1 - \tilde{n}_m)\omega_a - \omega_d \langle e^{k_m|y|/f_d} \rangle \tilde{n}_m \tag{16}$$

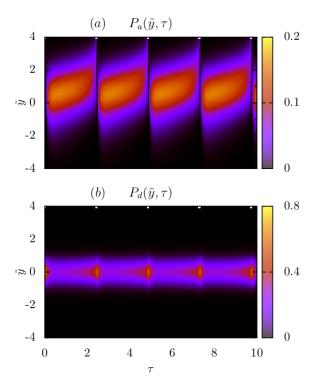


FIG. 6: (color online) Kymographs show the time evolution of the probability distributions of MPs with extension \tilde{y} : (a) in the attached state (P_a) , and (b) in the detached state (P_d) . The color-box describes the values of probability distributions. The relaxation dynamics are determined by dimensionless diffusion constants $\tilde{D}_a = 15.6$, $\tilde{D}_d = 17.4$, and the relaxation rate $\tilde{\nu} = 80$. Other parameter values used are $\tilde{v}_0 = 50$, $\tilde{f}_d = 4$, $\tilde{k}_m = 4.57$, $\tilde{\omega} = 5$, $\tilde{f}_s = 14.29$, N = 160 and $\tilde{K}_T = 4.57$.

By Jensen's inequality $\langle e^{k_m|y|/f_d} \rangle \geq e^{k_m \bar{y}/f_d}$ with $\bar{y} = \langle |y| \rangle$ denoting the mean extension in the attached state. Thus the actual relaxation $d\tilde{n}_m/dt$ is slower than that assumed in Eq.(4).

The active extension and relaxation dynamics in terms of $P_{a,d}(y,t)$ is shown in Fig.6 by numerically integrating the Fokker-Planck equations (Eq.(12) and (13)) along with the evolution of filament position x (Eq.(2)), and mean extension of MPs following

$$\dot{y} = v_m(y) + \dot{x}.\tag{17}$$

In the above equation the piecewise linear form of v_m is used from Eq.(1) replacing the load force $f_l = k_m y$. In Eq.(2), at this point, we use $F_m = -k_m \sum_{i=1}^{n_m} y^i = -k_m \int_{-\infty}^{\infty} dy \, y \, P_a(y,t)$. This reduces Eq.(2) to

$$\gamma_f \dot{x} = -K_T x - k_m \int_{-\infty}^{\infty} dy \, y \, P_a(y, t). \tag{18}$$

In plotting this graph we used $D_d = k_B T/\gamma_f$, $D_a < D_d$, and $\nu = k_m/\gamma_f$. The last choice maintains $\nu \gg \omega_a$.

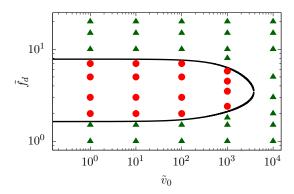


FIG. 7: (color online) Phase diagram for harmonically trapped microtubule-kinesin assay in $\tilde{v}_0 - \tilde{f}_d$ plane, at fixed $\tilde{k}_m = 4.57$, $\tilde{\omega} = 5$, $\tilde{f}_s = 14.29$, N = 160 and $\tilde{K}_T = 4.57$. The points denote the two phases characterized by decaying oscillations (\triangle) and stable limit cycle oscillations (\bigcirc) obtained from numerical simulations. The solid black line identifies the boundary of supercritical Hopf-bifurcation predicted by Eq.(11).

The parameters used correspond to kinesin-microtubule system (Table-I). Here the unit of length is chosen to be $l_0 = 8$ nm, the dimer-size of microtubules [2]. The unit of force is $f = k_B T/l_0 = 0.525$ pN, and time is $\omega_d^{-1} = 1$ s. As before, we express dimensionless extensions $\tilde{y} = y/l_0$, $\tilde{x} = x/l_0$ and dimensionless time $\tau = \omega_d t$.

Due to the directed nature of the MP extension and the loading acting against MPs on an average, the mean extension \tilde{y} of the attached fraction remains positive (Fig.6(a)), a fact used in replacing $\exp(k_m|\tilde{y}|/f_d)$ by $\exp(k_m \tilde{y}/f_d)$ in the mean field description of Eq.(4). As Fig.6(a) shows in terms of the evolution of $P_a(\tilde{y},\tau)$, the mean extension grows slowly up to a maximum, before relaxing back rapidly to zero in a time periodic manner. This is a characteristic of relaxation oscillators [36]. With detachment of one MP, the shared load on other attached MPs increases, increasing the effective rate of detachment. This mediates an avalanche of MP detachment leading to the rapid relaxation. Once relaxed, MPs reattaches, maintaining oscillations. The avalanche in detachments lead to an associated rapid increase in $P_d(\tilde{y},\tau)$ (Fig.6(b)). Note that the distribution $P_d(\tilde{y},\tau)$ maintains a maximum at $\tilde{y}=0$, and is always symmetric around $\tilde{y} = 0$, vindicating the simplification used in Eq.(14).

C. Numerical simulations

Finally, we perform a numerical simulation of the MP-filament model described in Sec. II. We consider a kinesin-microtubule system with the microtubule held in its positive end using a harmonic trap of strength K_T . A similar experimental system of myosin- F-actin with

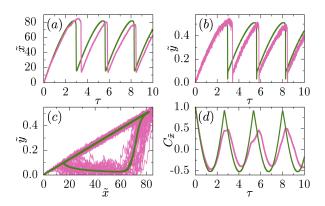


FIG. 8: (color online) Time evolutions of (a) microtubule displacement $\tilde{x}(\tau)$, and (b) mean kinsein extension $\tilde{y}(\tau)$. (c) A parametric plot of $\tilde{x}(\tau)$ and $\tilde{y}(\tau)$ shows stable limit cycle. (d) Correlation function of microtubule displacement $C_{\tilde{x}}(\tau)$ in the time-periodic steady state. The red (green) lines in all these figures correspond to numerical simulations (solutions to the Fokker-Planck based mean field equations). We used $\tilde{v}_0 = 50$, $\tilde{f}_d = 4$. All other parameter values are the same as in Fig. 7.

the F-actin held by an elastic load provided by a laser tweezer was considered before in Ref. [12]. We model the microtubule as a connected rigid string of $\sigma = 8$ nm segments. In this section, we use $l_0 = \sigma$ as the unit of length, which sets the unit of force $f=k_BT/l_0=0.525\,\mathrm{pN}$. We still use the unit of time $\omega_d^{-1}=1\,\mathrm{s}$. The *i*-th kinesin can attach to a microtubule segment within the cutoff radius $r_c/l_0 = 1.0$ stochastically with rate ω_a . When attached, the MP extends towards the plus end of microtubule stochastically with a rate v_m/l_0 . The instantaneous load force on the i-th MP is $f_l^i = k_m y^i$, expressed in terms of the extension $y^{i}(t)$. We use Eq.(1) for the load dependence of the extension rate. The MPs detach from the filament stochastically with the rate $\omega_d \exp(k_m|y^i|/f_d)$. This system along with Eq.s(2) and (3) for the position of the filament x and MP extension y_i are integrated numerically using Euler-Maruyama integration with time steps small enough so that probability of each event stays less than one. We simulate a large number of MPs N = 160.

Performing the numerical simulations over a range of activity \tilde{v}_0 and detachment force \tilde{f}_d we obtain the phase diagram in Fig. 7(a). It shows two phases, one is characterized by decaying oscillations corresponding to stability (\triangle), and the other displays stable limit cycle oscillations (\bigcirc). The phase boundary obtained from numerical simulations of the full stochastic model shows good agreement with analytic result, Eq.(11).

The dynamics deep inside the stable limit cycle phase is illustrated in Fig. 8, using results from numerical simulations. We plot the evolutions of filament position $\tilde{x}(\tau)$ and mean extension of kinesins $\tilde{y}(\tau)$ deep inside the limit cycle phase ($\tilde{v}_0 = 50$, $\tilde{f}_d = 4$). They show anharmonic oscillations with a well defined periodicity (Fig. 8(a), (b)). The slow extension (increase in \tilde{x}) followed by rapid re-

laxation is a typical relaxation oscillatior behavior [36]. Here the rapid relaxation is associated with an avalanche of MP detachments. When presented as a parametric plot in \tilde{x} - \tilde{y} plane, they clearly show a stable limit cycle, although with a spread in the trajectories due to their inherent stochastic nature (Fig. 8(c)). Similar spread has been observed in *in vitro* experiments [12]. The limit cycle oscillations are independent of initial conditions. Using the time series in Fig. 8(a), we obtain the two-time correlation function of the microtubule displacement $C_{\tilde{x}}(\tau) = \langle \tilde{\delta}x(\tau)\tilde{\delta}x(0)\rangle/\langle \tilde{\delta}x^2(0)\rangle$ where $\delta x = x - \langle x\rangle$ is measured in the time-periodic steady state (Fig. 8(d)).

We compare the simulation results with the Fokker Planck description developed in the previous subsection. For that we use the equation for $P_a(\tilde{y},\tau)$ using Eq.s(12), (13) and the expression of $P_d(\tilde{y},\tau)$ from Eq.(14). To determine $\tilde{n}_d(\tau) = 1 - \tilde{n}_m(\tau)$ we use Eq.(16). These equations are solved numerically along with Eq.(17) and (18). The comparisons are displayed in Fig. 8, and show semi-quantitative agreement.

IV. CONCLUSIONS

We have studied dynamics of a cytoskeletal filament in a motor protein (MP) assay under external loading. We used a mean field description along with linear stability analysis to determine various phase boundaries. Under constant loading, the system shows a transition from stable to unstable behavior. We have shown that the overdamped active system under harmonic loading displays an emergence of spontaneous oscillations via a supercritical Hopf bifurcation. In linear stability analysis this appears as a boundary between a stable and unstable spiral phase. The non-linearities makes the boundary between unstable spiral and linear instability irrelevant, with the system showing stable limit cycle oscillations in both of them. The increase in the critical number of MPs required at the onset of stable limit cycle oscillations with increase in the stiffness of elastic loading may be utilized by spreading cells for sensing the stiffness of extra-cellular matrix. Using a Fokker-Planck description, we analyzed

the limitations of the mean field equations used. Finally, we performed numerical simulations involving stochastic dynamics of individual MPs and the filament. The resulting phase diagram shows good agreement with the mean field prediction. While the stochastic dynamics displays characteristic spread of trajectories, they reproduce the limit cycle behavior in an average sense. We obtained semi-quantitative agreement between the mean field prediction for time evolution with stochastic trajectories.

In our numerical analysis we used parameters corresponding to microtubule-kinesin system. While our method is equally applicable to other filament-MP systems, e.g., filamentous actin-myosin, the quantitative results presented here are amenable to direct experimental verifications in gliding assay setups of microtubule and kinesin molecules. The parameter values used in Fig. 8 correspond to kinesin extension rate $0.4 \,\mu\text{m/s}$, and a detachment force 2 pN. We used a trapping potential of strength $K_T = 0.3 \,\mathrm{pN/nm}$, which can be controlled in experiments [12, 30, 34]. The amplitude and frequency of oscillations of the microtubule in the limit cycle phase shown in Fig. 8 correspond to $0.4 \,\mu\mathrm{m}$ and $0.5 \,\mathrm{Hz}$, respectively. On the other hand, for MPs while the frequency remains around 0.5 Hz, the amplitude of oscillation is $\sim 4\,\mathrm{nm}$.

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