

Last time poroelastic \rightarrow electrokinetic

$$\frac{\partial u_1}{\partial t} = Hk' \frac{\partial^2 u_1}{\partial x_1^2} + U_0 + \frac{k_{12}}{k_{22}} J_0$$

- neglect inertia - solid $L/\lambda \gg 1$
- fluid $Re \ll 1$

$$k' \equiv k_{11} - \frac{k_{21}k_{12}}{k_{22}} > 0 \text{ positive diffusivity}$$



- (1) $\sigma_{11} = (2G + \lambda) \epsilon_{11} - p$
- (2) $\frac{\partial \sigma_{11}}{\partial x_1} = 0$
- (3) $U_1 = -\frac{\partial u_1}{\partial t} + U_0$
- (4) $U_1 = -k_{11} \frac{\partial p}{\partial x_1} + k_{12} \frac{\partial v}{\partial x_1}$
- (5) $J_1 = k_{21} \frac{\partial p}{\partial x_1} - k_{22} \frac{\partial v}{\partial x_1}$
- (6) $\frac{\partial J_1}{\partial x_1} = 0$

Stacy

Ohm

charge relaxation is $\tau \sim ns$

- new effects - electroosmotic flows - flows generated by ΔV
- streaming potentials - ΔV generated by flow
- electrokinetics \Rightarrow fixed charge density (cartilage, cornea, skin, artery, extracellular matrix ECM, ...)
- e.g. $[NH_3^+]$, $[COO^-]$, $[SO_3^-]$ from GAGs
- not covered: osmotic pressure, dependence of k, λ, G on concentration

Today: transition to cell mechanics

<p>molecules (nm, pN)</p> <p>statistical mechanics</p> <p>single molecules</p>	<p>tissues (cm, mN-N)</p> <p>continuum mechanics</p> <p>poro/visco-elastic materials</p>
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- cells
 - molecular motors
 - ion channels
 - cytoskeletal filaments
- continuum mechanics
 - CSK as a continuum
 - CSK filaments

- mechanics: why cells?
- cartilage: its cell content doesn't really affect its stiffness, its mechanical behavior - but cells respond to mechanical deformation. "cells go along for the ride"
- arterial wall: mechanotransduction (response of endothelial cells to fluid shear stress \rightarrow atherosclerosis)
- smooth muscle cells (contractile, can redistribute the flow)
- muscle: cardiac / skeletal muscle cells, smooth muscle cells
- airway wall: mechanotransduction (epithelial cells)

- Force / biology interaction
- Prototypical example: muscle
 - anatomy / macroscopic behavior
 - activation / contraction
 - cross-bridge dynamics (Huxley)

see slides

• Macroscopic functions of muscles

- tetanus = maximum contracted state

- exponentially stiffening behavior $\frac{d\sigma}{d\varepsilon} = \alpha (\sigma + \beta) \Rightarrow \sigma = C \exp(\alpha \varepsilon) + \beta$



contractile + viscoelastic

- how much force can be generated depending on the velocity or force development?
zero if too fast

Hill's equations

$$\frac{v}{v_{max}} = \frac{1 - \frac{F}{F_{max}}}{1 + C \frac{F}{F_{max}}}$$

purely empirical, { true for all muscles
normalized

efficiency $\eta = \frac{\text{mechanical work}}{\text{chemical energy use}} \approx 25\%$ (comparable to car)

- $\Delta G \approx 25 k_B T$ per molecule

step size (myosin on actin) $\Delta x = 5 \text{ nm}$ and $F \approx 3-4 \text{ pN}$

• Activation / contraction

- depolarization (by nervous cell) conducted transversely $\Rightarrow \text{Ca}^{2+}$ ions released \Rightarrow contraction

after contraction is completed, Ca^{2+} sequestration in SR.

\hookrightarrow from sarcoplasmic reticulum all around

- sliding filament model: thick filament = myosin, walks along the actin
thin filament = actin



dark = overlap (A band) + light = actin only (I band)

A Huxley - Niedergerke

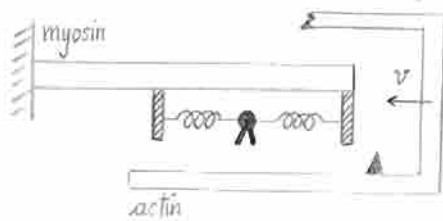
H. Huxley - Hanson

simultaneously in Nature in 1954

ATP-dependent conformational change \Rightarrow power stroke and displacement toward the (+) end of actin

Ca^{2+} removes the { tropomyosin
troponin } hindrance \Rightarrow walk possible

Model by Jonathan Howard (following article by Pate, 1993)



● myosin head

▲ actin binding site

$n(x, t)$ probability of binding

$$\frac{dn}{dt} = \frac{\partial n}{\partial t} + \frac{\partial n}{\partial x} v$$

0 at steady state

$$\frac{dn}{dt} = \frac{dx}{dt} \frac{\partial n}{\partial x} = -v \frac{\partial n}{\partial x} \quad \text{for different } x \text{ regions}$$

