New Journal of Physics

The open-access journal for physics

Inelastic mechanics of sticky biopolymer networks

Lars Wolff^{1,3}, Pablo Fernandez² and Klaus Kroy¹

¹ Institute for Theoretical Physics, University of Leipzig, Vor dem Hospitaltore 1, 04102 Leipzig, Germany
² Department of Physics, TU München, James Franck Strasse 1, 85748 Garching, Germany
E-mail: lars.wolff@itp.uni-leipzig.de

New Journal of Physics **12** (2010) 053024 (17pp) Received 6 January 2010 Published 14 May 2010 Online at http://www.njp.org/ doi:10.1088/1367-2630/12/5/053024

Abstract. We propose a physical model for the nonlinear inelastic mechanics of sticky biopolymer networks with potential applications to inelastic cell mechanics. It consists of a minimal extension of the glassy wormlike chain (GwLC) model, which has recently been highly successful as a quantitative mathematical description of the viscoelastic properties of biopolymer networks and cells. To extend its scope to nonequilibrium situations, where the thermodynamic state variables may evolve dynamically, the GwLC is furnished with an explicit representation of the kinetics of breaking and reforming sticky bonds. In spite of its simplicity, the model exhibits many experimentally established nontrivial features such as power-law rheology, stress stiffening, fluidization and cyclic softening effects.

³ Author to whom any correspondence should be addressed.

Contents

1.	. Introduction . Theory		2	
2.			3	
	2.1.	The glassy wormlike chain	3	
	2.2.	Effective interaction potential and bond kinetics	5	
3.	Results and discussions		7	
	3.1.	Viscoelastic properties	7	
	3.2.	Stress stiffening versus rate-dependent yielding	8	
	3.3.	Transient remodelling and cyclic softening	10	
	3.4.	Introducing multiple length scales	12	
4.	Conc	lusions and outlook	13	
Acknowledgments			14	
Ар	Appendix. Obtaining force–displacement curves and solving for <i>v</i>			
Re	References			

1. Introduction

In many studies of cell mechanics and dynamics, the cell is characterized as a viscoelastic body [1]-[3]. It is an intriguing question to what extent such mechanical behaviour can be rationalized in terms of comparatively simple polymer physics models. In this respect, the comparison of cell rheological data and minimalistic in vitro reconstituted constructs of the cytoskeleton, such as pure actin solutions [4]-[8] or cross-linked actin networks [9]-[14], has recently provided many new insights. Important progress has also been made in the development of phenomenological mathematical descriptions. This includes approaches related to the tube model [15]–[18], tensegrity-based approaches [19]–[21], effective-medium models [22]–[24] and some others [25, 26]. In particular, the glassy wormlike chain (GWLC) model [27], a phenomenological extension of the standard wormlike chain (WLC) model of semiflexible polymers [28], has been successful in describing a plethora of rheological data for polymer solutions [7, 29] and living cells [30] over many decades in time with a minimum of parameters. However, all these studies were primarily concerned with viscoelastic behaviour, while the latest investigations have underscored the glassy [31]-[33], fragile [34, 35] and inelastic [34]-[36] character of the mechanical response of living cells. Even for biopolymer networks in vitro, experiments operating in the nonlinear regime had so far to resort to special protocols that minimize plastic flow [8, 37] for making contact with dedicated theoretical models.

The aim of the present paper is to overcome this restriction by extending the GWLC to situations involving inelastic deformations. As a first step, we concentrate on *reversible inelastic* behaviour, where the deformation does not alter the microscopic ground state. The protocol applied by Trepat *et al* [34] provides a paradigmatic example. Cells are subjected to a transient stretch such that, after some additional waiting time in the unstretched state, the (linear) material properties of the initial state are recovered. The simplification for the theoretical modelling results from the assumption that not only the macro-state but also the micro-state of the system may, to a good approximation, be treated as reversible under such conditions; that is, we assume that the complete conformation of the polymer network, including the transiently broken bonds between adjacent polymers, is constrained to eventually return to its original equilibrium state.

2

For the time-delayed hysteretic response of the network to such protocols, one could thus still speak of a viscoelastic ('anelastic') response in an operational sense, but we refrain from doing so in view of the fundamentally inelastic nature of the underlying stochastic process—in contrast to the reversible softening effects observed in [38], for example. Indeed, by simply allowing bonds to reform in new conformational states, the model developed below can readily be extended to arbitrary irreversible plastic deformations, as will be demonstrated elsewhere [45]. Before entering the discussion of our model, we would also like to point out that the proposed (inelastic) extension of the GWLC is strongly constrained by identifying the newly introduced parameters with those of the original (viscoelastic) GWLC model, where possible. Despite its increased complexity, the extended model will therefore enable us to subject the underlying physical picture to a more stringent test than hitherto possible by comparing its predictions to dedicated experiments. Moreover, unlike current state-of-the-art simulation studies [25], it is not limited to rod networks but is firmly rooted in a faithful mathematical description of the underlying Brownian polymer dynamics.

This paper is organized as follows. First, we review some basic facts about the GwLC in section 2.1. Next, in section 2.2, we introduce our extended reversible inelastic version, which we formulate using the notion of an effective interaction potential as in the original construction of the GwLC in [27]. (A preliminary account of the basic procedure and some of its cell-biological motivation including reversible bond-breaking kinetics has recently been given in a conference proceedings [39].) Sections 3.1 and 3.2 explain the physical mechanism underlying the mechanical response under pulsed and periodically pulsed loadings, whereas section 3.3 illustrates its phenomenology. We demonstrate that the model exhibits the hallmarks of nonlinear cell mechanics: strain/stress stiffening, fluidization and cyclic softening [34]–[36]. Section 3.4 investigates the relevance of the lately quantified structural heterogeneities in networks of semiflexible polymers [40] for the mechanical properties; finally, we conclude and close with a brief outlook.

2. Theory

2.1. The glassy wormlike chain

The GWLC is a phenomenological extension of the WLC model, the well-established standard model of semiflexible polymers. A broad overview of WLC and GWLC dynamics can be found elsewhere [28]. The WLC describes the mechanics of an isolated semiflexible polymer in an isothermal viscous solvent. In the weakly bending rod approximation, a solution of the stochastic differential equations of motion for the WLC is possible via a mode decomposition ansatz for the transverse displacement of the polymer contour from the straight ground state. The individual modes labelled by an index *n* are independent of each other and decay exponentially with rates τ_n^{-1} . For convenience, we set the thermal energy $k_{\rm B}T = 1$, so that the bending rigidity can be identified with the persistence length $\ell_{\rm p}$, in the following. Using this convention, the WLC expression for the transverse susceptibility of a polymer of contour length *L* (with respect to a point force) reads

$$\alpha_{\rm WLC}(\omega) = \frac{L^3}{\ell_{\rm p}\pi^4} \sum_{n=1}^{\infty} \frac{1}{(n^4 + n^2 f/f_L)(1 + \mathrm{i}\omega\tau_n)}.$$
(1)



Figure 1. Cartoon of a test polymer (differently coloured) in a network. The trapping of the test polymer is schematically represented by means of an effective interaction (figure 2) indicated by the potential wells at sticky entanglement points, which are on average separated by a contour length L_e . The test polymer can bind/unbind by overcoming an energy barrier of height \mathcal{E} . The average contour length between the closed bonds is Λ .

Here, *f* is an optional backbone tension and $f_L \equiv \pi^2 \ell_p / L^2$ is the Euler buckling force for a hinged rod of length *L*. The different powers of *n* in the denominator show the competition between bending and stretching forces.

In the GwLC, interactions of the polymer with the surrounding network (e.g. excluded volume interactions or sticky contacts) are reflected in an altered WLC mode relaxation spectrum [27, 29]. The intuitive—albeit not fully microscopic—picture underlying the formulation of the GwLC is illustrated in figure 1, which depicts a test polymer reversibly bound to the background network via potential wells at all topological contacts, the so-called entanglement points. Consider now a generic point somewhere along the polymer backbone (not an entanglement point). It can relax freely until the constrictions are being felt, which slow down the contributions from long-wavelength bending modes. The GwLC translates this intuition into a simple prescription for the mode spectrum: the short-wavelength modes are directly taken over from the wormlike chain model, whereas modes of wavelength λ_n longer than the typical contour length Λ between adjacent bonds are modified to account for the slowdown. Motivated by the physical picture illustrated in figure 1, the slowdown of the relaxation of a wavelength λ_n is expressed by an Arrhenius factor exp[$\mathcal{E}(\lambda_n/\Lambda - 1)$] for breaking ($\lambda_n/\Lambda - 1$) potential energy barriers of height \mathcal{E} simultaneously. Accordingly, the phenomenological recipe to turn a WLC into a GWLC reads

$$\tau_n \to \tilde{\tau}_n = \begin{cases} \tau_n, & \lambda_n < \Lambda, \\ \tau_n \exp\left[\mathcal{E}(\lambda_n/\Lambda - 1)\right], & \lambda_n \ge \Lambda. \end{cases}$$
(2)

Upon inserting this into (1), one obtains the GWLC susceptibility $\alpha_{GWLC}(\omega)$. The microscopic 'modulus' for transverse point excitations of a generic backbone element on a test polymer is then defined as the inverse susceptibility, $g_{GWLC}(\omega) = 1/\alpha_{GWLC}(\omega)$. An approximate expression for the macroscopic shear modulus is obtained along similar lines [22, 27].

In the original equilibrium GWLC theory, Λ was assumed to be a constant of the order of the entanglement length of the network, $\Lambda \gtrsim L_e$. Note, however, that L_e is a geometric quantity



Figure 2. Sketch of the schematic effective interaction potential for a test segment against the background. A polymer segment can either be 'bound' by a transient crosslinker or a sticky patch on the backbone of another polymer (left potential well) or 'unbound' and merely confined by the surrounding network to a tube-like cage (right well). When considering an ensemble of contacts, the fraction of closed bonds depends on the free energy difference U between both states. The transition rates k_{-} and k_{+} between the bound and unbound states depend on the barrier heights \mathcal{E} and $(\mathcal{E} - U)$, respectively. An externally applied force can tilt the potential and favour binding or unbinding events.

(which is determined by the polymer concentration and the persistence length), whereas the contour length Λ between closed bonds clearly depends on the state of the bond network. One therefore has to allow for an increase of Λ with the number of broken bonds in non-equilibrium applications. This issue is explored in the following.

2.2. Effective interaction potential and bond kinetics

All mechanical quantities calculated within the GWLC model crucially depend on the interaction length Λ . In previous applications of the model [7, 27, 30], it was assumed that Λ remains constant—equal to its equilibrium value and unaffected by the deformation of the sample. In other words, the equilibrium theory allowed for statistical bond fluctuations but not for a dynamical evolution of the parameters characterizing the thermodynamic state of the bond network. An obvious starting point for generalizations of the model to non-equilibrium situations is therefore to consider the number of closed bonds, and therefore also Λ , as *dynamic variables*, dependent on the strain- and stress-history of the network. We now provide a meanfield description to account for such a dynamically evolving bond network. For clarity, we return to the intuitive picture underlying the GWLC, where the (possibly crosslinker- or molecularmotor-mediated) complex interactions between the polymers are summarized into an effective interaction potential for a test segment against the background, as sketched in figure 2. The same idea has also been used earlier in many related situations (e.g. [26, 41, 42]). The generic potential exhibits three essential features: a well-defined bound state, a well-defined unbound 6

state and a barrier in between (figure 2). The confining well corresponding to the unbound state represents the tube-like caging of the test polymer within the surrounding network [15, 17, 43]. For ease of notation, we further introduce the dimensionless fraction of closed bonds, or *bond fraction*

$$\nu(t) \equiv \Lambda_0 / \Lambda(t), \tag{3}$$

which is simply the ensemble-averaged fraction of sticky contacts that are actually in the bound state. The minimum bond distance Λ_0 is typically of the order of L_e , but may be somewhat larger in situations where the bonds are mediated by crosslinker molecules or by partially sterically inaccessible sticky patches (e.g. for helical molecules [12, 44]).

A quantitatively fully consistent way of calculating the dynamics of ν would involve solving the full Fokker–Planck equation for a WLC–WLC contact—a formidable programme to be pursued elsewhere [45]. Here, for the sake of our qualitative purposes, we chose to concentrate on the physical essence of the discussion, and keep the mathematical structure as simple and transparent as possible. We therefore approximate the dynamics by a simple exponential relaxation as is familiar from the standard example of reacting Brownian particles, conventionally described by Kramers theory with a Bell-like force dependence [46, 47]. Using this simplification and assuming a schematic interaction potential as depicted in figure 2, the value of ν is determined by the competition between bond breaking and bond formation with reaction rates k_{-} and k_{+} , respectively. Both rates are represented in the usual adiabatic approximation (meaning that the equilibration *inside* the wells is much faster than the barrier crossing and external perturbations) by the following [46]:

$$k_{-}\tau_{0} = e^{-\mathcal{E} + (x_{t} - x_{b})f}, \quad k_{+}\tau_{0} = e^{-\mathcal{E} + U - (x_{u} - x_{t})f},$$
(4)

where τ_0 is an intrinsic characteristic Brownian time scale for bond breaking and formation⁴, and *f* is the force pulling on the bond. Noting that the fraction of open bonds is $1 - \nu$, we can then write down the following rate equation for the fraction of closed bonds:

$$\dot{\nu}(t)\tau_0 e^{\mathcal{E}} = -(e^{-(x_u - x_t)f(t) + U} + e^{(x_t - x_b)f(t)})\nu(t) + e^{-(x_u - x_t)f(t) + U}.$$
(5)

The time dependence of v(t) leads via (3) to an implicit time dependence of the GwLC parameter $\Lambda(t)$ and thereby of all observables derived from the GwLC. The time-dependent force f(t) in (5) may be thought to result from an externally imposed stress protocol or from internal dynamical elements such as molecular motors setting the network under dynamic stress. Hence, via an appropriately chosen set of slowly changing state parameters f(t), U(t), $\mathcal{E}(t), \ldots$, the model can in principle accommodate the active biological remodelling in living cells and tissues [39]. (For a discussion of the relation between the microscopic f and the macroscopic stress, see [29].) Note that for constant force, the stationary value of v,

$$\nu_{\text{stat}} = (1 + \exp[-U + (x_u + x_b) f_{\text{stat}}])^{-1}, \qquad (6)$$

obtained by setting $\dot{\nu} \equiv 0$, does not depend on \mathcal{E} , as it should (the steady state is independent of the transition state).

⁴ Strictly speaking, the fluctuations in the potential wells at x_b and x_u are characterized by different Brownian time scales depending on the width of the potential wells. At the present stage, we do not bother to distinguish these time scales, nor to fine-tune their numerical values, and identify them, for the sake of simplicity, with the entanglement time scale τ_{L_e} in our numerical calculations.



Figure 3. Dependence of the microrheological 'modulus' g_{GWLC} on the bond fraction ν for various values of \mathcal{E} (brown, $\mathcal{E} = 0.1$: red, $\mathcal{E} = 1$: blue, $\mathcal{E} = 5$: green, $\mathcal{E} = 10$: black, $\mathcal{E} = \infty$) at a frequency $\omega_0 \approx 1/10\tau_{\Lambda_0}^{-1}$ and a force $f = f_{\Lambda_0}/2$ (τ_{Λ_0} is the relaxation time of a mode of length Λ_0 and f_{Λ_0} is the Euler buckling force for a rod of length Λ_0); (a) normalized real part g'; (b) loss angle.

3. Results and discussions

3.1. Viscoelastic properties

Whenever the bond kinetics can be disregarded ($\dot{\nu} = 0$), the viscoelastic properties are simply those of the bare GwLC [7, 27, 29, 30], which can basically be characterized as WLC shorttime dynamics followed by a slow, highly stretched logarithmic relaxation resembling powerlaw rheology with a non-universal exponent within the typical experimental time windows. Strong perturbations (stress or strain) then give rise to a pronounced stiffening due to the steeply nonlinear response of semiflexible polymers under elongation [2]. This well-established behaviour can be understood as the canvas against which we aim to discern the characteristic mechanical signatures of the bond kinetics.

An observable that will be of particular interest in the following is the complex microscopic 'modulus' $g \equiv g_{\text{GWLC}}(\omega_0, \Lambda_0/\nu, f)$, introduced in the previous section. It is used to determine the linear as well as (via a nonlinear update scheme, see appendix) the nonlinear force response of the system. To understand how the time dependence (5) of ν affects this important quantity, it is instructive to first examine the dependence of g on ν . To this end, we formally consider ν temporarily as an independent variable instead of determining it from (5). Note that this approach nevertheless makes sense, as for a fixed set of parameters, ν can still take any value between zero and one. This is due to the freedom in choosing an initial state, which can be imagined to have evolved from the prior deformation history.

For other fixed parameters, both the real part g' (figure 3(a)) and the imaginary part g''of g increase monotonically with v. We emphasize that g'' is not simply proportional to g'and therefore the loss angle $\delta = \arctan(g''/g')$ also depends on v (figure 3(b)). For small v, the loss angle is large, corresponding to fluid-like behaviour. With increasing v, the system becomes more solid-like. Increasing the barrier height \mathcal{E} makes the dependence of g on v more pronounced (figure 3). Conversely, as can be expected, the dependence on the barrier height \mathcal{E} vanishes with decreasing bond fraction v. Note that due to the boundary conditions, the limit of a Newtonian fluid ($\delta = \pi/2$) is not recovered when formally taking the limit of vanishing bond fraction $(\nu \to 0, \Lambda \to \infty)$. Finally, the potential difference U solely determines the dynamics of ν via (5) and therefore influences the modulus g solely through the equilibrium value for ν .

After these general considerations, we now concentrate on the nonlinear and nonequilibrium dynamic response of the extended inelastic GwLC model, which results from the coupled relaxation of the viscoelastic polymer network *and* the transient bond network.

3.2. Stress stiffening versus rate-dependent yielding

A convenient way to characterize the mechanical properties of an inelastic material is a force-displacement curve. For a perfectly linear elastic (Hookean) body, it would simply consist of a straight line, whereas a perfectly plastic body would feature as a rectangle delineated by a yield force f_y and an arbitrary plastic strain value. For our qualitative purpose, we identify the average transverse displacement of the test polymer segment (which is used to determine the force response, see appendix) with the reaction coordinate x of the schematic potential sketched in figure 2; and we identify the force f entering the reaction rates with the backbone tension of the test polymer. As a characteristic length scale for the transverse displacements, we use the width $(x_u - x_t)$ of the effective confinement potential, which is a measure of a typical mean-square displacement of the polymer in the unbound state. Using this convention, we now consider the effect of a time-symmetric displacement pulse on the evolution of the force f(t) and bond fraction v(t). For technical convenience, we use a Gaussian shape for the displacement pulses, but the qualitative conclusions to be drawn are largely independent of the precise shape. The duration of the displacement pulse, which sets the time scale for the dynamic response, is used as the unit of time in the following. Here, we are not interested in short-time tension propagation and single-polymer dynamics [48]; hence a lower bound for pertinent pulse durations is provided by the interaction time scale $\tau_{\Lambda_0} \simeq \tau_0$. On the other hand, for pulse durations longer than $\tau_0 e^{\mathcal{E}}$, the system deformation would be quasi-static so that no genuinely dynamic effects of the bond kinetics could be observed.

For a small Gaussian displacement pulse of relative amplitude 0.9 (cf figure 4(a) and appendix), the shape of the force-displacement curve predicted by our inelastic GwLc model shows all the features of a viscoelastic medium (figure 4(b), blue dashed curve). It starts with a nearly linear regime for relative displacements \lesssim 0.4, where a weak stiffening sets in. Owing to dissipation in the medium, the path back to the initial state takes its course at lower force, which causes a weak hysteresis. This is essentially the viscoelastic response that already the bare GWLC model would have predicted. For a larger relative deformation amplitude of 2.8, however, the predictions of the bare GWLC and the inelastic GWLC diverge. The bare GWLC predicts a strong stiffening (figure 4(b), red dotted curve), whereas the full model exhibits an initial stiffening regime followed by a pronounced softening (figure 4(b), solid red curve). A very interesting observation is that 'softening' in this case not only means a decrease of the slope of the force-extension curve, but that the slope actually becomes negative over an extended parameter region, once an operationally defined threshold force f_y is exceeded. This 'flow state' bears more resemblance to plastic flow than to viscoelastic relaxation. The reason for this qualitatively new phenomenon is the force-induced bond breaking. Theoretically, this is best exemplified by the time dependence of the (experimentally not easily accessible) fraction v(t) of closed bonds (figure 4(c)). A glance at the bond fraction during the weak deformation scenario (figure 4(c), blue dashed curve) reveals how the limit of a bare GWLC is recovered, namely whenever the deformation is not sufficiently violent and persistent to significantly





Figure 4. Transition from a stiffening to a softening response at a rate-dependent, operationally defined yield force f_y : Gaussian deformation pulses (a) with relative displacement amplitudes of 0.9 (dashed) and 2.8 (solid); corresponding force–displacement curves and bond fraction evolutions (b, c); dotted curve in (b): response of a GWLC without bond breaking); order-of-magnitude estimate for the rate dependence of the yield force, equation (7) with const. = 2.2, versus the numerically obtained maximum forces of the force–displacement curves for various mean force rates (d); the minimum bond fraction (d, inset) depends non-monotonically on the force rate; $\mathcal{E} = 10.5$, U = 2, $\tau_0 = 0.004$, $(x_t - x_b)/(x_u - x_t) = 0.07$, $\Lambda_0 = 22.3$.

decrease the fraction of closed bonds. For the large deformation scenario, in contrast, the bond fraction stays only initially constant (figure 4(c), solid red curve). As a consequence of the large strain, the force rises steeply, as can be seen from the strong stiffening in figure 4(b) (solid red curve). At the yield force, the bond fraction suddenly decreases to nearly half of its initial value during a very short time. The decrease in the bond fraction is accompanied by a somewhat slower drop in the force, which is reflected in the sudden softening of the force–displacement curve. The bond fraction eventually recovers on a much slower time scale, which is roughly given by $\tau_0 \exp(\mathcal{E} - U)$. Note that even though the effects of the inelastic response dominate the stress–strain curve, the viscoelastic relaxations from the underlying GwLC model are still present. They could hardly be disentangled from the inelastic contributions, however, without an underlying faithful model of the viscoelastic response at hand.

In summary, we observe a competition between force-amplitude-dependent stressstiffening and force-rate-dependent yielding events. If the backbone force stays much smaller than the yield force, the adaptation of the bond network requires an adiabatically long time, on the order of the bond lifetime $\tau_0 e^{\varepsilon}$, so that plain viscoelastic GWLC behaviour is observed on the pulse time scale. In contrast, if the backbone tension f reaches the yield force f_y , the bond fraction declines sharply, thereby switching the response from the WLC/GWLC-typical stiffening to a pronounced softening.

The rate dependence of the yield force f_y can be estimated by setting the time it takes to reach the yield force, f_y/\dot{f} , equal to the force-dependent time scale of bond opening, $k_{-1}^{-1}(f)$, from (4),

$$f_{\mathbf{y}}\dot{f}^{-1} \simeq k_{-}^{-1}(f_{\mathbf{y}}) \quad \Rightarrow \quad (x_t - x_b)f_{\mathbf{y}} \approx \mathrm{LW}\left(\mathrm{const.} \times (x_t - x_b)\tau_0 \mathrm{e}^{\mathcal{E}}\dot{f}\right).$$
(7)

Here, LW(x) is the positive real branch of the Lambert W function. In figure 4(d), this estimate is compared with results from numerical evaluations for Gaussian displacement pulses at different average rates. Apart from the numerical errors, the slight deviations from the estimate may be attributed to the fact that the force rate is not constant for the Gaussian protocol. They can be mostly eliminated by using force ramp protocols of various slopes instead of the Gaussian displacement pulses. For not too low rates, the rate dependence can be approximated by the even simpler relation

$$(x_t - x_b) f_y \approx (\mathcal{E} + \ln f \tau_0 + \text{const.}), \tag{8}$$

where the force rate has to be normalized by a suitable force scale.

The minimum bond fraction reached during the application of the pulse, which we interpret as a measure of the degree of fluidization, depends *non-monotonically* on the rate (figure 4(d), inset). Qualitatively, this can be understood by noticing that two different factors influence the fluidization, namely the maximum force attained during deformation and the time over which the force is applied. The maximum force is simply the yield force f_y , with the rate dependence in (7) and figure 4(d), while the reciprocal rate itself sets the time scale. For high rates $(x_u - x_t)\dot{f} \gtrsim 1$ (in our example) the rate dependence of the maximum force wins and the minimum ν reached decreases with increasing rate. For low rates, where the rate dependence of the maximum force is much weaker (figure 4(d)), the minimum ν decreases with increasing pulse duration, namely decreasing rate. Note that for slow pulses (low rates), the bond fraction after the pulse may be quite different from the minimum ν , as significant recovery may already take place during the pulse.

3.3. Transient remodelling and cyclic softening

The substantial changes in the material properties accompanying bond breaking can be exemplified by monitoring the linear elastic modulus $g'(\omega_0)$ (measured at a fixed frequency ω_0) in response to a strain pulse (figure 5(a)). Apart from the usual WLC/GwLC-typical stress stiffening below the threshold force f_y , the modulus is seen to drop *below* the value it had before the deformation pulse, where it apparently saturates (remember that the deformation vanishes for t > 1 and that the recovery takes roughly a time $\tau_0 e^{\mathcal{E}-U} \gg 1$). We thus observe what we call a 'passive', 'physical' remodelling of the bond network as opposed to the 'active', 'biological' remodelling of the cytoskeleton of living cells in response to external stimuli. These passive remodelling effects have recently been observed for human airway smooth muscle cells [34]. (A more quantitative discussion of the relation to the experimental observations will be given elsewhere.) The fact that the deformation pulse leads to a decrease in g' and an increase in the loss angle δ (figure 5(c)) suggests the notion of *fluidization* [34].



Figure 5. The internal force f and the bond fraction v largely determine the storage modulus g' and the loss angle δ : normalized storage modulus g'_n (a) and loss angle δ (c) during displacement pulses of large and small amplitudes, 2.8 (solid) and 0.9 (dashed); the stronger deformation pulse causes a strong and persistent overshoot of the loss angle, indicating *fluidization*; slow recovery after the pulse (b, d); parameters as in figure 4, $\omega_0 = 3$.

The passive remodelling described here is a *transient* phenomenon, because, after cessation of the external perturbations, the bond fraction will eventually recover its equilibrium value. This indicates that also the change in the system properties is a transient effect, which is demonstrated by the recovery from fluidization in figures 5(b) and (d). Thus, while the fluidization resembles a plastic process on short time scales, on long time scales, the phenomenology is more similar to pseudo- and superelasticity as observed in shape-memory alloys [49].

To demonstrate that the transient passive remodelling also affects the nonlinear material properties, we next consider a series of three pulses (figure 6). The force response to such a protocol is depicted in figure 6(b). The force–displacement curve for the second stretching (second left branch of the solid curve) is less steep than for the first stretching (leftmost branch of the solid curve) and the yield force is lower. This indicates a cyclic softening, or viscoelastic shake-down effect, closely related to the fluidization of the network by strain. The strength of the shake-down depends on the fraction of transiently broken bonds and hence on the rate as well as on the amplitude of the applied deformation (cf figure 6(c), solid red curve). Upon repeated application of deformation pulses the force–displacement curves settle on a limit curve that is essentially preconditioned by the initial deformation pulse and the inelastic work it performed on the sample. For more gentle protocols that only break a smaller fraction of bonds, the initial fluidization would not be as pronounced and one would obtain a gradual shake-down, which converges to a limit curve only after many cycles.



Figure 6. Protocol (a) and response (b, c) for three subsequent Gaussian strain pulses; force–displacement curves (b) for a single minimum interaction distance ($\Lambda_0 = 22.3$; solid) and for a distribution of Λ_0 according to Glaser *et al* [40] with mean $\bar{\Lambda}_0 = 22.3$ (dashed); the bond fraction $\nu(c)$ for $\Lambda_0 = 0.5 \bar{\Lambda}_0$ (dash-dotted), $\bar{\Lambda}_0$ (solid) and $2\bar{\Lambda}_0$ (dashed); other parameters are as in figure 4.

3.4. Introducing multiple length scales

So far, we have assumed that there is one well-defined characteristic length scale Λ_0 for the polymer interactions, which is of the order of the entanglement length L_e and interpreted as the minimum average contour distance between adjacent bonds of the test polymer with the background network. This is clearly a mean-field assumption. Recent combined experimental and theoretical studies have established that the local tube diameter and entanglement length in pure semiflexible polymer solutions actually exhibit a skewed leptocurtic distribution with broad tails [40]. To include this information in the above analysis is not straightforward, since the elastic interactions between regions of different entanglement lengths are not known, a priori. For the sake of a first qualitative estimate, the simplest procedure seems to be to average the above results over a distribution of Λ_0 , corresponding to a parallel connection of independent entanglement elements. Qualitatively, this renders the abrupt transition from stiffening to softening somewhat smoother (cf figure 6(b), dashed curve). Moreover, the initial stiffening is slightly more pronounced. This is due to the contributions of $\Lambda_0 < \overline{\Lambda}_0$, which exhibit both stronger stiffening and fluidization. Nevertheless, all qualitative features—like stiffening and remodelling effects and a 'flow state' indicating fluidization or inelastic shakedown—can still be well discerned. This is consistent with the assumption that pure semiflexible polymer solutions are at least qualitatively well described by a single entanglement length [17, 18, 22, 27, 50].

4. Conclusions and outlook

We have presented a theoretical framework for a polymer-based description of the inelastic nonlinear mechanical behaviour of sticky biopolymer networks. We represented the polymer network on a mean-field level by a test polymer described by the phenomenologically highly successful GwLc model. Beyond the equilibrium statistical fluctuations captured by the original model, we additionally allowed for a dynamically evolving thermodynamic state variable $\Lambda(t)$ characterizing the network of thermo-reversible sticky bonds, which is updated dynamically upon bond breaking according to an appropriate rate equation. Within our approximate treatment we could obtain a number of robust and qualitatively interesting results, which are not sensitive to the precise parameter values chosen, nor to the choice of the transverse rather than the longitudinal susceptibility in (1). For sufficiently strong deformations, changes in the mean fraction of closed bonds, reflected in $\Lambda(t)$, can influence material properties to a degree at which the material behaves qualitatively different from the usual viscoelastic paradigm. In particular, our theory predicts a pronounced fluidization response, which develops upon strong deformations on top of the intrinsic viscoelastic stiffening response provided by the individual polymers. After cessation of loading, the system slowly recovers its initial state. These observations are in qualitative agreement with recently published experimental results obtained for living cells [34], and a more quantitative comparison with dedicated measurement results for cells and in vitro biopolymer networks will be the subject of our future work. We found the yield force for the onset of the fluidization to be sensitive to the deformation rate. Moreover, the fluidization response was shown to be accompanied by a cyclic softening or shake-down effect. Taking into account the spatial heterogeneity of biopolymer solutions by a distribution of entanglement lengths leads to a smoothing of the force-displacement curves without affecting their qualitative characteristics. One may still expect qualitatively new effects in situations with unusually broad distributions of entanglement lengths, such as for strongly heterogeneous (e.g. phase separating) networks.

A parameter that was found to be very important for cells [51] but has not been discussed much in this contribution is the prestressing force f_0 (see appendix), which is present in adhering cells even in the absence of external driving. Experimentally, it was established that increasing prestress is correlated with a higher stiffness of adhering cells [51], which in turn is correlated with a higher stiffness of the substrate [52]. When naively treating the prestressing force as just an additive contribution to the overall force, our model predicts the opposite: the additional force breaks more bonds and the network becomes softer and more fluid-like. To reconcile these apparently contradicting trends, one can appeal to the notion of homeostasis, which essentially amounts to postulating that the cell actively adapts its structure such that a certain set of thermodynamic state variables remain in a physiologically sensible range [39]. This basically implies that the cell will respond to stiff substrates or persistent external stresses by a biological remodelling that corresponds to an increase of \mathcal{E} and U, and possibly to a decrease of Λ_0 . By adapting also the internal prestress f_0 , the cell may then avoid, apart from the structural collapse, an equally undesirable loss of flexibility. We note, however, that internal stresses are indeed observed to disrupt the cytoskeleton, as implied by the simple physical theory presented here, if they are not permanently balanced by a substrate [53].

Even though the nonlinear effects presented in this contribution depend crucially on the bond kinetics, it should not be overlooked that the GWLC is an equally indispensable ingredient of the complete theory. Firstly, it provides the constitutive relation that gives the force in

response to an infinitesimal displacement, which in turn governs the bond kinetics. Secondly, the time-dependent linear susceptibility strongly filters the dynamic remodelling of the bond network in practical applications, in which one rarely has access to the microstate of the underlying bond network. In summary, the extended model thus integrates experimentally confirmed features of the GWLC response such as slow relaxation, power-law rheology, viscoelastic hysteresis and shear stiffening with a simple bond kinetics scheme, which allows less intuitive complex nonlinear physical remodelling effects like fluidization and inelastic shake-down to be addressed.

Finally, we want to point out that it is straightforward to extend the above analysis in a natural way to account for irreversible plastic deformations. To this end, one has to allow for the possibility that the transiently broken bonds reform somewhere else than at their original sites (as always assumed in the foregoing discussion) after strong deformations with finite residual displacement. This can be included by accommodating an additional term acting as a 'slip rate' in (5) [45].

Acknowledgments

We acknowledge many fruitful discussions with J Glaser, S Sturm and K Hallatschek and financial support from the Deutsche Forschungsgemeinschaft (DFG) through FOR 877 and, within the German excellence initiative, the Leipzig School of Natural Sciences—Building with Molecules and Nano-Objects.

Appendix. Obtaining force–displacement curves and solving for v

The deformation pulse protocol used in this paper is

$$\Gamma(t) = \gamma_0 \exp\left(-\frac{(t-\tau)^2}{\sigma^2}\right). \tag{A.1}$$

To obtain the full nonlinear response of the (extended) GwLC to the displacement protocol (A.1), we make use of the superposition principle: we know that the Fourier transform of $g_{GWLC}(\omega, \Lambda, f)/(i\omega)$ is the linear response to a small displacement step. After decomposing a finite displacement into infinitesimal displacement steps and a partial integration, we write

$$f_{\rm GWLC}(t) = \int_{-\infty}^{\infty} dt' \int_{-\infty}^{\infty} \frac{d\omega}{\sqrt{2\pi}} g_{\rm GWLC}(\omega, \Lambda(t), f_{\rm GWLC}(t)) e^{-i\omega(t-t')} \Gamma(t').$$
(A.2)

Note that the right-hand side of (A.2) depends on $f_{GWLC}(t)$, rendering it a highly nonlinear implicit equation. Using (A.1) and integrating by parts, we obtain

$$f_{\rm GWLC}(t) = \sqrt{2}\gamma_0 \sigma \int_0^\infty d\omega \, \mathrm{e}^{-\mathrm{i}\omega(t-\tau)} g_{\rm GWLC}(\omega, \Lambda(t), f_{\rm GWLC}(t)) \mathrm{e}^{-(\sigma^2 \omega^2)/4}. \tag{A.3}$$

The next step is to establish a connection between the force in (5) and $f_{GwLC}(t)$. A finite prestressing force f_0 is introduced mainly for technical reasons, namely to avoid the unphysical region of negative (i.e. compressive) backbone stress, which would buckle the polymers. Whereas the physics of the prestressed network can essentially be mapped back to that of a network without prestress by a renormalization of the parameters \mathcal{E} and U, prestress may also be seen as an important feature: indeed, the cytoskeleton of adhered cells is well known to be

under permanent prestress, and suspended cells seem prone to spontaneous shape oscillations that could be indicative of a propensity of the cells to set themselves under prestress [39]. In the context and on the longer time scales of the biological remodelling of the cytoskeleton it would therefore make sense to think of f_0 as a dynamic force generated by molecular motors and polymerization forces. For the following, however, we take the prestress to be constant so that the force entering (5) is

$$f(t) = f_0 + f_{\text{GWLC}}(t).$$
 (A.4)

We now face the problem that (A.2) is an implicit equation for $f_{GWLC}(t)$, which depends on v(t), which in turn depends via f(t) on $f_{GWLC}(t)$. We therefore use a two-step Euler scheme: as initial values for f and v, we choose the prestressing force, $f(t = 0) = f_0$, and the steady-state value under the prestressing force, $v(t = 0) = v(t \rightarrow \infty | f = f_0)$, respectively. We then choose a sufficiently small time step Δt and apply the following iterative rule:

$$f(k\Delta t) \approx f_0 + f_{\text{GWLC}} \left(k\Delta t | f((k-1)\Delta t, \nu((k-1)\Delta t)) \right), \tag{A.5}$$

$$\dot{\nu}(k\Delta t) \approx \dot{\nu}(k\Delta t | f(k\Delta t)),$$
(A.6)

where $k \in \mathbb{N}_+$. In the limit $\Delta t \to 0$, this procedure converges to the exact solution.

References

- [1] Fung Y 1993 Biomechanics: Mechanical Properties of Living Tissues (Berlin: Springer)
- [2] MacKintosh F C, Käs J and Janmey P 1995 Elasticity of semiflexible biopolymer networks *Phys. Rev. Lett.* 75 4425
- [3] Kruse K, Joanny J F, Jülicher F, Prost J and Sekimoto K 2004 Asters, vortices and rotating spirals in active gels of polar filaments *Phys. Rev. Lett.* **92** 78101
- [4] Gisler T and Weitz D A 1999 Scaling of the microrheology of semidilute F-actin solutions *Phys. Rev. Lett.* 82 1606–9
- [5] Mason T G, Gisler T, Kroy K, Frey E and Weitz D A 2000 Rheology of F-actin solutions determined from thermally driven tracer motion J. Rheol. 44 917
- [6] Gardel M L, Valentine M T, Crocker J C, Bausch A R and Weitz D A 2003 Microrheology of entangled F-actin solutions *Phys. Rev. Lett.* 91 158302
- [7] Semmrich C, Storz T, Glaser J, Merkel R, Bausch A R and Kroy K 2007 Glass transition and rheological redundancy in F-actin solutions *Proc. Natl Acad. Sci. USA* 104 20199–203
- [8] Semmrich C, Larsen R J and Bausch A R 2008 Nonlinear mechanics of entangled F-actin solutions Soft Matter 4 1675–80
- [9] Gardel M L, Nakamura F, Hartwig J, Crocker J C, Stossel T P and Weitz D A 2006 Stress-dependent elasticity of composite actin networks as a model for cell behavior *Phys. Rev. Lett.* 96 88102
- [10] Tharmann R, Claessens M and Bausch A R 2007 Viscoelasticity of isotropically cross-linked actin networks *Phys. Rev. Lett.* 98 88103
- [11] Luan Y, Lieleg O, Wagner B and Bausch A R 2008 Micro- and macrorheological properties of isotropically cross-linked actin networks *Biophys. J.* 94 688–93
- [12] Claessens M, Semmrich C, Ramos L and Bausch A R 2008 Helical twist controls the thickness of F-actin bundles *Proc. Natl Acad. Sci. USA* 105 8819
- [13] Kasza K E, Nakamura F, Hu S, Kollmannsberger P, Bonakdar N, Fabry B, Stossel T P, Wang N and Weitz D A 2009 Filamin a is essential for active cell stiffening but not passive stiffening under external force *Biophys. J.* 96 4326–35

- [14] Koenderink G H, Dogic Z, Nakamura F, Bendix P M, MacKintosh F C, Hartwig J H, Stossel T P and Weitz D A 2009 An active biopolymer network controlled by molecular motors *Proc. Natl Acad. Sci. USA* 106 15192
- [15] Käs J, Strey H and Sackmann E 1994 Direct imaging of reptation for semiflexible actin filaments *Nature* 368 226–9
- [16] Morse D C 1999 Viscoelasticity of concentrated isotropic solutions of semiflexible polymers. 3. Nonlinear rheology *Macromolecules* 32 5934–43
- [17] Morse D C 2001 Tube diameter in tightly entangled solutions of semiflexible polymers Phys. Rev. E 63 31502
- [18] Hinsch H, Wilhelm J and Frey E 2007 Quantitative tube model for semiflexible polymer solutions *Eur. Phys.* J. E 24 35–46
- [19] Canadas P, Laurent V M, Oddou C, Isabey D and Wendling S 2002 A cellular tensegrity model to analyse the structural viscoelasticity of the cytoskeleton J. Theor. Biol. 218 155–73
- [20] Sultan C, Stamenović D and Ingber D E 2004 A computational tensegrity model predicts dynamic rheological behaviors in living cells Ann. Biomed. Eng. 32 520–30
- [21] Cañadas P, Wendling-Mansuy S and Isabey D 2006 Frequency response of a viscoelastic tensegrity model: structural rearrangement contribution to cell dynamics J. Biomech. Eng. 128 487
- [22] Gittes F and MacKintosh F C 1998 Dynamic shear modulus of a semiflexible polymer network *Phys. Rev.* E 58 1241–4
- [23] Storm C, Pastore J J, MacKintosh F C, Lubensky T C and Janmey P A 2005 Nonlinear elasticity in biological gels Nature 435 191
- [24] Broedersz C P, Storm C and MacKintosh F C 2008 Nonlinear elasticity of composite networks of stiff biopolymers with flexible linkers *Phys. Rev. Lett.* 101 118103
- [25] Åström J A, Kumar P B S, Vattulainen I and Karttunen M 2008 Strain hardening, avalanches and strain softening in dense cross-linked actin networks *Phys. Rev.* E 77 51913
- [26] Kollmannsberger P and Fabry B 2009 Active soft glassy rheology of adherent cells Soft Matter 5 1771-4
- [27] Kroy K and Glaser J 2007 The glassy wormlike chain New J. Phys. 9 416
- [28] Kroy K 2008 Dynamics of wormlike and glassy wormlike chains Soft Matter 4 2323-30
- [29] Glaser J, Hallatschek O and Kroy K 2008 Dynamic structure factor of a stiff polymer in a glassy solution *Eur. Phys. J.* E 26 123–36
- [30] Kroy K and Glaser J 2009 Rheological redundancy—from polymers to living cells AIP Conf. Proc. 1151 52–5
- [31] Fabry B and Fredberg J J 2003 Remodeling of the airway smooth muscle cell: are we built of glass? *Respir*. *Physiol. Neurobiol.* 137 109–24
- [32] Deng L, Trepat X, Butler J P, Millet E, Morgan K G, Weitz D A and Fredberg J J 2006 Fast and slow dynamics of the cytoskeleton *Nat. Mater.* 5 636–40
- [33] Zhou E H, Trepat X, Park C Y, Lenormand G, Oliver M N, Mijailovich S M, Hardin C, Weitz D A, Butler J P and Fredberg J J 2009 Universal behavior of the osmotically compressed cell and its analogy to the colloidal glass transition *Proc. Natl Acad. Sci. USA* **106** 10632
- [34] Trepat X, Deng L, An S S, Navajas D, Tschumperlin D J, Gerthoffer W T, Butler J P and Fredberg J J 2007 Universal physical responses to stretch in the living cell *Nature* 447 592
- [35] Krishnan R et al 2009 Reinforcement versus fluidization in cytoskeletal mechanoresponsiveness PLoS ONE 4 e5486
- [36] Fernandez P and Ott A 2008 Single cell mechanics: stress stiffening and kinematic hardening *Phys. Rev. Lett.* 100 238102
- [37] Xu J, Tseng Y and Wirtz D 2000 Strain hardening of actin filament networks. Regulation by the dynamic cross-linking protein alpha-actinin J. Biol. Chem. 275 35886
- [38] Chaudhuri O, Parekh S H and Fletcher D A 2007 Reversible stress softening of actin networks Nature 445 295
- [39] Wolff L and Kroy K 2009 Mechanical stability: a construction principle for cells Diffus. Fundam. 11 56

New Journal of Physics 12 (2010) 053024 (http://www.njp.org/)

16

- [40] Glaser J, Chakraborty D, Kroy K, Lauter I, Degawa M, Kirchgessner N, Hoffmann B, Merkel R and Giesen M 2009 Tube width fluctuations in F-actin solutions arXiv:0910.5864
- [41] Bell G I 1978 Models for specific adhesion of cells to cells Science 200 618-27
- [42] Schwarz U, Erdmann T and Bischofs I 2006 Focal adhesions as mechanosensors: the two-spring model BioSystems 83 225–32
- [43] Dichtl M A and Sackmann E 2002 Microrheometry of semiflexible actin networks through enforced singlefilament reptation: frictional coupling and heterogeneities in entangled networks *Proc. Natl Acad. Sci. USA* 99 6533
- [44] Grason G and Bruinsma R 2008 Chirality and equilibrium biopolymer bundles Phys. Rev. Lett. 99 98101
- [45] Wolff L, Bullerjahn J and Kroy K unpublished
- [46] Evans E and Ritchie K 1997 Dynamic strength of molecular adhesion bonds Biophys. J. 72 1541-55
- [47] Feneberg W, Westphal M and Sackmann E 2001 Dictyostelium cells' cytoplasm as an active viscoplastic body *Eur. Biophys. J.* 30 284–94
- [48] Hallatschek O, Frey E and Kroy K 2007 Tension dynamics in semiflexible polymers. I. Coarse-grained equations of motion *Phys. Rev.* E 75 031905
- [49] Delaey L, Krishnan R V, Tas H and Warlimont H 1974 Thermoelasticity, pseudoelasticity and the memory effects associated with martensitic transformations J. Mater. Sci. 9 1521–35
- [50] Bausch A R and Kroy K 2006 A bottom-up approach to cell mechanics Nat. Phys. 2 231-8
- [51] Wang N, Tolic-Norrelykke I M, Chen J X, Mijailovich S M, Butler J P, Fredberg J J and Stamenovic D 2002 Cell prestress. Stiffness I and prestress are closely associated in adherent contractile cells Am. J. Physiol. Cell Physiol. 282 C606–16
- [52] Discher D, Janmey P and Wang Y 2005 Tissue cells feel and respond to the stiffness of their substrate Science 310 1139–43
- [53] Salbreux G, Joanny J F, Prost J and Pullarkat P 2007 Shape oscillations of non-adhering fibroblast cells *Phys. Biol.* 4 268–84