

Cytoskeletal system. Motor proteins. Molecular mechanisms of biological motion.



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Lecture outline

TB. pages 346-356.

- cytoskeleton
 - history
 - polymer mechanics
 - measurement methods
 - polymerization
 - actin
- 10 min pause
- microtubuli
- intermediate filaments
- motor proteins
 - types
 - duty cycle
 - animations

POSSIBLE final exam questions

The cytoskeletal system.
Polymerization of cytoskeletal filaments.

Actin filament system. Actin-dependent biological motion.

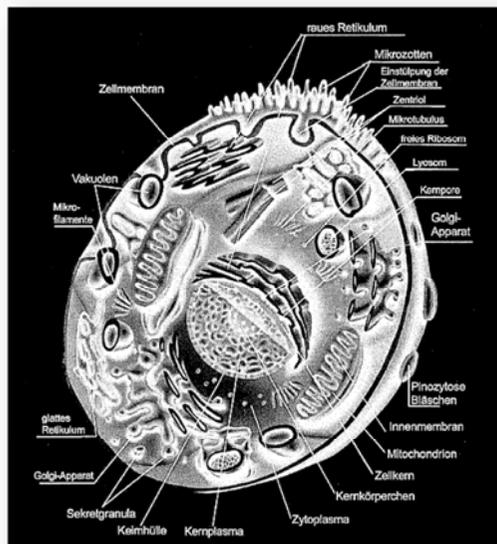
Microtubular system. Microtubule-dependent biological motion.

Intermediate filament system.

Motor proteins. Duty cycle.
Processivity.

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Historical overview



old approach:

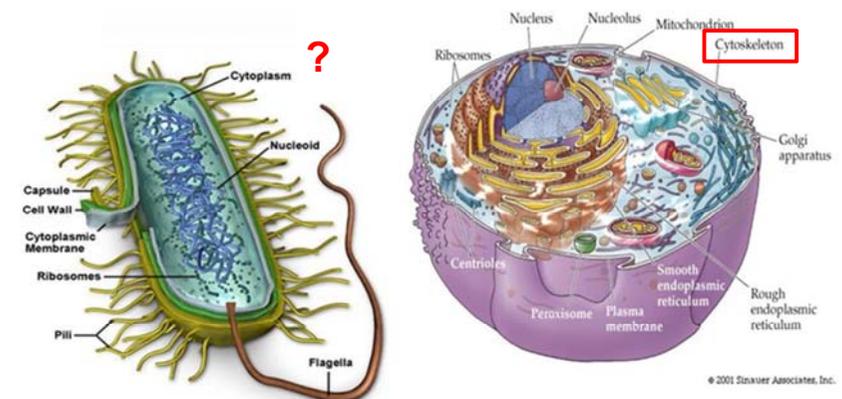
- cell: colloidal system
- membrane-bound structures, organelles
- cytoskeleton: 1-2% volume percent of the cell

„big names“:

- Nikolai Koltsov (1903): the shape of cells is determined by a network of tubules
- Rudolph Peters (1929): protein mosaic coordinates cytoplasmic biochemistry
- Paul Wintrebert (1931):
- *cytosquelette*

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Prokaryotes → Eukaryotes



prokaryotic „cytoskeleton“: analogous to the eukaryotic cytoskeleton

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The cytoskeleton

- dynamic skeleton of eukaryotic cells
- long end-to-end distance (= „filament“):
- network building at low concentrations

three main filament classes:

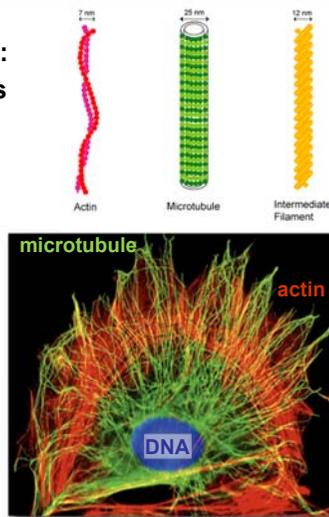
- A. thin (actin) (d ~ 7 nm)
- B. intermediate (d ~ 10 nm)
- C. microtubule (d ~ 25 nm)

+ associated proteins

polymerization: monomer subunits

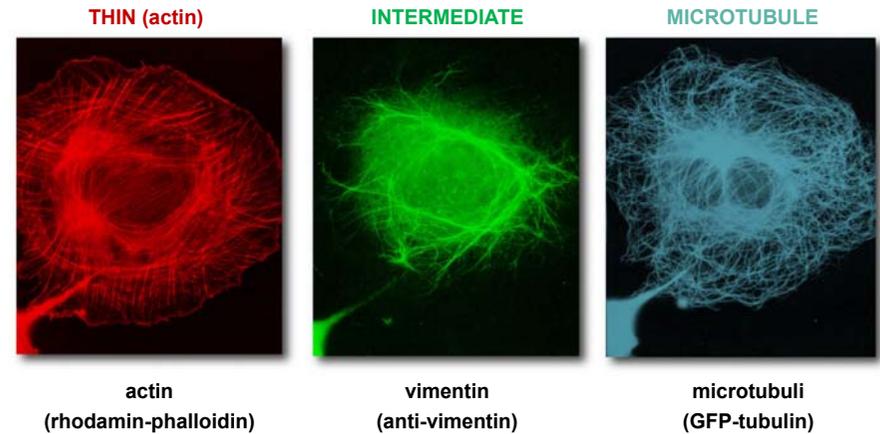
role:

- A. motility, metamorphosis
- B. mechanical stabilization
- C. cell division, transport



The filaments of the cytoskeleton

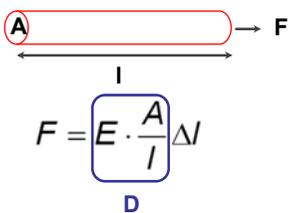
(visualized by fluorescence microscopy)



Polymer mechanics: Hookean elasticity

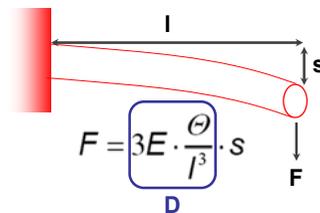
- the stiffness or spring constant ($D = F/\Delta l$) is not only material-dependent
- the stiffness (D) depends on the shape of the body and the direction of the force

Longitudinal stiffness:



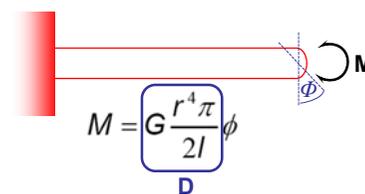
E: elastic modulus (Young modulus)

Bending stiffness:



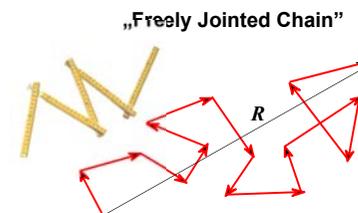
θ : flexural modulus (bending modulus)

Torsional stiffness:



G: shear modulus (modulus of rigidity)

Polymer mechanics: FJC, WLC



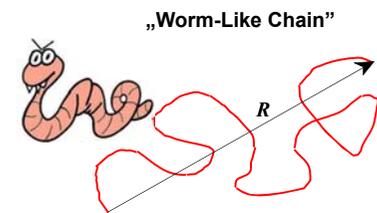
polymer built up from elementary vectors (N pieces, one elementary vector maintains its orientation within the chain)

l = correlation length

(average length of one elementary vector)

$Nl = L$ = contour length

R = end-to-end distance

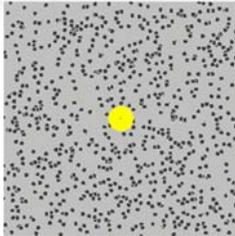


L_p = persistence length (measure of the bending rigidity; the ability of the polymer to maintain its direction)

$$\langle \cos \varphi(s) \rangle = e^{-\frac{s}{L_p}}$$

- if $s \ll L_p$: $\cos(\varphi) \sim 1$, and $\varphi(s) \sim 0^\circ$
- if $s \gg L_p$: $\cos(\varphi) \sim 0$, thus $\varphi(s)$ is between 0° and 360°

Excursion: Brownian motion



$$\frac{1}{2}mv^2 = \frac{3}{2}k_B T \quad (\text{for ideal gases})$$

The small (black) particles move randomly with different velocities and impart momentum to the bigger one (yellow). As a consequence, the yellow particle performs random walk.

Excursion: Entropy

characterizes the disorder / thermodynamic probability of a state in a system



„thermal“ excitation

investing work



Polymer mechanics: „thermal“ elasticity

$$L_p = \frac{E\Theta}{k_B T}$$

Θ : flexural modulus (second moment of inertia)
 – for a rod with circular cross-section:
 $\Theta = r^4 \pi / 4$

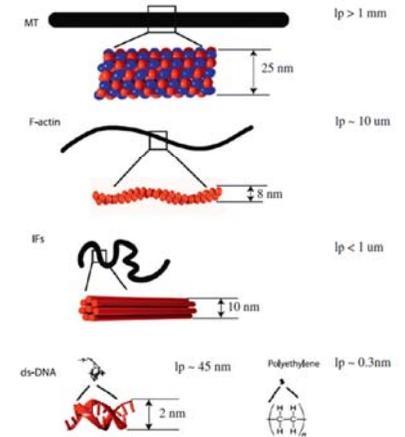
rigid chain
 $L_p \gg L$



semiflexible chain
 $L_p \sim L$

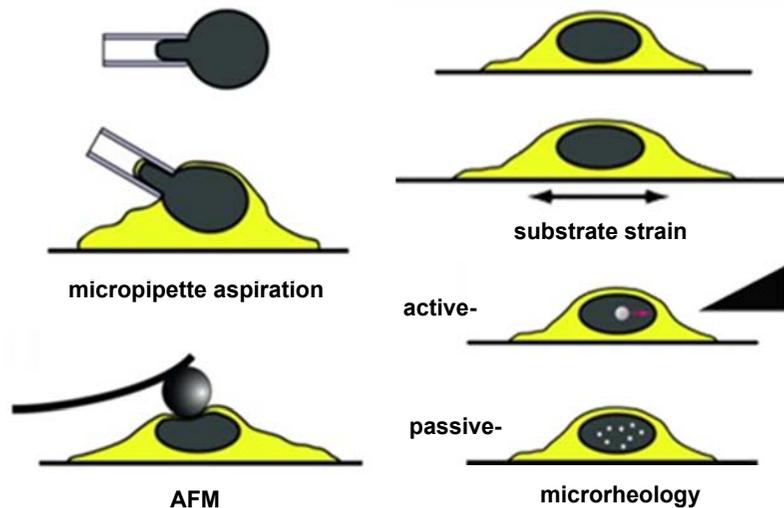


flexible chain
 $L_p \ll L$



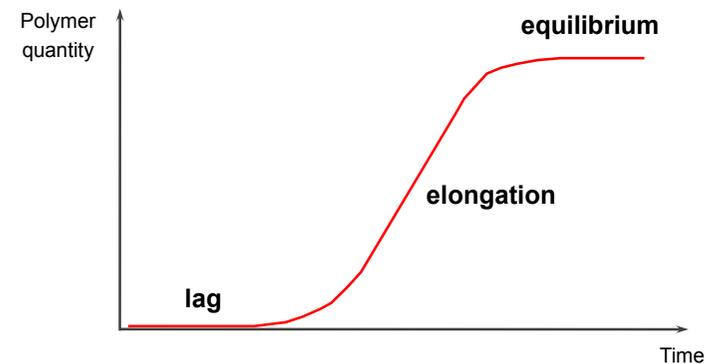
Mechanical measurement methods

(applicable to the nucleus or the whole cell)



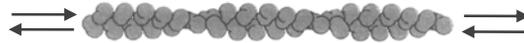
Phases of polymerization

1. Lag phase: nucleation
2. Elongation
3. Equilibrium



Polymerization equilibria

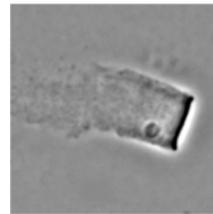
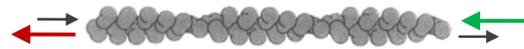
1. real equilibrium:



2. dynamic instability: continuous, slow elongation followed by a catastrophic depolymerization



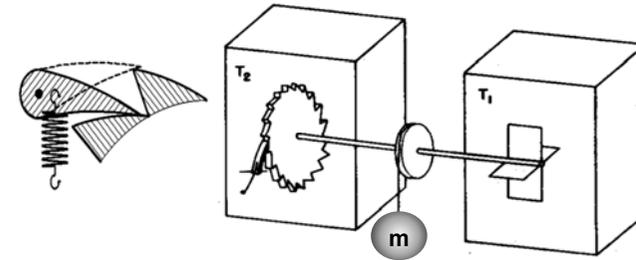
3. treadmilling:



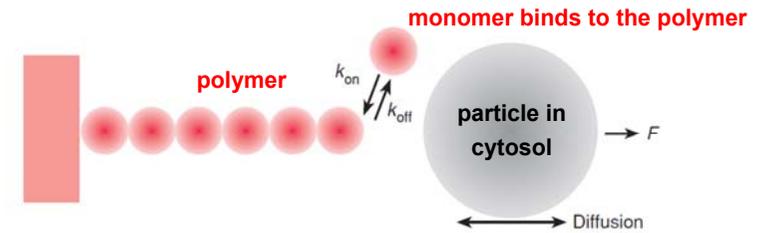
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Force generation by polymerization

(Brownian ratchet mechanism)



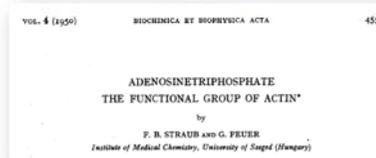
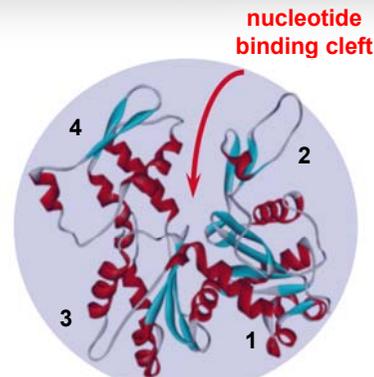
only works if $T_1 > T_2$



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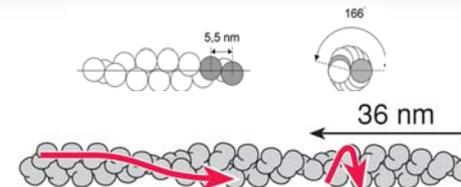
Actin monomer (G-actin)

- actin: 5% of all proteins present in eukaryotic cells,
- monomer unit: globular (G-) actin,
- 43 kDa molecule weight, 4 subdomains (covalent bonds),
- intracellular concentration: 2-8 mg/mL (50-200 μ M): corresponds to ~25 nm average intermolecular distance in solution...
- 1 bound adenosine nucleotide / monomer (ATP or ADP).



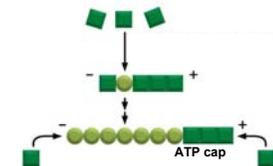
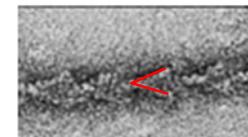
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Actin filament (F-actin)



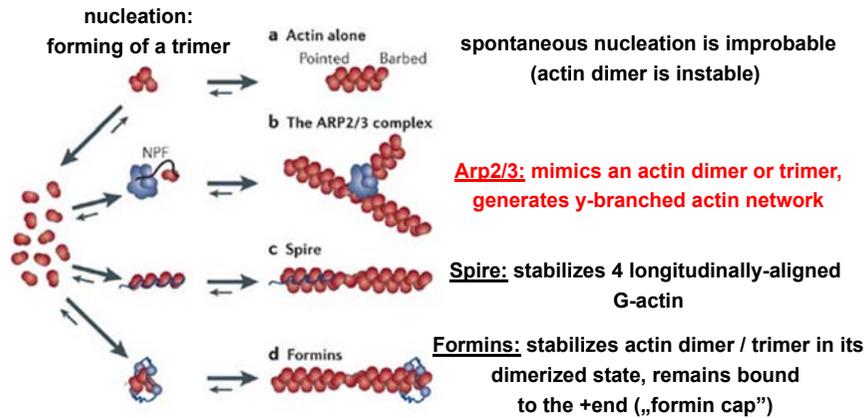
Structure, properties:

- ~7 nm diameter, *in vitro* length: several 10 μ m, *in vivo* length ~1-2 μ m; right-handed long-pitch helix; left-handed short-pitch helix,
- semiflexible polymer (L_p : ~10 μ m),
- structural polarity:
 - barbed end: +,
 - pointed end: -.
- asymmetric polymerization: ATP cap (at the + end).



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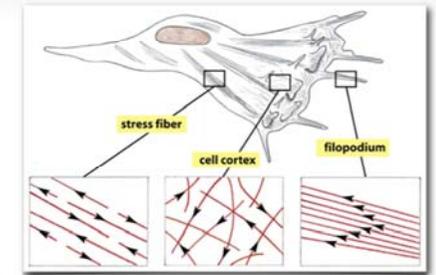
Nucleation factors (associated proteins) of actin



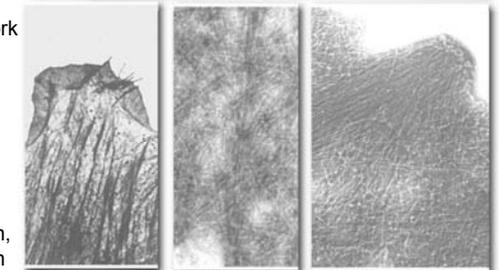
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Occurrence of actin filaments

- **cortex** (periphery of the cell),
- **stress fibers**: cross-linked actin network with myosin-II motors (originally: tension initiates its formation); provides mechanical force for cell adhesion

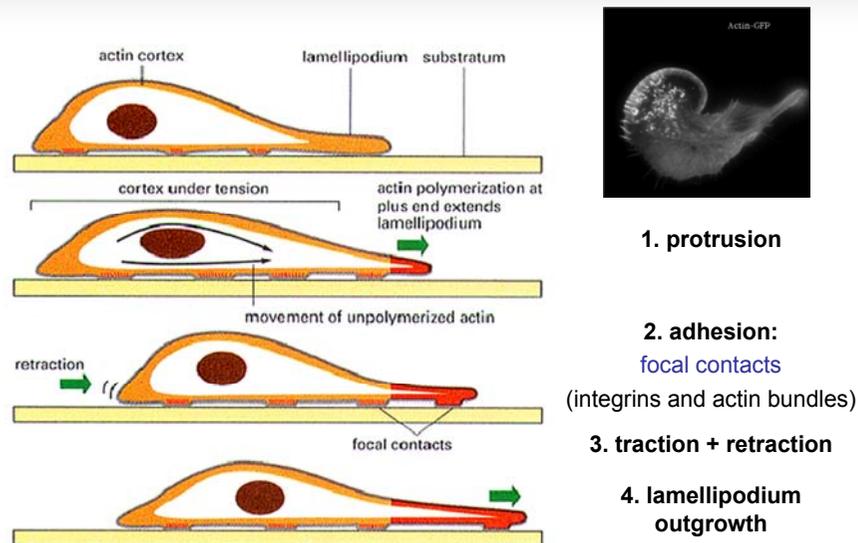


- **cytoplasmic projections**:
 - **lamellipodium**: 2D actin network on the mobile edge of the cell, propels the cell forward
 - **filopodium**: (microspikes) forms focal adhesion with the substrate,
 - **microvillus**: increases the surface area of cells; absorption, secretion, mechanotransduction



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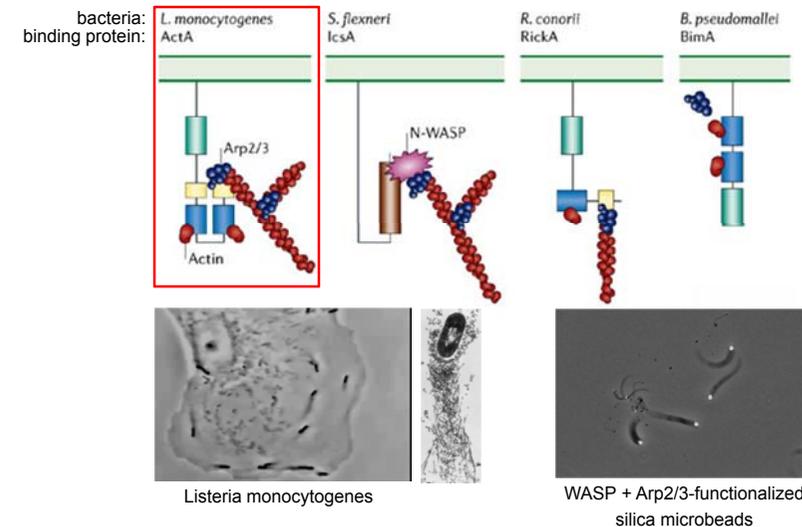
Steps of cell locomotion



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Intracellular locomotion of bacteria

(they use the actin network for „travelling”)

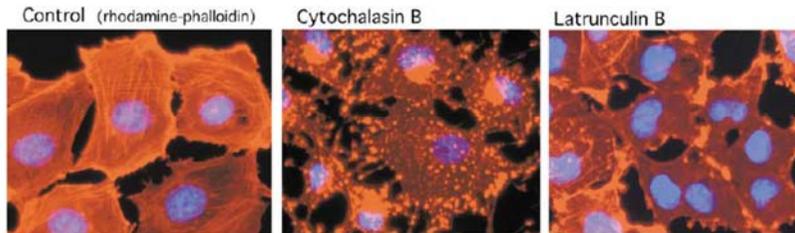


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Actin as target in medicine

(mechanism of effect: inhibition of actin dynamics)

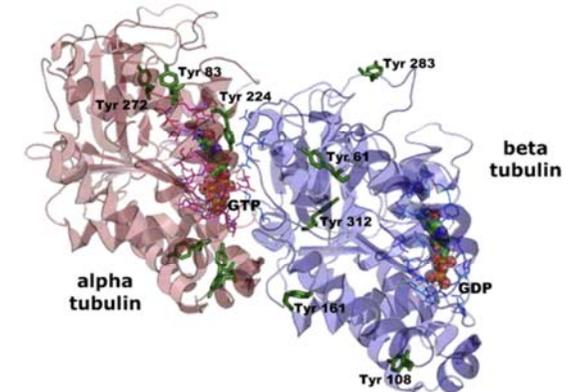
substance	target	disease / application
phalloidin (death cap)	binds to F-actin, inhibits polymerization	mushroom poisoning
cytochalasin (chalysis: relaxation)	inhibits monomer incorporation	experimental tumor therapy
latrunculin	actin monomer (nucleotide-binding cleft)	experimental tumor therapy



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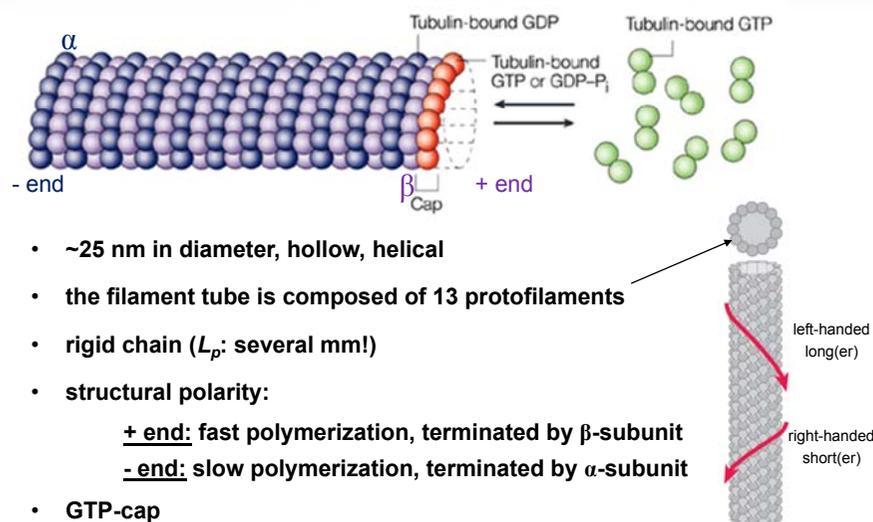
The microtubular system

- subunit: **tubulin** (α - and β -tubulin), ~ 50 kDa
- in neural tissue: up to 10-20% of the total protein
- 1 guanosine nucleotide (GTP or GDP) / subunit
- the nucleotide might be exchangeable (β) or not exchangeable (α)



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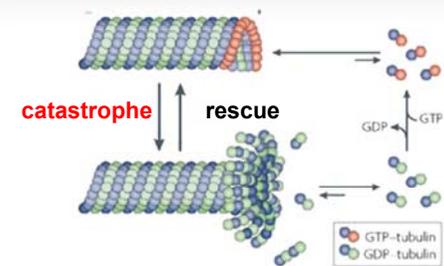
The microtubule



- ~25 nm in diameter, hollow, helical
- the filament tube is composed of 13 protofilaments
- rigid chain (L_p : several mm!)
- structural polarity:
 - + end: fast polymerization, terminated by β -subunit
 - end: slow polymerization, terminated by α -subunit
- GTP-cap

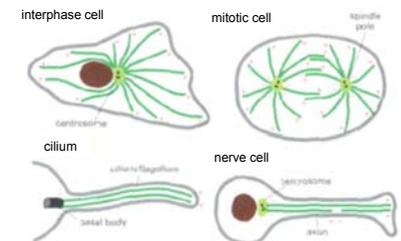
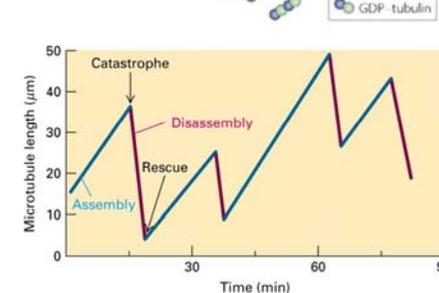
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Microtubular dynamics, occurrence



Where does it occur in eukaryotes?

- cytoplasm of interphase cells,
- axon,
- cilium, flagellum,
- mitotic spindle of mitotic cells.

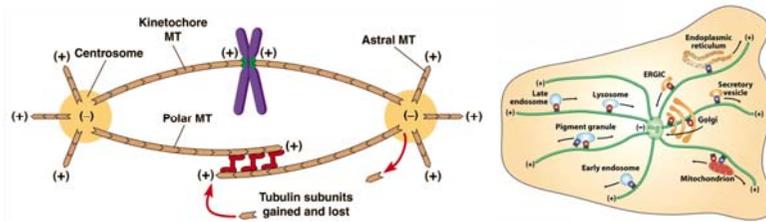


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Localization and function of microtubules

Specific polarity pattern inside the cell:

- in the centrosome: - end, on the periphery: + end,
- **centrosome**: 2 centrioles, centrosome matrix containing γ -tubulin,
- maintaining the polarity of the cell with the help of associated proteins.



Functions:

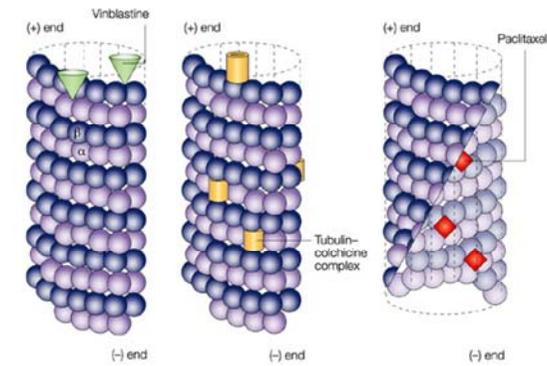
1. „motorways” for motor proteins,
2. senses, finds and monitors the geometric centre of the cell,
3. motility (cell division).

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Microtubuli as targets in medicine

(mechanism of effect: inhibition of microtubule dynamics)

substance	target	disease / application
vinblastin	+ end of the microtubule	cancer therapy
colchicine	tubulin dimer	gout
paclitaxel	interior surface of MT	cancer therapy



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Intermediate filaments

- diameter: 8-10 nm,
- chemically resistant,
- fibrous monomer (non-globular), polymerization does not require ATP/GTP
- tissue-specific monomers differ in their tail sequence / structure:

epithelium	keratins (α , β)
muscle	desmin
connective tissue	vimentin
glia	glial fibrillary acidic protein (GFAP)
nerve	neurofilaments (L, M, H)

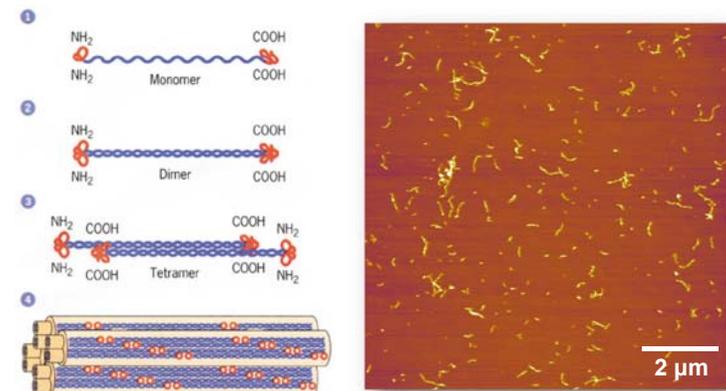
Intermediate filament dimer:



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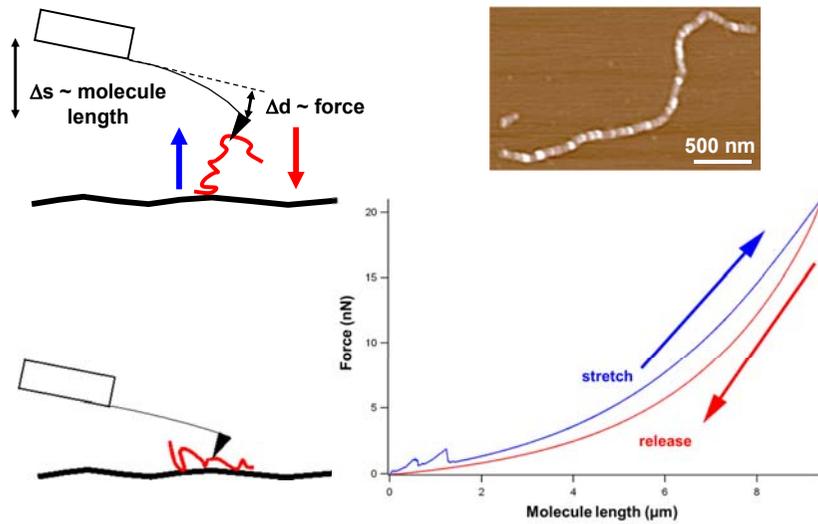
Polymerization of intermediate filaments

- in cells they are usually present in their fully-polymerized state,
- no polymerization-depolymerization dynamism,
- depolymerization is regulated via phosphorylation.



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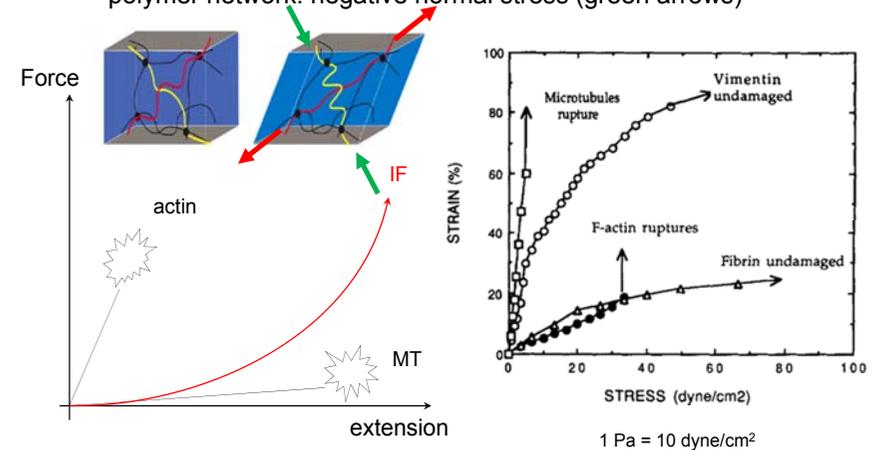
Stretching a single IF (desmin) with AFM



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Hypothetic role of IFs: mechanical stabilization

- nonlinear elasticity: strain hardening at high forces
- polymer network: negative normal stress (green arrows)

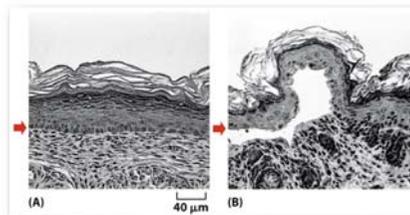


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Tissue-specific role of intermediate filaments

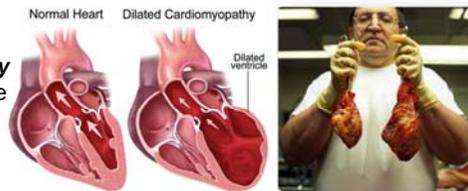
in epithelial cells:

- **clinical picture:** *epidermolysis bullosa simplex*. Defect in anchoring between the epidermis and dermis, resulting in friction and skin fragility: blisters in the skin and mucosal membranes.
- **cause:** keratin gene mutation



in cardiac tissue:

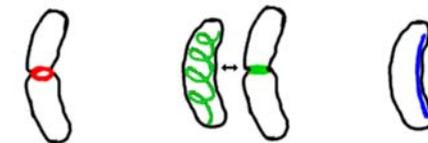
- **clinical picture:** *cardiomyopathy* (deterioration of the function of the myocardium = the heart muscle)
- **cause:** desmin gene mutation



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Point of interest: prokaryotic cytoskeleton

	cell division	cell polarity	cell shape
eukaryotes	tubulin	actin	intermediate filaments
prokaryotes	FtsZ	MreB	CreS



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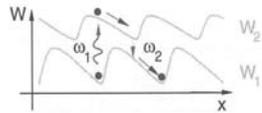
Motor

a cyclic working device capable of converting energy into movement (force and thus displacement)

$$E_x \rightarrow W$$

E_x :

- mechanical energy
- **chemical energy**
- electric energy
- thermal energy



Motor proteins

- non-equilibrium systems (the displacement coupled to the chemical reaction is strictly directed)
- velocities: 0.01-100 $\mu\text{m/s}$
- step sizes: 0.3-40 nm
- forces: 1-60 pN
- fuel: hydrolysis of ATP or similar macroergic compound, (e.g.: coupled to polymerization): 54 kJ/mol energy = $22 k_b T$
- efficiency: 50-100% (!!!)

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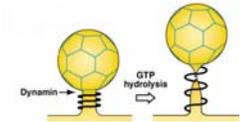
Types of motor proteins

1. Actin-based:

- **myosins**: conventional (myosin-II) and non-conventional;
- myosin superfamily (classes I-XXIV), move towards plus end.

2. Microtubule-based:

- **dyneins**: ciliary (flagellar) and cytoplasmic dyneins;
 - move towards the minus end.
- **kinesins**: kinesin superfamily: conventional and non-conventional;
 - move towards the plus end.
- **dynamins**: MT-dependent GTPase activity;
 - biological role: vacuolar protein sorting



3. DNA-based mechanoenzymes:

- DNA and RNA **polymerases**, virus capsid **packaging motor**;
- produce force and displacement along the DNA strand.

4. Rotary motors:

- F1F0-ATP synthase,
- bacterial flagellar motor.

5. Mechanoenzyme complexes

- ribosome

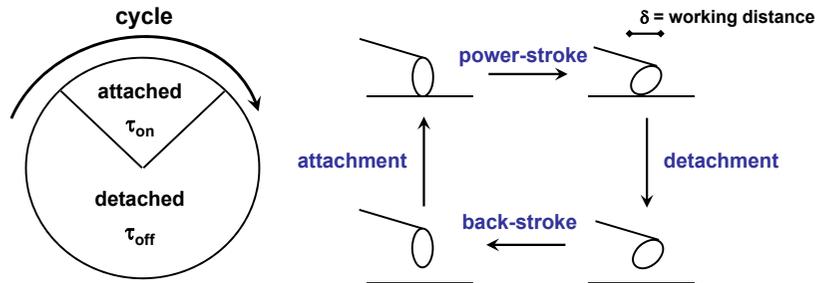
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Duty cycle of motor proteins 1.

C-terminus: functional binding site (e.g.: cargo)

N-terminus: globular head: motor domain (binds and hydrolyzes nucleotide), binding site for the respective cytoskeletal filament

ATP hydrolysis cycle **tight coupling: 1 ATP is being hydrolyzed per 1 cycle**



motor protein conformational change: $\sim \mu\text{s}$

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Duty cycle of motor proteins 2.

duty ratio (r):

$$r = \frac{\tau_{\text{on}}}{\tau_{\text{on}} + \tau_{\text{off}}} = \frac{\tau_{\text{on}}}{\tau_{\text{total}}}$$

sliding velocity:

$$v = \frac{\delta}{\tau_{\text{on}}} \xrightarrow{\text{attached time:}} \tau_{\text{on}} = \frac{\delta}{v} \quad \left. \vphantom{\tau_{\text{on}} = \frac{\delta}{v}} \right\} r = \frac{\delta V}{v}$$

cycle time:

$$\tau_{\text{total}} = \frac{1}{V}$$

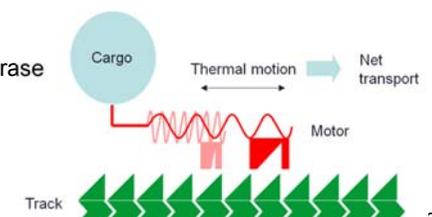
δ =working distance or step size; V =ATP-ase rate; sliding velocity

Processive motor: $r \sim 1$

- e.g.: kinesin, DNA, RNA-polymerase
- carries its load by itself

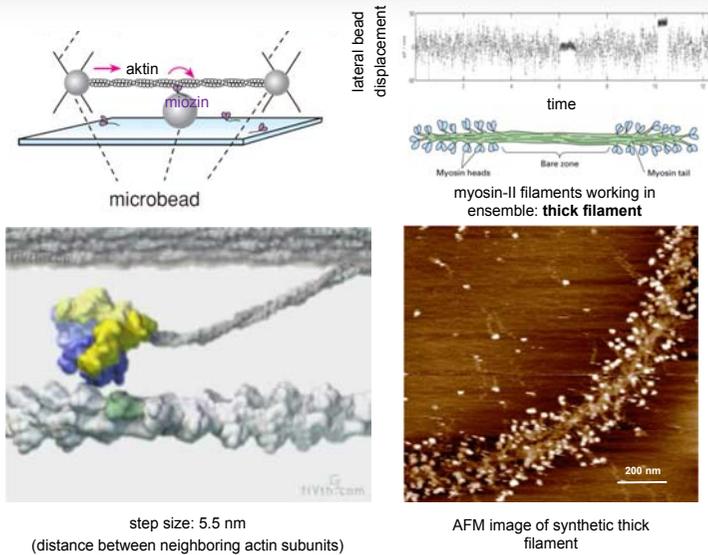
Non-processive motor: $r \sim 0$

- e.g.: myosin-II
- works in ensembles

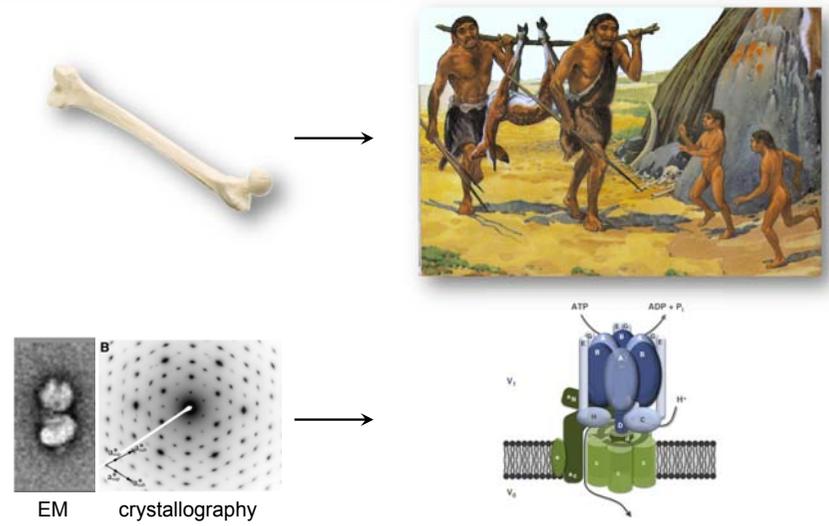


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Non-processive motors: myosin-II



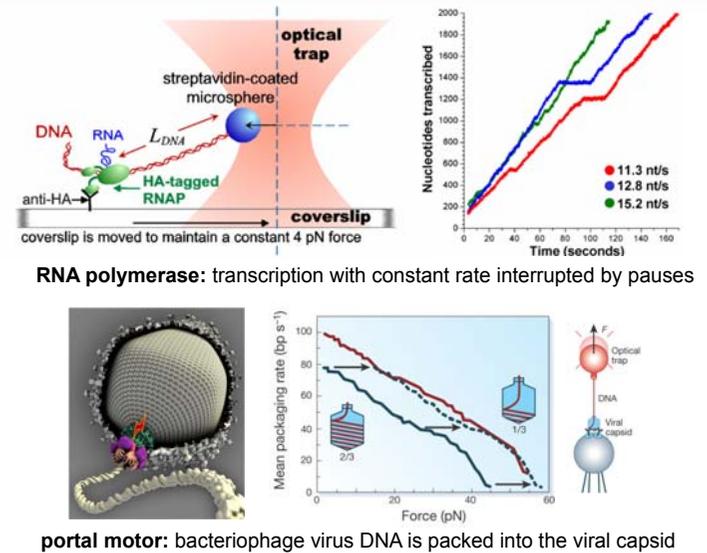
Relaxation: how were the animations made?



Processive motors

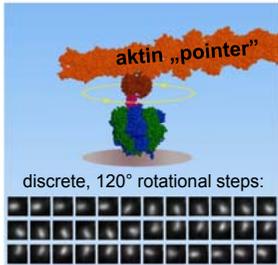
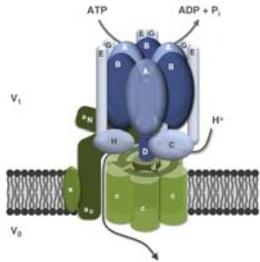


Nucleic acid-based motors

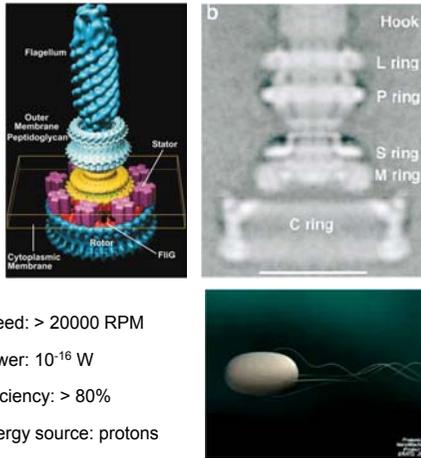


Rotary motors

F1F0 ATP-synthase



Bacterial flagellar motor



- speed: > 20000 RPM
- power: 10^{-16} W
- efficiency: > 80%
- energy source: protons

Thank You for Your attention!

