

Nanomechanics of hierarchical biological materials

Lecture 6

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Introduction to Mechanics of Materials 1. Basic concepts of mechanics, stress and strain, deformation, strength and fracture Monday Jan 8, 09-10:30am Introduction to Classical Molecular Dynamics 2. Introduction into the molecular dynamics simulation; numerical techniques Tuesday Jan 9, 09-10:30am **Mechanics of Ductile Materials** 3. Dislocations; crystal structures; deformation of metals Tuesday Jan 16, 09-10:30am The Cauchy-Born rule 4. Calculation of elastic properties of atomic lattices Friday Jan 19, 09-10:30am **Dynamic Fracture of Brittle Materials** 5. Nonlinear elasticity in dynamic fracture, geometric confinement, interfaces Wednesday Jan 17, 09-10:30am **Mechanics of biological materials** 6. Monday Jan. 22, 09-10:30am Introduction to The Problem Set 7. Atomistic modeling of fracture of a nanocrystal of copper. Wednesday Jan 22, 09-10:30am Size Effects in Deformation of Materials 8. Size effects in deformation of materials: Is smaller stronger? Friday Jan 26, 09-10:30am



- Topic: Elasticity in biological materials: Entropic versus energetic contributions
- **Examples:** Deformation of collagen, vimentin, ...: Protein mechanics
- Material covered: Covalent bonding and models, chemical complexity, reactivity, molecular potentials: CHARMM
- Important lesson: Models for bonding in proteins, entropic vs. energetic elasticity
- **Historical perspective:** AFM, single molecule mechanics





Very brief review:

Material covered in last 2-3 lectures



Figure by MIT OCW.

Ductile vs. brittle Depends on atomic behavior at crack tip





Figure by MIT OCW.

Dislocations are the <u>discrete</u> entities that carry plastic (permanent) deformation; measured by "Burgers vector"

http://www.people.virginia.edu/~lz2n/mse209/Chapter7.pdf



Brittle or ductile?











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1. Buehler, M. J., et al. "The dynamical complexity of work-hardening: a large scale molecular dynamics situation." *Acta Mechanica Sinica* 21, no. 2 (2005): 103-111.

2. Buehler, M. J., et al. "Atomic plasticity: description and analysis of a one-billion atom simulation of ductile materials failure." *Computer Methods in Applied Mechanics and Engineering* 193, no. 48-51 (2004): 5257-5282.







$$F = k_s \Delta u$$





1st law of TD



 $\frac{\mathrm{d}S}{\mathrm{d}t} \geq \frac{\delta Q}{T}$ Change in entropy is always greater or equal than the entropy supplied in form of heat; difference is due to internal dissipation

$$\frac{\mathrm{d}D}{\mathrm{d}t} = T\frac{\mathrm{d}S}{\mathrm{d}t} - \delta Q \ge 0 \qquad \qquad \delta Q = \mathrm{d}U/\mathrm{d}t - \delta W$$

Dissipation rate

Dissipation rate after consider 1st law of TD:

 $\frac{\mathrm{d}D}{\mathrm{d}t} = \delta W - \frac{\mathrm{d}}{\mathrm{d}t} \left(U - TS \right)$

Dissipation rate=External work rate -change in usable energy U-TS`

F = U - TS is defined as free energy or Helmholtz energy,





Elastic deformation (no dissipation by definition):

$$\frac{\mathrm{d}D}{\mathrm{d}t} = 0 \qquad \qquad \delta W - \frac{\mathrm{d}F}{\mathrm{d}t} = 0 \qquad \text{Assume only internal energy change}$$
$$\delta W = \dot{x}F_e$$
$$\dot{x}\left(F_e - \frac{\mathrm{d}F}{\mathrm{d}x}\right) = 0. \qquad \qquad \underset{\text{dU/dt} = \text{dU/dx dx/dt}}{\overset{\text{Expand equation}}{\overset{\text{du}}{\text{d}t}} \dot{x}F_e - \frac{\mathrm{d}F}{\mathrm{d}x}\frac{\mathrm{d}x}{\mathrm{d}t} = 0$$

Therefore: If applied force equals change in free energy of the system, have elastic deformation`

$$F_e = \frac{\mathrm{d}F}{\mathrm{d}x}$$
 $F_e = \frac{\mathrm{d}U}{\mathrm{d}x}$

With strain energy density: $\Psi = F/V$ $\Psi = U/V$

$$\sigma_{ij} = \frac{\mathrm{d}\Psi}{\mathrm{d}\varepsilon_{ij}} \quad c_{ijkl} = \frac{\mathrm{d}^2\Psi}{\mathrm{d}\varepsilon_{ij}\mathrm{d}\varepsilon_{kl}}$$





- Idea: Use thermodynamic approach to develop expression of elasticity of a atomic microstructure, provided a specific interatomic potential
- Assuming locally homogeneous deformation, take a unit cell representation (typically PBCs) and express free energy density as a function of strain tensor
- This provides direct link between interatomic potential and constitutive behavior
- For example: FCC crystal and EAM potential can calculate elastic properties directly





Idea: Express elastic energy (strain energy density) for a atomistic representative volume element as a function of macroscopic applied strain

$$\Phi = \frac{1}{\Omega_0} \int\limits_{\Omega_0^*} U(l) \, D_\Omega \, d\Omega \qquad \mbox{U(I) a function of deformation} \\ \mbox{gradient}$$

- D_{Ω} : Mapping function, e.g. $D(l_0, \theta, \phi) = \delta_D(l_0 l_0^*)D_{\theta\phi}(\theta, \phi)$
- Impose macroscopic deformation gradient on atomistic volume element, then calculate atomic stress – this corresponds to the macroscopic stress
- Strictly valid only far away from defects in periodic lattice (homogeneous deformation, perfect lattice, amorphous solid-average)
- Allows direct link of potential to macroscopic continuum elasticity



Impose homogeneous strain field on 1D string of atoms

• Then get $\sigma_{ij} = c_{ijkl} \varepsilon_{kl}$ from that



 $r_0 \cdot D$ Atomic volume





Detailed write-up on 1D / 2D Cauchy-Born rule is included on line









Mechanics of biological materials

(a) Brittle



(b) Ductile



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Example: Stretching of proteins

. . .





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Physical origin? Soft matter Nonlinear Viscoelastic (time dependent)



Additional Reading



Book Chapters

T. Courtney, Mechanical Behavior of Materials Chapter 2

D.I. Bower An Introduction to Polymer Physics (Chapter 6 [+7 and 3])

D. Boal Mechanics of the Cell (Chapters 1+2)

D. Whitford Proteins – Structures and Function (Chapters 1-4)



- Interactions between atoms, molecules and/or particles govern the elastic response of materials
- For a fundamental understanding of elasticity, need to consider atomic bonding (chemistry or quantum mechanics)
- Elastic response is governed by change of free energy as a function of deformation (yields stress versus strain); second derivative of free energy with respect to strain yields Young's modulus

F(T,V) = U - TS

- Polymers
- Biological structures and materials (proteins, DNA, ...)



 $\lambda_i = 1 + \Delta_i$

$$\mathcal{E}_{ii} = \mathcal{E}_i = \Delta_i$$

In small-strain elasticity

$$\lambda_i^2 = 1 + 2\Delta_i + \Delta_i^2$$

 $\Delta_i \rightarrow 0$ Leads to (small strains)

 $\lambda_i^2 = 1 + 2\Delta_i$

 λ_i

Extension ratios in three directions; pure tensile stress state (directions of principal stress)





Phenomenological theory that uses the concept of strainenergy functions

$$U = f(\lambda_i)$$

- Shear strains are assumed to be zero (coordinate system of principal stresses)
- U is (physically) not equivalent to the thermodynamic internal energy function; instead it is a function that maps changes in entropy and internal energy into a mathematical function (phenomenological model) free energy density!
- **Examples**: Rivlin, Neo-Hook, ...



$$U = C\left(\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - 3\right)$$

Express "strain energy density" as a function of extension ratios λ_i

C is a constant (parameter) that is related to the Young's modulus (will be derived later)

The SED function is used to calculate the stress for a given deformation state (note: 2nd derivatives=modulus=not constant!!)





Incompressible (retain volume during deformation)

 $U = C(\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - 3)$ $\sigma_1 = \sigma_2 = 0 \quad \sigma_3 = \sigma$

Orthogonal to pulling direction:

$$\lambda_{1}\lambda_{2}\lambda_{3} = 1 \qquad \lambda_{1} = \lambda_{2} \qquad \lambda_{3} = \lambda$$
$$\lambda_{3} = \lambda_{2} = 1/\sqrt{\lambda}$$
$$\longrightarrow \lambda_{3} \qquad U = C(\lambda^{2} + 2/\lambda - 3) \text{ for uniaxial tension}$$

$$U/d\lambda = 2C(\lambda - 1/\lambda^2) = \sigma$$





Then, the relation between extension ratio and stress can be written as

$$\sigma = 2C(\lambda^2 - 1/\lambda)$$

Consider the nominal stress force/unit area of the undeformed medium:

$$\sigma = 2C(\lambda - 1/\lambda^2) \quad \text{For small strains:} \quad \lambda_i = 1 + \varepsilon_i$$

$$\sigma = 2C((1 + \varepsilon)^2 - 1/(1 + \varepsilon)) \approx 2C[\chi + 2\varepsilon - (\chi - \varepsilon) + O(0)]$$

$$\sigma = 6C\varepsilon = E\varepsilon \quad \text{Hooke's law (linear elasticity)} \quad C = E/6$$





High entropy



Low entropy





Figure by MIT OCW.



 Based on the assumption that the rubber entropy S can be calculated in terms of

$$\lambda = l / l_0$$

• For example, it can be shown that

$$U = C(\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - 3) \qquad C = E/6$$
$$\sigma_n = C(\lambda - 1/\lambda^2)$$

This is the result for a neo-Hookean solid

(see Courtney, p. 64)



Note: No change in elastic energy of molecules



Figure by MIT OCW.

Needed to understand elasticity: Expression of free energy as a function of the applied strain!

Here: Entropic elasticity – therefore change in entropy



Figure by MIT OCW. $r^2 = \langle r_b^2 \rangle = n \cdot l^2$ RMS length of the chain (no force applied)

 $\sqrt{n} \cdot l \rightarrow nl$ Maximum extension due to force

Physical meaning of *I*: Length at which molecular bending is uncorrelated





Freely jointed Gaussian chain with *n* links and length *l* each (same for all chains in rubber)

 $S = c - kb^2r^2$ where $b^2 = \frac{3}{2nl^2}$ r distance of chain





Figure by MIT OCW.





$$\Delta S = -kb^{2} \sum_{N_{b}} (\lambda_{1}^{2} - 1)x^{2} + (\lambda_{2}^{2} - 1)y^{2} + (\lambda_{3}^{2} - 1)z^{2}$$

$$\Delta S = -kb^{2} N_{b} [(\lambda_{1}^{2} - 1) < x^{2} > + (\lambda_{2}^{2} - 1) < y^{2} > + (\lambda_{3}^{2} - 1) < z^{2} >]$$

<...> Average values over all N_b chains – need multiply by N_b to get total energy

Isotropic solid: End-to-end distances of the Nb chains are directed equally in all directions; therefore

$$< x^{2} > = < y^{2} > = < z^{2} > = \frac{1}{3} < r_{b}^{2} >$$







The length $< r_b^2 >$ in the unstressed state is equal to the mean square length of totally free chains.

It can be shown that

$$r_{RMS} = \sqrt{n} \cdot l = \sqrt{\langle r_b^2 \rangle}$$

$$< r_b^2 \ge n \cdot l^2$$

$$< x^2 \ge < y^2 \ge < z^2 \ge \frac{1}{3}n \cdot l^2 = \frac{1}{2b^2}$$

$$\Delta S = -kN_b / 2 [(\lambda_1^2 - 1) + (\lambda_2^2 - 1) + (\lambda_3^2 - 1)]$$
No explicit dep.
on *b* any more

$$U = -T\Delta S = \frac{1}{2}N_b kT (\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - 3)$$

$$C = E / 6$$

$$U = C (\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - 3)$$

$$\sigma = (E / 3)(\lambda^2 - 1 / \lambda)$$





For SED: Free energy density Young's modulus $U = -T\Delta S = \frac{1}{2}N_b kT \left(\lambda_1^2 + \lambda_2^2 + \lambda_3^2 - 3\right)$ C = E/6Predictions:

$$E = 3N^*kT \quad N^* = N_b / V$$

Stiffness is proportional to temperature

$$E \sim T$$

Stiffness is proportional to degree of cross-linking (for ideal network, N^* equals twice the cross-link density) $E \sim N^*$





- Developed rigorous link extension ratio and elastic properties of rubber-like materials
- Based on statistical theory and by considering changes in entropy due to deformation, arrived at an expression for the Neo-Hookeian solid
- This enables to link cross-linking density and temperature with Young's modulus



The length at which a filament is capable of bending significantly in independent directions, at a given temperature.

This is defined by a autocorrelation function which gives the characteristic distance along the contour over which the tangent vectors **t(s)** become uncorrelated



Bending deformation (R=radius, EI=flexural rigidity of the rod) - energy $E_{bend} = EI \frac{L}{2R^2}$

Thermal (kinetic) energy per molecule (kinetic theory of gases)
 energy
 3

$$E_{kin,mol} = \frac{3}{2}kT$$

- Example: kT~4E-21 J at room temperature
- Persistence length is defined as $\xi_p = \frac{EI}{kT}$

(unit: length)



The contour length of a molecule is the total length in the stretched configuration, denoted as L

• When $L << \xi_p$

a filament appears relatively straight.

When $L >> \xi_p$

No energetic interactions!



a filament adopts more convoluted shapes

• To pull a highly convoluted molecule apart ($L >> \xi_p$), a force is necessary; define effective spring constant

$$k_{sp} = \frac{3kT}{2L\xi_p} \qquad F \sim k_{sp}x \qquad x \ll L$$



Worm-like chain model



Freely-jointed rigid rods



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Continuously flexible ropes

Worm like chain model







This spring constant is only valid for small deformations from a highly convoluted molecule, with length far from its contour length

$x \ll L$

- A more accurate model (without derivation) is the Worm-like chain model (WLC) that can be derived from the Kratky-Porod energy expression (see D. Boal, Ch. 2)
- A numerical, approximate solution of the WLC model:

$$F = \frac{kT}{\xi_p} \left(\frac{1}{4} \frac{1}{(1 - x/L)^2} - \frac{1}{4} + x/L \right)$$

Marko and Siggia, 1995





- An important building block in biological systems are proteins
- Proteins are made up of amino acids
- 20 amino acids carrying different side groups (R)
- Amino acids linked by the amide bond via condensation
- Proteins have four levels of structural organization: primary, secondary, tertiary and quaternary

Protein structure



- Primary structure: Sequence of amino acids
- Secondary structure: Protein secondary structure refers to certain common repeating structures found in proteins. There are two types of secondary structures: <u>alpha-helix</u> and <u>beta-pleated</u> sheet.
- Tertiary structure: Tertiary structure is the full 3-dimensional folded structure of the polypeptide chain.
- Quartenary Structure: Quartenary structure is only present if there is more than one polypeptide chain. With multiple polypeptide chains, quartenary structure is their interconnections and organization.

A A S X D X S L V E V H X X

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Images removed due to copyright restrictions. Table of amino acid chemical structures. See similar image: http://web.mit.edu/esgbio/www/lm/proteins/aa/aminoacids.gif.



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Collagen features hierarchical structure

Goal: Understand the scale-specific properties and cross-scale interactions

Macroscopic properties of collagen depend on the finer scales

Material properties are scale-dependent

(Buehler, JMR, 2006)

Elasticity of tropocollagen molecules







Covalent bonds (directional) Electrostatic interactions H-bonds vdW interactions

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Model for covalent bonds



$$V(R) = E_{bonded} + E_{non-bonded}$$
$$E_{bonded} = E_{bond-stretch} + E_{angle-bend} + E_{rotate-along-bond}$$

Bonding between atoms described as combination of various terms, describing the angular, stretching etc. contributions

Courtesy of the EMBnet Education & Training Committee. Used with permission. Images created for the CHARMM tutorial by Dr. Dmitry Kuznetsov (Swiss Institute of Bioinformatics) for the EMBnet Education & Training committee (http://www.embnet.org)









Review: CHARMM potential



Chemical type	K _{bond}	b _o
С-С	100 kcal/mole/Å ²	1.5 Å
C=C	200 kcal/mole/Å ²	1.3 Å
C≡C	400 kcal/mole/Å ²	1.2 Å

Bond Energy versus Bond length



Different types of C-C bonding represented by different choices of b_0 and k_b ;

Need to retype when chemical environment changes

$$V_{bond} = K_b (b - b_o)^2$$



Review: CHARMM potential



$$E_{non-bonded} = E_{van-der-Waals} + E_{electrostatic}$$

$$E_{van-der-Waals} = \sum_{\substack{nonbonded \\ pairs}} \left(\frac{A_{ik}}{r_{ik}^{12}} - \frac{C_{ik}}{r_{ik}^{6}} \right)$$

Image removed for copyright restrictions. See the graph on this page:

http://www.ch.embnet.org/MD_tutorial/pages/MD.Part2.html

$$E_{electrostatic} = \sum_{\substack{nonbonded \\ pairs}} \frac{q_i q_k}{Dr_{ik}}$$

Nonbonding interactions vdW (dispersive) Coulomb (electrostatic) H-bonding





- Can handle complete periodic table
- Force constants derived using general rules of element, hybridization and connectivity

$$E_{\rm R} = \frac{1}{2}k_{\rm IJ}(r - r_{\rm IJ})^2$$

Features:

- Atom types=elements
- Chemistry based rules for determination of force constants

 $r_{\rm IJ} = r_{\rm I} + r_{\rm J} + r_{\rm BO} + r_{\rm EN}$

Pauling-type bond order correction

 $r_{\rm BO} = -\lambda(r_{\rm I} + r_{\rm J}) \ln (n)$

$$r_{\rm EN} = r_{\rm I} r_{\rm J} (\sqrt{\chi_{\rm I}} - \sqrt{\chi_{\rm J}})^2 / (\chi_{\rm I} r_{\rm I} + \chi_{\rm J} r_{\rm J})$$

$$k_{\rm IJ} = \left(\frac{\partial^2 E_{\rm r}}{\partial R^2}\right)_0 = 2G \, \frac{Z_{\rm I}^* Z_{\rm J}^*}{R^3} = 664.12 \frac{Z_{\rm I}^* Z_{\rm J}^*}{r_{\rm IJ}^3}$$







http://www.pharmacy.umaryland.edu/faculty/amackere/force_fields.htm http://amber.scripps.edu/





Hydrogen bonding e.g. between O and H in H_2O Between N and O in proteins...

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http://www.columbia.edu/cu/biology/courses/c2005/ima ges/3levelpro.4.p.jpg



Figure by MIT OCW. After Ackbarow and Buehler, 2007.







Titin I27 domain: Very resistant to unfolding due to parallel Hbonded strands

Figure by MIT OCW.

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Three-point bending test: Tropocollagen molecule





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Source: Buehler, M. J., and S. Y. Wong. "Entropic Elasticity Controls Nanomechanics of Single Tropocollagen Molecules." *Biophys J* 93 (2007): 37-43.



Three-point bending test: Tropocollagen molecule





Deformation Rate m/sec

MD: Calculate bending stiffness; consider different deformation rates

Result: Bending stiffness at zero deformation rate (extrapolation)

Yields: Persistence length – between 3 nm and 25 nm (experiment: 7 nm)

Source: Buehler, M. J., and S. Y. Wong. "Entropic Elasticity Controls Nanomechanics of Single Tropocollagen Molecules." *Biophys J* 93 (2007): 37-43.





Figure by MIT OCW. Source: Buehler, M. J., and S. Y. Wong. "Entropic Elasticity Controls Nanomechanics of Single Tropocollagen Molecules." *Biophys J* 93 (2007): 37-43.