

# Stable length distributions in co-localized polymerizing and depolymerizing protein filaments

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## Abstract

A model for the dynamics of the length distribution in co-localized groups of polar polymer filaments is presented. It considers nucleation, polymerization at plus-ends, and depolymerization at minus-ends and is derived as a continuous macroscopic limit from a discrete description. Its main feature is a nonlinear coupling due to competition of the depolymerizing ends for the limited supply of a depolymerization agent. The model takes the form of an initial-boundary value problem for a one-dimensional nonlinear hyperbolic conservation law, subject to a nonlinear, nonlocal boundary condition. Besides existence and uniqueness of entropy solutions, convergence to a steady state is proven. Technical difficulties are caused by the fact that the prescribed boundary data are not always assumed by entropy solutions.

## 1 Introduction

The motility of a number of species of living cells is driven by polymerization and depolymerization of the protein *actin* in the *lamellipodium*, a thin sheet-like protrusive structure [SSVR02]. During phases of steady motion, which can be rather long for certain types of cells (e.g. the fish keratocyte), the shape of the lamellipodium does not change much, indicating a stable, stationary length distribution of actin filaments (polymerized actin) with an average filament length of several  $\mu\text{m}$ . In recent mathematical models of lamellipodium dynamics [ÖS09], knowledge of the length distribution is required as input datum.

Actin filaments are polar. Almost all so called *barbed (or plus-)ends* are oriented towards the leading edge of the lamellipodium, and polymerization dominates there. A stationary length distribution therefore requires a decomposition mechanism. A candidate is filament severing promoted by the proteins *gelsolin* or *ADF/cofilin*, leading to stable length distributions in mathematical models [EKE98], [EEK98], [EKE00], [EKE01], [Gov07], [RBM<sup>+</sup>08]. An alternative is depolymerization at the minus-ends, known to be up-regulated by ADF/cofilin [CLS<sup>+</sup>97]. However, without coordination between polymerization and depolymerization velocities this either leads to unbounded growth of filaments or to almost extinction of the filament population, leaving an exponential length distribution with a mean filament length of at most a few tens of nm [EKE98]. On the other hand, the computation of metastable length distributions in [HMO07] indicates that not only stationary states might be relevant.

It is the purpose of this work to describe a mechanism based on minus-end depolymerization, which

stabilizes the length distribution. The new ingredient is an interaction between minus-ends of filaments of the same length by competition for a depolymerization agent (DA) (as suggested in [Fla08]). A mathematical model will be derived under the following assumptions:

1. Ensembles of filaments are considered, whose plus-ends are aligned as well as the minus-ends of filaments of the same length.
2. At the plus-ends, polymerization occurs with the same rate for all filaments.
3. At the minus-ends, depolymerization is triggered by DA and occurs in two steps: A) one free molecule of DA binds to a free minus-end; B) one monomer is removed from the filament, and the DA molecule and the minus-end become free again. The main modeling assumption is that this process is limited by the availability of free DA. The sum of the densities of free and bound DA molecules is a prescribed constant.

Among these assumptions, the last one is questionable and definitely a simplification, since it neglects diffusion of free DA molecules. For the other extreme of very fast diffusion of DA, the competition for DA would be global, and the resulting model would not have the desired properties.

Since depolymerization at the minus-ends might be faster than polymerization at the plus-ends, filaments might be completely depolymerized, reducing the total number of filaments. We assume, however, that at the position of the plus-ends a fixed number of nucleation agents (NAs) is available, such that, whenever the filament number becomes smaller, free NAs nucleate new filaments. For actin filaments in lamellipodia, a candidate for the role of NA is the Arp2/3 complex occurring at the leading edge [SXPM99]. Although in this situation the nucleation process involves branching from an already existing filament, the nucleation rate will be assumed to be limited only by the availability of the NA.

In the following section, a discrete model is presented, where the distribution of polymer filaments with respect to the number of monomers is accounted for. A macroscopic continuous limit leads to a nonlinear hyperbolic conservation law for the density of minus-ends, as a function of distance from the plus-ends. The non-standard feature of the problem is a nonlocal boundary condition describing nucleation of new polymers. In Section 3, the theory of LeFloch [LeF88] is employed to deduce existence and uniqueness of an entropy solution of the initial-boundary value problem. The main result is large time convergence to a steady state, which is independent from the initial data and which corresponds to a linearly decreasing length distribution with compact support. A slight generalization of the model, where the total number of filaments can be above a preferred equilibrium, leads to a situation, where the boundary condition has to be interpreted in the generalized sense of Bardos [BLN79]. Existence and uniqueness of a solution is proven in Section 4 by a fixed point iteration. It is also shown that, after finite time, the boundary condition is assumed in the classical sense, which again allows to deduce convergence to the steady state. In Section 5, results of numerical experiments are presented, showing rather complex transient behaviour depending on the initial data.

## 2 Derivation of the model

The one-dimensional variable  $x \geq 0$  stands synonymously for filament length as well as for the position along the main filament direction, where  $x = 0$  denotes the common position of the plus-ends. Denoting the number of monomers per filament length by  $1/l_m$ ,  $l_m$  can be interpreted as the 'length' of a monomer. Possible filament lengths (or minus-end positions) are then given by  $x_j = jl_m$ ,  $j \geq 1$ . The number of filaments of length  $x_j$  at time  $t$  is denoted by  $l_m u_j(t)$ .

The number of filaments with length at least  $x_j$  is given by

$$\eta_j(t) = \sum_{i \geq j} l_m u_i(t)$$

and has the form of a Riemann sum with integrand  $u_i$ ; hence  $u_i$  can be interpreted as the number of filament ends per unit length. Accordingly, the total number of filaments is given by

$$N(t) = \eta_1(t).$$

The minus-end density  $u_j = u_{f,j} + u_{b,j}$  is split into the density  $u_{f,j}(t)$  of free minus-ends and the density  $u_{b,j}(t)$  of minus-ends bound to a DA molecule. We also introduce the density  $\rho_{f,j}(t)$  of free DA molecules at position  $x_j$  and time  $t$ , and note that  $u_{b,j}(t)$  can also be interpreted as the density of bound DA molecules. According to the assumption C) of the introduction, we require

$$\rho_{f,j}(t) + u_{b,j}(t) = \bar{\rho} \quad (1)$$

with a fixed (positive) constant  $\bar{\rho}$ . The dynamics of the minus-end densities is governed by the system

$$\frac{du_{b,j}}{dt} = k_b u_{f,j} \rho_{f,j} - k_d u_{b,j} + k_p (u_{b,j-1} - u_{b,j}), \quad (2)$$

$$\frac{du_{f,j}}{dt} = -k_b u_{f,j} \rho_{f,j} + k_d u_{b,j+1} + k_p (u_{f,j-1} - u_{f,j}), \quad (3)$$

The first terms on the right hand sides describe the binding of free DA molecules and free minus-ends with rate  $k_b$ . The second terms describe the reverse reaction, which also involves depolymerization by one monomer with rate  $k_d$ . The third terms model the elongation of filaments by polymerization by one monomer at the plus-ends with rate  $k_p$ .

We introduce a nondimensionalization of (1)–(3), where  $\bar{\rho}$  is used as reference value for the densities  $\rho_{f,j}$ ,  $u_{b,j}$ , and  $u_{f,j}$ . A macroscopic length scale  $l$ , much bigger than the monomer length  $l_m$ , is used as reference length. The reference time is  $l/(l_m k_d)$ . Using the same symbols for the dimensionless variables, we arrive at the scaled system

$$\rho_{f,j} + u_{b,j} = 1, \quad (4)$$

$$\varepsilon \frac{du_{b,j}}{dt} = \alpha u_{f,j} \rho_{f,j} - u_{b,j} + \varepsilon v_p \frac{u_{b,j-1} - u_{b,j}}{\varepsilon}, \quad (5)$$

$$\varepsilon \frac{du_{f,j}}{dt} = -\alpha u_{f,j} \rho_{f,j} + u_{b,j} + \varepsilon \frac{u_{b,j+1} - u_{b,j}}{\varepsilon} + \varepsilon v_p \frac{u_{f,j-1} - u_{f,j}}{\varepsilon}, \quad (6)$$

with the dimensionless parameters

$$\alpha = \frac{k_b \bar{\rho}}{k_d}, \quad v_p = \frac{k_p}{k_d}, \quad \varepsilon = \frac{l_m}{l} \ll 1. \quad (7)$$

Note that, for the scaled minus-end positions,  $x_{j+1} - x_j = \varepsilon$  holds. The scaling assumption that the rates of the binding reaction, of the depolymerization reaction, and of the polymerization reaction are of the same order of magnitude, implies that the scaled binding reaction  $\alpha$  and the scaled polymerization speed  $v_p$  take moderate values.

As a preparation for the continuous limit  $\varepsilon \rightarrow 0$ , we state the equation

$$\frac{du_j}{dt} + v_p \frac{u_j - u_{j-1}}{\varepsilon} - \frac{u_{b,j+1} - u_{b,j}}{\varepsilon} = 0, \quad (8)$$

for the total minus-end density

$$u_j = u_{b,j} + u_{f,j}. \quad (9)$$

Assuming that there are smooth functions  $\rho_f(t, x)$  and  $u(t, x) = u_b(t, x) + u_f(t, x)$  such that  $\rho_{f,j}(t)$  approximates  $\rho_f(t, x_j)$  for small  $\varepsilon$  and analogously for the other variables, the formal limit  $\varepsilon \rightarrow 0$  of (4), (9), (5), (8) leads to the problem

$$\begin{aligned} \rho_f + u_b &= 1, & u_f + u_b &= u, & u_b &= \alpha u_f \rho_f, \\ \partial_t u + \partial_x (v_p u - u_b) &= 0. \end{aligned}$$

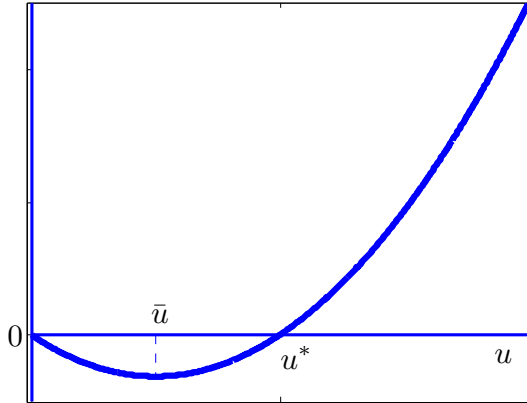


Figure 1: The graph of the flux function  $f$ . It is smooth, convex, with two roots  $0, u^*$  and a minimum  $\bar{u} \in (0, u^*)$ .

Computing  $u_b$  as a function of  $u$  from the first three equations, the problem can be reduced to a nonlinear hyperbolic conservation law:

$$\partial_t u + \partial_x f(u) = 0, \quad \text{with } f(u) = v_p u - \frac{1}{2} \left( u + \frac{1}{\alpha} + 1 - \sqrt{\left( u + \frac{1}{\alpha} + 1 \right)^2 - 4u} \right). \quad (10)$$

The flux function  $f$  has the properties

$$f'' > 0, \quad f'(0) = v_p - \frac{\alpha}{\alpha + 1}, \quad f(u) = v_p u - 1 + O(u^{-1}) \text{ as } u \rightarrow \infty,$$

and a typical graph is presented in Figure 1.

The last property shows that polymerization dominates for large minus-end densities. On the other hand, we assume that depolymerization wins for small values of  $u$ :

$$v_p < \frac{\alpha}{\alpha + 1}. \quad (11)$$

In this case,

$$f(0) = f'(\bar{u}) = f(u^*) = 0 \quad \text{for } 0 < \bar{u} = 1 - \frac{1}{\alpha} + \frac{1 - 2v_p}{\sqrt{\alpha v_p(1 - v_p)}} < u^* = \frac{\alpha - v_p(1 + \alpha)}{\alpha v_p(1 - v_p)}. \quad (12)$$

*Remark 1.* Before proceeding, we note that due to (7), relation (11) yields a condition on the dimensional rates  $k_p, k_d, k_b$  that govern the competitive dynamics of polymerization and depolymerization. For this discussion we refer to the last paragraph of this work.

*Remark 2.* The microscopic details of the derivation of the model could be skipped and replaced by the ansatz  $f(u) = v_p u - v_d(u)u$  for the flux function, where the depolymerization speed  $v_d(u)$  is a decreasing function of the minus-end density, with the assumptions that  $f$  is strictly convex and that  $v_d(0) > v_p$ .

**Nucleation – auxiliary conditions:** The total number of filaments is given by

$$N(t) = \int_0^\infty u(t, x) dx.$$

The plus-ends are assumed to be tied to nucleation agents (NA), whose total number  $\bar{N} > 0$  is fixed, implying  $N(t) \leq \bar{N}$ . We assume that free NA nucleate new filaments with a rate  $1/\tau_n$ , which can be translated as a flux of minus-ends at  $x = 0$ . In the nondimensionalization above, the choice of the length scale  $l$  has not been fixed so far. We make up for this by requiring the characteristic time  $l/(l_m k_d)$  to be equal to  $\tau_n$ , i.e.  $l = l_m k_d \tau_n$ . The small parameter can now be written as  $\varepsilon = 1/(k_d \tau_n)$  with the interpretation that the nucleation process is much slower than the depolymerization reaction. In the nondimensionalized setting, our considerations result in the nonlocal boundary condition

$$f(u(t, 0)) = \bar{N} - \int_0^\infty u(t, x) dx.$$

Since the right hand side is nonnegative and  $f : [u^*, \infty) \rightarrow [0, \infty)$  is strictly increasing and possesses the inverse  $\varphi$ , the boundary condition can be written as

$$u(t, 0) = \varphi\left(\bar{N} - \int_0^\infty u(t, x) dx\right) \geq u^*, \quad t > 0. \quad (13)$$

The problem (10), (13) is completed by prescribing initial data:

$$u(0, x) = u_0(x) \geq 0, \quad x > 0. \quad (14)$$

Assuming that  $u_0$  has compact support, the same is true for  $u(t, \cdot)$  for  $t > 0$  by the finite propagation speed. By integration of the conservation law (10) with respect to  $x$  (for a solution with compact support) and by using the boundary condition, the simple ODE

$$\frac{dN}{dt} = \bar{N} - N \quad (15)$$

is derived. Together with the initial condition (14) this implies

$$N(t) = \bar{N} + \left(\int_0^\infty u_0(x) dx - \bar{N}\right) e^{-t}. \quad (16)$$

### 3 Existence, uniqueness, and convergence to a steady state

If the formal computations at the end of the previous section are justified, then the problem reduces to a standard initial-boundary value problem with prescribed boundary data

$$u(t, 0) = u_b(t) := \varphi(\bar{N} - N(t)), \quad (17)$$

with  $N(t)$  given by (16) and, consequentially,  $u_b(t) \rightarrow u^*$  exponentially fast as  $t \rightarrow \infty$ .

The justification of (17) is nontrivial, however, since solutions of hyperbolic conservation laws cannot be expected to assume given boundary values in general. The generally accepted solution concept, which can be justified by a vanishing viscosity approach, is due to [BLN79]. It states that the boundary condition should be replaced by

$$\begin{aligned} u(t, 0+) &= b_m(t) := \max\{\bar{u}, u_b(t)\}, \\ \text{or} \quad f'(u(t, 0+)) &\leq 0 \quad \text{and} \quad f(u(t, 0+)) \geq f(b_m(t)), \end{aligned}$$

where  $u(t, 0+)$  denotes the trace of  $u$  at  $x = 0$ . We note that the max that appears in the first line does not constitute a restriction, and that the second line describes the situation, where the boundary condition  $b_m(t)$  is not assumed by the solution.

With the properties of the flux function  $f$  and of the boundary datum  $u_b$  given in the previous section, the above conditions always reduce to (17). This is easily seen, since  $u_b > u^*$  so  $b_m = u_b$ ; hence  $f(b_m) \geq 0$  (cf Figure 1) and the second line of the boundary condition is not valid.

By the boundedness and continuity of  $u_b$ , the results of [LeF88] imply existence and uniqueness of a global entropy solution of (10), (14), (16), (17) and, thus, of (10), (13), (14).

As a preparation for the long time asymptotics, we show that  $u$  can be obtained as the solution of a Cauchy problem. However, this process typically cannot be started at  $t = 0$ , but only at a large enough time  $T_1$ . The backward characteristic

$$x = (t - \tau)f'(u_b(\tau))$$

through the point  $(t, x) = (\tau, 0)$ ,  $\tau > T_1$ , carrying the value  $u_b(\tau)$  intersects the line  $t = T_1$  at

$$x = Y(\tau) := (T_1 - \tau)f'(u_b(\tau)) < 0.$$

The derivative  $Y'(\tau) = -f'(u_b(\tau)) + (T_1 - \tau)f''(u_b(\tau))u'_b(\tau)$  can be estimated by

$$Y'(\tau) \leq -f'(u_b(\tau)) + c\tau|u'_b(\tau)| \xrightarrow{\tau \rightarrow \infty} -f'(u^*) < 0,$$

because of the exponential decay of  $u'_b$ . The constant  $c$  is a bound for  $f''$  on  $[\bar{u}, u^*]$ . This implies that  $Y$  is strictly monotonically decreasing on  $[T_1, \infty)$  for large enough  $T_1$ .

The appropriate definition for the Cauchy data therefore is

$$u_C(x) := \begin{cases} u_b(Y^{-1}(x)) & \text{for } x < 0 \\ u(T_1, x) & \text{for } x > 0. \end{cases}$$

In other words, the solution of the initial-boundary value problem for  $t \geq T_1$  can be computed by solving the Cauchy problem

$$\partial_t u + \partial_x f(u) = 0, \quad \text{for } t > T_1, \quad u(T_1, x) = u_C(x), \quad (18)$$

and by restricting the solution to  $x > 0$ .

Since  $u_b(t) - u^* = O(e^{-t})$  as  $t \rightarrow \infty$ ,

$$Y(t) \approx -tf'(u^*) \quad \text{as } t \rightarrow \infty.$$

As a consequence,

$$u_C(x) - u^* = O\left(\exp\left(\frac{x}{f'(u^*)}\right)\right) \quad \text{as } x \rightarrow -\infty. \quad (19)$$

The study of long time asymptotics for nonlinear hyperbolic conservation laws has a long history (see, e.g., [Liu77]). A typical result for Cauchy problems is the following.

**Lemma 1.** *Let  $f'' > 0$ ,  $f(0) = f(u^*) = 0$  with  $0 < u^*$ , let*

$$u_\infty(x) = \begin{cases} u^* & \text{for } x < x_\infty, \\ 0 & \text{for } x > x_\infty, \end{cases} \quad (20)$$

let  $u_I - u_\infty \in L^1(\mathbb{R}) \cap L^\infty(\mathbb{R})$ , and let

$$\int_{\mathbb{R}} (u_I - u_\infty) dx = 0.$$

Then the solution  $u$  of (18) for  $u_C = u_I$  satisfies

$$\lim_{t \rightarrow \infty} \|u(t, \cdot) - u_\infty\|_{L^1(\mathbb{R})} = 0.$$

Since the restriction of  $u$  to  $x > 0$  solves the initial-boundary value problem (10)–(14),  $x_\infty$  can be determined from the requirement

$$\bar{N} = N(\infty) = \int_0^\infty u_\infty dx = u^* x_\infty.$$

We collect the results of this section:

**Theorem 2.** *Let (11) hold, let  $0 \leq u_0 \in L^1(\mathbb{R}^+) \cap L^\infty(\mathbb{R}^+)$  have compact support and satisfy*

$$\int_0^\infty u_0 dx \leq \bar{N}. \quad (21)$$

*Let  $\varphi$  be the inverse of  $f : [u^*, \infty) \rightarrow [0, \infty)$ . Then the problem (10)–(14) has a global unique entropy solution  $u$  satisfying*

$$\lim_{t \rightarrow \infty} \|u(t, \cdot) - u_\infty\|_{L^1(\mathbb{R}^+)} = 0,$$

*with  $u_\infty$  defined in (20) with  $x_\infty = \bar{N}/u^*$ .*

## 4 A generalized model

In this section, we consider the more general case, in which  $N(t)$  might be bigger than a preferred number of filaments. Such cases of oversaturation allow for situations in which, differently from above, the prescribed boundary conditions are not assumed at all times. This complicates the analysis significantly.

Consider the problem

$$\partial_t u + \partial_x f(u) = 0, \quad x, t > 0, \quad (22)$$

subject to the boundary condition

$$u(t, 0+) = u_b(t) := \psi \left( \int_0^\infty u(t, x) dx \right), \quad (23)$$

$$\text{or } f'(u(t, 0+)) \leq 0 \quad \text{and} \quad f(u(t, 0+)) \geq f(u_b(t)), \quad t > 0, \quad (24)$$

and to the initial condition

$$u(0, x) = u_0(x), \quad x > 0, \quad (25)$$

where the data satisfy

$$f \text{ smooth, } f'' > 0, \quad f(0) = f'(\bar{u}) = f(u^*) = 0, \quad f'(0) < 0 \quad \text{with } 0 < \bar{u} < u^*, \quad (26)$$

$$\psi : [0, \infty) \rightarrow [\bar{u}, \infty) \quad \text{Lipschitz, non-increasing,}$$

$$C^1 \text{ in a neighborhood of } \bar{N} > 0, \quad \psi(\bar{N}) = u^*, \quad \psi'(\bar{N}) < 0, \quad (27)$$

$$0 \leq u_0 \in L^\infty(\mathbb{R}^+) \quad \text{with compact support.} \quad (28)$$

### Existence and uniqueness

The essential difference to the previous section, where  $\bar{N}$  was the maximum number of filaments the cell can sustain, in this section  $\bar{N}$  represents the preferable number of filaments, i.e. we do not assume (21) any more. This in turn allows for  $u_b(t) < u^*$  and, thus, makes the alternative (24) of the boundary condition possible. Note that, if (23) holds, the total number of filaments solves the ODE

$$\frac{dN}{dt} = f(\psi(N)), \quad N(t) = \int_0^\infty u(t, x) dx, \quad (29)$$

which – similarly to (15) – has the unique, globally attractive steady state  $N = \bar{N}$ . However, since (23) cannot be guaranteed,  $N(t)$  cannot be computed a priori by solving (29), so the existence and uniqueness of a solution is not a simple consequence of [LeF88] any more.

**Theorem 3** (Global existence and uniqueness). *Let (26)–(28) hold. Then the problem (22)–(25) has a unique entropy solution  $u \in C([0, \infty); L^1(\mathbb{R}^+))$ , such that  $u(t, \cdot)$  has compact support for every  $t \geq 0$ , and*

$$0 \leq u(t, x) \leq \max \left\{ \psi(0), \sup_{\mathbb{R}^+} u_0 \right\}, \quad \forall t, x > 0.$$

*Proof.* It suffices to prove local existence and uniqueness of a solution with the properties stated in the theorem, which imply that, for  $t > 0$ , the solution satisfies the assumptions (28) on the initial data and therefore can be continued.

For a given  $T > 0$  we consider the Banach space

$$\mathcal{X}_T = C([0, T], L^1(\mathbb{R}^+)), \quad \text{equipped with} \quad \|u\|_{\mathcal{X}_T} = \max_{t \in [0, T]} \|u(t, \cdot)\|_{L^1(\mathbb{R}^+)},$$

and its closed subset

$$\mathcal{S}_T = \left\{ u \in \mathcal{X}_T \mid 0 \leq u \leq u_{max} \right\}, \quad u_{max} := \max \left\{ \psi(0), \sup_{\mathbb{R}^+} u_0 \right\}.$$

We define a fixed point operator  $\mathcal{P} : \mathcal{S}_T \rightarrow \mathcal{S}_T$  by  $\mathcal{P}(v) = u$ , where  $u$  solves (22)–(25) with the definition of the boundary data in (23) replaced by

$$u_b[v](t) := \psi \left( \int_0^\infty v(t, x) dx \right).$$

By (27) and  $v \in \mathcal{X}_T$ ,  $u_b[v]$  is continuous and bounded. This, together with (26)–(28) suffices for the application of Theorems 2.1 and 2.2 of [LeF88], guaranteeing that  $\mathcal{P}(v) \in \mathcal{X}_T$  is well defined and that  $\mathcal{P}$  maps  $\mathcal{S}_T$  into itself.

Theorem 2.2 of [LeF88] also provides the stability estimate

$$\|\mathcal{P}(v_1)(t, \cdot) - \mathcal{P}(v_2)(t, \cdot)\|_{L^1(\mathbb{R}^+)} \leq \int_0^t |f(u_b[v_1](s)) - f(u_b[v_2](s))| ds \leq TL_f L_\psi \|v_1 - v_2\|_{\mathcal{X}_T},$$

where  $L_f$  is the Lipschitz constant of  $f$  on  $[\bar{u}, \psi(0)]$  and  $L_\psi$  is the Lipschitz constant of  $\psi$ . This proves that  $\mathcal{P}$  is a contraction for small enough  $T$  and, thus, completes the existence and uniqueness proof.

Finally, by the finite speed of propagation, if  $\text{supp}(u_0) \subset [0, \bar{x}]$  then

$$\text{supp}(u(t, \cdot)) \subset [0, \bar{x} + f'(u_{max})t].$$

□

## Convergence to the steady state

For proving decay to equilibrium, it will suffice to show that the boundary data are assumed for large enough times, i.e. that there exists  $T > 0$  such that (23) holds for  $t > T$ . Then the strategy of the previous section can be applied.

We shall need more details concerning the solution of the initial-boundary value problem. The basic idea of [LeF88] is to use the explicit representation of the entropy solution due to Lax [Lax57] for an equivalent Cauchy problem as derived in the previous section. Complications arise, however, due to the different alternatives in the boundary conditions.



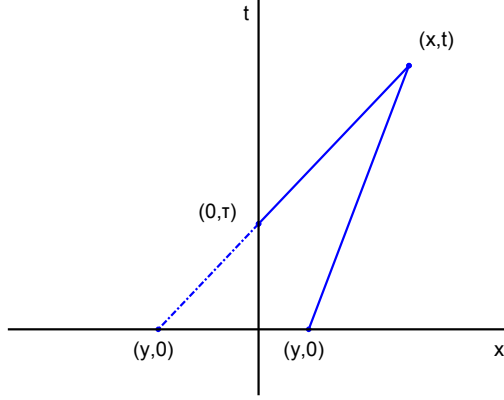


Figure 2: For a given  $(x, t)$  the minimizer  $y$  of  $G(y, t, x)$  can be  $y \geq 0$  or  $y \leq 0$ . In both cases the slope of the line connecting  $(x, t)$  with  $(y, 0)$  is given by the solution formula:  $u(t, x) = (f')^{-1}\left(\frac{x-y}{t}\right)$ ,  $y \in \mathbb{R}$ , and reads  $\frac{t}{x-y} = \frac{1}{f'(u(t, x))}$ . In the case  $y \leq 0$ ,  $\tau$  represents the intersection point of the straight line  $(x, t) - (y, 0)$  with the boundary  $x = 0$ ; that is  $\frac{x-y}{t} = \frac{x}{t-\tau}$ .

We shall need the Legendre transform

$$g(v) = \sup_{u \in \mathbb{R}} (uv - f(u)) = vb(v) - f(b(v)), \quad b := (f')^{-1},$$

of the flux function. As a consequence of (26),  $g$  has the properties

$$g \text{ smooth}, \quad g'' > 0, \quad g(c) = g'(c) = 0, \quad \text{for } c := f'(0) < 0, \quad g'(v) = b(v). \quad (30)$$

The solution of (22)–(25) can then be written as ([LeF88], see also [JG91])

$$u(t, x) = b\left(\frac{x - y(t, x)}{t}\right), \quad \text{with } y(t, x) = \operatorname{argmin}_{y \in \mathbb{R}} G(y; t, x),$$

where

$$G(y; t, x) = \begin{cases} \int_0^y u_0(\xi) d\xi + t g\left(\frac{x-y}{t}\right), & y \geq 0, \\ -\int_0^{yt/(y-x)} \mathcal{Y}(s) ds + \frac{xt}{x-y} g\left(\frac{x-y}{t}\right), & y \leq 0. \end{cases}$$

The definition of the function  $\mathcal{Y}$  involves the solution of a variational inequality. Its details will not be needed in the following. It is interesting to note, however, that  $\mathcal{Y}(t) = f(u(t, 0+))$  holds for the solution.

It is easily seen that, for continuous  $u_0$  and  $y(t, x) > 0$ ,  $u(t, x) = u_0(y(t, x))$  holds. Similarly, if the boundary data are assumed (i.e. (23) holds) and continuous then, for  $y(t, x) < 0$ ,  $u(t, x) = u_b(\tau(t, x))$  holds with

$$\frac{x - y(t, x)}{t} = \frac{x}{t - \tau(t, x)}.$$

For a fixed  $(t, x)$  we produce a curve  $C_{t,x} = \{(\tau, y_\tau(t, x)) : 0 \leq \tau \leq t\}$  connecting  $y_0(t, x) = y(t, x)$  and  $y_t(t, x) = x$  by replacing  $t$  by  $t - \tau$  and  $u_0$  by  $u(\tau, \cdot)$  in the definition of  $G$  and computing the minimizers, and we refer to Figure 2 for a graphical explanation.

We call  $C_{t,x}$  the *backward characteristic* through  $(t, x)$  emanating from  $y(t, x)$ . For smooth solutions backward characteristics are classical characteristics with equation  $x = y + tf'(u_0(y))$  if  $y > 0$ , or  $x = (t - \tau)f'(u_b(\tau))$  if  $y < 0$ .

We start by proving that for large enough  $t$  and for arbitrary  $x > 0$ ,  $C_{t,x}$  cannot emanate from the support of  $u_0$ .

**Lemma 4.** *Let (26)–(28) hold and  $\text{supp}(u_0) \subset [0, \bar{x}]$ . Then there exists  $T_2 > 0$ , such that  $y(t, x) \notin (0, \bar{x})$  for all  $x > 0$ ,  $t > T_2$ .*

*Proof.* Let  $t, x > 0$  be such that  $0 \leq \eta := y(t, x) < \bar{x}$ . Then, in view of (30),

$$\frac{x - \eta}{t} > -\frac{\bar{x}}{t} > c,$$

for  $t$  large enough. Since, again by (30),  $g$  is increasing on  $[c, \infty)$  and  $g(0) = -f(\bar{u}) > 0$ , there exists  $\bar{t} > 0$  (independent of  $x$ ) such that

$$g\left(\frac{x - \eta}{t}\right) \geq g\left(-\frac{\bar{x}}{t}\right) \geq -\frac{f(\bar{u})}{2} \quad \text{for every } t \geq \bar{t}.$$

This in turn implies that for every  $t \geq \bar{t}$

$$G(\eta; t, x) \geq -\frac{f(\bar{u})}{2} t, \tag{31}$$

since  $\eta = y(t, x) \geq 0$ . Moreover, since  $x - tc > 0$  and since  $\eta$  is the minimizer of  $G(\cdot; t, x)$ ,

$$G(\eta; t, x) \leq G(x - tc; t, x) = \int_0^{x-tc} u_0(\xi) d\xi + tg(c) \leq K, \tag{32}$$

where the last inequality stems from (28), using  $g(c) = 0$  and  $K = \text{diam}(\text{supp}(u_0)) \|u_0\|_{L^\infty}$ .

Combination of (31) and (32) implies that we arrive at a contradiction for  $t > T_2 := \max\{\bar{t}, -2K/f(\bar{u})\}$ , which completes the proof.  $\square$

**Lemma 5.** *Let the assumptions of Lemma 4 hold. Then (23) holds for  $t > T_2$  with  $T_2 > 0$  from Lemma 4.*

*Proof.* We distinguish between two cases:

**Case 1:** Assume for a  $t \geq 0$ ,  $N(t) < \bar{N}$ . Then, by the continuity of  $N(t)$ , this remains true at least for a small time interval. Since, in this interval,  $u_b(t) > u^*$ , (23) and, consequently, the ODE (29) hold there. This in turn implies that  $N(t) < \bar{N}$  and, thus, (23) remains true for all times. This is actually the setting of the previous section.

**Case 2:** Assume that for a  $t > T_2$ ,  $N(t) \geq \bar{N}$  holds. For  $x \in (0, \bar{N}/u_{max})$  (where  $u_{max}$  is given in Theorem 3 and is an upper bound of the solution  $u$ ) assume  $y(t, x) > \bar{x}$ . Then, since

$$G(y; t, x) = \int_0^\infty u_0(\xi) d\xi + tg\left(\frac{x - y}{t}\right)$$

in a neighbourhood of  $y = y(t, x)$ , the minimum satisfies  $dG/dy = 0$ ; hence

$$u(t, x) = b\left(\frac{x - y(t, x)}{t}\right) = 0.$$

By the monotonicity of  $y(t, x)$  with respect to  $x$  (Proposition 2.4 in [LeF88]),  $u(t, x') = 0$  holds for  $x' \geq x$ . As a consequence  $N(t) \leq xu_{max} < \bar{N}$  in contradiction to the basic assumption of this case. Therefore,  $y(t, x) \leq 0$  follows from Lemma 4, implying  $\frac{x-y(t,x)}{t} > 0$  and since  $b \nearrow$  in  $[0, \infty)$ , it reads

$$u(t, x) = b\left(\frac{x-y(t, x)}{t}\right) > b(0) = \bar{u} \quad \text{for } x < \bar{N}/u_{max}.$$

The consequence  $u(t, 0+) \geq \bar{u}$  excludes (24), completing the proof.  $\square$

For proving convergence to the steady state it suffices to repeat the argument of the preceding section with a  $T_1$  satisfying the additional requirement  $T_1 \geq T_2$ .

**Theorem 6.** *Let (26)–(28) hold. Then the entropy solution  $u$  of (22)–(25) satisfies*

$$\lim_{t \rightarrow \infty} \|u(t, \cdot) - u_\infty\|_{L^1(\mathbb{R}^+)} = 0,$$

with  $u_\infty$  defined in (20) with  $x_\infty = \bar{N}/u^*$ .

*Remark 3.* The steady state solution is independent from the initial conditions. With respect to the flux function, it depends only on its root, and with respect to the boundary condition it depends only on the ‘preferred’ number of filaments  $\bar{N}$ .

## 5 Numerical experiments

For the numerical tests we note that the finite speed of propagation and the convergence of the solution  $u$  to a steady state with bounded support guarantee that  $\mathcal{S} = \bigcup_{t \geq 0} \text{supp}(u(t, \cdot)) \subset \mathbb{R}^+$  is finite. So, for the needs of the numerical tests we shall consider a finite computational domain  $[0, x_{num}]$  satisfying  $[0, x_{num}] \supset \mathcal{S}$ .

We discretize the spatial computational domain  $[0, x_{num}]$  by a fixed-in-time, uniform-in-space grid with  $N_x$  points:

$$\mathcal{D}_x = \left\{ x_i = (i-1)\Delta x \mid i = 1, \dots, N_x, \Delta x = \frac{x_{num}}{N_x - 1} \right\}.$$

The temporal domain is discretized as

$$\mathcal{D}_t = \left\{ t^n = t^{n-1} + \Delta t^n \mid n = 1, \dots, N_t, t^0 = 0 \right\},$$

where the time steps  $\Delta t^n$  are chosen such that the CFL condition is satisfied. The maximum time step  $N_t$  is chosen sufficiently large so that the problem under consideration reaches its steady state.

The numerical solution at the time step  $t^n$  is denoted by

$$U^n = \left\{ u_i^n \mid i = 1, \dots, N_x \right\},$$

where  $u_i^n$  are discrete approximations of the cell averages of the exact solution, i.e

$$u_i^n \approx \frac{1}{\Delta x} \int_{x_i}^{x_{i+1}} u(x, t^n) dx, \quad i = 1, \dots, N_x, \quad n = 1, \dots, N_t.$$

The numerical solution is computed as follows:

- For  $i = 1$ , the computation of the numerical solution, depends on the boundary conditions and is described in the paragraphs that follow.

- For  $i = 2, \dots, N_x - 1$ , the numerical solution is computed using an explicit in time conservative Finite Volume scheme

$$u_i^{n+1} = u_i^n - \frac{\Delta t^n}{\Delta x} \left( F_{i+1/2}^n - F_{i-1/2}^n \right).$$

For the computation of the numerical flux we use the superbee flux limiter:

$$\begin{aligned} F_{i+1/2}^n &= F_{i+1/2}^{n,LxF} - \rho(r_i^n) \left( F_{i+1/2}^{n,LxF} - F_{i+1/2}^{n,Rchtmr} \right) \\ F_{i-1/2}^n &= F_{i-1/2}^{n,LxF} - \rho(r_{i-1}^n) \left( F_{i-1/2}^{n,LxF} - F_{i-1/2}^{n,Rchtmr} \right) \end{aligned}$$

where the flux limiter function  $\rho$  is defined by

$$\begin{aligned} r_i^n &= \frac{u_i^n - u_{i-1}^n}{u_{i+1}^n - u_i^n} \\ \rho(r) &= \max\{0, \min\{2r, 1\}, \min\{r, 2\}\} \end{aligned}$$

and the numerical fluxes used as building block are the Lax-Friedrichs flux and the Richtmyer flux

$$\begin{aligned} F_{i+1/2}^{n,LxF} &= \frac{f(u_i^n) + f(u_{i+1}^n)}{2} - \frac{\Delta x}{2\Delta t^n} (u_{i+1}^n - u_i^n), \\ F_{i+1/2}^{n,Rchtmr} &= f \left( \frac{u_i^n + u_{i+1}^n}{2} - \frac{\Delta t^n}{2\Delta x} (f(u_i^n) - f(u_{i+1}^n)) \right). \end{aligned}$$

For more details on flux limiters, we refer to [LeV04].

- For  $i = N_x$ , we use that the computational domain satisfies  $[0, x_{num}] \supset \mathcal{S}$  and we set

$$u_{N_x}^{n+1} = u_{N_x}^n.$$

In the following numerical tests, the spatial mesh is discretized using  $N_x = 2000$  grid points, and the time steps  $\Delta t^n$  are chosen such as the Courant number is 0.9

$$\frac{\Delta t^n}{\Delta x} \max_i |f'(u_i^n)| = 0.9.$$

**Numerical test 1.** This numerical test refers to the initial model as in Theorem 2. In this case the evolution of the total mass  $N(t)$  is given by (16), and the boundary condition by (17). So at every time step  $t^{n+1}$  we compute

$$N^{n+1} = N(t^{n+1}),$$

and  $u_1^{n+1}$  by solving the nonlinear problem

$$f(u_1^{n+1}) = \bar{N} - N^{n+1}.$$

So the numerical scheme is

$$\text{Sch1} :: \begin{cases} f(u_1^{n+1}) = \bar{N} - N^{n+1}, & i = 1, \\ u_i^{n+1} = u_i^n - \frac{\Delta t}{\Delta x} \left( F_{i+1/2}^n - F_{i-1/2}^n \right), & i = 2, \dots, N_x - 1 \\ u_{N_x}^{n+1} = u_{N_x}^n, & i = N_x \end{cases} .$$

The parameters used in this case are

$$\text{Par1} :: \begin{cases} u_0(x) = 0, & x > 0, \\ f(u) = u(u - 2), & (u^* = 2) \\ \bar{N} = 30 \end{cases} .$$

We refer to Figure 3 for a graphical representation of this test case.

**Numerical test 2.** This numerical test refers to the generalized model case as summarized in Theorem 6.

In this case the boundary condition is given by (23) or (24). At every time step  $t^n$  we compute the boundary function  $u_b$  by using (23) and the trace  $u_{tr}$  by extrapolation of the values  $u_3^n, u_4^n$  to the boundary, i.e

$$u_b(t^n) = \psi \left( \Delta x \sum_{i=1}^{N_x} u_i^n \right),$$

$$u_{tr}(t^n) = 3u_3^n - 2u_4^n$$

respectively. We then use (24) for  $u_b(t^n)$  and  $u_{tr}(t^n)$  to decide whether the boundary function value is attained, and we set

$$u_1^{n+1} = \begin{cases} u_b(t^n), & \text{if } u_b(t^n) \text{ is assumed} \\ u_1^n - \frac{\Delta t^n}{\Delta x} (f(u_2^n) - f(u_1^n)), & \text{if } u_b(t^n) \text{ is not assumed} \end{cases}.$$

The numerical scheme is

$$\text{Sch2} :: \begin{cases} u_1^{n+1}, & i = 1, \\ u_i^{n+1} = u_i^n - \frac{\Delta t}{\Delta x} (F_{i+1/2}^n - F_{i-1/2}^n), & i = 2, \dots, N_x - 1 \\ u_{N_x}^{n+1} = u_{N_x}^n, & i = N_x \end{cases}.$$

The parameters used are

$$\text{Par2} :: \begin{cases} u_0(x) = 0, & x > 0, \\ f(u) = e^u - \frac{e^2 - 1}{2} u - 1, & (u^* = 2) \\ \psi = 2 \exp \left( \frac{\bar{N} - \int_0^\infty u(t,x) dx}{20} \right), \\ \bar{N} = 10 \end{cases}.$$

We refer to Figure 4 for a graphical representation of this test case.

**Numerical test 3.** In this numerical test, we examine the dependence of the steady state solutions on the polymerization/depolymerization rate ratio  $v_p$ .

We use (7), and assume that the binding/depolymerization ratio  $\alpha = \frac{k_b \bar{p}}{k_d}$  is constant, while we allow for variable  $v_p = \frac{k_p}{k_d}$ . The numerical scheme that we use is the same as in the Numerical test 1 case, with flux function given by the modeling approach (10)

$$f(u) = v_p u - \frac{1}{2} \left( u + \frac{1}{\alpha} + 1 - \sqrt{\left( u + \frac{1}{\alpha} + 1 \right)^2 - 4u} \right).$$

The flux functions that we consider satisfy the conditions (12), more specifically their root  $u^*$  depends on  $\alpha$  and  $v_p$  as

$$u^* = \frac{\alpha - v_p(1 + \alpha)}{\alpha v_p(1 - v_p)}.$$

The respective steady state profiles are given by (20) and Theorem 2, namely

$$u_\infty(x) = \begin{cases} u^* & \text{for } x < x_\infty, \\ 0 & \text{for } x > x_\infty, \end{cases},$$

with  $x_\infty = \frac{\bar{N}}{u^*}$ .

We also note, for the curve  $(x_\infty, u^*)$ , that  $\lim_{v_p \rightarrow \frac{\alpha}{\alpha+1}} u^* = 0$ ; hence  $\lim_{v_p \rightarrow \frac{\alpha}{\alpha+1}} x_\infty = \infty$ , and that  $\lim_{v_p \rightarrow 0} u^* = \infty$ ; hence  $\lim_{v_p \rightarrow 0} x_\infty = 0$ .

For this test, we choose  $\bar{N} = 100$ ,  $\alpha = 1$  and we compute the steady state solutions for various values of  $v_p$  that satisfy the restriction (11)

$$v_p < \frac{\alpha}{\alpha + 1} = \frac{1}{2}.$$

The results of this test are presented in Figure (5).

**Numerical test 4.** In this numerical test we examine the dependence of the steady state solution on the binding/depolymerization rate ratio  $\alpha$ , while keeping  $v_p$  fixed. The computational setting (numerical scheme, flux function) is as described in the paragraph **Numerical test 3**.

We note, due to (11), that

$$0 < v_p < 1, \quad \alpha > \frac{v_p}{1 - v_p},$$

and due to (12) that

$$u^* = \frac{1}{v_p} - \frac{1}{1 - v_p} \frac{1}{\alpha}$$

So

$$u^* \longrightarrow \begin{cases} \frac{1}{v_p}, & \text{as } \alpha \rightarrow +\infty \\ 0, & \text{as } \alpha \rightarrow \frac{v_p}{1 - v_p} \end{cases}$$

and the maximum length of filaments

$$x_\infty = \frac{\bar{N}}{u^*} \longrightarrow \begin{cases} \bar{N} v_p, & \text{as } \alpha \rightarrow +\infty \\ +\infty, & \text{as } \alpha \rightarrow \frac{v_p}{1 - v_p} \end{cases}$$

For this test, we choose  $\bar{N} = 100$ ,  $v_p = \frac{1}{4}$  and we compute the steady state solutions and respective length distributions, for various values of  $\alpha$  that satisfy

$$\alpha > \frac{v_p}{1 - v_p} = \frac{1}{3}.$$

The results of this test are presented in Figure (6).

## 6 Conclusions

With the model derived in Section 2, the dimensional version of the stationary length distribution

$$\eta_\infty(x) = \int_x^\infty u_\infty(x) dx$$

is given by

$$\eta_\infty(x) = \max \left\{ \bar{N} - x \frac{k_d [k_b \bar{\rho} k_d - k_p (k_d + k_b \bar{\rho})]}{k_b k_p (k_d - k_p)}, 0 \right\}. \quad (33)$$

The model is valid under the assumption (11) on the parameters, which can be rewritten as the condition

$$\frac{1}{k_p} > \frac{1}{k_b \bar{\rho}} + \frac{1}{k_d},$$

on the relaxation time for polymerization on the one hand, and the sum of the relaxation times for binding of free DA and for depolymerization on the other hand. Note however that for the binding reaction,  $1/(k_b\bar{\rho})$  is the minimal relaxation time since  $\bar{\rho}$  is the maximal density of free DA molecules. For smaller densities of free DA molecules, the relation can be reversed. This competition between polymerization and depolymerization is the essence of our model. Note that if a limit is approached, where the above inequality becomes an equality, the maximal filament length

$$x_\infty = \frac{\bar{N}k_b k_p (k_d - k_p)}{k_d [k_b \bar{\rho} k_d - k_p (k_d + k_b \bar{\rho})]}$$

(and therefore also the mean filament length  $x_\infty/3$ ) tends to infinity by the dominance of the polymerization effect.

A mathematical question left open in this work is the validity of the macroscopic and fast reaction limit of Section 2. Some evidence has been collected by using a time discretization of the microscopic model (4)–(8) as an alternative numerical method. Without presenting the detailed results, we report that both numerical methods produced virtually the same results for several test problems.

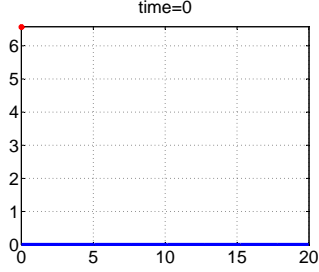
A result like (33) can in principle be used as input, when modeling actin dynamics in lamellipodia [ÖS09] under the assumption that the dynamic processes described here are fast compared to other effects in the cytoskeleton. Of course, several effects have been neglected (see, e.g. [MEK02]), like other disassembly mechanisms and the details and limitations of the polymerization process. On the other hand, a very precise modeling of in vivo length distributions seems not to be feasible anyway, since both the available experimental data and our theoretical knowledge are rather incomplete. An example of the former is Fig. 12 in [EKE01], where the fluctuations in the experimental data do not seem to permit a conclusive answer as to which theoretical model is correct. Thus, our main conclusion is only a qualitative one: In the presence of competition for a depolymerization agent, a balance between plus-end polymerization and minus-end depolymerization is possible, and stable length distributions can also be achieved without filament severing.

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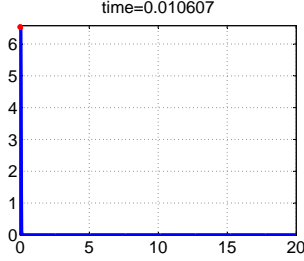
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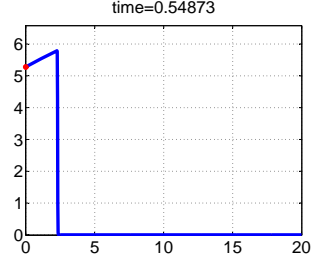




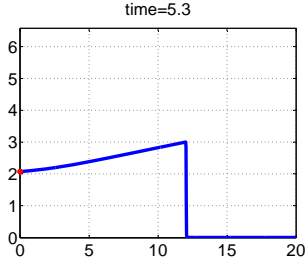
(a) Zero initial conditions; zero initial mass. The boundary condition attains the maximum possible value,  $u_b(0) = \phi(\bar{N}) > u^*$ .



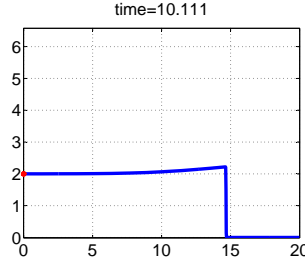
(b) Since  $u_b(0) > u^*$ , the boundary condition is instantly assumed, and remains assumed for every  $t > 0$ . So there is influx at the boundary.



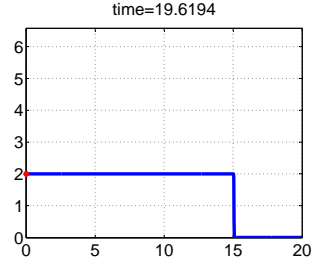
(c) The influx at the boundary, leads to mass production (16), and to boundary condition decrease (13). A propagating shock has formed, followed by a rarefaction wave.



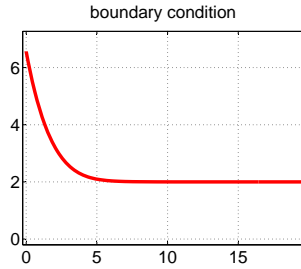
(d) The decrease of the boundary condition and the increase of the mass slow down, due to the (16) and (13). The rarefaction consumes the magnitude of the shock.



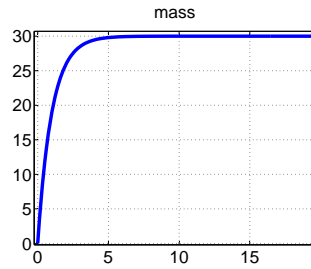
(e) The mass  $N(t)$  and the boundary condition  $u_b(t)$  have almost reached their limiting values  $\bar{N}$  and  $u^*$ . The speed of the shock has decreased since its end values are close to  $u^*$  and 0.



(f) Steady state of  $u$ . The shock speed is  $\sigma = \frac{f(u^*) - f(0)}{u^* - 0} = 0$ .



(g) Time evolution of the boundary condition. Since  $u_b(t) > u^*$ ,  $t > 0$  the boundary condition is assumed for every  $t > 0$ . Note the exponential convergence as  $t \rightarrow \infty$ .



(h) Time evolution of the mass. Since  $u_b(t) > u^*$ ,  $t > 0$  is always assumed and the, the mass evolves according to (16). Note again the exponential convergence.

Figure 3: Numerical test 1. In this case  $\bar{N} = 30$ , the flux function  $f(u) = u(u - 2)$ , with  $u^* = 2$  and the initial conditions are zero. As the Theorem 2 predicted, the boundary condition and the mass converge as  $u_b(t) \rightarrow u^*$  and  $N(t) \rightarrow \bar{N}$  respectively, and the asymptotic profile of  $u$  is a single steady shock at  $x_\infty = \frac{\bar{N}}{u^*}$  with jump from  $u^*$  to 0.

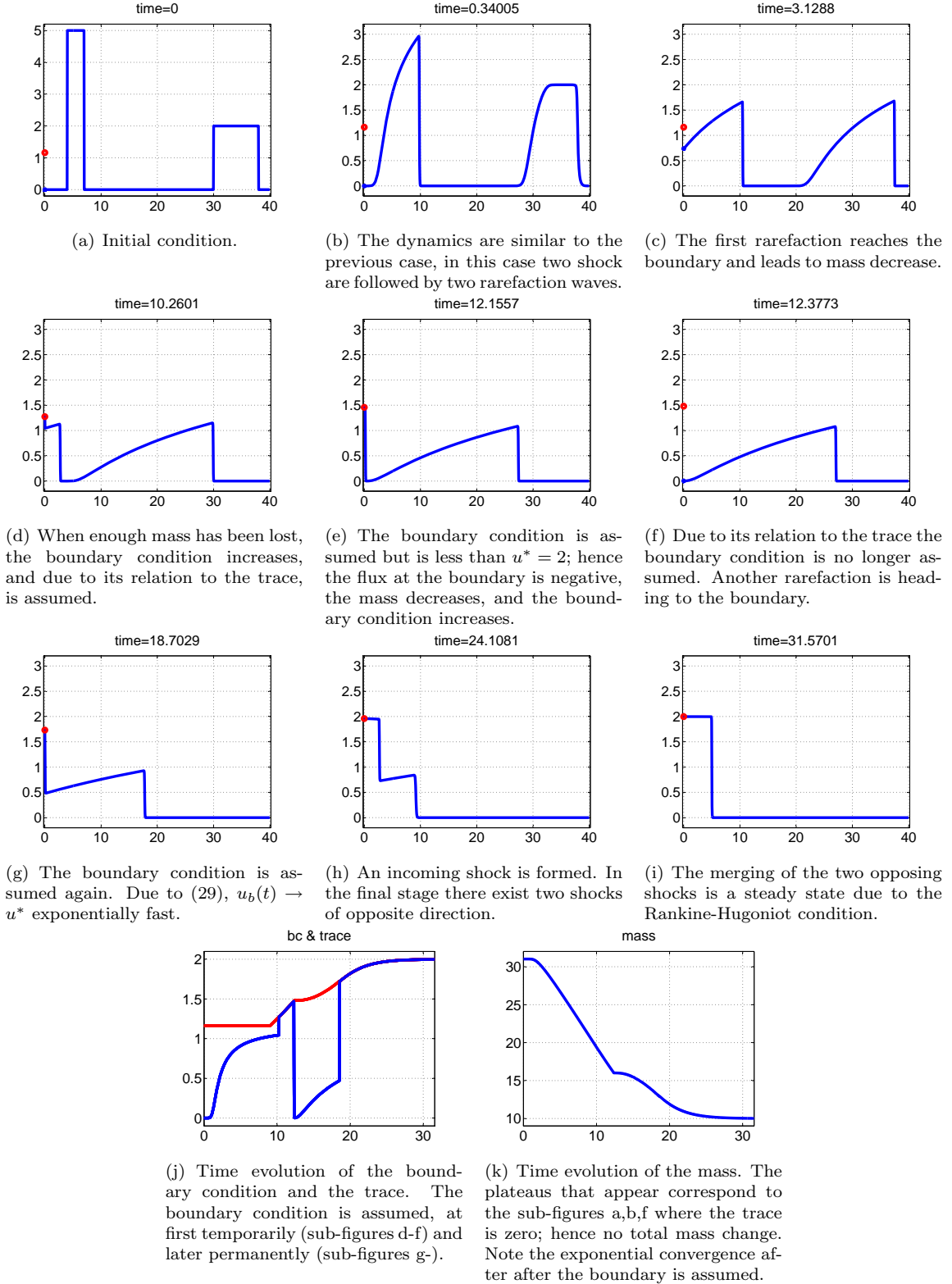
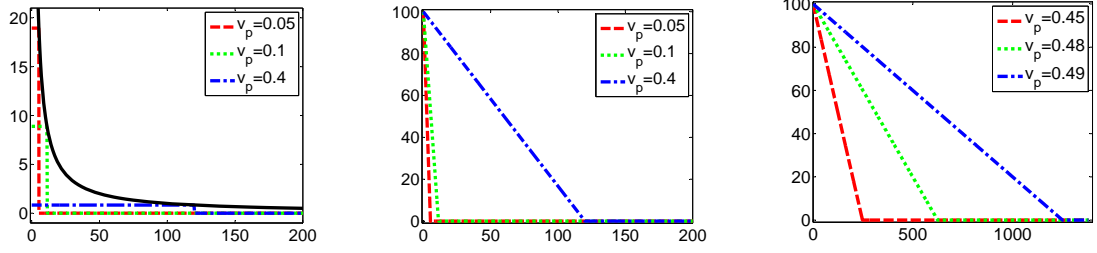


Figure 4: Numerical test 2. In this case  $\bar{N} = 10$ , and  $f(u) = e^u - \frac{e^2-1}{2}u - 1$  with  $u^* = 2$ . The initial mass is larger than  $\bar{N}$ ; in this case the generalized model and the Theorem 6 predict the evolution.

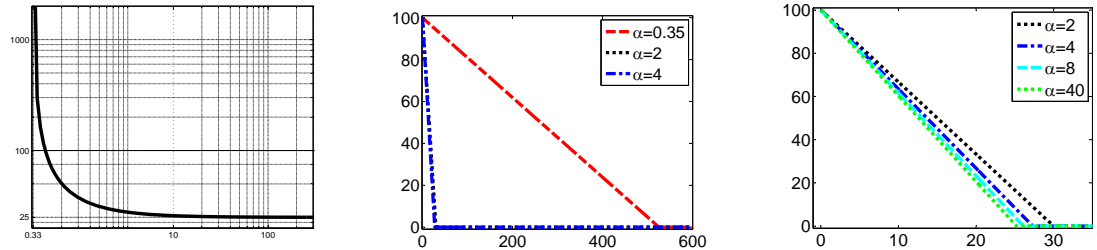


(a) We present the steady state density profiles  $u_\infty$  for  $\alpha = 1$  and for various values of  $v_p$  (dashed lines), as a function of  $x$ . We also present the curve  $(x_\infty, u^*)$  (solid curve) as a function of  $v_p$ , with  $\alpha = 1$ . We note that the discontinuities of the steady state shock wave solutions, coincide with the the  $(x_\infty, u^*)$  curve.

(b) The steady state length distribution of the filament ends as functions of  $x$ . The total number of filaments is the same in all cases, coinciding with  $\bar{N} = 100$ , and the length of the filaments is distributed uniformly between  $x = 0$  and  $x = x_\infty$ .

(c) The maximum length of filaments  $x_\infty$  tends to  $\infty$  as  $v_p$  tends to is upper bound, which is  $\frac{\alpha}{\alpha+1} = \frac{1}{2}$  in this case.

Figure 5: Numerical test 3. We study in this test, the steady state of the density and of the length distribution of the filament end points, as functions of  $v_p$ . We consider  $\alpha = 1$  and  $v_p = 0.05, 0.1, 0.4, 0.45, 0.48, 0.49$ . We conclude that the higher the polymerization/depolymerization ratio  $v_p$  is, the larger the maximum filament lengths  $x_\infty$  are, more specifically we verify numerically that  $\lim_{v_p \rightarrow \frac{\alpha}{\alpha+1}} x_\infty = \infty$ .



(a) We present the graph of  $x_\infty$  as a function of  $\alpha > \frac{1}{3}$  in logarithmic scales. We note the convergence of  $x_\infty$  at the endpoints its domain.

(b) We present the length distribution of the filament ends, for  $\alpha = 0.35, 2, 4$ , as functions of  $x$ . We note the large value of  $x_\infty$  for  $\alpha = 0.35$ . The total number of filaments is  $\bar{N} = 100$ . The length of the filaments is distributed uniformly between  $x = 0$  and  $x = x_\infty$ .

(c) The same graph for other values of  $\alpha$ . We note the convergence of  $x_\infty$  as  $\alpha$  increases.

Figure 6: Numerical test 4. We study in this test, the steady state of the length distribution of the filament end points, for fixed  $v_p = \frac{1}{4}$  and variable  $\alpha$ . We conclude that the smaller the  $\alpha$ , the longer the filaments are, i.e  $\lim_{\alpha \rightarrow \frac{1}{3}} x_\infty = \infty$ , on the contrary, the maximum filament length is bounded for large values of  $\alpha$ , i.e  $\lim_{\alpha \rightarrow \infty} x_\infty = 25 = \bar{N} v_p$ .