

Front speed and pattern selection of a propagating chemical front in an active fluidClara del Junco *Department of Chemistry and Department of Sociology, University of Chicago, Chicago, Illinois 60637, USA
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Spontaneous pattern formation in living systems is driven by reaction-diffusion chemistry and active mechanics. The feedback between chemical and mechanical forces is often essential to robust pattern formation, yet it remains poorly understood in general. In this analytical and numerical paper, we study an experimentally motivated minimal model of coupling between reaction-diffusion and active matter: a propagating front of an autocatalytic and stress-generating species. In the absence of activity, the front is described by the well-studied Kolmogorov, Petrovsky, and Piskunov equation. We find that front propagation is maintained even in active systems, with crucial differences: an extensile stress increases the front speed beyond a critical magnitude of the stress, while a contractile stress has no effect on the front speed but can generate a periodic instability in the high-concentration region behind the front. We expect our results to be useful in interpreting pattern formation in active systems with mechanochemical coupling *in vivo* and *in vitro*.

DOI: [10.1103/PhysRevE.105.014602](https://doi.org/10.1103/PhysRevE.105.014602)**I. INTRODUCTION**

The emergence of form in the living embryo involves a series of out-of-equilibrium chemical and physical processes [1]. Among them, two mechanisms stand out: reaction-diffusion (RD) [2–6], which creates well-defined concentration patterns, and active matter (AM), mediated by molecules that convert chemical energy into mechanical forces generating hydrodynamic flows [7–16]. RD dynamics have also been used to model a wide range of situations in biology ranging from spatiotemporal protein dynamics in cells [17] to the migration of a cell monolayer [18,19]. In these contexts, the species undergoing RD dynamics may regulate or control active forces in the living system. Indeed, the importance of interactions between chemistry and mechanics in modeling living systems has long been recognized [4], yet RD and AM systems have generally been examined separately, whether theoretically [4,8] or in *in vivo* [5,20] and *in vitro* [21,22] experiments. More recently, however, mechanochemical couplings have been investigated theoretically [23,24], demonstrating the ability of such systems to display novel patterns even in the absence of chemical reactions [23]. When combined with chemical reactions, the coupling modifies the patterning behavior of RD systems [23], and it can also generate spatiotemporal oscillations [24]. Their importance for *in vivo* assays has been established [1,7], and chemical

reactions have also been implicated in the active context in spontaneously creating stable droplets of a finite size and the formation of protocells [25]. However, how active flows affect RD dynamics in general remains sparsely understood.

To the best of our knowledge, couplings between RD and AM have not yet been engineered *in vitro*. However, recent experimental developments suggest that mechanochemical self-organization could be investigated in well-controlled *in vitro* experiments in the near future [26]. Such *in vitro* systems, which we will call reaction-diffusion active matter (RDAM) systems, would be particularly interesting for quantitatively testing theoretical predictions concerning the propagation of a reacting and diffusing activity regulator—a situation that is likely to be ubiquitous in *in vivo* settings.

In this paper, we present an examination of how active flows affect the propagation of a reacting and diffusing chemical species that controls the strength of activity, using a highly simplified model. We describe the simplest, nontrivial RDAM system that could be reasonably assembled *in vitro* with current knowledge (Sec. II). It is an RD system generating a traveling front that is coupled to a compressible active fluid. We model it with a dynamical equation for the concentration of the reactant and a constitutive equation for the velocity field of the active fluid [23] (Sec. III). A linear stability analysis of these equations about a homogeneous state shows that there is a spatial instability for a contractile RDAM system

(Sec. IV). However, the linear theory falls short of being able to predict the behavior of the system when symmetry is broken by a traveling front. In particular, a linear theory incorrectly predicts that the front speed is insensitive to the active stresses. Moreover, it does not provide a prediction of how the spatial instability propagates or how its selected wavelength is shifted in the presence of the front. Numerical simulations of the two dynamical equations provide the answers to these questions (Sec. VA). The simulations show that, in extensile fluids, fronts propagate faster as the active stress increases and spatial instabilities are never observed. We show that the front speed picked out in simulations is the minimum physically allowed speed, consistent with a marginal stability mechanism that shows up in much simpler systems, but which does not hold in our highly nonlinear model. In turn, contractile fluids generate fronts whose velocity is independent of the active stress, and they display spatial instabilities whose group velocity is not equal to the speed of the front and whose wave number is shifted to smaller values compared to the linear prediction.

II. IN VITRO REACTION-DIFFUSION ACTIVE MATTER

In this section, we will describe an RDAM system that could be assembled *in vitro* with the current experimental state of the art. We will take into account known experimental constraints to motivate the theoretical description undertaken in the rest of the paper. Although a wide array of AM experimental systems exist, we will restrain ourselves to molecular active matter because we believe that it has the highest chance to be combined with molecular RD systems. In this category, extracts of cytoskeletal filaments and motors, either actomyosin or kinesin/microtubule gels [26,27], are particularly suitable. Depending on the situation, these gels are locally contractile or extensile and are composed of polar, nematic, or isotropic particles [28]. A wide variety of dissipative behaviors have been observed in these gels *in vitro*, such as aster formation [22], global contractions [29,30], chaotic flows [31], active nematics [31], and active film buckling [32]. These systems function in aqueous solution, at neutral pH and room temperature.

Chemical systems that generate RD patterns have historically relied on strongly acidic reactions [21] which are incompatible with cytoskeletal extracts. However, in the past 15 years, biochemical RD systems working at neutral pH have been developed [33–37]. Among them, DNA-only [35] and DNA/enzyme [36] reaction schemes are particularly interesting because the modularity of the DNA hybridization reaction facilitates the engineering of mechanochemical couplings with active fluids. In particular, one may envision that the single-stranded (ssDNA) species A produced by these reactions dimerizes DNA-motor conjugates [38,39], or releases them from a reservoir [40]. Indeed, a number of active fluids become active only in the presence of dimers of motors [22,31], and thus, with this scheme, active stresses would be generated only in the presence of species A .

With such a design, we will consider an autocatalytic reaction of ssDNA species A , which has been reported to generate RD fronts propagating at constant velocity in a thin microchannel [41] and that is compatible with kinesin-microtubule active matter [42]. In the following, we will

suppose that this autocatalytic reaction takes place in a solution that is in contact with a thin film of an active fluid composed of filaments and motor-DNA conjugates. We note that active thin films, as well as motor-DNA conjugates, have been obtained both for myosin [43,44] and kinesin [31,38,45–47]. Such a thin film of DNA-responsive active fluid could be obtained at a water-oil interface in the presence of a depletion agent, such as polyethyleneglycol, that pushes the filaments against that interface [27,43].

III. ACTIVE TRAVELING FRONT MODEL

In the situation just described, the thin film of active fluid can exchange both momentum and matter (i.e., the filaments can diffuse into the fluid above) with the solution where DNA/enzyme reactions take place. At a given point in space, as the concentration of A increases in the bulk solution, motor dimers will form in the film and will generate active flows (Fig. 1). To model this three-variable (DNA A , motors M , and filaments F) biphasic system as a one-variable monophasic system, we will make two sets of assumptions. First, we will suppose that the timescales of motor-motor dimerization via DNA hybridization, $\tau_{M,A}^+$, of motor-motor dissociation, $\tau_{M,A}^-$, of motor binding to filaments, $\tau_{M,F}^+$, and of motor-filament dissociation, $\tau_{M,F}^-$, are fast compared with the timescale of DNA autocatalysis, τ_A . Then, one can model such a system with a single average species C , of concentration c , that could be seen as the concentration of motor dimers *bound to filaments* that undergo an autocatalytic reaction (referred to as autocatalytic motors in the following). Note that the effective diffusion coefficient D of this species is an average of the diffusion coefficient of free A , D_A , and of A

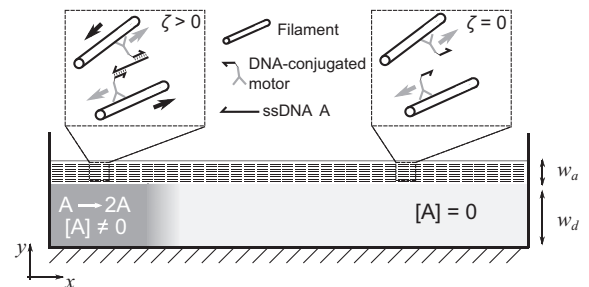


FIG. 1. Scheme of a plausible experimental reaction-diffusion active matter system that can be described using Eqs. (1)–(3). A chamber is filled with a solution containing an RD and an AM subsystem. The RD subsystem is an autocatalytic DNA/enzyme reaction of the form $A \rightarrow 2A$, where A is a ssDNA. The AM subsystem is made of filaments and DNA-conjugated motors. Due to depletion forces, the AM subsystem forms a film of height w_a (segmented pattern) above the RD solution of height w_d . A small quantity of A is introduced on the left side of the chamber (dark gray). In the absence of activity (i.e., $[ATP] = 0$ and $\zeta = 0$), A is expected to generate a traveling front of constant velocity [41]. In the presence of ATP, A will diffuse on the active region, generating motor dimers that will move along the filaments (gray arrows) generating a flow of filaments (black arrows) and thus induce $\zeta > 0$. At the right side of the front, $[A] = 0$, the motors are monomeric and thus they move along the filaments without generating flow $\zeta = 0$.

bound to M , D_M , weighted by the time spent in each state [41]. Second, we will suppose that the thickness of both the active film, w_a , and the DNA/enzyme solution, w_d , is small enough such that the corresponding diffusion time $\tau_D = [\max(w_a, w_d)]^2/2D_A \ll \tau_A$, with D_A the diffusion coefficient of A . As C is advected by active flows in the thin active film, this transport may in turn influence the concentration of C in the bulk by the rapid diffusion of C across both phases.

To estimate $\tau_{M,A}^+$, we suppose that motor-DNA conjugates dimerize through DNA hybridization with second-order kinetics. Thus $\tau_{M,A}^+ = 1/(\kappa_{M,A}^+ M_0)$, where we take $\kappa_{M,A}^+ \sim 10^4\text{--}10^6 \text{ M}^{-1} \text{ s}^{-1}$ as a typical value of DNA association rate [48] and $M_0 \sim 1\text{--}100 \text{ nM}$ the typical concentration of kinesin in active matter experiments [32]. $\tau_{M,A}^- = 1/\kappa_{M,A}^-$, where the dissociation rate of a DNA double strand $\kappa_{M,A}^-$ is controlled by its length and can vary in the range $\kappa_{M,A}^- \sim 10^{-2}\text{--}10^{-5} \text{ s}^{-1}$. $\tau_{M,F}^+ = 1/\kappa_{M,F}^+$, and we take $\kappa_{M,F}^+ = 5 \text{ s}^{-1}$ from Leduc *et al.* [49]. $\tau_{M,A}^- = 1/\kappa_{M,F}^-$, and we calculate $\kappa_{M,F}^- = v/\ell_r$, where v is the velocity of kinesin along microtubules and ℓ_r is the run length before detachment. Furuta *et al.* [46] measured v and ℓ_r for DNA-promoted kinesin clusters with different numbers of kinesins and found $v = 800 \text{ nm/min}$ and $\ell_r = 2 \times 10^3 \text{ nm}$, $4 \times 10^3 \text{ nm}$, and $15 \times 10^3 \text{ nm}$, respectively, for one-, two- or four-kinesin clusters. τ_A has been measured by Zadorin *et al.* [41] to be in the range $10^2\text{--}10^4 \text{ s}$. Finally $D_A \approx D_M = 10^{-10} \text{ m}^2 \text{ s}^{-1}$, and w_a and w_d are in the range $10\text{--}100 \mu\text{m}$. We thus get $\tau_{M,A}^+ \sim 10\text{--}10^5 \text{ s}$, $\tau_{M,A}^- \sim 10^2\text{--}10^5 \text{ s}$, $\tau_{M,F}^+ \sim 0.2 \text{ s}$, $\tau_{M,F}^- \sim 2\text{--}20 \text{ s}$, $\tau_A \sim 10^2\text{--}10^4 \text{ s}$, and $\tau_D \sim 1\text{--}100 \text{ s}$. As a result, provided that the motor-DNA system is designed such that M_0 and $\kappa_{M,A}^-$ are on the upper side of the aforementioned ranges, τ_A is larger than all the other characteristic times as required by our model.

Beyond the specific and explicit *in vitro* scenario described here, the RDAM model we will construct is relevant to a wide range of *in vivo* systems. Indeed, our theory describes all active thin films of uniform thickness h , where the thickness of the film is maintained by the exchange of material with the surrounding medium [50,51], and in which activity is controlled by an activity regulator C , which in the absence of active flows would have the following RD dynamics: an unstable lower concentration state of the regulator is changed to a higher concentration state at a front moving at a well-defined speed. Stated in this general form, our theory may be relevant for understanding the propagation of a cell layer in a wound-healing assay [52], where C models the cells or for the dynamics of a chemical active stress regulator in the cellular cortex.

Having established the experimental relevance of an RD system coupled to an active flow, we now construct a model for understanding two key features of such a system: (i) whether active flows enhance or inhibit the speed of the RD front, and (ii) whether the profile of the component C behind the front remains homogeneous or forms a patterned state. For this, we consider a system that is perfectly homogeneous in \hat{y} and has a moving front of C along \hat{x} at which a low (zero) concentration state of C changes to a high (nonzero) concentration state. We take c to be the concentration of the species C and $v \equiv v_x \hat{x}$ to be the flow velocity.

With these assumptions, the general equation of motion for the concentration field is

$$\partial_t c = D \partial_{xx} c - \partial_x(vc) + f(c), \quad (1)$$

where D denotes the diffusivity, the second term on the right-hand side accounts for the advection of the concentration by the flow, and $f(c)$ is a generic reaction term with the following properties: (i) it destabilizes a steady state with $c_{ss} = 0$, and (ii) it saturates at finite value c_0 . For most of the paper, we will use a simple form for $f(c)$:

$$f(c) = rc(1 - c/c_0), \quad (2)$$

but we have checked that our results do not depend qualitatively on this particular form. Note that, in the absence of the advection by the velocity field, (1) constitutes the well-known equation for a traveling wave introduced independently by Fisher [53] and by Kolmogorov, Petrovsky, and Piskunov [54], which we refer to as the KPP equation. The absence of activity can be experimentally realized in two ways (Fig. 1). In the absence of ATP, A can still bind to the DNA-conjugated motors, and thus the average species C will still have the diffusion coefficient D . In the absence of motors, instead, the relevant diffusion coefficient would be D_A .

The overdamped equation of motion for the velocity field, relevant for slow flows encountered in biological and biomimetic systems, has the form

$$(\gamma - \eta \partial_{xx})v = \partial_x[\zeta(c)], \quad (3)$$

where γ is a friction coefficient that accounts for relative motion with respect to the surrounding medium, η is the viscosity of the active layer, and $\zeta(c)$ is a stress that depends on the regulator concentration. This stress may have a passive component arising from an equation of state for c , but we take it to be primarily active with $\zeta(c) > 0$ implying a contractile system and $\zeta(c) < 0$ an extensile system. For much of this paper, we will assume a form for $\zeta(c)$: $\zeta(c) = \alpha c$, where α is a constant. However, we checked that our results do not depend qualitatively on the form of ζ by also considering two alternate forms of the stress coupling: a quadratic form, $\zeta_c = \alpha c^2/c_0$, and a saturating form, $\zeta(c) = \alpha c/(1 + c/c_0)$ (see Appendix B and Fig. 7). Equations (1) and (3) fully characterize the effectively one-dimensional RDAM system.

To facilitate numerical calculations, we now nondimensionalize (1) and (3) by rescaling $x \rightarrow x/\ell$, $t \rightarrow t/\tau$, and $c \rightarrow c/c_0$, where $\ell = \sqrt{\eta/\gamma}$ is the hydrodynamic correlation length, which is proportional to the thickness of the active layer, and $\tau = 1/r$ is the zero-wave-number inverse relaxation (growth) rate of concentration fluctuations. The rescaled equations of motion are

$$\partial_t c = \frac{1}{\text{Da}} \partial_{xx} c - \partial_x(vc) + c(1 - c), \quad (4)$$

$$(1 - \partial_{xx})v = \Omega(c) \partial_x c, \quad (5)$$

where c , x , t , and v are now nondimensional, and $\text{Da} = r\eta/D\gamma$ —the second Damköhler number—and $\Omega(c) = \partial_c \zeta(c)/r\eta$ are nondimensional parameters of the model. Da and Ω express ratios, respectively, of characteristic timescales of reaction to diffusion, and of active transport to reaction. Thus, Ω can also be expressed as $\Omega = \text{Pe}/\text{Da}$, where Pe is the

more familiar Péclet number expressing the ratio of active to diffusive timescales, given in our case by $\text{Pe}(c) = \partial_c \zeta(c)/D\gamma$.

To understand the behavior of this model, we first consider the linear stability of a state with a homogeneous concentration implied by (4) and (5), and then we numerically analyze how activity modifies the propagation of a concentration front into a state with $c = 0$.

IV. LINEAR THEORY

We start by examining the linear stability of a state at zero average concentration, i.e., $c_{ss} = 0$, with boundary conditions $c(\pm\infty) = 0$, $v(\pm\infty) = 0$. Linearizing (4) and (5) about $c_{ss} = 0$ and Fourier-transforming in space and time, we get

$$-i\sigma c_{q,\sigma} = \left(1 - \frac{1}{\text{Da}} q^2\right) c_{q,\sigma}, \quad (6)$$

$$|q| \in \left\{ \frac{\sqrt{-[1 + \text{Da}(1 - \Omega_1)]} - \sqrt{[1 + \text{Da}(1 - \Omega_1)]^2 - 4\text{Da}}}{\sqrt{2}}, \frac{\sqrt{-[1 + \text{Da}(1 - \Omega_1)]} + \sqrt{[1 + \text{Da}(1 - \Omega_1)]^2 - 4\text{Da}}}{\sqrt{2}} \right\} \quad (8)$$

when

$$\Omega_1 > \frac{1 + 2\sqrt{\text{Da}} + \text{Da}}{\text{Da}} = \Omega_c \quad (9)$$

[or $\text{Pe}_1 > (1 + \sqrt{\text{Da}})^2$, where $\text{Pe}_1 \equiv \text{Pe}|_{c=1}$], as sketched in Fig. 2. Since this requires $\Omega_1 > 0$, it can only be achieved in *contractile* systems. The maximum growth rate is at the wave number $q_c = \sqrt{-1 + \sqrt{\Omega_1 \text{Da}}} = \sqrt{-1 + \sqrt{\text{Pe}_1}}$. The instability starts at finite wave number in our model, in contrast to the zero-wave-number instability predicted in [23], due to the reaction term that damps concentration fluctuations at zero wave number.¹

We now consider the case in which a front moving at a constant speed k interpolates between these two states, such that $c(-\infty) = 1$ and $c(\infty) = 0$, and $v(\pm\infty) = 0$. The steady-state front must then be described by a function $c(\xi)$, where $\xi = x - kt$, which, in the absence of flow-advection due to activity, solves the equation

$$0 = \frac{1}{\text{Da}} \partial_{\xi\xi} c + k \partial_{\xi} c + c(1 - c). \quad (10)$$

The speed of the $c = 1$ front in the absence of activity is expected to be, in dimensional form, $k = 2(Dr)^{1/2}$, using a marginal stability argument originally due to [55]. Contractile

where σ and q are the (nondimensional) frequency and wave number of the perturbation, respectively. The dispersion relation is $\sigma = i(1 - q^2/\text{Da})$, which implies that the amplitude of a small wave-number fluctuation grows, destabilizing the $c_{ss} = 0$ state. Next, linearizing about $c_{ss} = 1$ [$c(\pm\infty) = 1$, $v(\pm\infty) = 0$] and defining $\delta c = c - 1$ and $\Omega|_{c=1} \equiv \Omega_1$, we get

$$-i\sigma \delta c_{q,\sigma} = -\left(1 + \frac{1}{\text{Da}} q^2 - \frac{\Omega_1 q^2}{1 + q^2}\right) \delta c_{q,\sigma}. \quad (7)$$

This implies that a state with $c_{ss} = 1$ is stable in the long-wavelength $q \rightarrow 0$ limit where $\sigma = -i$. Equation (7), however, suggests an activity-induced *nonhydrodynamic*, *finite* wave-number instability of the homogeneous state for wave numbers in the range

or extensile activity qualitatively modifies the KPP concentration dynamics via the advection by the hydrodynamic velocity. Formally solving (5) in terms of the concentration field leads to an integrodifferential equation for the concentration, which is not easy to deal with analytically. We will therefore examine the effect of advection due to active flow on the front propagation numerically. However, we can first gain some qualitative insight by assuming that $v \gg \partial_{xx} v$ in (5), which is generically the case when $\ell \rightarrow 0$, and replace the velocity field in (4) with $v = \Omega(c) \partial_x c$ to obtain

$$0 = \left(\frac{1}{\text{Da}} - c\Omega\right) \partial_{\xi\xi} c + [k - (c\Omega' + \Omega) \partial_{\xi} c] \partial_{\xi} c + c(1 - c), \quad (11)$$

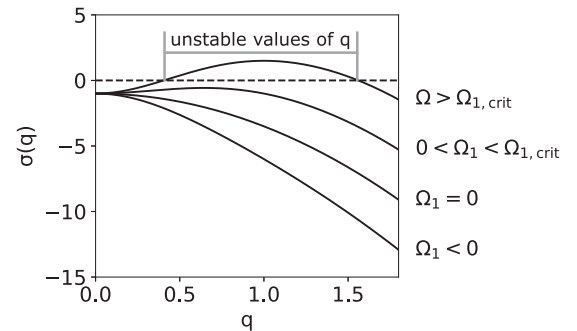


FIG. 2. Growth rate σ of wave number q at different values of the active stress parameter $\Omega_1 = \Omega|_{c=1}$, from linear stability analysis of (4) and (5) around $c = 1$ with a linear stress term $\zeta(c) = \alpha c$. The black dashed line is at $\sigma = 0$. For contractile stresses above a critical value of Ω_1 , given in (9), the reaction-diffusion active matter (RDAM) system is unstable to perturbations in a finite range of wave numbers. Extensile stresses (negative Ω) stabilize the system. In this figure, $\text{Da} = 0.4$.

¹A note of caution: A prediction of a finite wave-number instability within a hydrodynamic theory should not be taken to imply that all systems with the same symmetry, and the same sign of activity, will display this instability, in contrast to prediction for a zero-wave-number instability. Whether the growth rate turns positive at a nonzero wave number depends not only on the lowest order in wave-number terms retained within the hydrodynamic theory, but terms at all orders in wave numbers that are not. Such terms encode system-specific details and can eliminate a finite wave-number instability even when a theory retaining only the lowest order in wave-number terms suggests its presence.

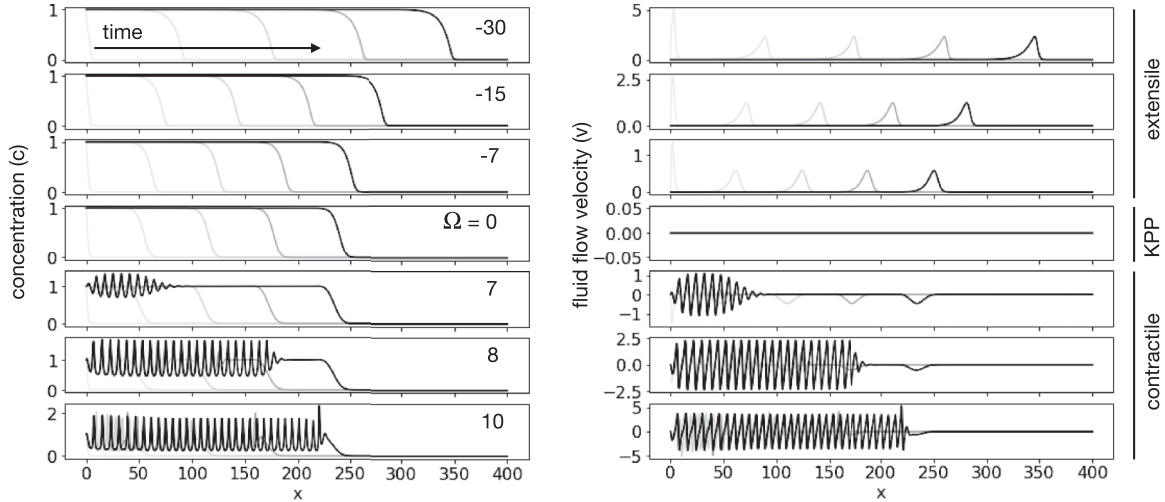


FIG. 3. Representative simulations of the RDM model [(4) and (5)] studied in this article. This model of mechanochemical coupling may represent, for example, a simplified 1D version of an experiment in a thin film where a reacting and diffusing chemical species A interacts with stress-generating molecules (for instance, molecular motors walking on filamentous proteins) such that the magnitude of the stress increases with increasing concentration of A . Positive values of Ω model contractile stresses, while negative values model extensile stresses. Similar to Ref. [23], we find that a periodic instability develops behind the front above a critical value of contractile stresses. Moreover, we note several other interesting features that we explore in this paper. First, there is an asymmetric effect of stresses on the speed of the traveling front, which is increased by extensile stresses but unaffected by contractile stresses. Second, the instability is modified by the boundary condition imposed by the front in three ways: the critical value of Ω is shifted to a higher value compared to an infinite system; there is a gap between the front and the instability that closes with increasing Ω ; and the instability selects a single wavelength from the spectrum of unstable wavelengths in the ideal system. Simulation parameters for all simulations in this figure: $L = 400$, $T = 100$, $dx = 0.01$, $dt = 0.1$, $Da = 0.4$.

where the prime denotes differentiation with respect to c . Therefore, the coupling to an active species leads to two distinct modifications of the usual KPP equation: (i) it reduces the (nondimensional) diffusivity at nonzero concentrations via the term $c\Omega$, and (ii) it leads to a new nonlinear term $(c\Omega' + \Omega)\partial_\xi c$ in the equation of motion. The former implies that a $c = 1$ concentration profile behind the front cannot be homogeneous for large values of Ω , as seen in the linear stability analysis. Note, however, that linearizing (11) suggests an incorrect value for the threshold of the instability of the $c_{ss} = 1$ state discussed earlier, the range of unstable wave numbers, and the wave number at which the growth rate of the instability is maximum. It predicts an instability for $\Omega_1 = 1/Da$ and, since there is no stabilizing mechanism at large wave numbers, no fastest growing mode. The reason for this is that we assumed the hydrodynamic correlation length ℓ to be 0, which amounts to replacing the factor $1/(1 + q^2)$ in the last term of the right-hand side in (7) by 1. If we expand $1/(1 + q^2)$ in a binomial series for $q \rightarrow 0$, we see that this approximation ignores even *linear* terms at higher wave numbers. Therefore, this implies that (11) becomes qualitatively different from the full dynamics when $\Omega \gtrsim 1/Da$ and cannot be used to obtain qualitative information about it beyond this point.

If instead of setting $\ell \rightarrow 0$ and ignoring the $\partial_{xx}v$ term in (5), we formally invert the differential operator $(1 - \partial_{xx})^{-1} \equiv \sum_{n=0}^{\infty} \partial_x^{2n}$, we find that the nonlocal character of the full velocity field leads to linear and nonlinear higher order in gradient terms in the concentration equation which should lead to wave-number selection behind the front when the homogeneous state is unstable.

Finally, even in the regime where (11) is predictive, the nonlinear modification of diffusivity and speed implies that results regarding the speed of the front derived using marginal stability analysis for KPP equations cannot be carried over to this case [55–57]. In particular, if (11) is linearized around $c = 0$, it simply reduces to the usual KPP equation (10) which would predict that the front speed would be unaffected by activity Ω . This turns out to be incorrect, as we explicitly demonstrate in the next section by numerically simulating (4) and (5).

V. NUMERICAL RESULTS

To explore the propagation of a concentration front in the RDM system, we numerically simulated (4) and (5). In Fig. 3 we show example trajectories of these simulations. In this section, we consider only stress linear in c , i.e., $\zeta(c) = \alpha c$, so that Ω is independent of c and we drop the subscript on Ω_1 . Further examples are shown in Fig. 7 with alternative forms of the stress term $\zeta(c)$ and the reaction term $f(c)$. We simulated the system using a custom python code, which is available along with the analysis scripts used in this paper [58]. The coupled equations were integrated in real space using an implicit Euler scheme. Discretization parameters (grid size dx , time step dt) were chosen to be small enough that the simulations were numerically stable and results did not depend on them. Box length L and total time T were chosen to be large enough that the simulation results reached their asymptotic values. Dirichlet boundary conditions were imposed on c and v with $c(0) = 1$, $c(L) = 0$, and $v(0) = v(L) = 0$, implying $\partial c_x(0) = \partial c_x(L) = 0$ due to Eq. (5). $Da = \frac{r\eta}{D\gamma}$ was chosen

to match experimentally realizable conditions with $r = \tau_A^{-1} = 10^{-2} - 10^{-4} \text{ s}^{-1}$, the rate of the DNA/enzyme autocatalytic reaction, and $D = D_A = 10^{-10} \text{ m}^2 \text{ s}^{-1}$, the diffusion coefficient of a 10-mer single-stranded DNA [41]. For a thin film of thickness h , $\eta/\gamma \sim h^2$, yielding $\text{Da} = 1 - 10^{-2}$ for $h = 100 \mu\text{m}$. We chose Da ranging from 0.2 to 0.8 in the following. Finally, the stress parameter Ω was varied in the range -30 to 12 to probe different regimes of contractile and extensile stresses. Specific parameter values are stated along with the results in what follows.

Figure 3 summarizes our central findings. It shows that with contractile stresses, the front speed does not change significantly and, above a critical value of Ω , an instability appears behind the front where the concentration is high, as predicted from the linear theory. In contrast, in the extensile case, the selected front speed k^* increases as a function of Ω but no spatial instability is observed. We now examine these results in more detail.

A. Front speed selection in the presence of activity

In this section, we examine how activity modifies the speeds of the concentration fronts. We defined the front location as the rightmost point where $c = 0.5$ (when the contractile instability sets in, there can be points behind the front with $c = 0.5$). We computed the location of the front from the simulations in Fig. 3 and found that it is a linear function of time after an initial transient resulting from the step-function initial conditions that we used. The transient happens because the speed of a KPP front and its shape are related; specifically, linearizing (10) about $c = 0$ and assuming that $c(x, t)$ is a traveling wave with the exponential form $c(x, t) \propto \exp(-a(x - kt))$ at the leading edge yields a dispersion relation that predicts that the front speed k decreases with increasing a , saturating at $k = 2\text{Dr}^{-1/2}$ for $a \geq 1$ [59]. As a result, simulations begun from an infinitely steep front must accelerate to their asymptotic speed. We therefore calculated the front speed from a linear fit of the front position versus time at late times after the speed had stopped changing. We verified our results by checking that we obtained the same front speed in simulations initialized with a shallow initial profile of the form $c(x, 0) = \tanh((x - x_0)/m)$ with m chosen to be large enough that the front speed converges to its final value from above. Additionally, we checked that our choices of lattice spacing ($dx = 0.5$) and time step ($dt = 0.1$) were small enough that further reducing them had a negligible effect on the calculated front speed.

We first consider the effect of extensile activity on the front speed. In our reduced model (11), if Ω is independent of c , the activity *nonlinearly* enhances the $\partial_{\xi\xi}$ term and reduces the ∂_{ξ} term compared to the standard KPP equation, making it quite different from the standard KPP equation (10). It follows from a simple physical argument that restricts solutions to have non-negative values of c that a KPP front can adopt any speed above a minimum allowed value [59], but it has been shown using marginal stability analysis that in fact the front asymptotically picks out this minimum speed [55]. While a marginal stability argument is not expected to yield the front speed in this case, we hypothesize that the selected front speed in full simulations (k^*) will still be given by the

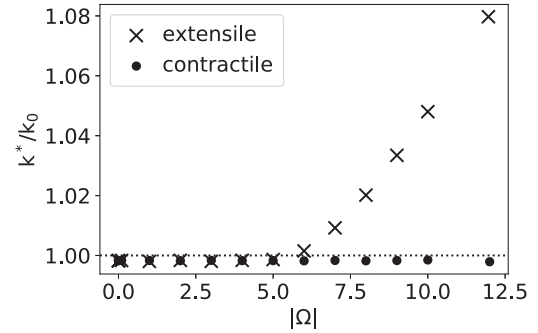


FIG. 4. The effect of extensile and contractile stresses on the selected front speed k^* . Extensile active stresses result in an increase in k^* above a critical value Ω^* of Ω , while contractile stresses have no effect. All y-axes are normalized by $k_0 = 2\text{Da}^{-1/2}$, the front speed at $\Omega = 0$. The line at $k^* = k_0$ is a guide for the eye. Simulation parameters for these plots: $T = 900$, $dt = 0.05$, $dx = 0.1$, $L = 3000$ for $\Omega > -7$ and $L = 2800 + 58|\Omega|$ for $\Omega \leq -7$. L was chosen such that the front would not reach the end of the box before $t = 900$.

minimal allowed value of k , k_{\min} , for which c is non-negative in simulations of (11). We numerically integrated (11) for each value of Ω using MATLAB's ode23s function for a range of values of k (code online [60]). The minimum allowed value of the front speed, k_{\min} , was identified as the smallest value of k for which $c \geq 0$ for all ξ . The k_{\min} obtained via this procedure is an increasing function of Ω [in contrast to the prediction from a naive linearization of (11) about the $c = 0$ state [55]]. This observation can be heuristically accounted for by the following argument. For (10), linearizing about $c = 0$ shows that, due to the constraint that $c \geq 0$, the only possible trajectories have $k \geq 2(\text{Dr})^{1/2}$ [59] (which is also the speed of the front [55]). Now, since c must always be positive and $\partial_{\xi}c$ is always negative in permitted trajectories where the front moves from left to right, in (11) a negative value of Ω (corresponding to extensile stresses) increases the diffusivity ($D_{\text{eff}} = 1/\text{Da} - c\Omega$). At the same time, the part of k_{eff} [which is the coefficient of $\partial_{\xi}c$ in (11)] that depends on $\partial_{\xi}c$ is $-\Omega\partial_{\xi}c$ and therefore decreases. Since k_{eff} is constrained to be greater than D_{eff} , and since D_{eff} increases and $-\Omega\partial_{\xi}c$ decreases as Ω becomes more negative (i.e., for increasing extensile stress), the value of k must increase to compensate. Thus, the allowed values of the front speed shift upwards.

We now compare this result with the full numerical simulation to test our hypothesis. Figure 4(a) shows that the front speed begins to increase in extensile simulations for $\Omega \approx -5$ (for $\text{Da} = 0.4$). In Fig. 5 we compare k^* to k_{\min} for three values of Da and $0 < |\Omega| < 25$. Figure 5(b) shows that k_{\min} is within 5% of k^* for the largest value of Da that we looked at ($\text{Da} = 0.8$), and is within 1% for the smallest value $\text{Da} = 0.2$. This is a significant finding for two reasons: (i) Equation (11) was obtained within an $\ell \rightarrow 0$ approximation and therefore ignored both linear and nonlinear terms of higher order in gradients. Despite this, the reduced model predicts the front speed reasonably accurately. (ii) This demonstrates that even in the presence of activity, which makes the effective concentration equation nonlocal due to the nonlocal dependence of the velocity field on the concentration, the selected speed is still given by the minimum allowed speed, as had been

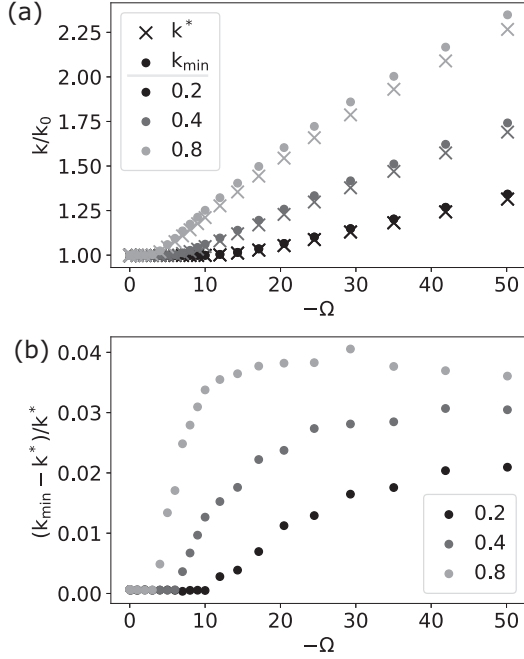


FIG. 5. The front selects the minimum allowed speed. (a) Comparing the front speed selected in simulations of the full model (k^*) to the minimum allowed front speed k_{\min} in a simplified model where we take $\ell \rightarrow 0$. Shades of gray represent different values of Da (0.2, 0.4, 0.8), \times 's are results from a simulation of the full system, and circles are results from the simplified model. The agreement is good over this range of parameters and improves with decreasing Da . (b) Relative difference between the data from full and simplified models in (a). Simulation parameters for these plots: $T = 900$, $dt = 0.05$, $dx = 0.1$. For $Da = 0.2$ and 0.4 , $L = 3000$ for $\Omega > -7$ and $L = 2800 + 58|\Omega|$ for $\Omega \leq -7$. For $Da = 0.8$, $L = 3000$ for $\Omega > -6$ and $L = 3000 + 75|\Omega|$ for $\Omega \leq -7$.

predicted by [55,56] in the context of much simpler models. The selected speed in both the full simulation and from (11) remains equal to the speed in the absence of activity up to a critical value $|\Omega^*|$ and then starts to increase with $|\Omega|$, as shown in Fig. 5(a). Figure 5(a) also demonstrates that Ω^* decreases with increasing Da . This feature may also be heuristically understood from (11): when $c Da |\Omega| \ll 1$, the effective diffusivity in (11) is effectively controlled by $1/Da$ and the speed is equal to the one in the absence of activity. However, as Da increases, $|\Omega| \sim 1/Da$ at smaller values of Ω , implying that the front speed is controlled by activity for $|\Omega| > |\Omega^*|$ that decreases with Da .

Comparing the results in Fig. 5 with the values of the experimental parameters summarized in Sec. III allows us to evaluate the magnitude of the increase in the front velocity expected in the experiment sketched in Fig. 1. Taking the thin-film approximation $\eta/\gamma \sim w_d^2$, we have $Da = w_d^2 \frac{1}{D_A \tau_A} \approx 2 \times (10^{-3} - 10^{-1})$ for a reasonable value of $w_d = 40 \mu\text{m}$. Kinesin/microtubule active fluids generate flows with velocities up to [61] $v_a = 600 \mu\text{m/s}$ associated with typical lengths $l_a = 100 \mu\text{m}$, yielding $Pe = v_a l_a / D_A \approx 6$ and thus $\Omega = Pe/Da \approx 6 \times (10^3 - 10)$. In these conditions, from Fig. 5, we expect a change in the front velocity of about 20% in the presence of activity, which could be reasonably measured experimentally.

We now consider the effect of contractile active stress on the front propagation. As discussed earlier, (11) is only expected to be predictive for $\Omega < 1/Da$. We find that, in this regime [i.e., when (11) does not imply an instability of the high concentration state], (11) implies that the minimum allowed value of k is essentially the same as the one for $\Omega = 0$. This is heuristically understandable: even for the extensile system, the front speed was only controlled by activity, and increased with it, when $|\Omega| \gtrsim 1/Da$. Therefore, since (11) is only valid for $\Omega < 1/Da$, we do not expect the front speed to decrease with contractility at least when the high concentration front is not itself unstable. Our results from the full numerical simulations confirm that the front speed is not modified when the active stress is contractile, as shown in Fig. 4. Surprisingly, we find that the front speed remains independent of Ω even at larger values of Ω when the concentration and velocity profiles behind the front develop a periodic pattern, which we discuss in the next section.

B. Patterns behind the front with contractile stresses

We now consider the dynamics at large Ω when the homogeneous profile behind the front is destabilized and a periodic pattern is formed. We measure Ω_c —the minimum value of Ω for which a periodic pattern emerges, and q_c —the wave number of the periodic pattern, from the simulation of the full model. We checked that in simulations with periodic boundary conditions, Ω_c and q_c are consistent with linear stability analysis predictions (see Appendix C for methods and results).

In simulations with a front, there is a small peak in the concentration just behind the front, which we identified using a standard peak-finding algorithm as the peak with the largest value of x . We call the location of this peak x_f as it tracks the location of the front. We define the leading edge of the instability x_{p1} to be the location of the second leftmost peak. We then Fourier-transformed the region from $x = 0$ to $x = x_f$, as shown in Fig. 6(a). Because x_{p1} is a function of time, the region of the Fourier transform changes over time and so do the possible wave numbers of the instability. We therefore approximated the height of the dominant wave number at long times (q_c) by the amplitude h_{\max} of the wave number with maximum amplitude (q_1) at any point in time. We plotted $\ln(h_{\max})$ versus t and fit the linear region of the curves to extract the growth rate $\sigma(q_1; \Omega)$. We then fit Ω versus $\sigma(q_1)$ to a line and extrapolated Ω_c as the value of Ω where $\sigma(q_1) = 0$.

Surprisingly, we find from this analysis that $\Omega_c \approx (1 + 2\sqrt{Da + Da})/Da$ as predicted by the linear stability analysis of a homogeneous $c_{ss} = 1$ state. Next, examining the time evolution of the dominant wave number q_1 , we find that it relaxes from below to a constant mean value q_c after $t \approx 1000$. In Fig. 6(b), we plot the mean ± 1 standard deviation of q_1 from $t = 1200$ to 1800 (i.e., $q_c \pm 1$ standard deviation), measured at intervals of $\Delta t_{\text{sample}} = 2$. The value of q_c is within 1% of the value predicted by the linear stability analysis when Ω is just above Ω_c , and diverges from it as Ω_c increases.

Finally, we measured the lag of the leading edge of the instability behind the front (i.e., the front speed k^* minus the periodic pattern's group velocity v_g). After a brief initial transient, the distance $\Delta x_{\text{instab}} = x_f - x_{p1}$ is a linear function of time. We calculated $k^* - v_g$ as the slope of the linear fit to

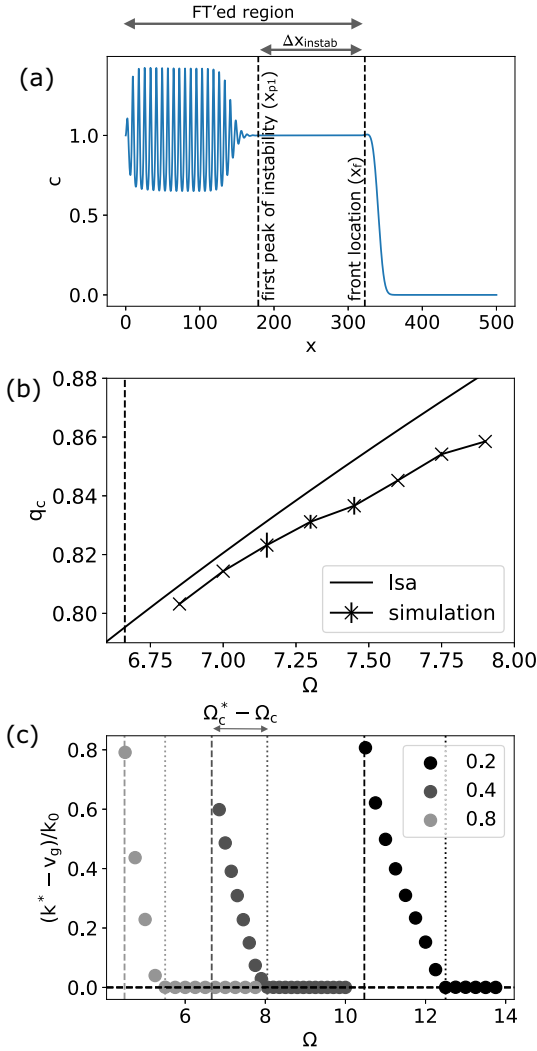


FIG. 6. Wave number and group velocity of the periodic pattern behind the front. (a) To measure the group velocity v_g and the fastest growing wave number q_c , we defined the location of the front x_f as the rightmost peak of the concentration, and the location of first peak of the instability x_{p1} as the location of the second rightmost peak. (b) q_c as a function of Ω , defined as the mean value over the final third of a long simulation of the wave number with maximum amplitude in the Fourier transform of the concentration between $x = 0$ and $x = x_f$. The vertical dashed line is located at Ω_c . Parameters for these simulations were $Da = 0.4$, $dt = 0.05$, $dx = 0.05$, $L = 6300$, and $T = 1800$. (c) Front speed k^* minus the group velocity of the periodic pattern v_g as a function of Ω at three values of Da (colors). Each simulation data point is obtained from the slope of the distance from the front to the first peak of the instability, $\Delta x_{\text{instab}} = x_f - x_{p1}$, as a function of time, which is a linear function at long times. The vertical dashed lines are located at Ω_c , and the dotted lines are located at Ω_c^* , showing decreasing $\Omega_c^* - \Omega_c$ with increasing Da . Parameters for these simulations were $dt = 0.05$, $dx = 0.05$, $L = 500$, and $T = 150$.

t versus Δx_{instab} . As shown in Fig. 6(c), $k^* - v_g$ goes from a positive value just beyond Ω_c to 0. That is, the pattern lags behind the traveling front until a second value Ω_c^* . Before this value, the front “outruns” the periodic pattern, implying that near the traveling edge, a homogeneous region emerges before

the patterned state sets in (see the supplemental movie [62]). Beyond Ω_c^* , the pattern “catches up” with the front, and for all higher values it moves with the front. The value of $\Omega_c^* - \Omega_c$ decreases with increasing Da (which depends on the ratio of the reaction coefficient and the bare diffusivity). This implies that in highly diffusive species, the patterns formed due to the contractile instability lag arbitrarily far behind the advancing front and will not be observed in experiments that track the front. Experimentally, microtubule active matter also displays a contractile behavior and could thus be used to test these predictions, either with multiheaded kinesins [63], clustered kinesins [42,64], or dyneins [30].

VI. CONCLUSION

In this paper, we examined the effect of active flows on the propagation of a reacting and diffusing activity regulator species. We demonstrated that neither contractile nor extensile active flows inhibit the propagation of the activity regulator. We further showed that extensile activity (corresponding to negative $\Omega = Pe/Da$ in our model) increases the propagation speed of the activity regulator relative to k_0 , the speed in the absence of activity, when the magnitude of Ω exceeds a critical value Ω^* , while contractile activity does not modify the front speed. On the other hand, contractile active flows above a critical value Ω_c destabilize the homogeneous state behind the front leading to a periodic pattern. For moderate Da and activity, however, a homogeneous state prevails near the front edge as the speed of the pattern lags behind the front speed.

RDAM systems may describe a wide range of *in vivo* biological systems. For instance, they may be relevant for the dynamics of activity regulators [23] in the cellular cortex, or they may describe the dynamics of cell monolayers [18]. They may also be synthesized *in vitro*. Indeed, we motivated our one-dimensional, one-variable model from the state-of-the-art *in vitro* experiments involving a DNA/enzyme autocatalytic RD front coupled with a DNA-responsive kinesin/microtubule active fluid. We believe that the qualitative features predicted by the theory presented here can be observed in *in vitro* RDM systems such as the one suggested here. Taken together, our results may prove useful in interpreting pattern formation in RDM systems *in vivo* and *in vitro* that are on the experimental horizon.

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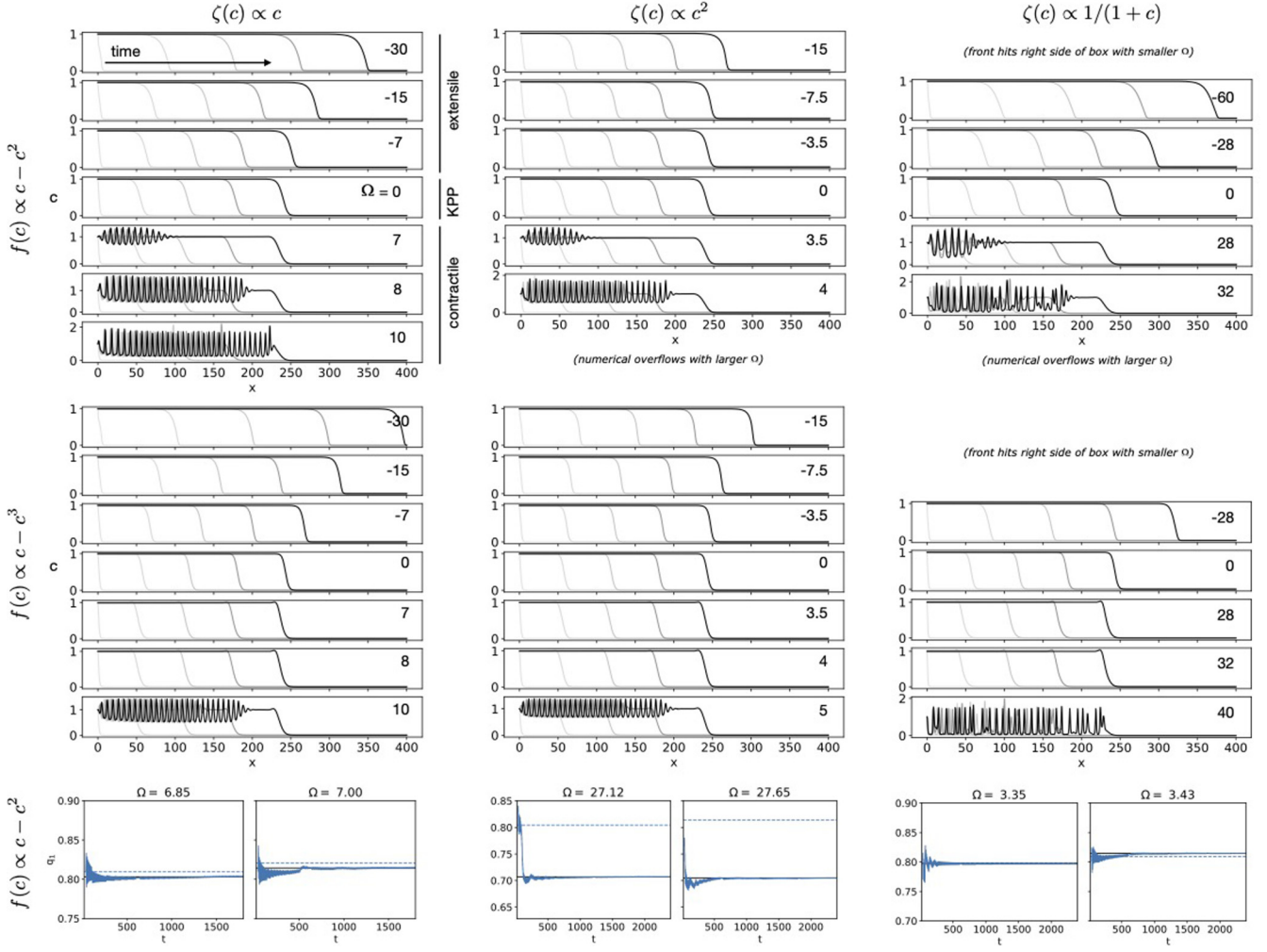


FIG. 7. Simulation results with alternative choices of stress and reaction terms. Top two rows: Representative simulations of the concentration as a function of time in the reaction-diffusion active matter (RDAM) model [(4) and (5)], with three alternative forms of the stress term $\zeta(c)$ and two forms of the reaction term $f(c)$. The upper-left panel is the same as in Fig. 3. Parameters: $Da = 0.4$, $L = 400$, $T = 100$ for all simulations; $dx = 0.05$ and $dt = 0.1$ for simulations with $\zeta(c) \propto c$ and $\zeta(c) \propto c^2$; $dx = 0.01$ and $dt = 0.05$ for simulations with $\zeta(c) \propto c/(1+c)$. Bottom row: Wave number with maximum amplitude in the Fourier transform of the concentration behind the front when $\Omega > \Omega_c$, as a function of time, computed as described in Sec. VB. Parameters: $Da = 0.4$ for all simulations; $T = 1800$, $L = 6300$, $dx = 0.05$, and $dt = 0.05$ for simulations with $\zeta(c) \propto c$; $T = 2400$, $L = 8400$, $dx = 0.05$, and $dt = 0.05$ for $\zeta(c) \propto c^2$; $T = 2400$, $L = 8400$, $dx = 0.01$, and $dt = 0.05$ for $\zeta(c) \propto c/(1+c)$.

APPENDIX A: DEMONSTRATION THAT THE RDM MODEL IS APPLICABLE BOTH FOR ALIGNED AND ISOTROPIC SYSTEMS

In the main text, we described a simplified RDM model in which we did not take orientational ordering of the active units into account. However, active matter systems, such as the kinesin-microtubule one described in the main text, are often aligned [32]. In this Appendix, we show that the RDM model is still applicable in this case as long as the speed of the KPP front is fast compared to the relaxation of the angular fluctuations in the aligned state.

We consider an aligned system (for instance, composed of microtubules or polarized cells) with the degree of alignment denoted by

$$\mathbf{Q} = \frac{S}{2} \begin{pmatrix} 1 & 0 \\ 0 & -1 \end{pmatrix}, \quad (\text{A1})$$

where S is the degree of order, with $S = 0$ denoting an isotropic organization and $S \neq 0$ denoting an aligned state. With this, the equation for the concentration field is modified to

$$\partial_t c = \bar{D} \nabla^2 c + \zeta_c \nabla \nabla : (\mathbf{Q}c) - \nabla \cdot (\mathbf{v}c) + f(c), \quad (\text{A2})$$

where \bar{D} is the bare diffusivity and $f(c)$ is a generic reaction term. The “curvature current” $\propto \zeta_c$ is a standard active concentration current in active, aligned systems [8]. The equation

TABLE I. Values of the constants μ and β in (B1) for each choice of $\zeta(c)$ and $f(c)$ shown in Fig. 7.

$\zeta(c)/\alpha \rightarrow$	c	c^2	$c/(1+c)$	$f(c)/r \rightarrow$	$c - c^2$	$c - c^3$
$\beta \rightarrow$	1	2	1/4	$\mu \rightarrow$	1	2

TABLE II. Critical values of Ω in nondimensional units, when $Da = 0.4$, for each choice of $\zeta(c)$ and $f(c)$ featured in Fig. 7 as predicted from a linear stability analysis.

$\Omega_{1,c}$	$\zeta(c) \propto c$	$\zeta(c) \propto c^2$	$\zeta(c) \propto c/(1+c)$
$f(c) \propto c - c^2$	6.66	3.33	26.65
$f(c) \propto c - c^3$	8.97	4.49	35.89

of the velocity field becomes

$$(\eta\nabla^2 - \gamma)\mathbf{v} = -\nabla\Pi - \nabla\bar{\zeta}_c(c) - \nabla \cdot [\bar{\zeta}(c)\mathbf{Q}], \quad (\text{A3})$$

where Π is the equilibrium pressure stemming from an equation of state for c , $\bar{\zeta}_c$ is the isotropic active stress, and $\bar{\zeta}$ is the active apolar stress that arises in aligned systems [8,65]. To complete the description of the model, we also need to specify the fluctuations of the order parameter \mathbf{Q} . However, we assume that the propagation of the chemical species happens at a timescale that is much faster than the alignment dynamics, and on that timescale the order-parameter \mathbf{Q} can be assumed to be frozen at a constant value. Therefore, we replace \mathbf{Q} by its value given by (A1) with $S = 0$ for an isotropic state and $S = 1$ for an aligned state. Further assuming that only variations of c and \mathbf{v} are along \hat{x} , we obtain the equations of motion described in the main text with $v \equiv v_x$, $D = \bar{D} + \zeta_c$ when $S = 1$ (i.e., an aligned state) and $D = \bar{D}$ when $S = 0$, $\zeta(c) = \bar{\zeta}(c) + \bar{\zeta}_c(c) - \Pi$ in the aligned state, and $\zeta(c) = \bar{\zeta}_c(c) - \Pi$ in the isotropic state.

APPENDIX B: ALTERNATIVE STRESS AND REACTION TERMS

Figure 7 demonstrates that the phenomena that we describe in this paper are also present with several alternative choices for the stress and reaction terms $\zeta(c)$ and $f(c)$. Specifically, for two reaction terms and three stress terms, we observe an increased front speed with extensile stresses and an instability behind the front that picks out a single wavelength at long times. Linear stability analysis predicts that instabilities arise when

$$\Omega_1 > (1 + 2\sqrt{\mu Da} + Da)/(\beta Da), \quad (\text{B1})$$

where μ and β are constants that depend on the choice of $\zeta(c)$ and $f(c)$ with values given in Table I. Predicted values of Ω_c for each value of μ and β are given in Table II. Figure 7(a) shows that the location of the instability is shifted in agreement with the linear stability prediction for all choices of $f(c)$ and $\zeta(c)$ considered here. While it is not clear from the short simulation in Fig. 7(a) that the model with saturating stress

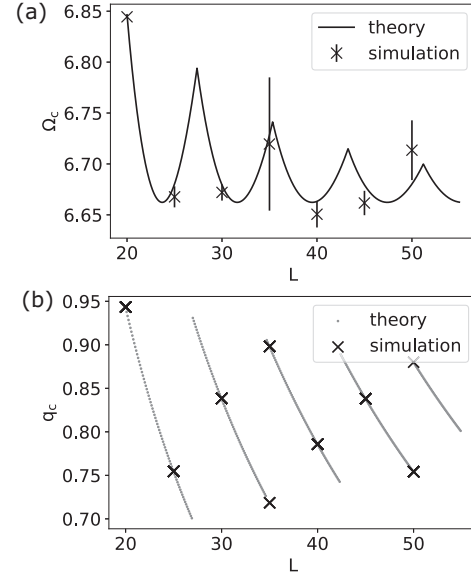


FIG. 8. Checking linear stability analysis predictions in simulations. (a) The critical value of the stress for instabilities to arise as a function of the box length. Error bars are ± 1 standard deviation about the mean from 10 simulations with different random initial conditions. (b) The most unstable wave number q_c at $\Omega = 6.95$ as a function of the box length. All of the points from 10 simulations with different random initial conditions are plotted. Simulation parameters for all data in this figure are $dx = 0.01$, $dt = 0.1$, $T = 100$, and $Da = 0.4$.

[$\zeta(c) \propto c/(1+c)$] picks out a single wavelength, the longer simulations in Fig. 7(b) show that at long times q_1 converges even in this case.

APPENDIX C: CONFIRMING LINEAR STABILITY ANALYSIS RESULTS IN SIMULATIONS

In this Appendix, we compare our linear analysis in Sec. IV to simulations with periodic boundary conditions. Beginning from a system with a high concentration plus a random initial perturbation, $c(x) = 1 + 0.001v(x)$, where $\langle v(x) \rangle = 0$, $\langle v(x)v(x') \rangle = \delta(x-x')$, we simulated a system with parameters $Da = 0.4$, $dx = 0.01$, $dt = 0.1$, $T = 100$, and a range of box sizes. Denoting the Fourier-transformed concentration as \hat{c} and assuming that fluctuations in the concentration grow as $\hat{c}(q) \propto \exp(\sigma(q)t)$, we measured the growth rate of the fastest growing wave number, $\sigma(q_c)$, as the slope of the best-fit line to $\ln(c(q_c))$ versus t . We then fit Ω versus $\sigma(q_c)$ to a line and extrapolated Ω_c as the value of Ω where $\sigma(q_c) = 0$. We find that the critical values of Ω_c [Fig. 8(a)] and q_c [Fig. 8(b)]—the wave number that dominates the spectrum of c at long times] are in good agreement with our linear stability analysis calculations.

[1] C. Collinet and T. Lecuit, Programmed and self-organized flow of information during morphogenesis, *Nat. Rev. Mol. Cell Biol.* **22**, 245 (2021).

[2] M. Cross and H. Greenside, *Pattern Formation and Dynamics in Nonequilibrium Systems* (Cambridge University Press, Cambridge, 2009).

- [3] M. C. Cross and P. C. Hohenberg, Pattern formation outside of equilibrium, *Rev. Mod. Phys.* **65**, 851 (1993).
- [4] A. M. Turing, The chemical basis of morphogenesis, *Philos. Trans. R. Soc. B* **237**, 37 (1952).
- [5] S. Kondo and T. Miura, Reaction-diffusion model as a framework for understanding biological pattern formation, *Science* **329**, 1616 (2010).
- [6] J. B. A. Green and J. Sharpe, Positional information and reaction-diffusion: two big ideas in developmental biology combine, *Development* **142**, 1203 (2015).
- [7] P. Gross, K. V. Kumar, and S. W. Grill, How active mechanics and regulatory biochemistry combine to form patterns in development, *Annu. Rev. Biophys.* **46**, 337 (2017).
- [8] M. C. Marchetti, J. F. Joanny, S. Ramaswamy, T. B. Liverpool, J. Prost, M. Rao, and R. A. Simha, Hydrodynamics of soft active matter, *Rev. Mod. Phys.* **85**, 1143 (2013).
- [9] J. Prost, F. Jülicher, and J.-F. Joanny, Active gel physics, *Nat. Phys.* **11**, 111 (2015).
- [10] F. Jülicher, S. W. Grill, and G. Salbreux, Hydrodynamic theory of active matter, *Rep. Prog. Phys.* **81**, 076601 (2018).
- [11] S. Ramaswamy, The mechanics and statistics of active matter, *Annu. Rev. Condens. Matter Phys.* **1**, 323 (2010).
- [12] J. Toner, Y. Tu, and S. Ramaswamy, Hydrodynamics and phases of flocks, *Ann. Phys.* **318**, 170 (2005).
- [13] S. Ramaswamy, Active matter, *J. Stat. Mech.: Theor. Expt.* (2017) 054002.
- [14] L. P. Dadhichi, A. Maitra, and S. Ramaswamy, Origins and diagnostics of the nonequilibrium character of active systems, *J. Stat. Mech.: Theor. Expt.* (2018) 123201.
- [15] A. P. Solon, M. E. Cates, and J. Tailleur, Active brownian particles and run-and-tumble particles: A comparative study, *Eur. Phys. J.: Spec. Top.* **224**, 1231 (2015).
- [16] M. E. Cates and J. Tailleur, Motility-induced phase separation, *Annu. Rev. Condens. Matter Phys.* **6**, 219 (2015).
- [17] J. Eliaš and J. Clairambault, Reaction-diffusion systems for spatio-temporal intracellular protein networks: A beginner's guide with two examples, *Comput. Struct. Biotechnol. J.* **10**, 12 (2014).
- [18] B. D. MacArthur, P. S. Stumpf, and R. O. C. Oreffo, From mathematical modeling and machine learning to clinical reality, in *Principles of Tissue Engineering*, 5th ed., edited by R. Lanza, R. Langer, J. P. Vacanti, and A. Atala (Academic Press, 2020), Chap. 2, pp. 37–51.
- [19] B. G. Sengers, C. P. Please, and R. O. C. Oreffo, Experimental characterization and computational modelling of two-dimensional cell spreading for skeletal regeneration, *J. R. Soc., Interface* **4**, 1107 (2007).
- [20] P. Müller, K. W. Rogers, B. M. Jordan, J. S. Lee, D. Robson, S. Ramanathan, and A. F. Schier, Differential diffusivity of nodal and lefty underlies a reaction-diffusion patterning system, *Science* **336**, 721 (2012).
- [21] I. R. Epstein and J. A. Pojman, *An Introduction to Nonlinear Chemical Dynamics: Oscillations, Waves, Patterns, and Chaos* (Oxford University Press, Oxford, 1998).
- [22] F. J. Nedelec, T. Surrey, A. C. Maggs, and S. Leibler, Self-organization of microtubules and motors, *Nature (London)* **389**, 305 (1997).
- [23] J. S. Bois, F. Jülicher, and S. W. Grill, Pattern Formation in Active Fluids, *Phys. Rev. Lett.* **106**, 028103 (2011).
- [24] K. V. Kumar, J. S. Bois, F. Jülicher, and S. W. Grill, Pulsatory Patterns in Active Fluids, *Phys. Rev. Lett.* **112**, 208101 (2014).
- [25] D. Zwicker, R. Seyboldt, C. A. Weber, A. A. Hyman, and F. Jülicher, Growth and division of active droplets provides a model for protocells, *Nat. Phys.* **13**, 408 (2017).
- [26] A. Senoussi, Y. Vyborna, H. Berthoumieux, J.-C. Galas, and A. Estevez-Torres, Learning from embryo development to engineer self-organizing materials, in *Out-of-Equilibrium Supramolecular Systems and Materials*, edited by N. Giuseppone and A. Walther (Wiley-VCH, 2021), Chap. 2, pp. 21–60.
- [27] R. Andorfer and J. D. Alper, From isolated structures to continuous networks: A categorization of cytoskeleton-based motile engineered biological microstructures, *Wiley Interdiscip. Rev.: Nanomed. Nanobiotechnol.* **11**, e1553 (2019).
- [28] D. Needleman and Z. Dogic, Active matter at the interface between materials science and cell biology, *Nat. Rev. Mater.* **2**, 17048 (2017).
- [29] P. M. Bendix, G. H. Koenderink, D. Cuvelier, Z. Dogic, B. N. Koeleman, W. M. Briehar, C. M. Field, L. Mahadevan, and D. A. Weitz, A quantitative analysis of contractility in active cytoskeletal protein networks, *Biophys. J.* **94**, 3126 (2008).
- [30] P. J. Foster, S. Fürthauer, M. J. Shelley, and D. J. Needleman, Active contraction of microtubule networks, *Elife* **4**, e10837 (2015).
- [31] T. Sanchez, D. T. N. Chen, S. J. DeCamp, M. Heymann, and Z. Dogic, Spontaneous motion in hierarchically assembled active matter, *Nature (London)* **491**, 431 (2012).
- [32] A. Senoussi, S. Kashida, R. Voituriez, J.-C. Galas, A. Maitra, and A. Estevez-Torres, Tunable corrugated patterns in an active nematic sheet, *Proc. Natl. Acad. Sci. USA* **116**, 22464 (2019).
- [33] M. Isalan, C. Lemerle, and L. Serrano, Engineering gene networks to emulate drosophila embryonic pattern formation, *PLoS Biol.* **3**, e64 (2005).
- [34] M. Loose, E. Fischer-Friedrich, J. Ries, K. Kruse, and P. Schwille, Spatial regulators for bacterial cell division self-organize into surface waves in vitro, *Science* **320**, 789 (2008).
- [35] S. M. Chirieleison, P. B. Allen, Z. B. Simpson, A. D. Ellington, and X. Chen, Pattern transformation with dna circuits, *Nat. Chem.* **5**, 1000 (2013).
- [36] A. Padirac, T. Fujii, A. Estévez-Torres, and Y. Rondelez, Spatial waves in synthetic biochemical networks, *J. Am. Chem. Soc.* **135**, 14586 (2013).
- [37] A. M. Tayar, E. Karzbrun, V. Noireaux, and R. H. Bar-Ziv, Propagating gene expression fronts in a one-dimensional coupled system of artificial cells, *Nat. Phys.* **11**, 1037 (2015).
- [38] A. J. M. Wollman, C. Sanchez-Cano, H. M. J. Carstairs, R. A. Cross, and A. J. Turberfield, Transport and self-organization across different length scales powered by motor proteins and programmed by dna, *Nat. Nanotechnol.* **9**, 44 (2014).
- [39] A. M. Tayar, M. F. Hagan, and Z. Dogic, Active liquid crystals powered by force-sensing dna-motor clusters, *Proc. Natl. Acad. Sci. USA* **118**, e2102873118 (2021).
- [40] Y. Vyborna, J.-C. Galas, and A. Estevez-Torres, Dna-controlled spatiotemporal patterning of a cytoskeletal active gel, *J. Am. Chem. Soc.* **143**, 20022 (2021).
- [41] A. S. Zadorin, Y. Rondelez, J.-C. Galas, and A. Estevez-Torres, Synthesis of Programmable Reaction-Diffusion Fronts using Dna Catalyzers, *Phys. Rev. Lett.* **114**, 068301 (2015).

- [42] A. Senoussi, J.-C. Galas, and A. Estevez-Torres, Programmed mechano-chemical coupling in reaction-diffusion active matter, *Sci. Adv.* **7**, eabi9865 (2021).
- [43] N. Kumar, R. Zhang, J. J. de Pablo, and M. L. Gardel, Tunable structure and dynamics of active liquid crystals, *Sci. Adv.* **4**, aat7779 (2018).
- [44] R. F. Hariadi, R. F. Sommese, A. S. Adhikari, R. E. Taylor, S. Sutton, J. A. Spudich, and S. Sivaramakrishnan, Mechanical coordination in motor ensembles revealed using engineered artificial myosin filaments, *Nat. Nanotechnol.* **10**, 696 (2015).
- [45] B. Martínez-Prat, J. Ignés-Mullol, J. Casademunt, and F. Sagués, Selection mechanism at the onset of active turbulence, *Nat. Phys.* **15**, 362 (2019).
- [46] K. Furuta, A. Furuta, Y. Y. Toyoshima, M. Amino, K. Oiwa, and H. Kojima, Measuring collective transport by defined numbers of processive and nonprocessive kinesin motors, *Proc. Natl. Acad. Sci. USA* **110**, 501 (2013).
- [47] Y. Sato, Y. Hiratsuka, I. Kawamata, S. Murata, and S.-i. M. Nomura, Micrometer-sized molecular robot changes its shape in response to signal molecules, *Sci. Robot* **2**, aal3735 (2017).
- [48] A. Estevez-Torres, T. Le Saux, C. Gosse, A. Lemarchand, A. Bourdoncle, and L. Jullien, Fourier transform to analyse reaction-diffusion dynamics in a microsystem, *Lab Chip* **8**, 1205 (2008).
- [49] C. Leduc, O. Campas, K. B. Zeldovich, A. Roux, P. Jolimaitre, L. Bourel-Bonnet, B. Goud, J.-F. Joanny, P. Bassereau, and J. Prost, Cooperative extraction of membrane nanotubes by molecular motors, *Proc. Natl. Acad. Sci. USA* **101**, 17096 (2004).
- [50] G. Salbreux, J. Prost, and J. F. Joanny, Hydrodynamics of Cellular Cortical Flows and the Formation of Contractile Rings, *Phys. Rev. Lett.* **103**, 058102 (2009).
- [51] A. Maitra, Stable long-range uniaxial order in active fluids at two-dimensional interfaces, [arXiv:2010.15044](https://arxiv.org/abs/2010.15044).
- [52] M. Poujade, E. Grasland-Mongrain, A. Hertzog, J. Jouanneau, P. Chavrier, B. Ladoux, A. Buguin, and P. Silberzan, Collective migration of an epithelial monolayer in response to a model wound, *Proc. Natl. Acad. Sci. USA* **104**, 15988 (2007).
- [53] R. A. Fisher, The wave of advance of advantageous genes, *Ann. Eugenics* **7**, 355 (1937).
- [54] A. Kolmogorov, I. Petrovskii, and N. Piskunov, A study of the diffusion equation with increase in the amount of substance, in *Selected Works of A. N. Kolmogorov I*, edited by V. M. Tikhomirov (Kluwer, Dordrecht, 1991), pp. 242–270, translated by V. M. Volosov from *Bull. Moscow Univ., Math. Mech.* **1**, 1 (1937).
- [55] G. Dee and J. S. Langer, Propagating Pattern Selection, *Phys. Rev. Lett.* **50**, 383 (1983).
- [56] W. van Saarloos, Front propagation into unstable states: Marginal stability as a dynamical mechanism for velocity selection, *Phys. Rev. A* **37**, 211 (1988).
- [57] W. van Saarloos, Front propagation into unstable states, *Phys. Rep.* **386**, 29 (2003).
- [58] <https://github.com/cdeljunco/rdam>.
- [59] J. D. Murray, *Mathematical Biology I* (Springer, New York, 2003).
- [60] <https://github.com/cdeljunco/rdam>.
- [61] T. E. Bate, E. J. Jarvis, M. E. Varney, and K.-T. Wu, Collective dynamics of microtubule-based 3d active fluids from single microtubules, *Soft Matter* **15**, 5006 (2019).
- [62] See Supplemental Material at <http://link.aps.org/supplemental/10.1103/PhysRevE.105.014602> for a movie of a simulation trajectory where a pattern emerges behind the travelling front.
- [63] T. Torisawa, D. Taniguchi, S. Ishihara, and K. Oiwa, Spontaneous formation of a globally connected contractile network in a microtubule-motor system, *Biophys. J.* **111**, 373 (2016).
- [64] L. Bezia, N. P. Mitchell, R. Subramanian, D. J. Needleman, and Z. Dogic, Active microphase separation in mixtures of microtubules and tip-accumulating molecular motors, [arXiv:2107.12281](https://arxiv.org/abs/2107.12281).
- [65] R. A. Simha and S. Ramaswamy, Hydrodynamic Fluctuations and Instabilities in Ordered Suspensions of Self-Propelled Particles, *Phys. Rev. Lett.* **89**, 058101 (2002).