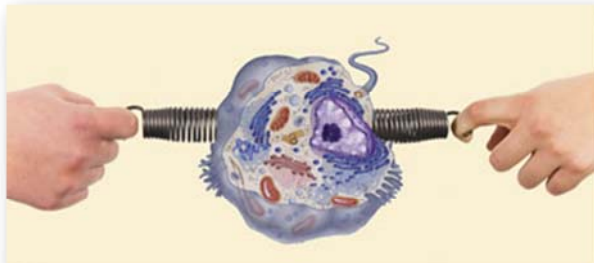


## Cytoskeletal system. Motor proteins. Molecular mechanisms of biological motion.



[http://www.rsc.org/images/b901714c-400-FOR-TRIDION\\_tcm18-152053.jpg](http://www.rsc.org/images/b901714c-400-FOR-TRIDION_tcm18-152053.jpg)



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20. February, 2014

### 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.

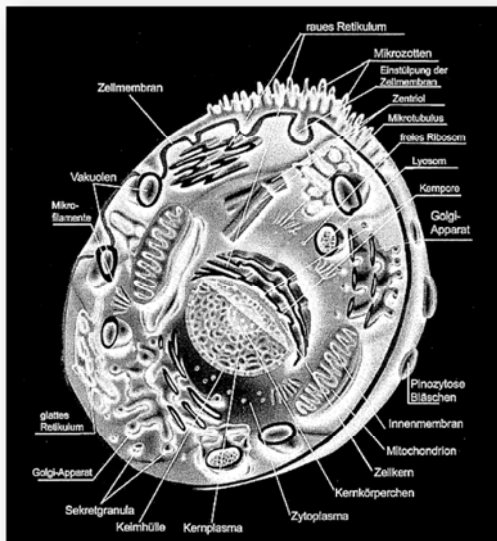
Microtubular system. Microtubule-dependent biological motion.

Intermediate filament system.

Motor proteins. Duty cycle. Processivity.

2

## Historical overview



### old approach:

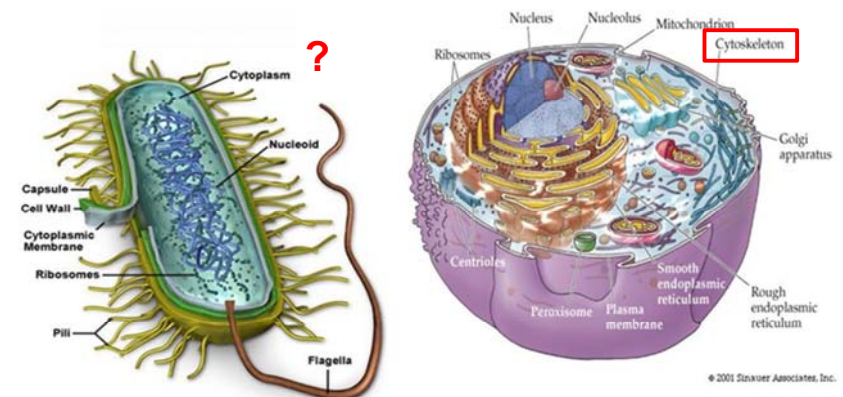
- cell: colloidal system
- membrane-bound structures, organelles
- cytoskeleton: 1-2% volume percent of the cell

### „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*

3

## Prokaryotes → Eukaryotes



prokaryotic „cytoskeleton”: analogous to the eukaryotic cytoskeleton

4

## The cytoskeleton

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

### three main filament classes:

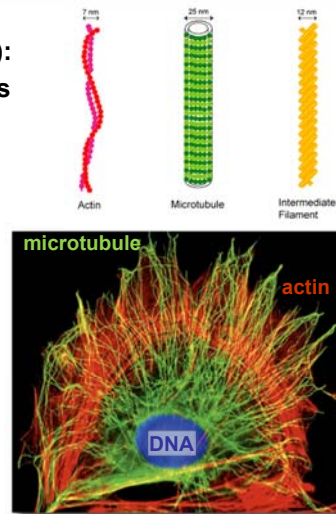
- thin (actin) (d ~ 7 nm)
- intermediate (d ~ 10 nm)
- microtubule (d ~ 25 nm)

+ associated proteins

polymerization: monomer subunits

role:

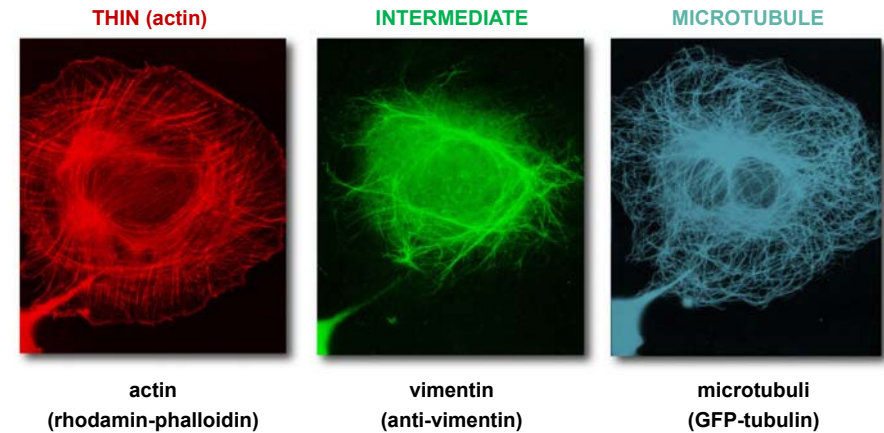
- motility, metamorphosis
- mechanical stabilization
- cell division, transport



5

## The filaments of the cytoskeleton

(visualized by fluorescence microscopy)

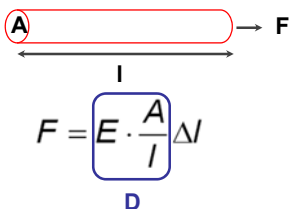


6

## Polymer mechanics: Hookean elasticity

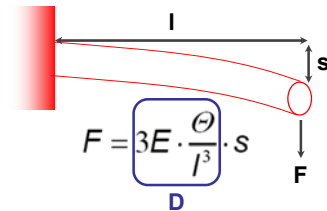
- the stiffness or spring constant ( $D = F/\Delta l$ ) is not only material-dependent
- the stiffness ( $D$ ) depends on the shape of the body and the direction of the force

### Longitudinal stiffness:



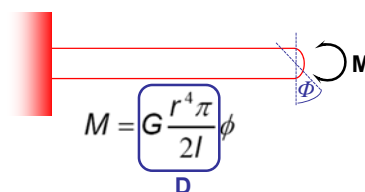
E: elastic modulus (Young modulus)

### Bending stiffness:



$I$ : flexural modulus (bending modulus)

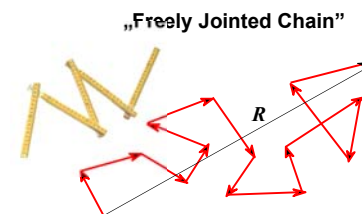
### Torsional stiffness:



G: shear modulus (modulus of rigidity)

7

## Polymer mechanics: FJC, WLC



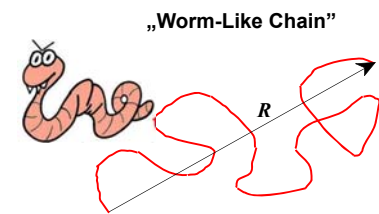
polymer built up from elementary vectors ( $N$  pieces, one elementary vector maintains its orientation within the chain)

$l$  = correlation length

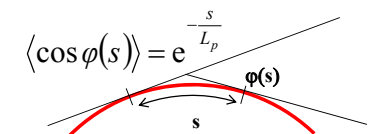
(average length of one elementary vector)

$Nl = L$  = contour length

$R$  = end-to-end distance



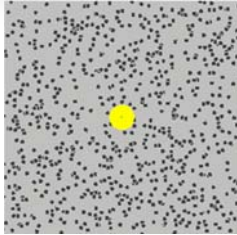
$L_p$  = persistence length (measure of the bending rigidity; the ability of the polymer to maintain its direction)



- 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  $0^\circ$  and  $360^\circ$

8

## 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



9

## Polymer mechanics: „thermal“ elasticity

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

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

rigid chain

$$L_p \gg L$$



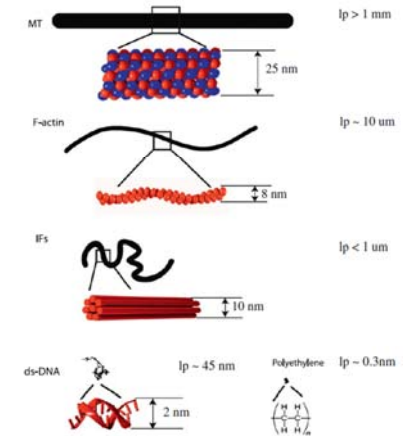
semiflexible chain

$$L_p \sim L$$



flexible chain

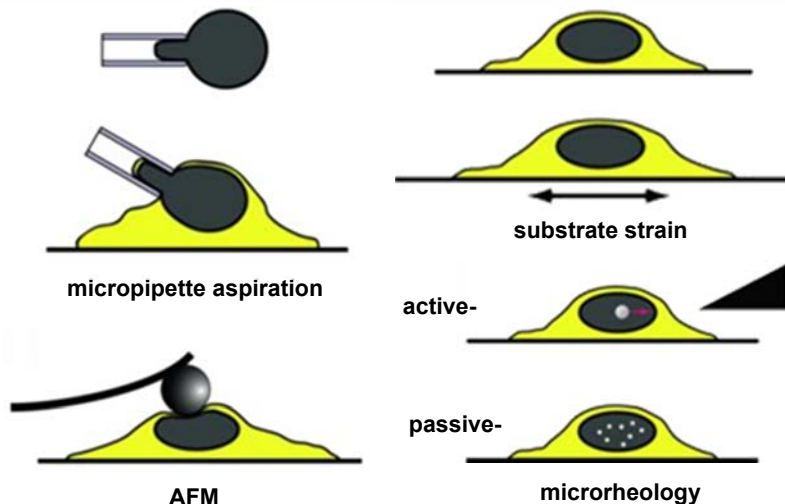
$$L_p \ll L$$



10

## Mechanical measurement methods

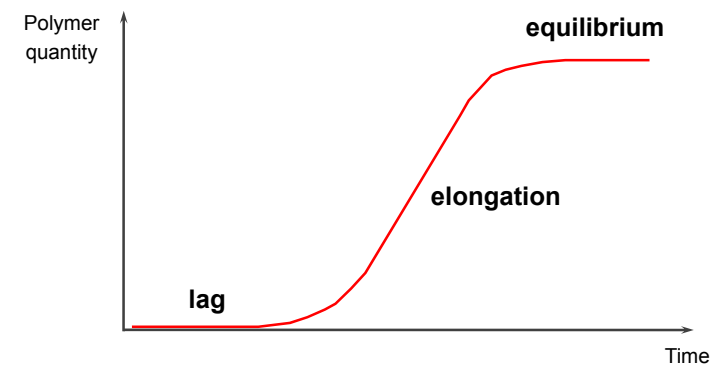
(applicable to the nucleus or the whole cell)



11

## Phases of polymerization

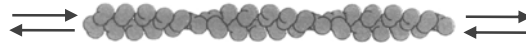
1. Lag phase: nucleation
2. Elongation
3. Equilibrium



12

## Polymerization equilibria

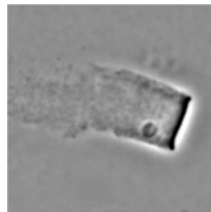
### 1. real equilibrium:



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



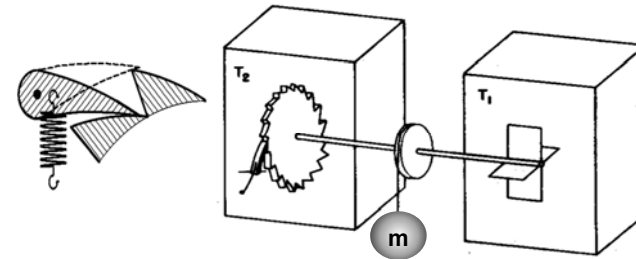
### 3. treadmilling:



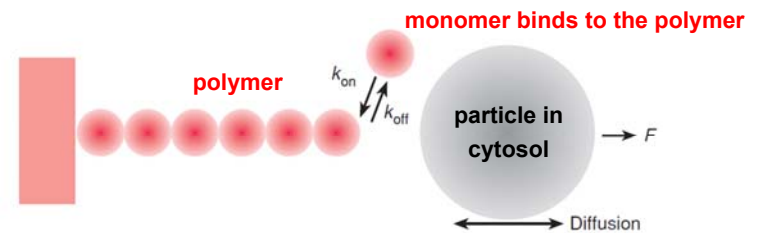
13

## Force generation by polymerization

(Brownian ratchet mechanism)



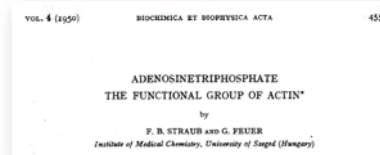
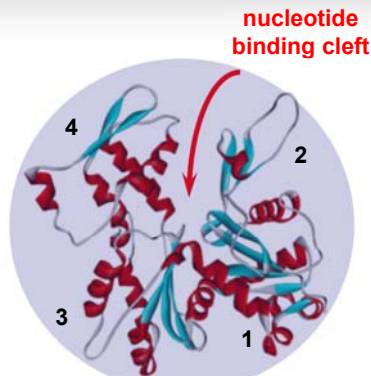
only works if  
 $T_1 > T_2$



14

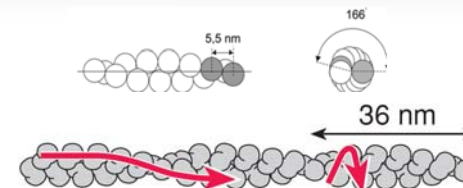
## Actin monomer (G-actin)

- 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  $\mu$ M): corresponds to ~25 nm average intermolecular distance in solution...
- 1 bound adenosine nucleotide / monomer (ATP or ADP).



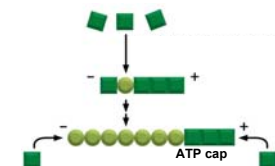
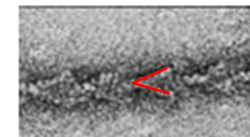
15

## Actin filament (F-actin)



### Structure, properties:

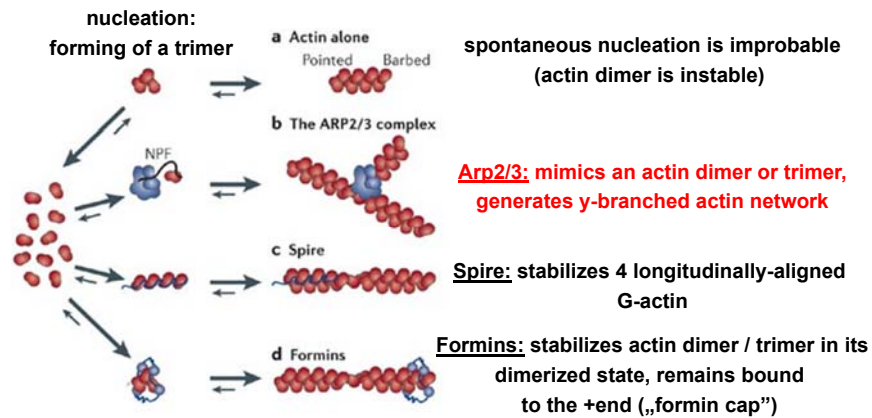
- ~7 nm diameter, *in vitro* length: several 10  $\mu$ m, *in vivo* length ~1-2  $\mu$ m; right-handed long-pitch helix; left-handed short-pitch helix,
- semiflexible polymer ( $L_p$ : ~10  $\mu$ m),
- structural polarity:
  - barbed end: +,
  - pointed end: -.
- asymmetric polymerization: ATP cap (at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



16



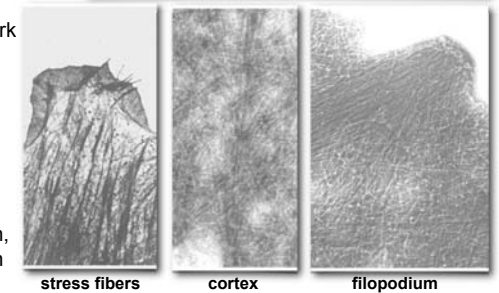
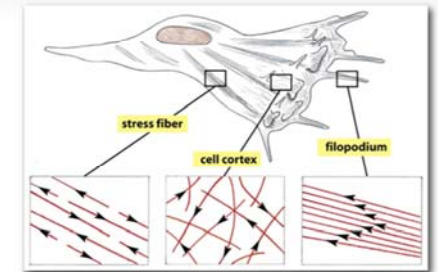
## Nucleation factors (associated proteins) of actin



17

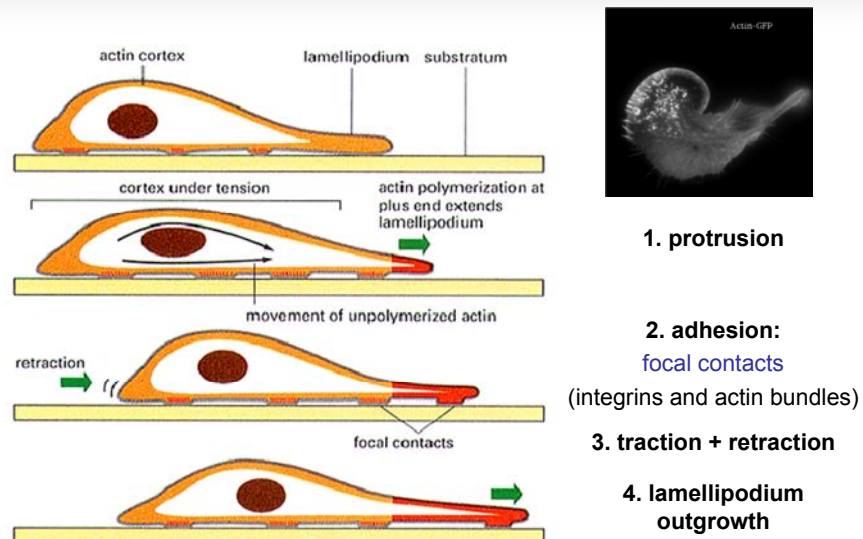
## Occurrence of actin filaments

- **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



18

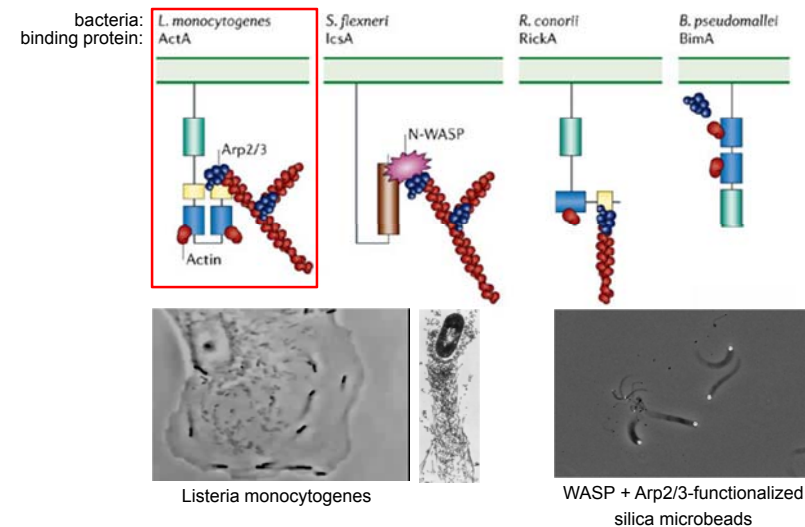
## Steps of cell locomotion



19

## Intracellular locomotion of bacteria

(they use the actin network for „travelling”)

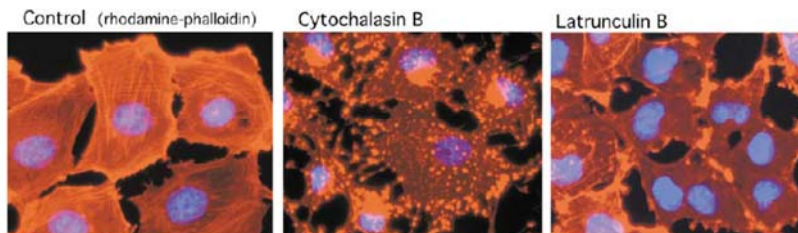


20

## Actin as target in medicine

(mechanism of effect: inhibition of actin dynamics)

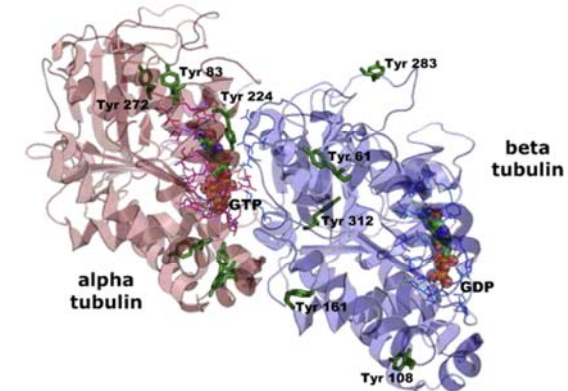
substance	target	disease / application
phalloidin (death cap)	binds to F-actin, inhibits polymerization	mushroom poisoning
cytochalasin (chalysis: relaxation)	inhibits monomer incorporation	experimental tumor therapy
latrunculin	actin monomer (nucleotide-binding cleft)	experimental tumor therapy



21

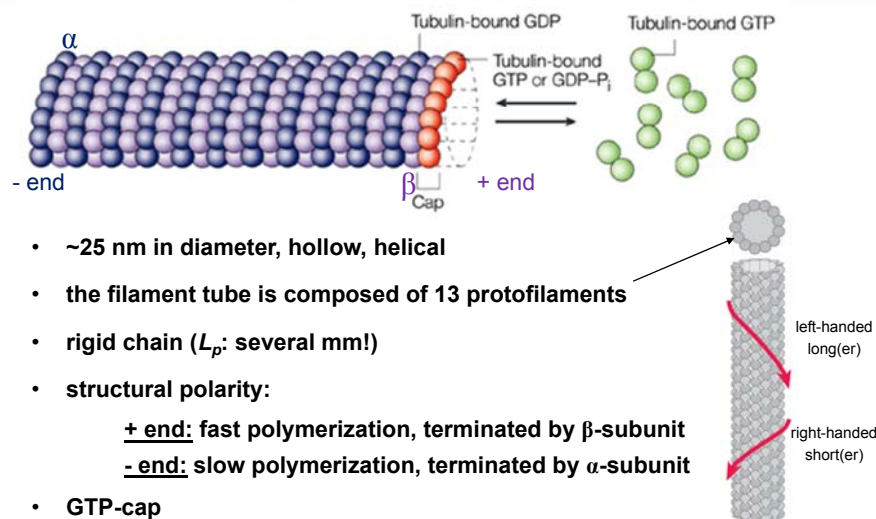
## The microtubular system

- subunit: **tubulin** ( $\alpha$ - and  $\beta$ -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 ( $\beta$ ) or not exchangeable ( $\alpha$ )



22

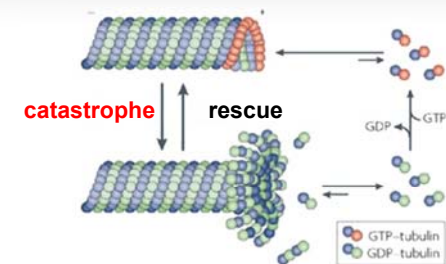
## The microtubule



- ~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  $\beta$ -subunit
  - end: slow polymerization, terminated by  $\alpha$ -subunit
- GTP-cap

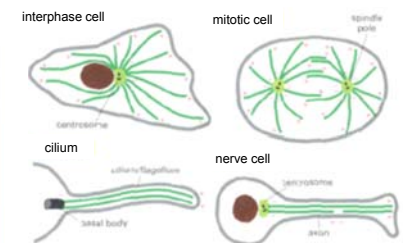
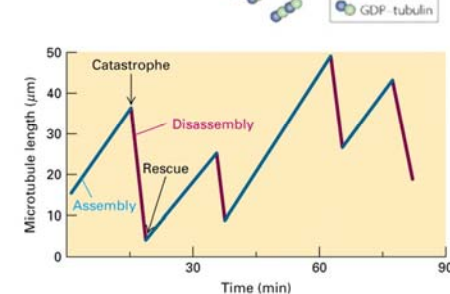
23

## Microtubular dynamics, occurrence



Where does it occur in eukaryotes?

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

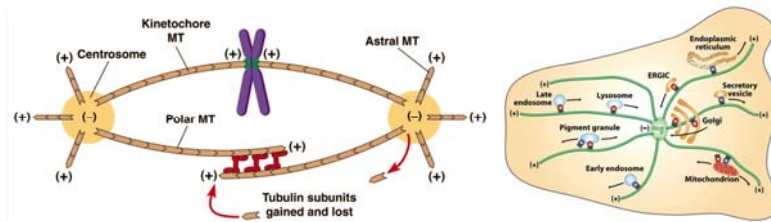


24

## 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  $\gamma$ -tubulin,
- maintaining the polarity of the cell with the help of associated proteins.



### Functions:

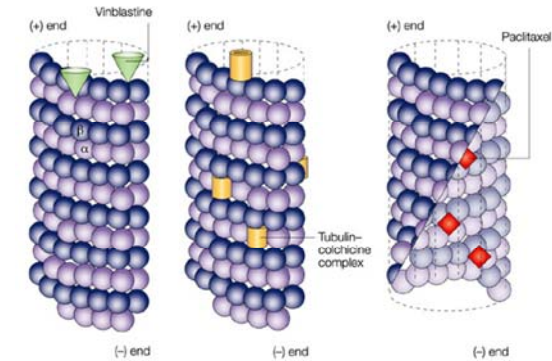
1. „motorways” for motor proteins,
2. senses, finds and monitors the geometric centre of the cell,
3. motility (cell division).

25

## 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



26

## 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 ( $\alpha$ , $\beta$ )
muscle	desmin
connective tissue	vimentin
glia	glial fibrillary acidic protein (GFAP)
nerve	neurofilaments (L, M, H)

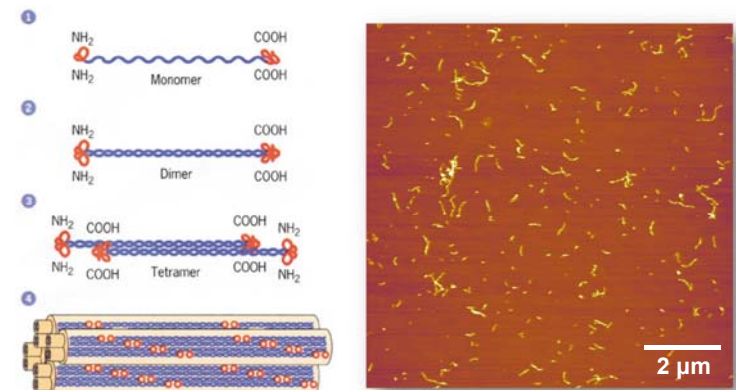
### Intermediate filament dimer:



27

## Polymerization of intermediate filaments

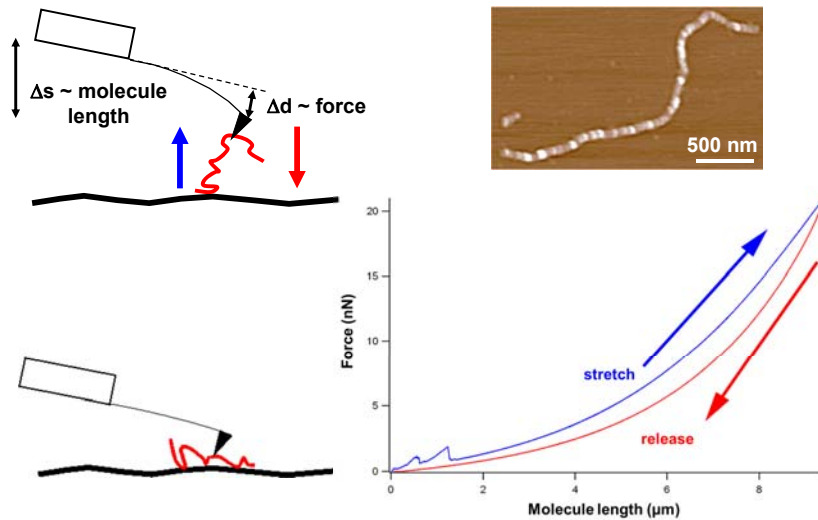
- in cells they are usually present in their fully-polymerized state,
- no polymerization-depolymerization dynamism,
- depolymerization is regulated via phosphorylation.



28



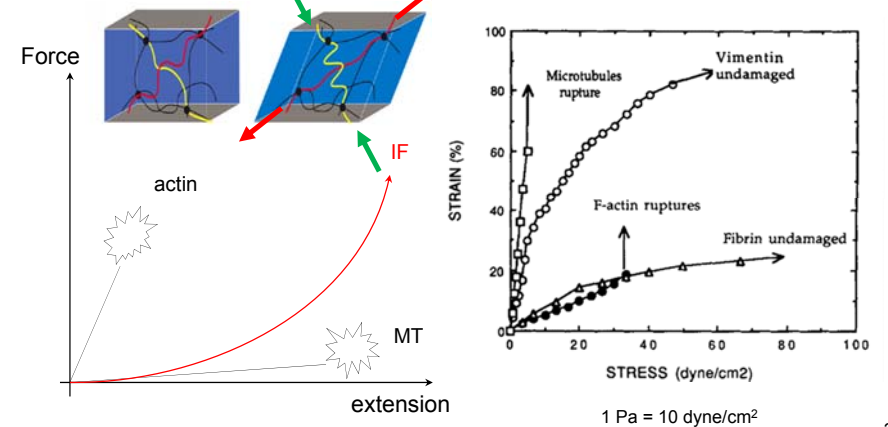
## Stretching a single IF (desmin) with AFM



29

## Hypothetic role of IFs: mechanical stabilization

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

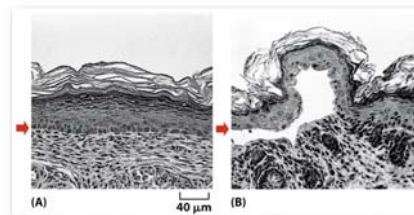


30

## Tissue-specific role of intermediate filaments

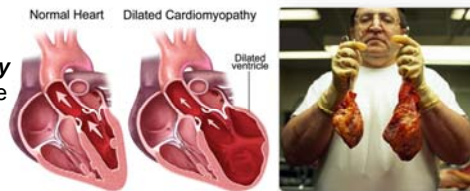
### in epithelial cells:

- 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:

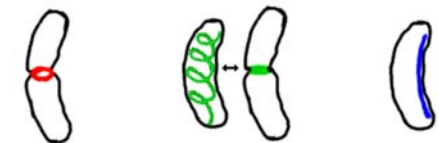
- clinical picture: **cardiomyopathy** (deterioration of the function of the myocardium = the heart muscle)
- cause: desmin gene mutation



31

## Point of interest: prokaryotic cytoskeleton

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



32



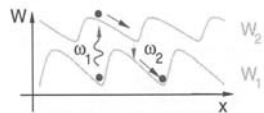
## Motor

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% (!!!)

33

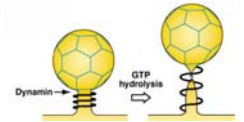
## Types of motor proteins

### 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

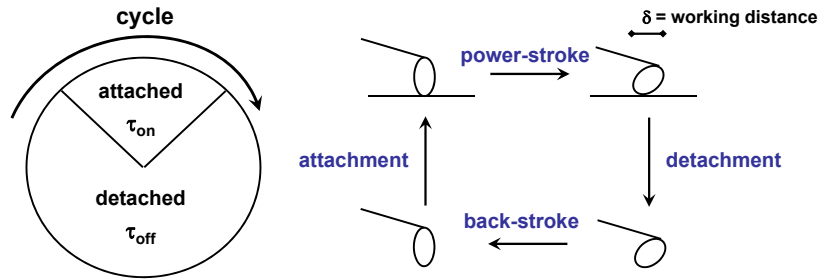
34

## 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:  $\sim \mu\text{s}$

35

## 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:

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

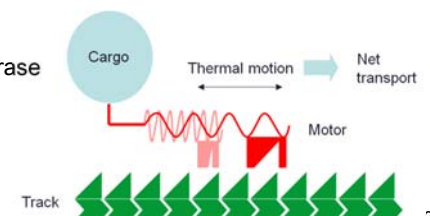
$\delta$ =working distance or step size;  $V$ =ATP-ase rate; sliding velocity

**Processive motor:  $r \sim 1$**

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

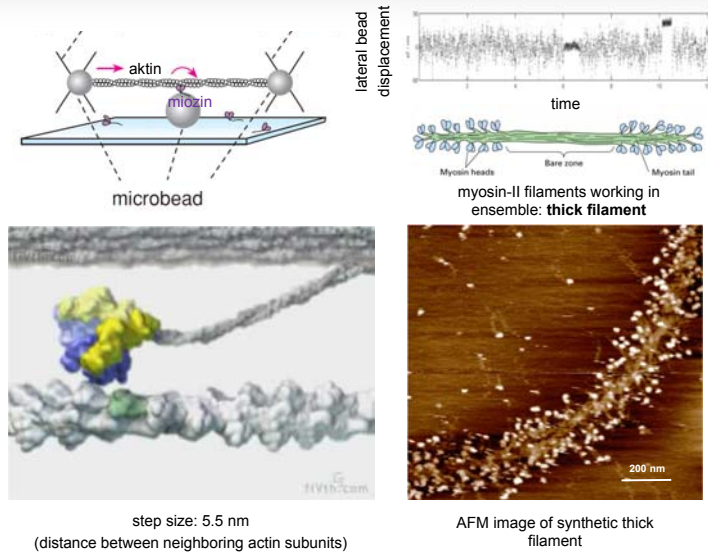
**Non-processive motor:  $r \sim 0$**

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



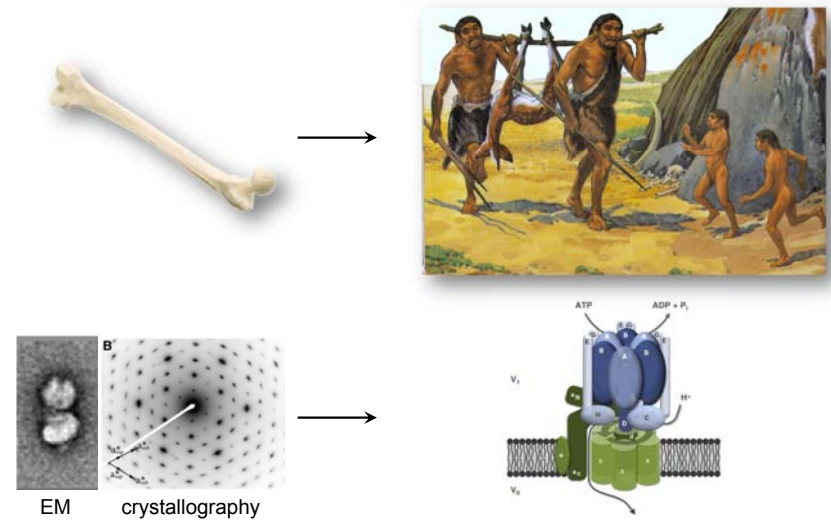
36

## Non-processive motors: myosin-II



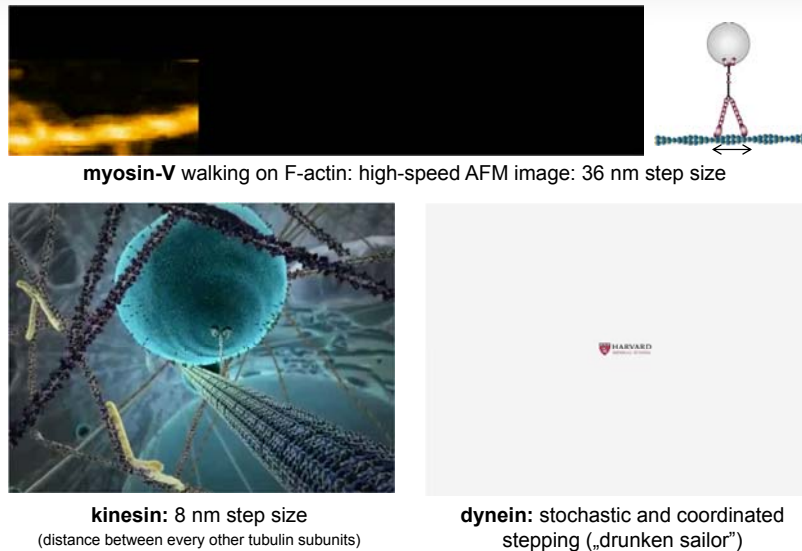
37

## Relaxation: how were the animations made?



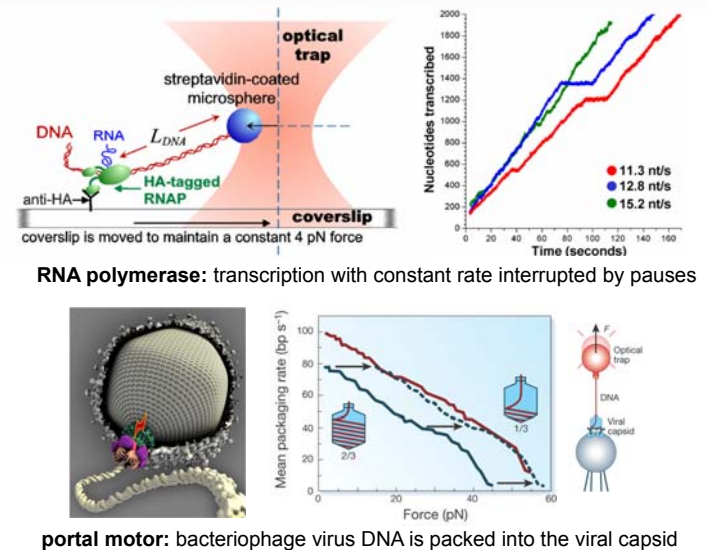
38

## Processive motors



39

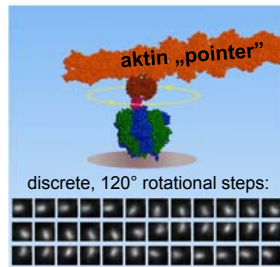
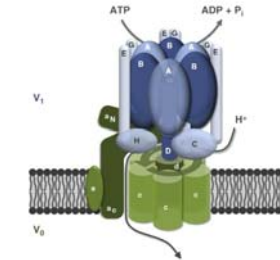
## Nucleic acid-based motors



40

## Rotary motors

F1F0 ATP-synthase



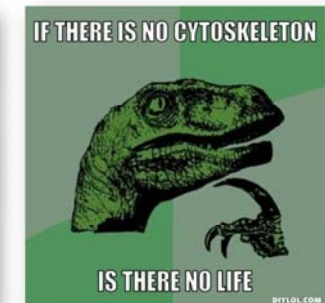
Bacterial flagellar motor



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

41

Thank You for Your attention!



42