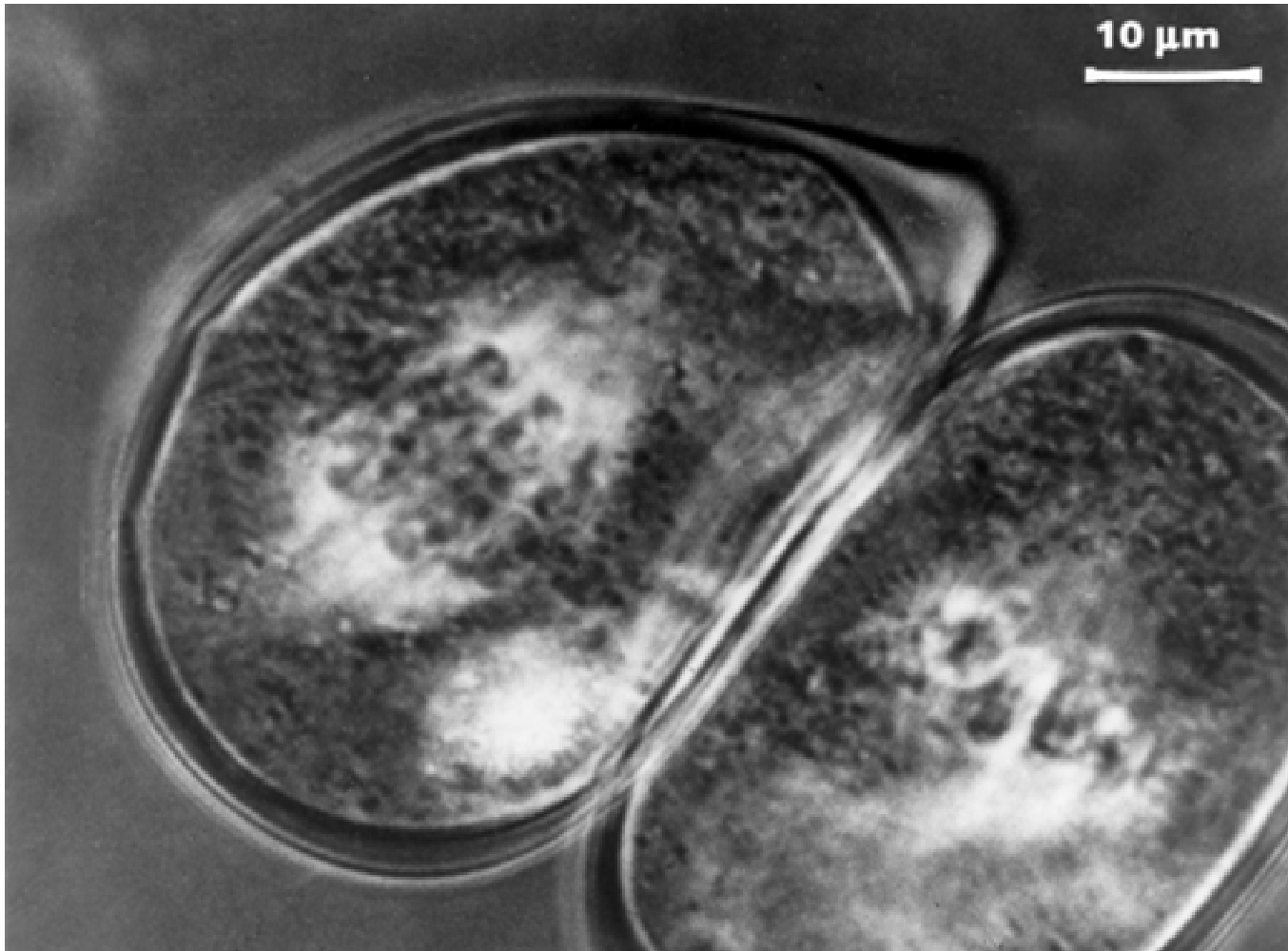


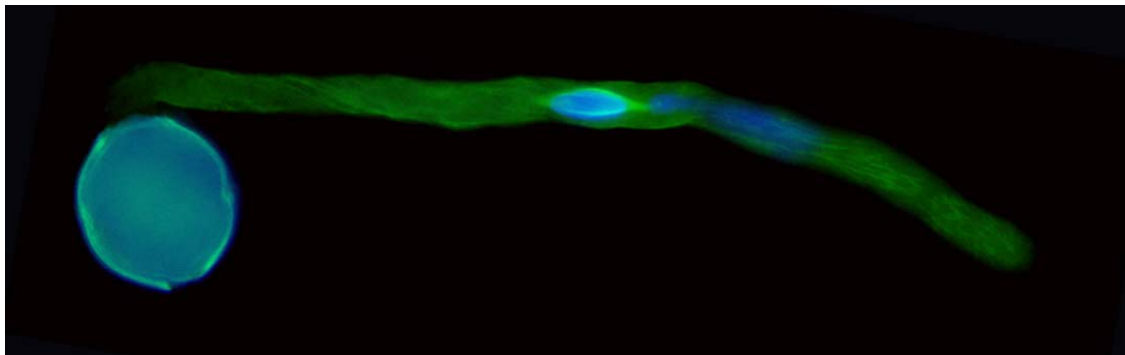
# Cytoskelet

# Co je vidět v buňkách?

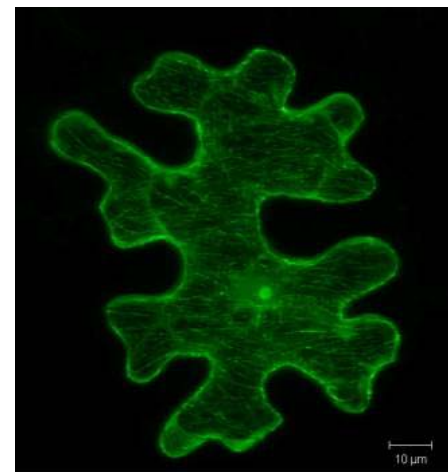
Inoue, 50. léta: „vláknité struktury“ (meióza mikrospor *Lilium longiflorum*)



# Aktin a tubulin

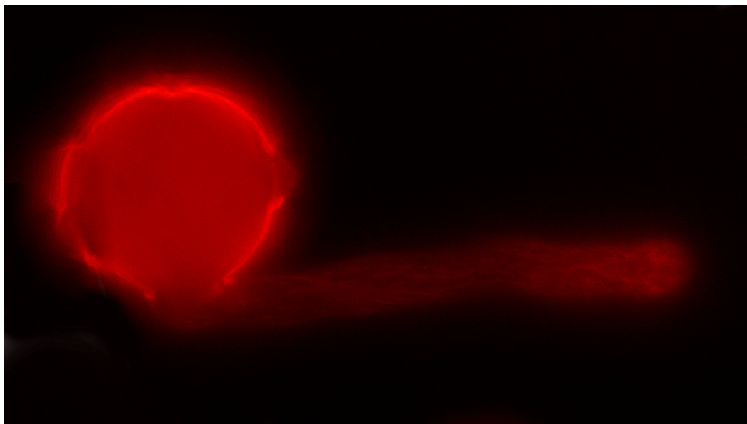


tubulin (a DNA) v pylové láčce tabáku

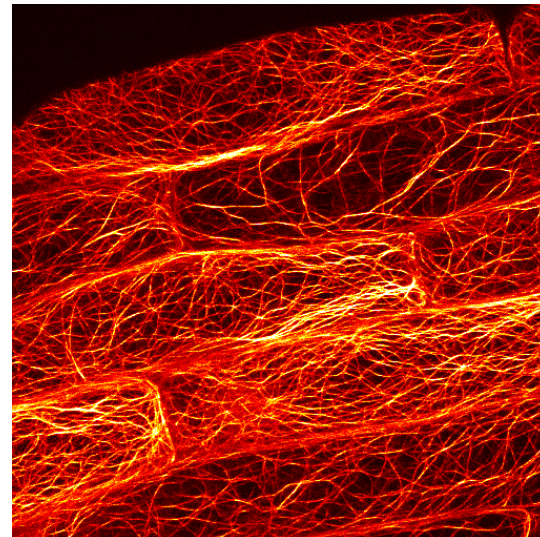


tubulin v epidermis listů tabáku

aktin v pylové láčce tabáku

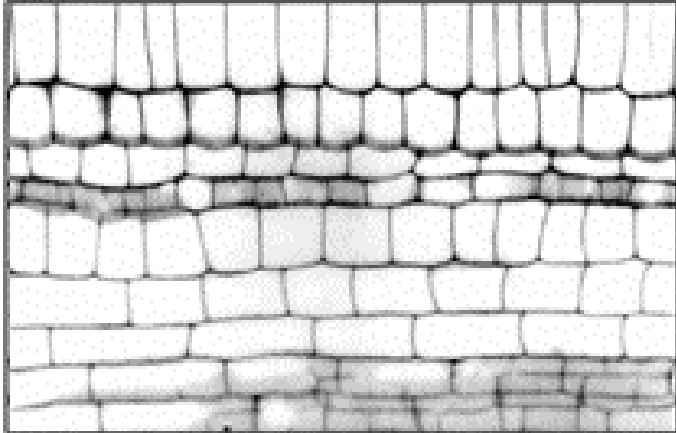


aktin v rhizodermis Arabidopsis



# Co je vidět v buňkách?

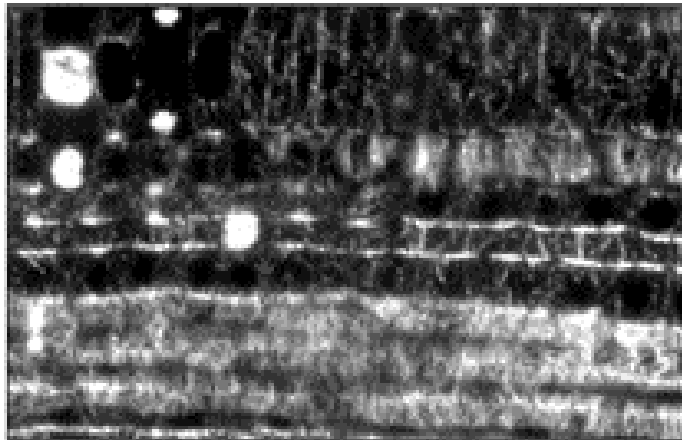
(A)



(B)



(C)



kořen Arabidopsis

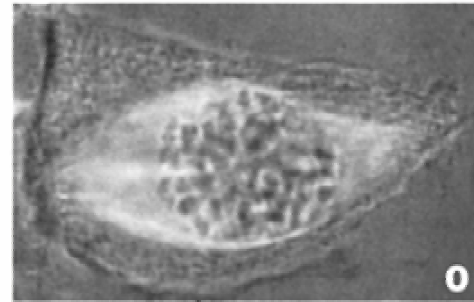
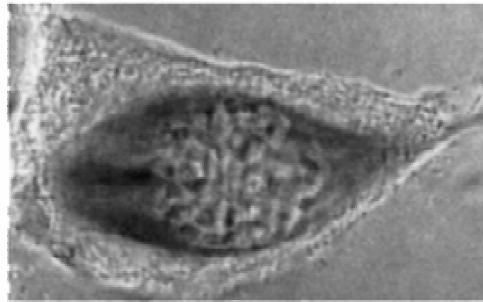
Ledbetter MC, Porter KR (1964):  
Morphology of Microtubules of Plant Cell.

(B)

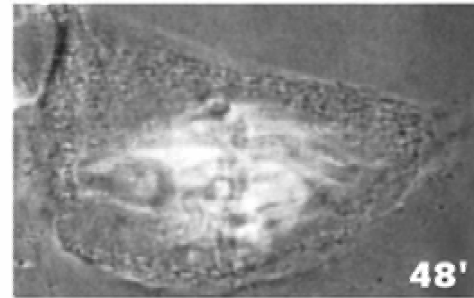
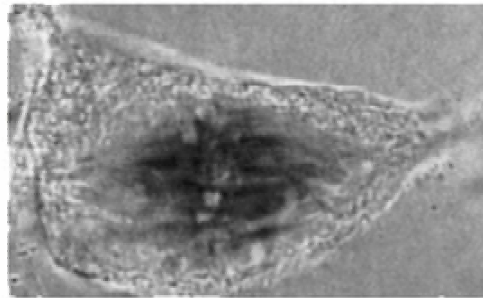
Subtractive contrast

Additive contrast

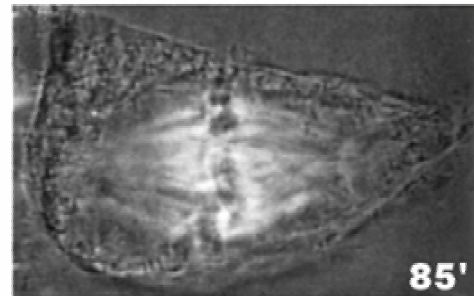
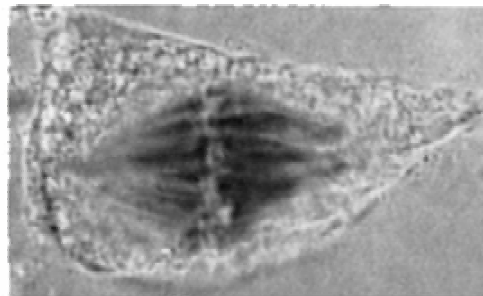
Prophase



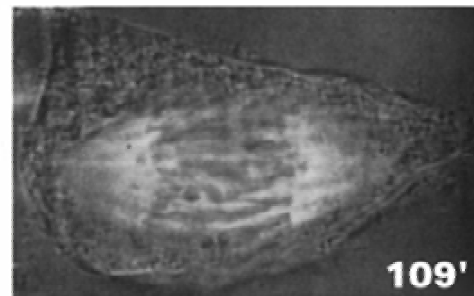
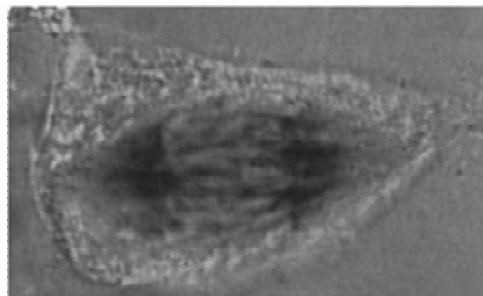
Prometaphase



Metaphase



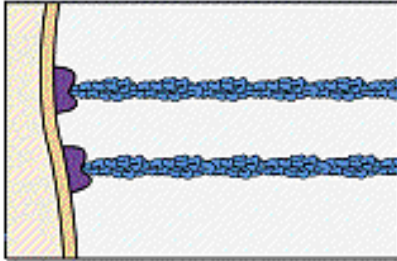
Anaphase



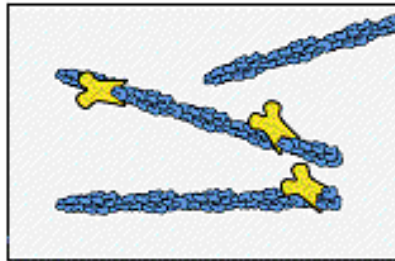
Endosperm,  
Haemanthus  
sp.

# Obecné rysy cytoskeletu

Nucleating protein



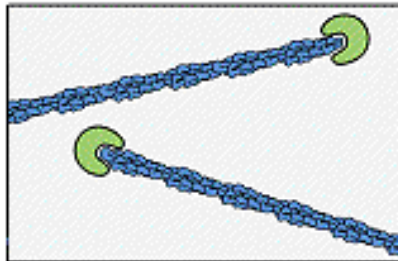
Severing protein



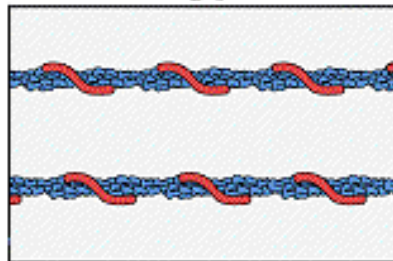
Cross-linking protein



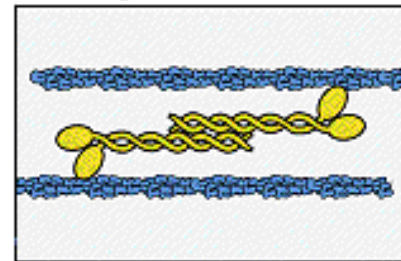
Capping (end-blocking) protein



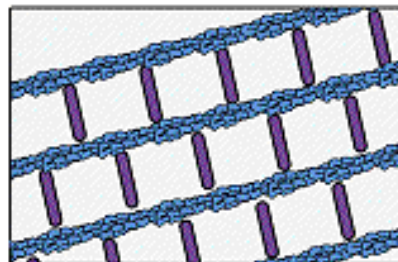
Side-binding protein



Motor protein



Bundling protein

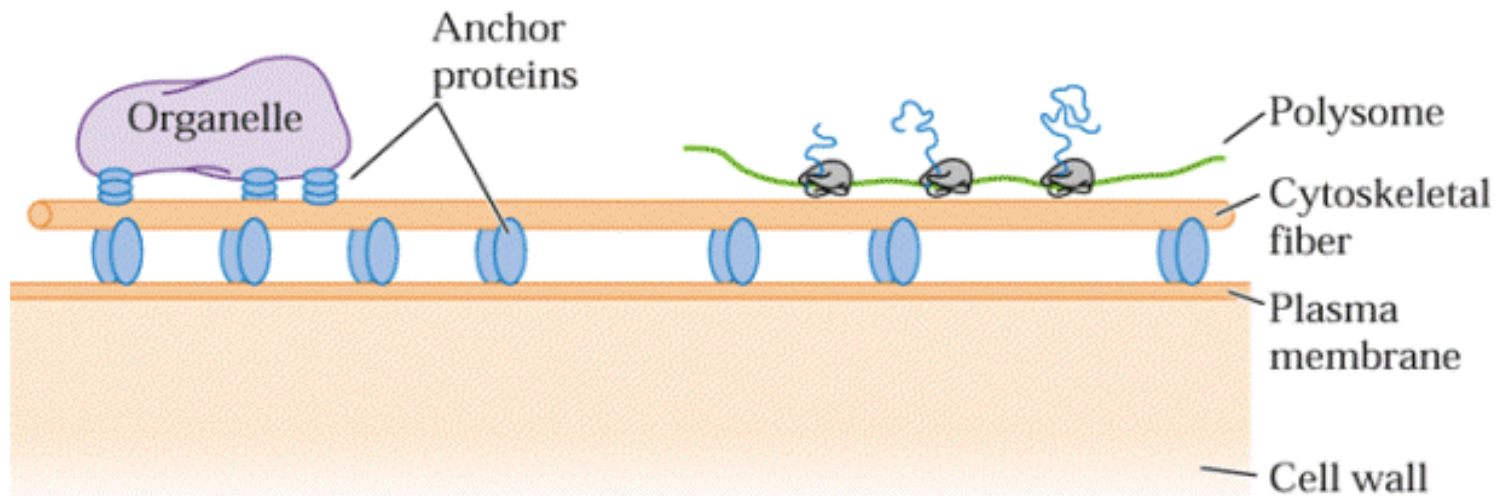


vlákná jsou polymery z několika málo typů podjednotek

diverzitu a nejrůznější funkce jim udělají asociované proteiny

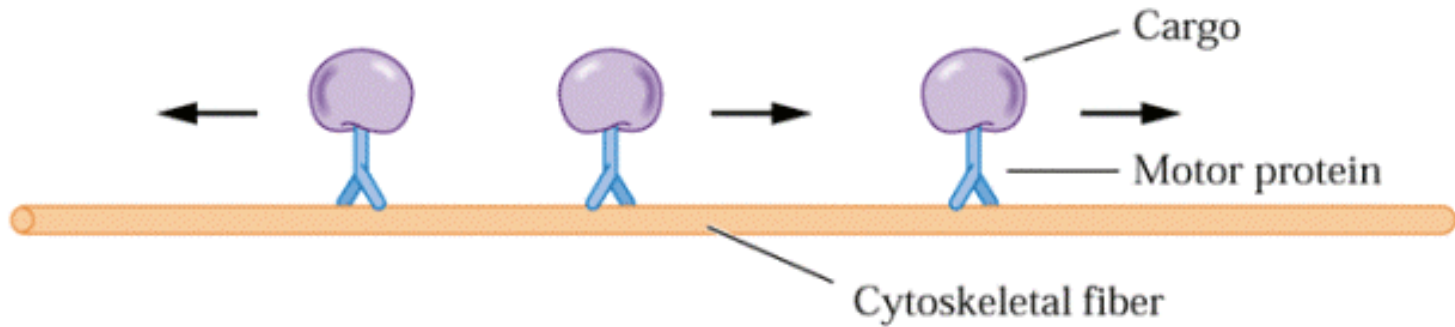
# Obecné funkce cytoskeletu

vazba organel a multimolekulárních komplexů

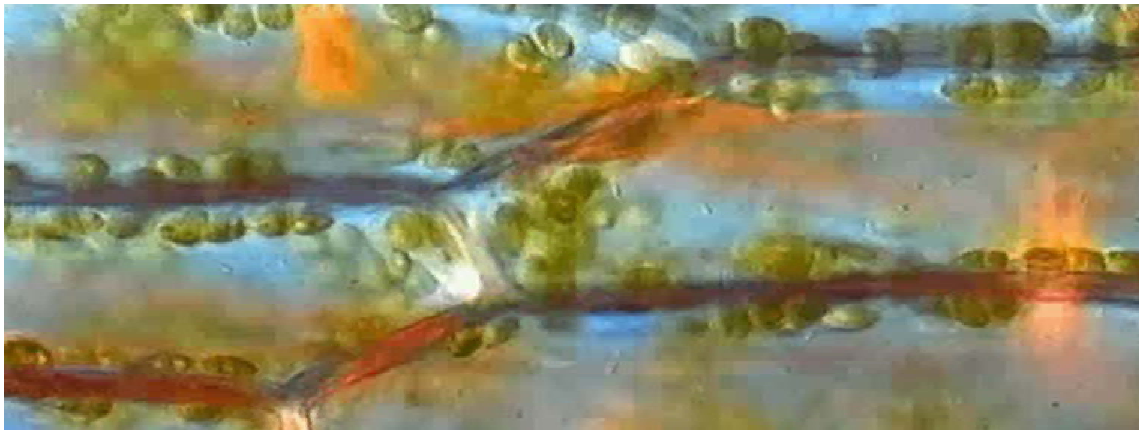


# Obecné funkce cytoskeletu

řízený vnitrobuněčný pohyb



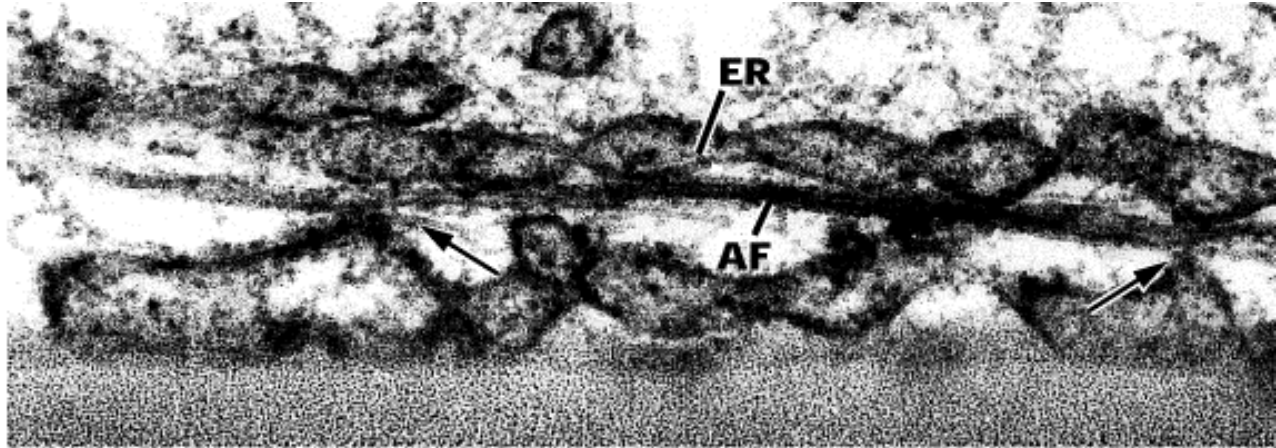
proudění cytoplazmy – *cytoplasmic streaming*



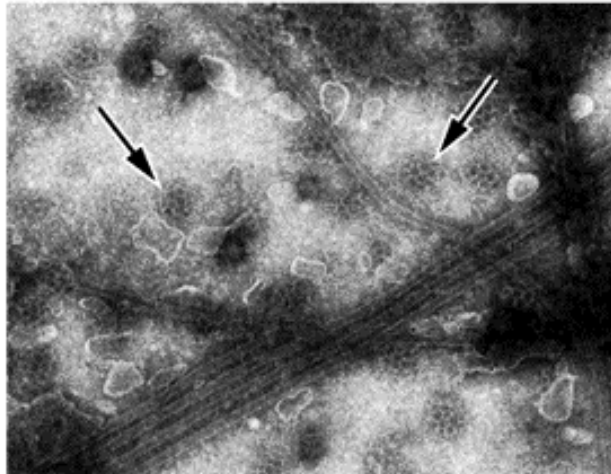


## řízený vnitrobuněčný pohyb

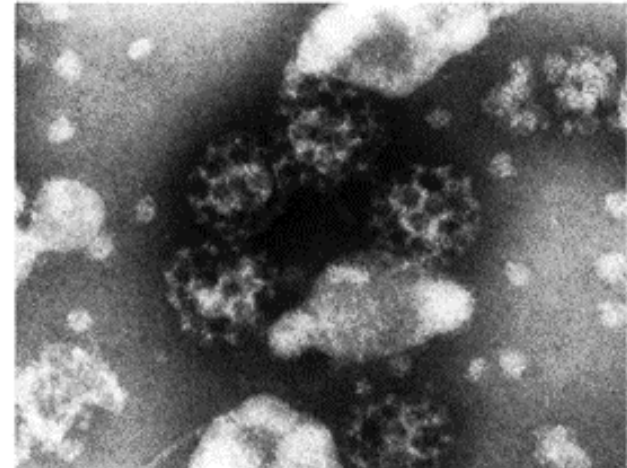
ER and  
clathrin-coated  
vesicles  
associated with  
AF and MT



(A)

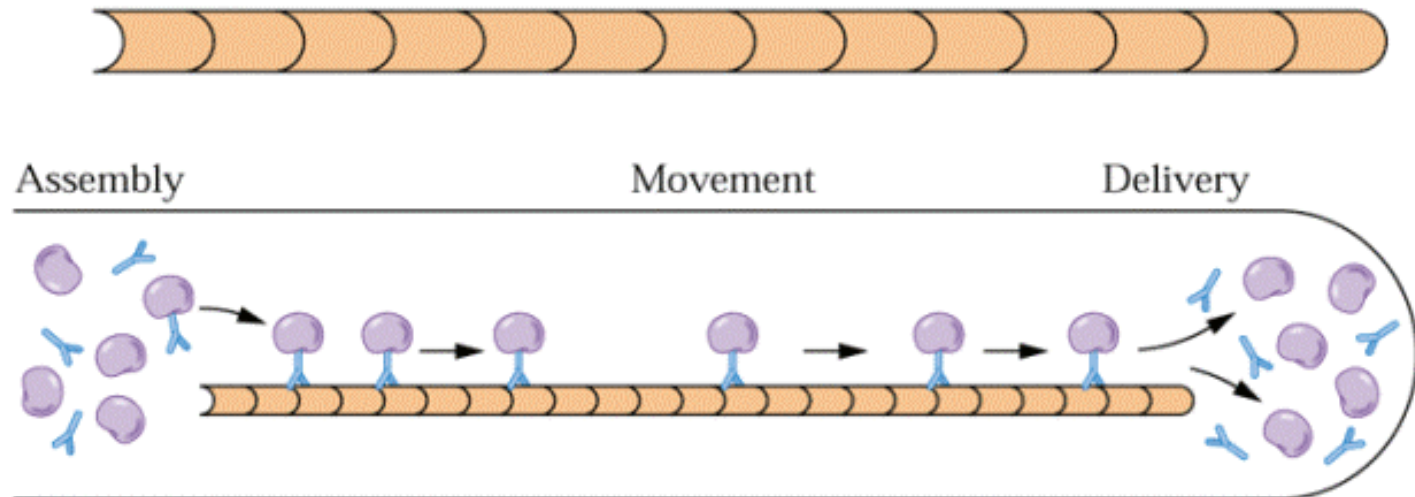


(B)



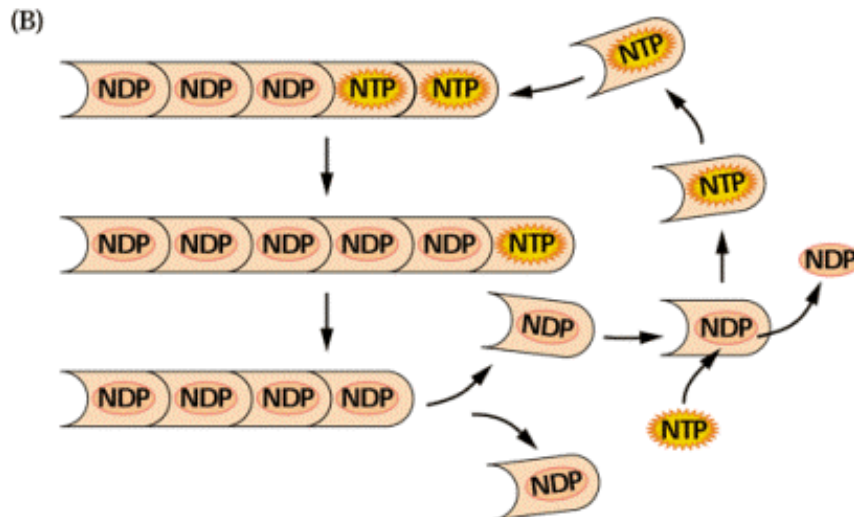
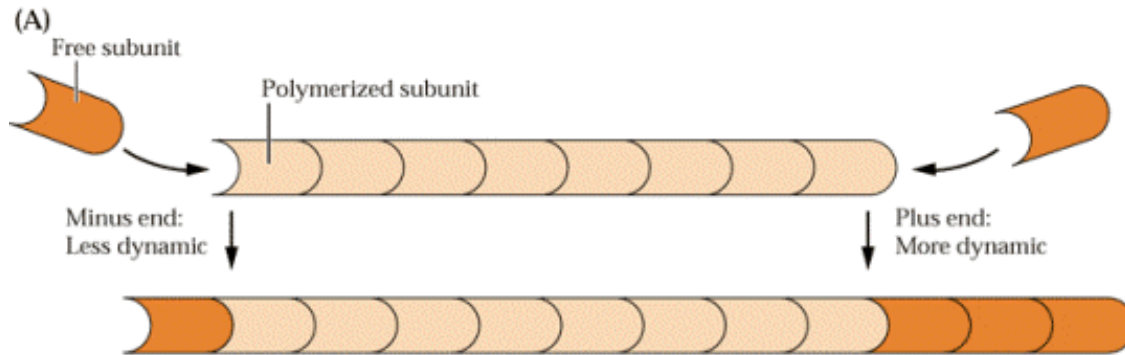
# Obecné funkce cytoskeletu

buněčná polarita

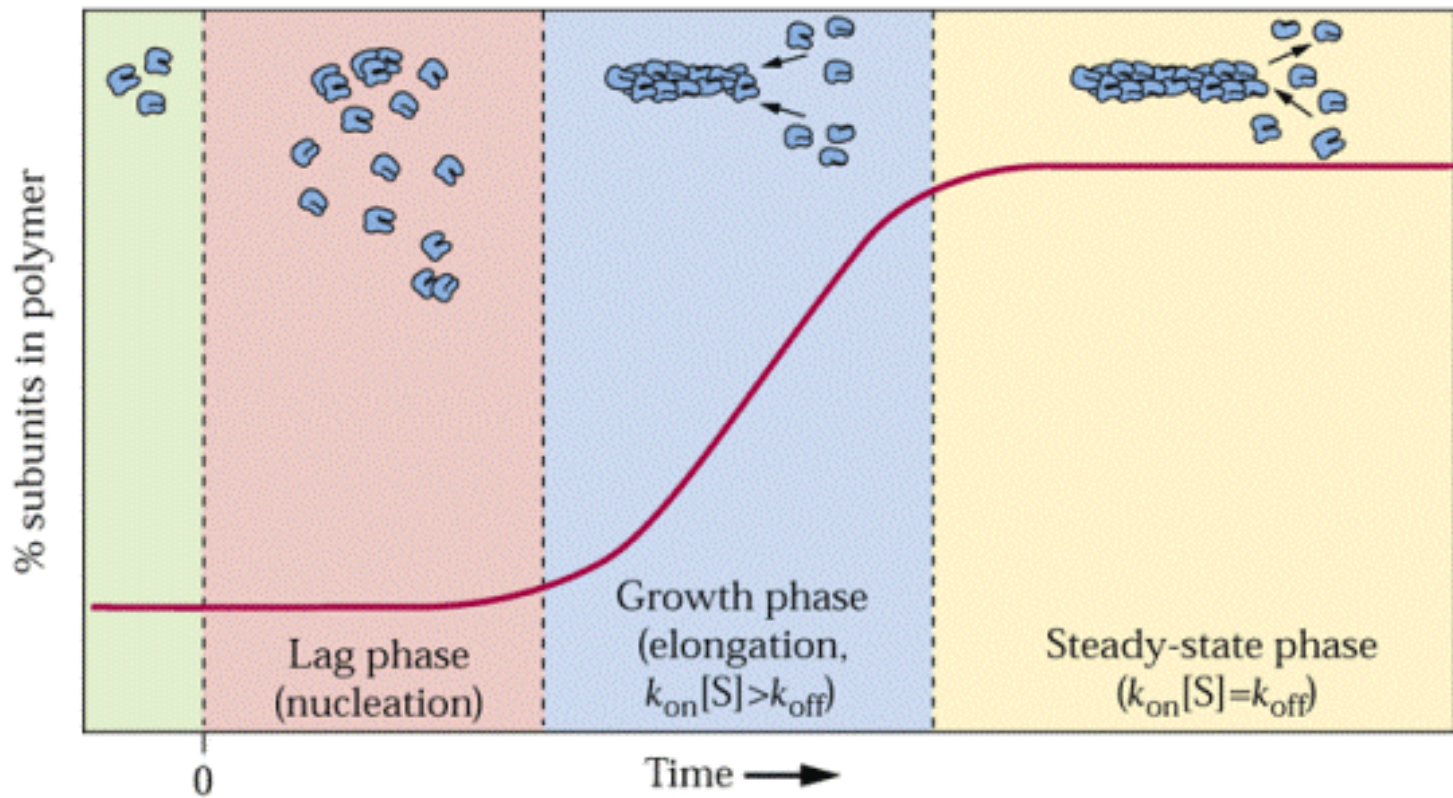


# Vznik polárního vlákna z monomerů

obecný princip u aktinu i tubulinu



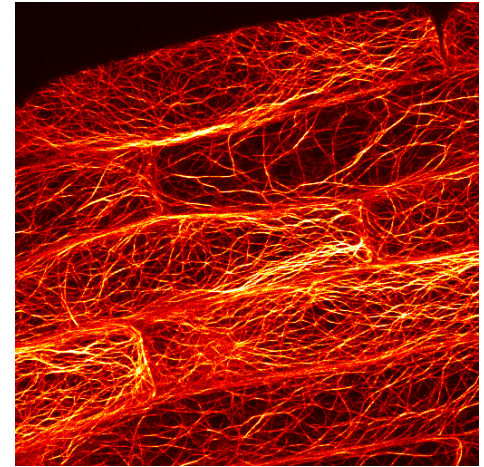
# Vznik polárního vlákna z monomerů



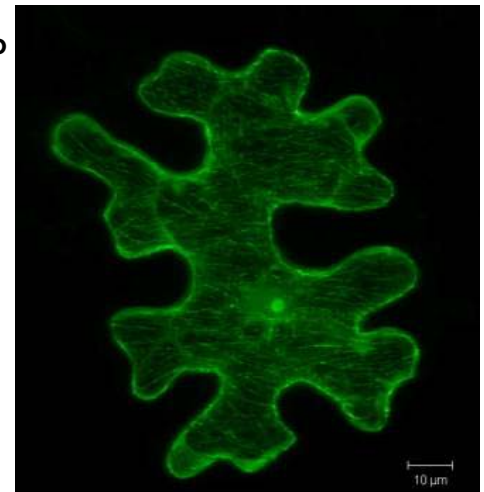
# Základní cytoskeletální systémy eukaryot

- Aktinový (mikrofilamenta)
  - všude (prokaryotní homolog MreB)
- Tubulinový (mikrotubuly)
  - všude (prokaryotní homolog FtsZ)
- Intermediální filamenta
  - „klasická“ zatím jen u živočichů
  - keratin, vimentin, neurofilamenta, laminy...

talin-RFP

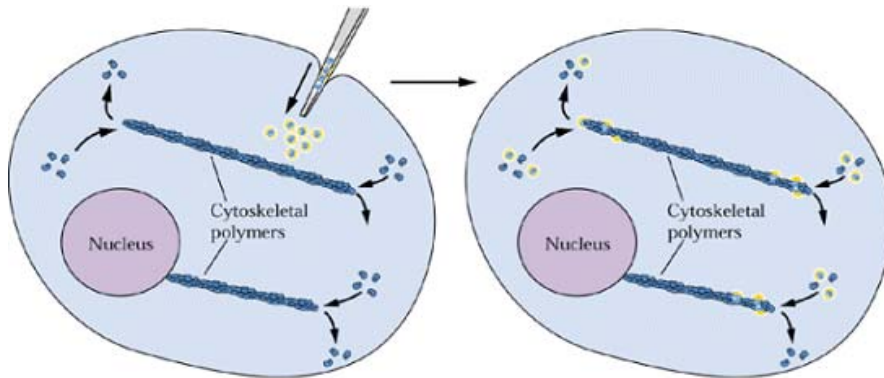


MAP4-GFP



# Metody studia cytoskeletu

mikroinjekce



nepřímá fluorescence

infiltrace agrobakterií do listů tabáku



exprese fúzních fluorescenčních proteinů

přímá fluorescence (faloidin)

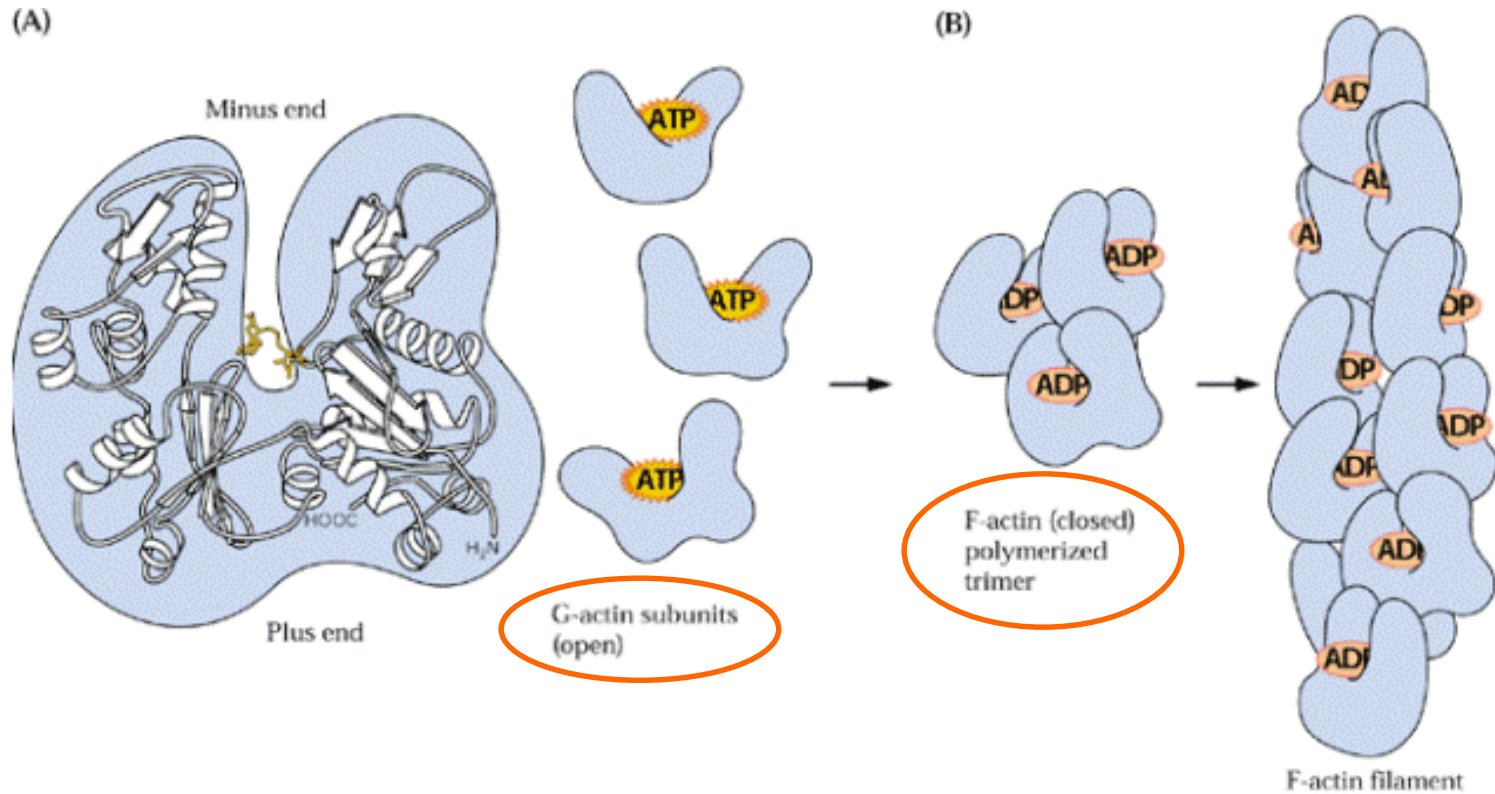
farmakologické působení (stabilizující x destabilizující látky)

tubulin-destabilizing	tubulin-stabilizing	actin-destabilizing	actin-stabilizing
oryzalin	taxol	latrunculin B	phalloidin
nocodazole		cytochalasin D	
propyzamid			

# Aktinový cytoskelet

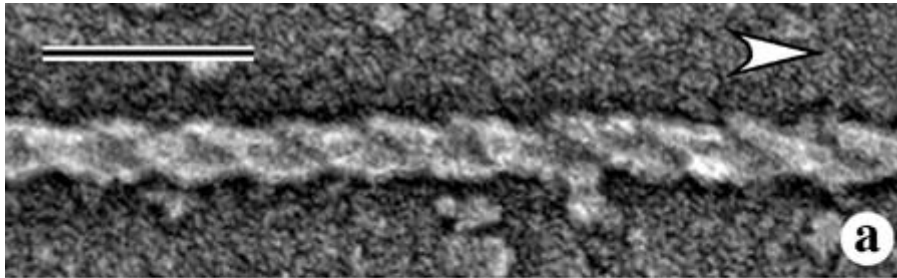
# Vznik polárního vlákna aktinu

aktin

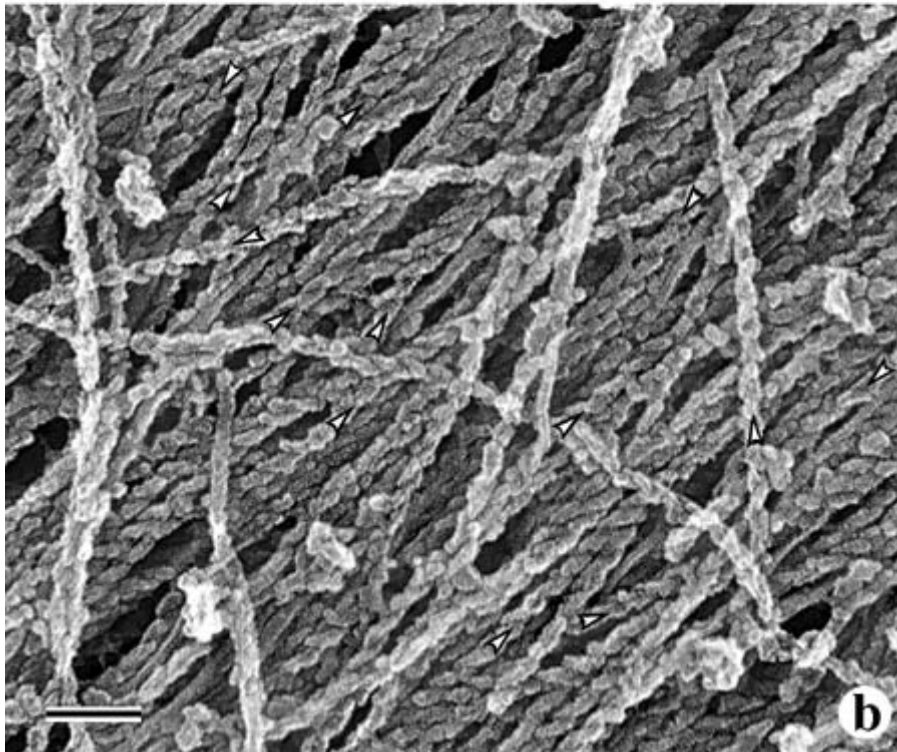




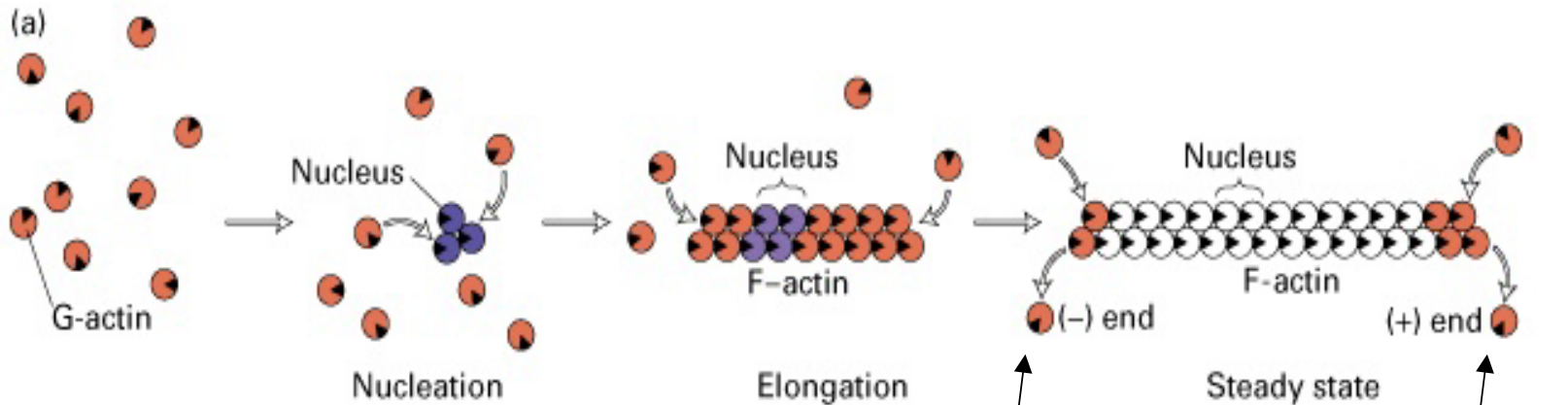
# Vznik polárního vlákna z monomerů



(a). S1 (myosin fragment)-decorated actin filament displays a helical rope-like appearance; the thicker part of a turn is directed to the pointed end of a filament (direction of the arrowhead). (b), part of an actin filament bundle from a REF-52 fibroblast after S1 decoration. Bars, 0.1  $\mu$ m

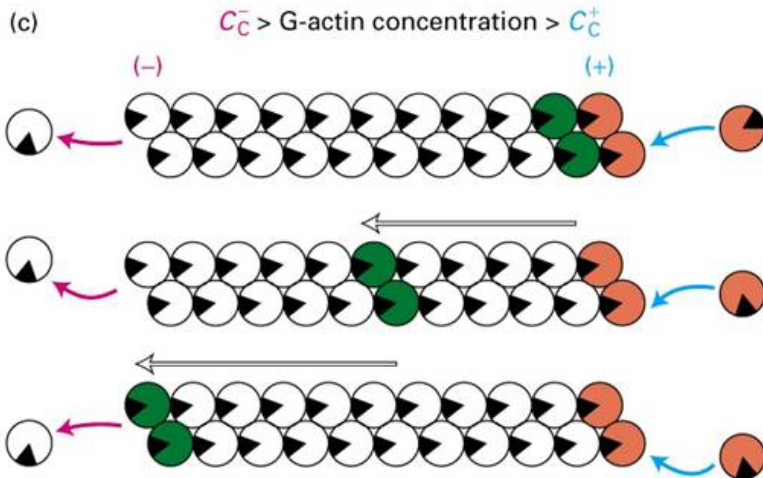


# Dynamika aktinových vláken: polymerace a treadmilling

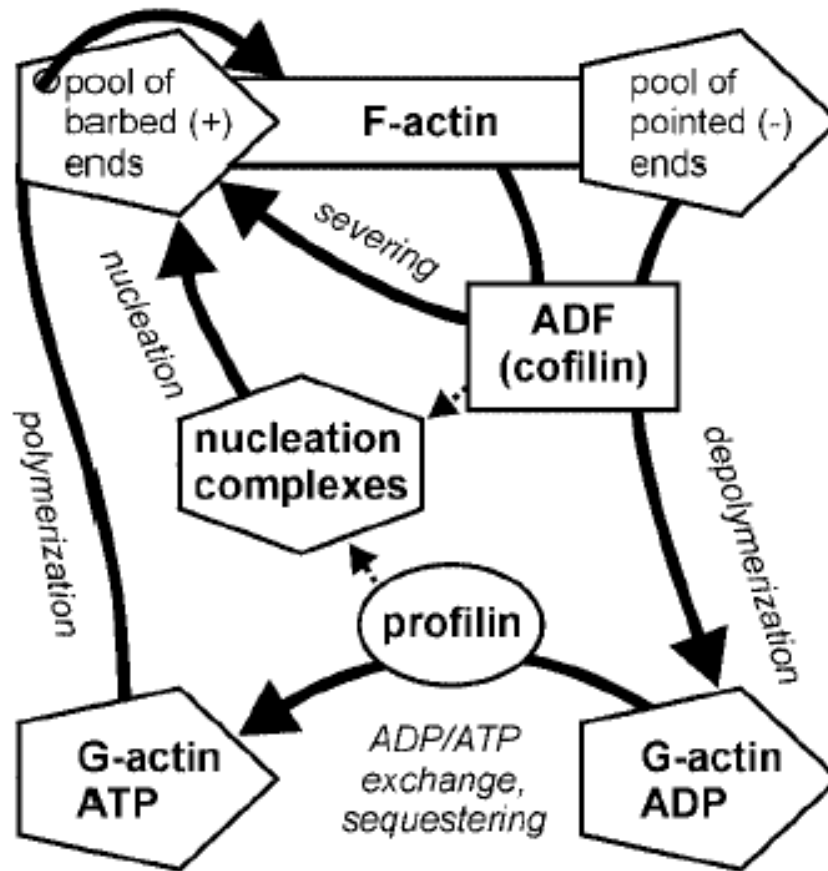


pointed end

barbed end

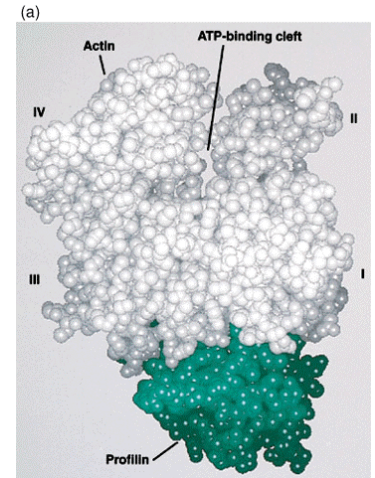
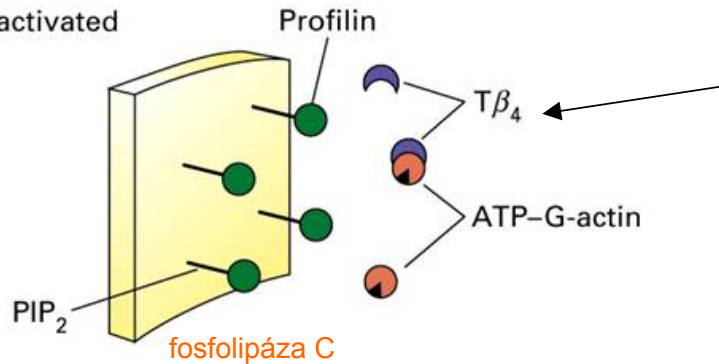


# Proteiny ovlivňující dynamiku aktinu

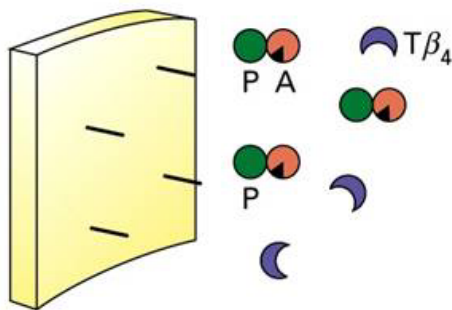


# Proteiny vázající G-aktin: regulace dostupnosti podjednotek

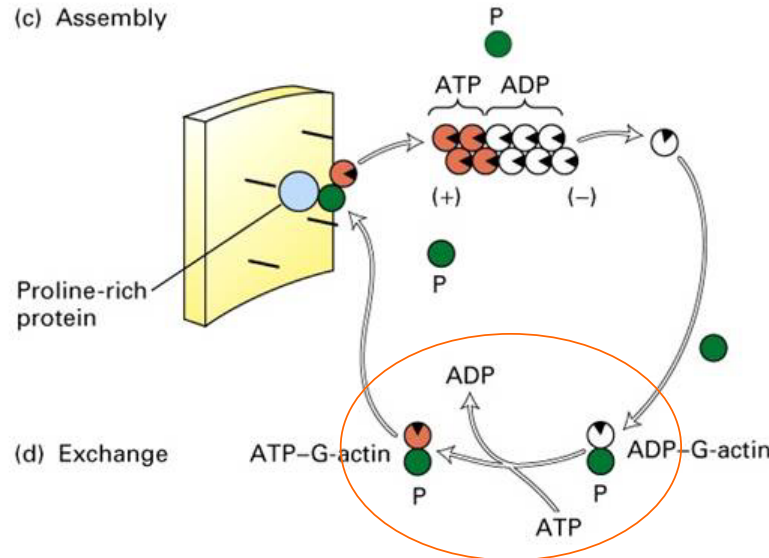
(a) Unactivated



(b) Activated



(c) Assembly



(d) Exchange

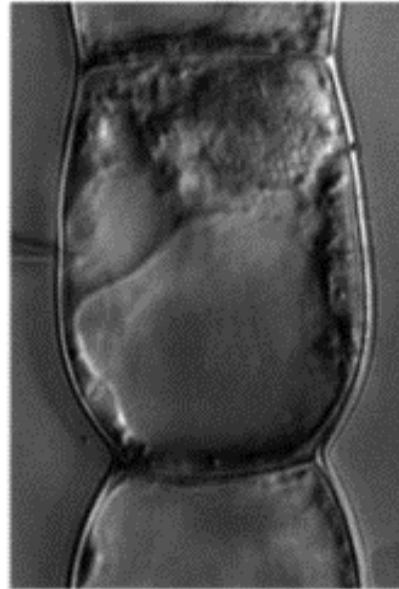
T $\beta_4$  a profilin vážou přednostně ATP-G-aktin – udržují vysokou pohotovostní hladinu aktinu pro rychlou polymeraci

## Profilin

(A)



Cytoplasmic streaming along transvacuolar strands is evident in an uninjected cell.



Ten minutes after profilin is microinjected, streaming has stopped and most of the transvacuolar strands have broken down.

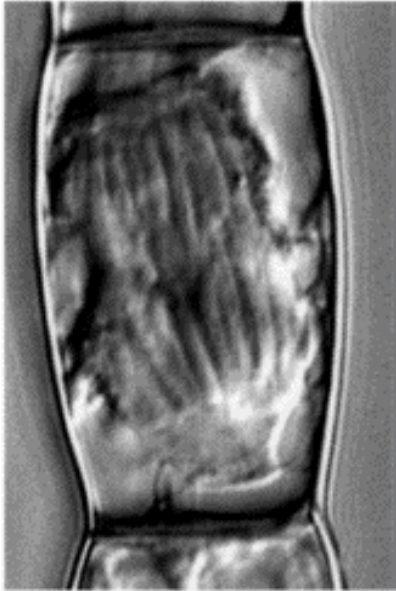


Streaming continues unabated 10 minutes after bovine serum albumin (BSA) is microinjected into a control cell.

Tradescantia stamen hairs

## Profilin

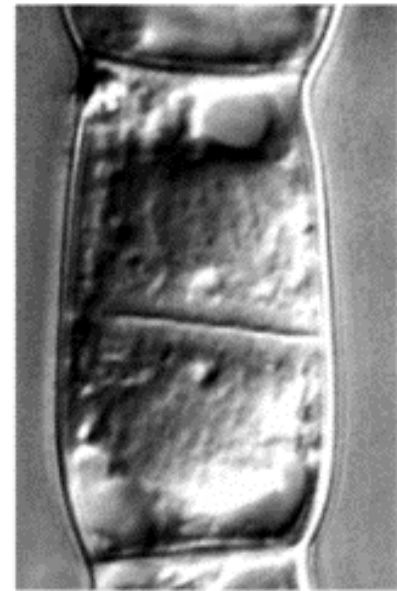
(B)



Anaphase cell undergoing chromosomal separation in an uninjected cell.

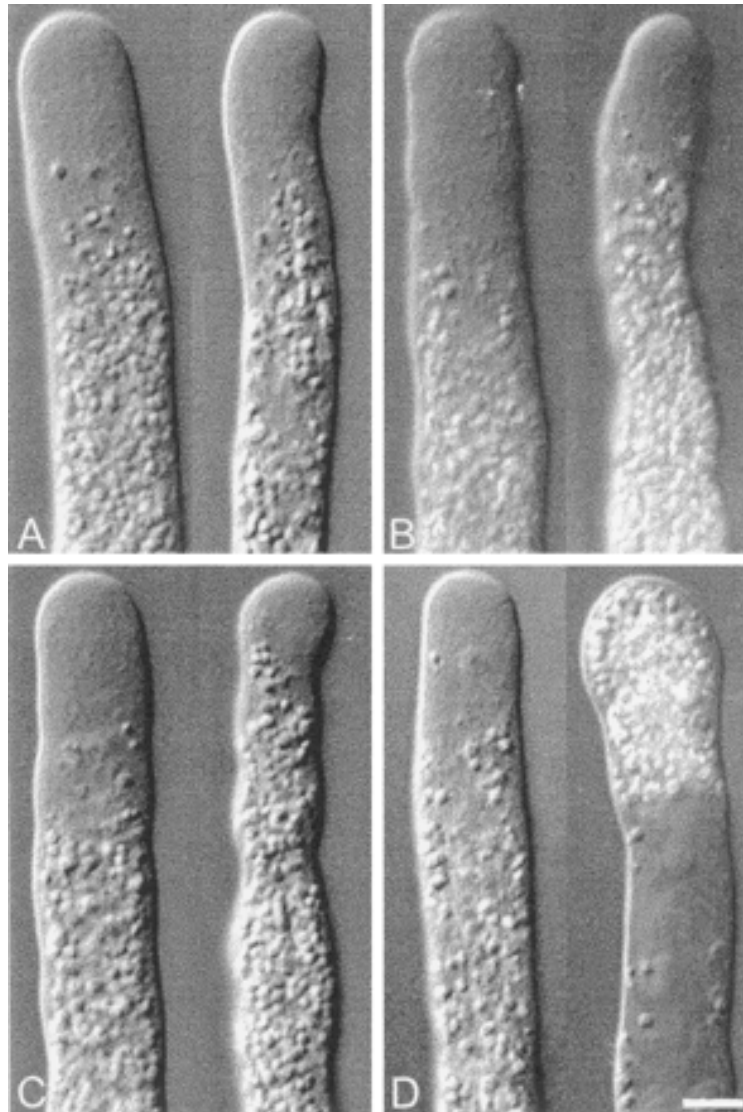


Fifty minutes after profilin is microinjected, chromosomal separation is complete, but a cell plate has not formed.

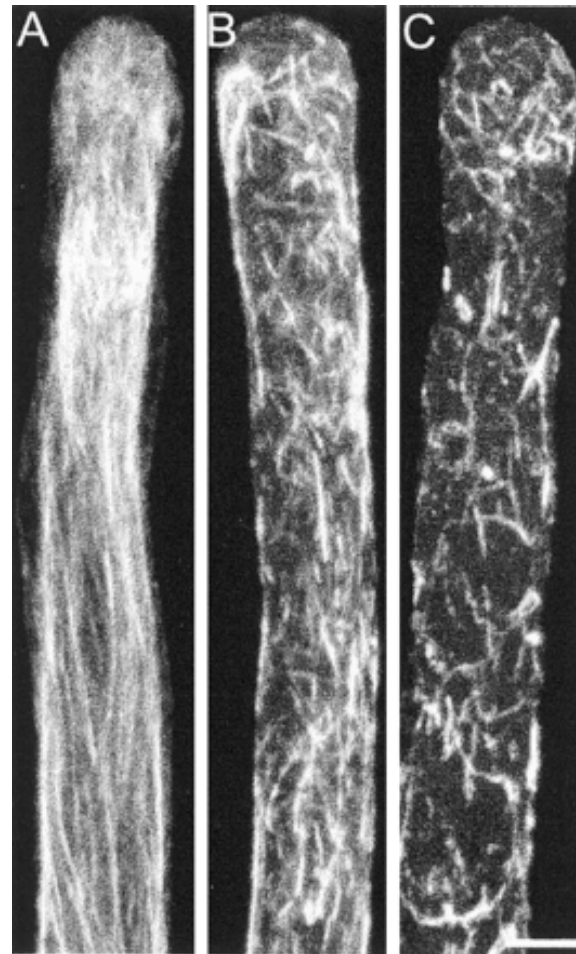


Twenty-five minutes after BSA is microinjected, a cell plate separates the control cell into two daughter cells.

## Injekce profilinu inhibuje růst pylových láček a narušuje strukturu aktinu



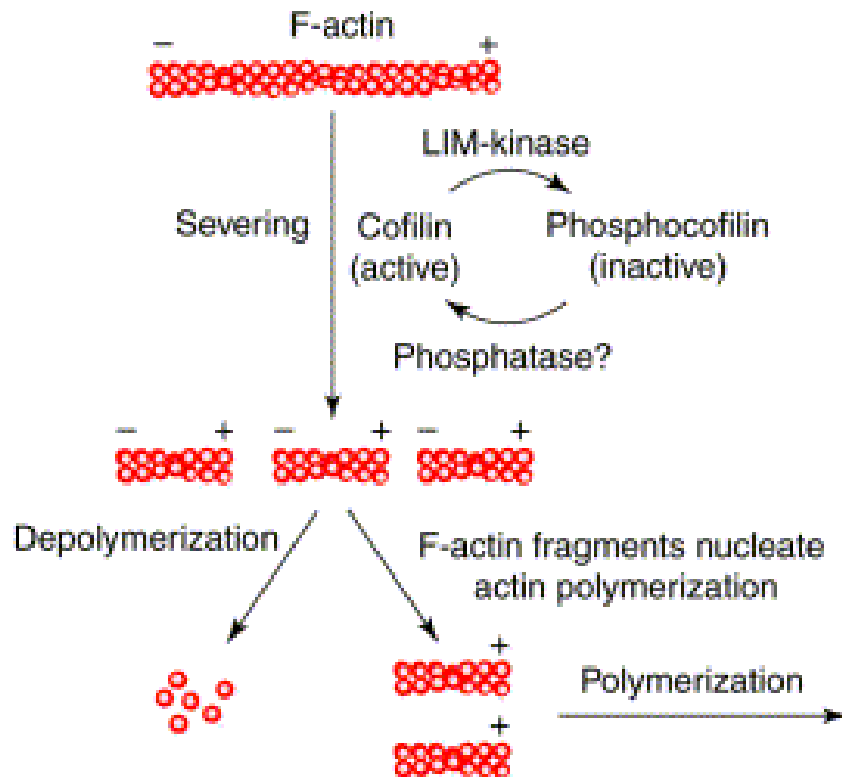
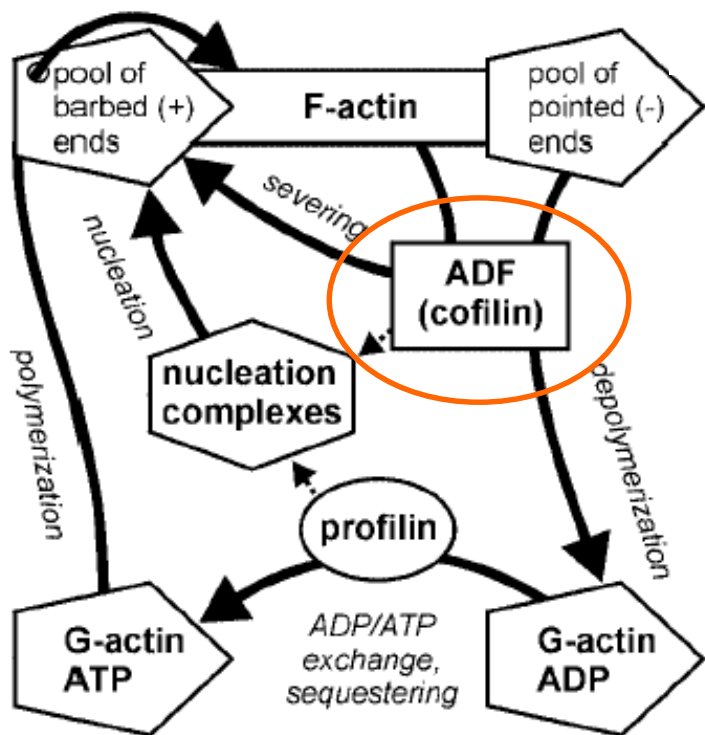
Pollen tubes injected with increasing doses of native pollen profilin. Leftmost cell - before injection, rightmost 20 min after injection. (A) 10  $\mu\text{M}$ , (B) 16  $\mu\text{M}$ , (C) 22  $\mu\text{M}$ , and (D) 62  $\mu\text{M}$ . Bar, 10  $\mu\text{m}$ .



F-actin distribution of profilin-injected cells after chemical fixation and Alexa-phalloidin staining. Twenty minutes after injection cells were chemically fixed and stained with 0.3  $\mu\text{M}$  Alexa-phalloidin. (A) Control cell. (B) Cell injected with a low concentration of profilin ( $\sim 30 \mu\text{M}$ ). (C) Cell injected with a higher concentration of profilin ( $>60 \mu\text{M}$ ).

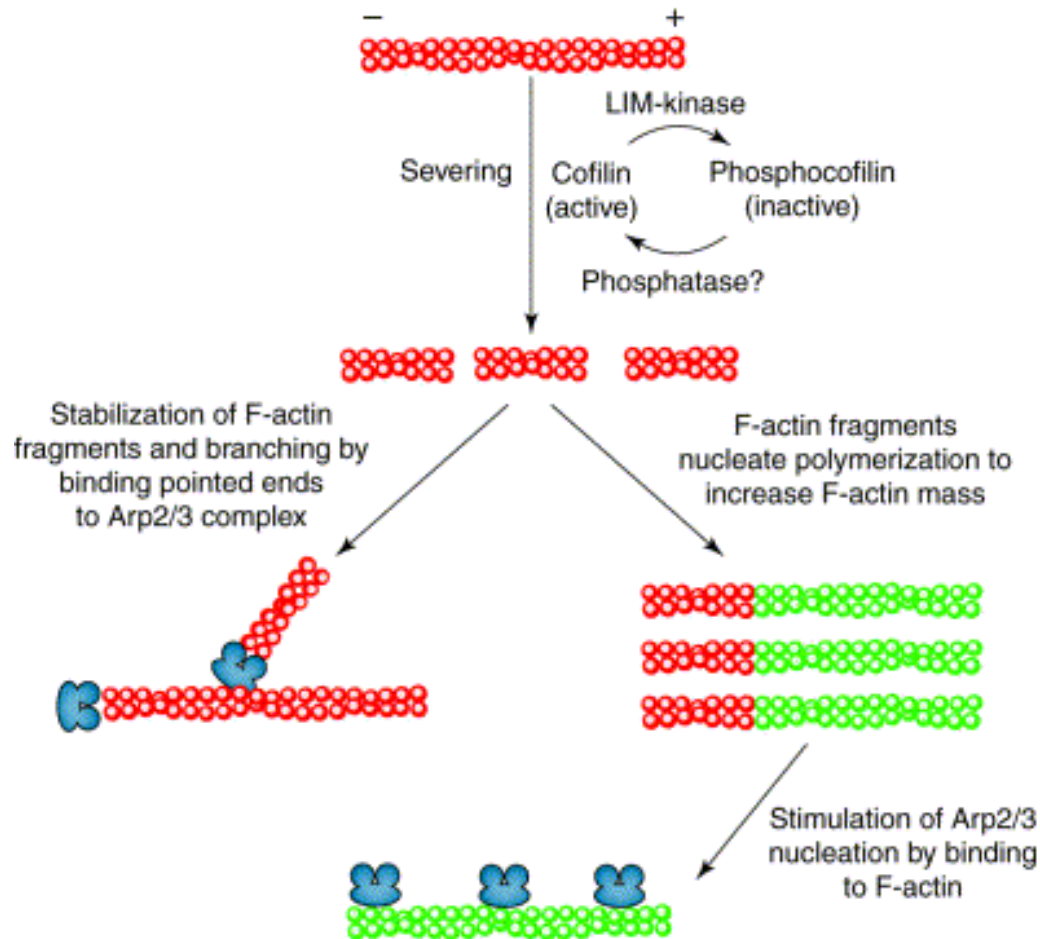
# „Střihání“ mikrofilament: ADF/cofilin

cofilin = Actin Depolymerizing Factor (ADF)

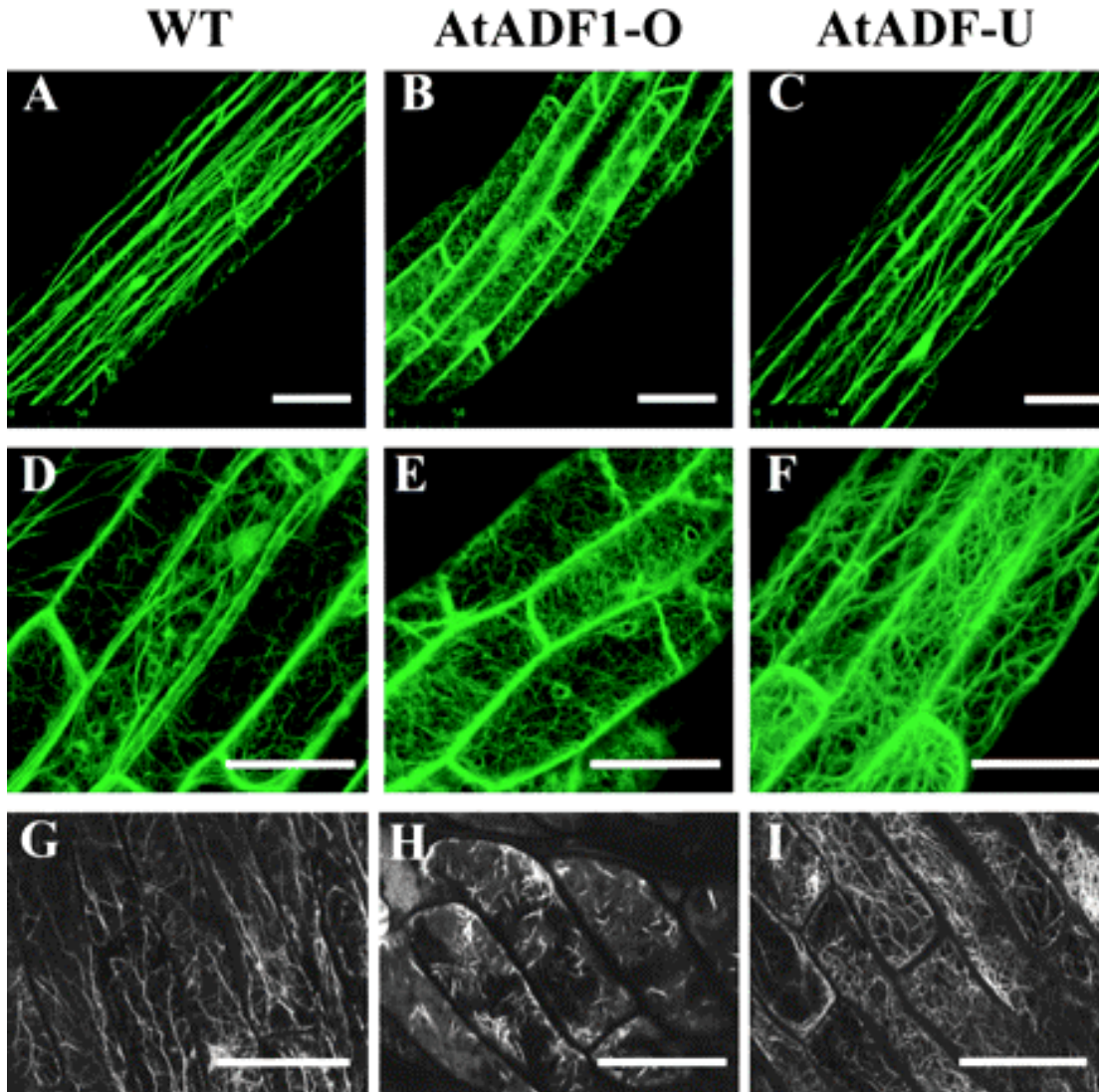




# „Stříhání“ mikrofilament: ADF/cofilin



## Změny hladiny ADF1 u Arabidopsis



overexpresse:

vymizení svazků  
mikrofilament (cables)

menší buňky

inhibice:

více svazků  
mikrofilament

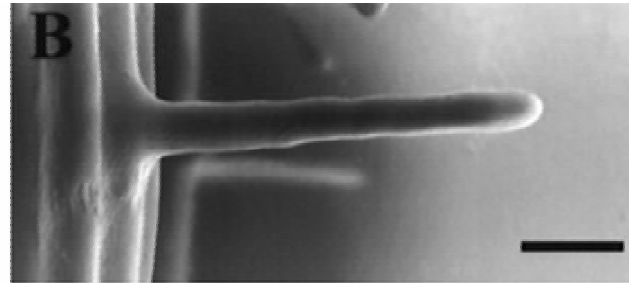
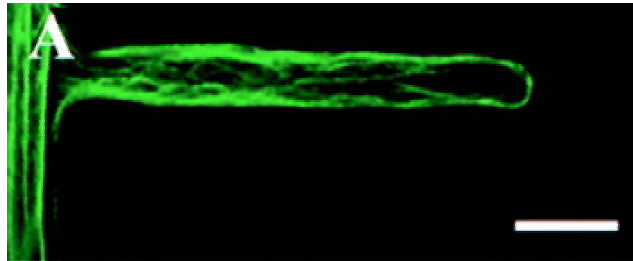
sense cDNA

anti-sense cDNA

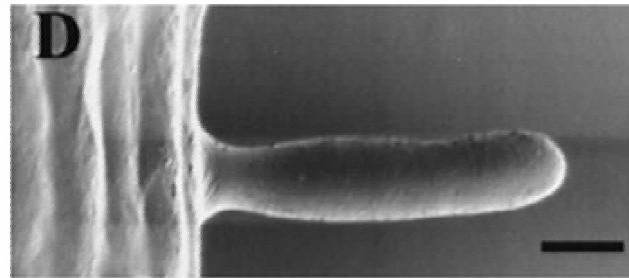
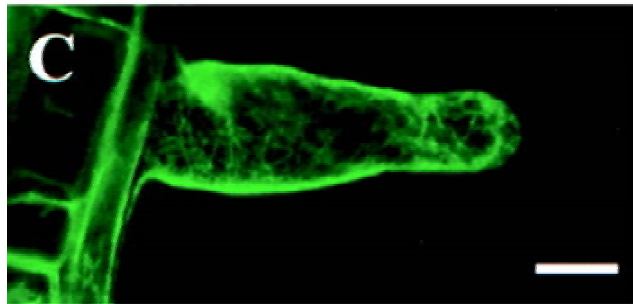
Dong et al. 2001

ADF1

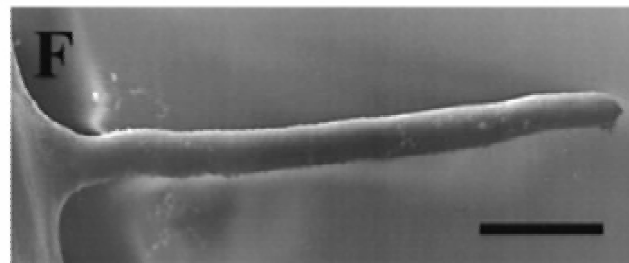
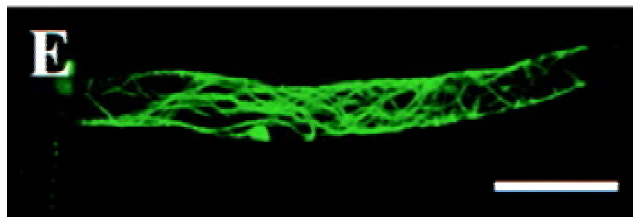
## Změny hladiny ADF1 u Arabidopsis



WT

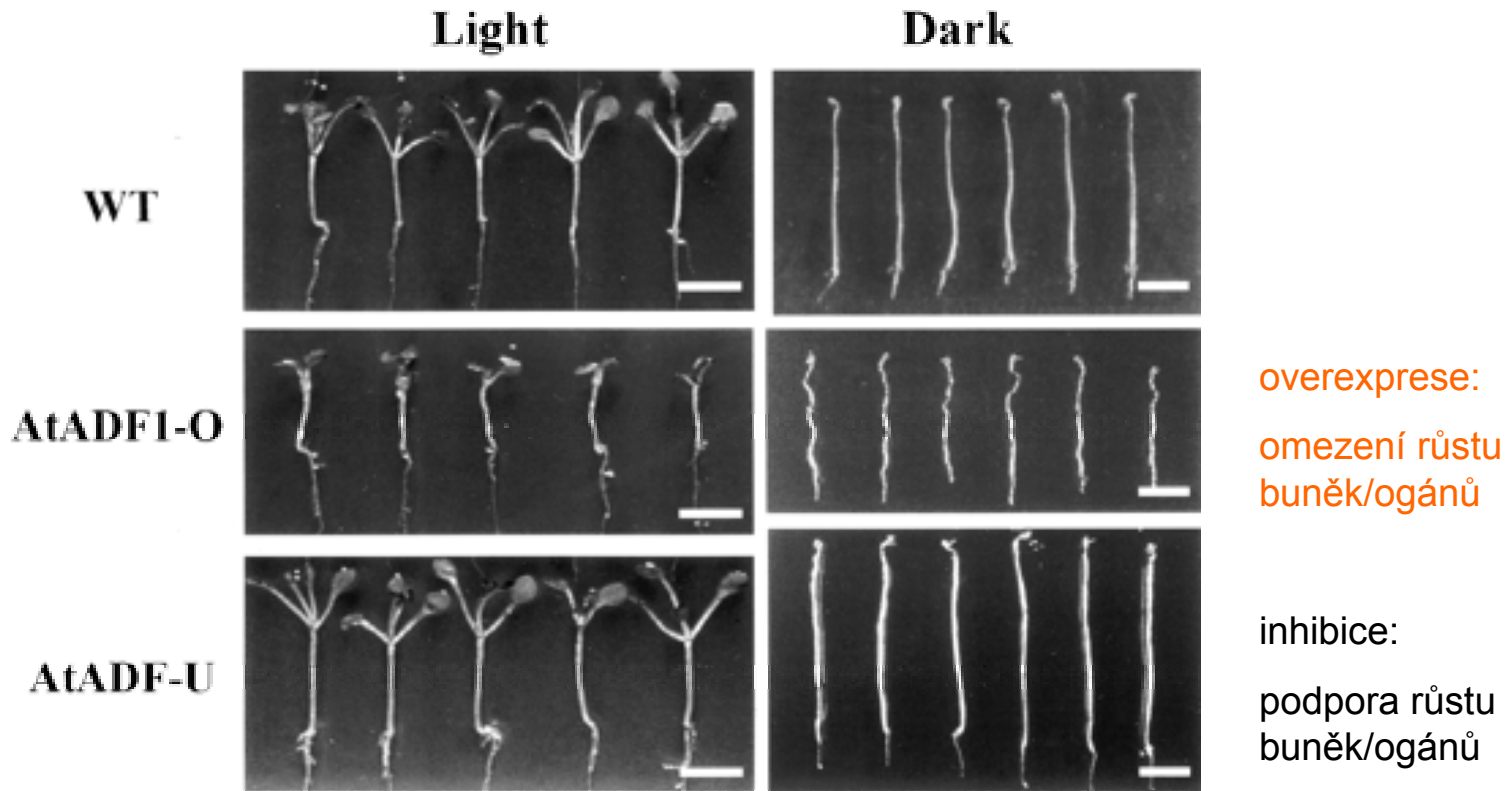


AtADF1-O



AtADF1-U

## Změny hladiny ADF1 u Arabidopsis

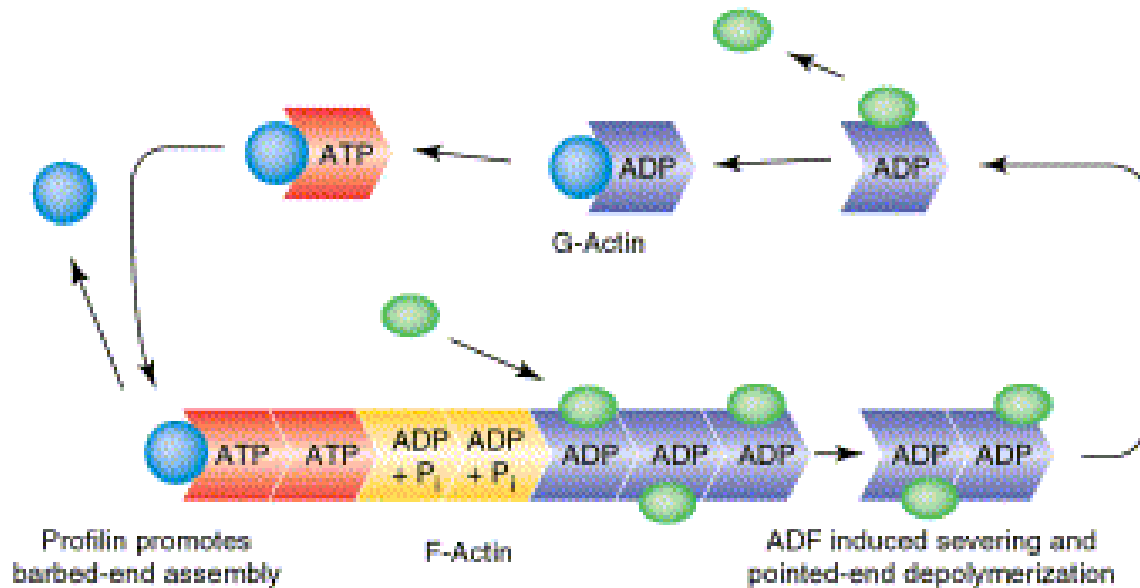


Počet listů v růžici před vykvetením (vliv na indukci kvetení):

WT:  $16,57 \pm 0,95$     **ADF1-O:  $16,46 \pm 1,10$**     AtADF1-U:  $25,61 \pm 1,90$

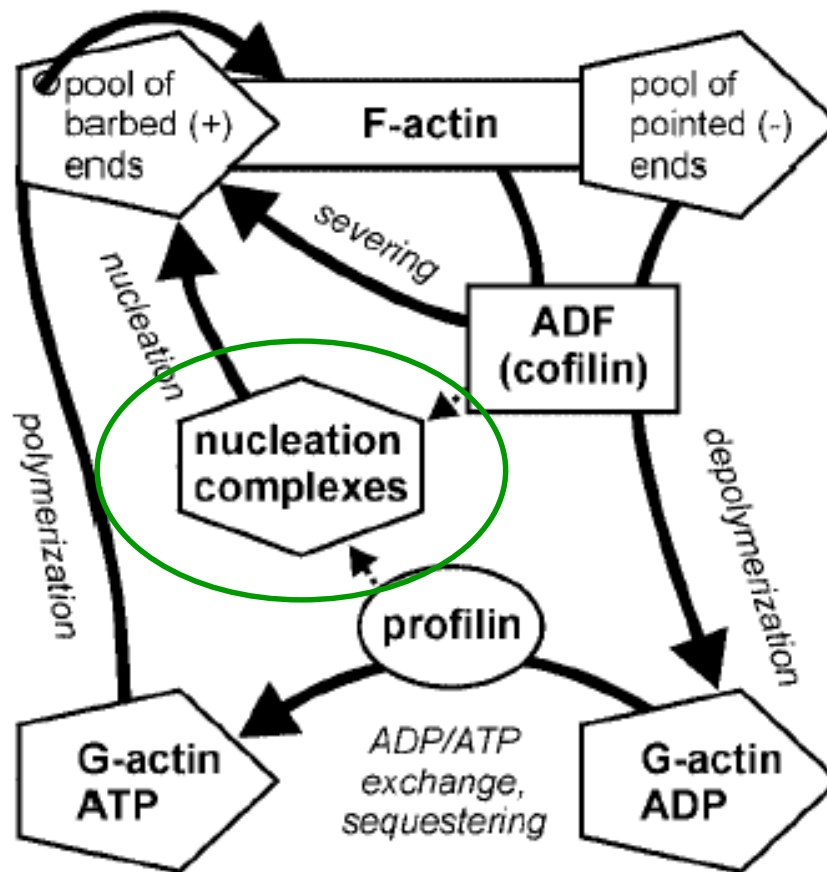
**ADF je jedním z klíčových regulátorů aktinového cytoskeletu!**

# Profilin versus Cofilin



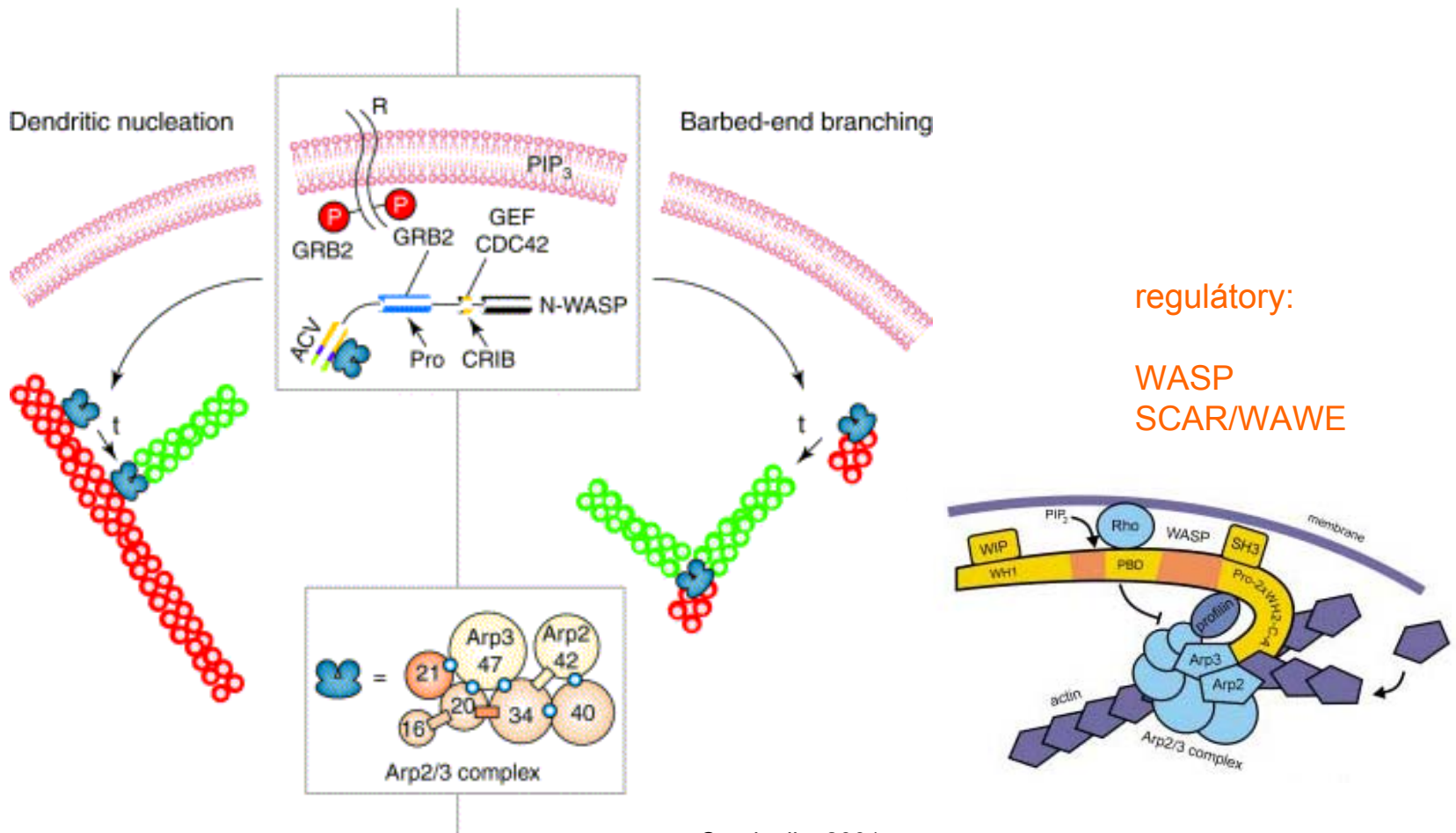
*TRENDS in Plant Science*

# Nukleace aktinových filament *de novo*

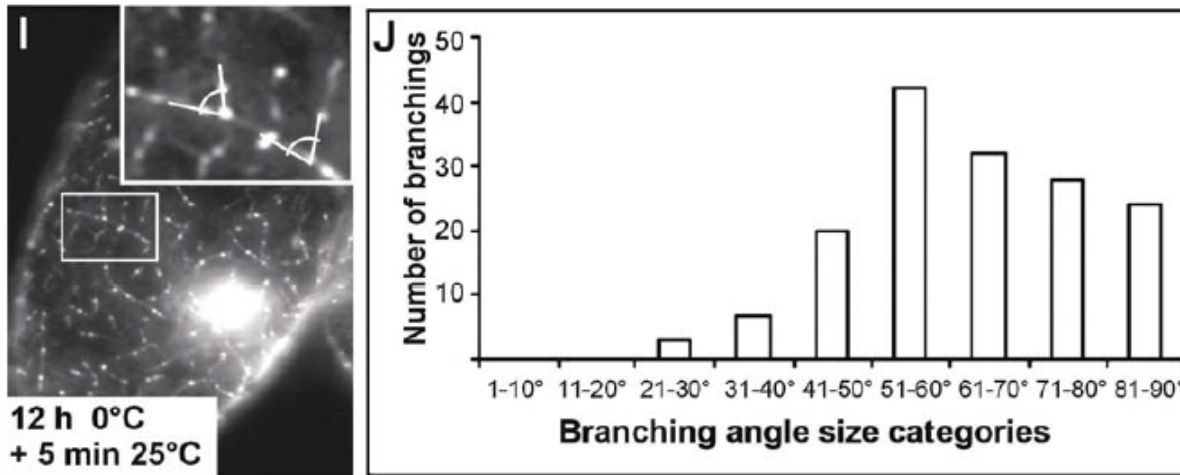


# Nukleace aktinových filament *de novo*

## Arp2/3 komplex



# Nukleace aktinových filament *de novo*



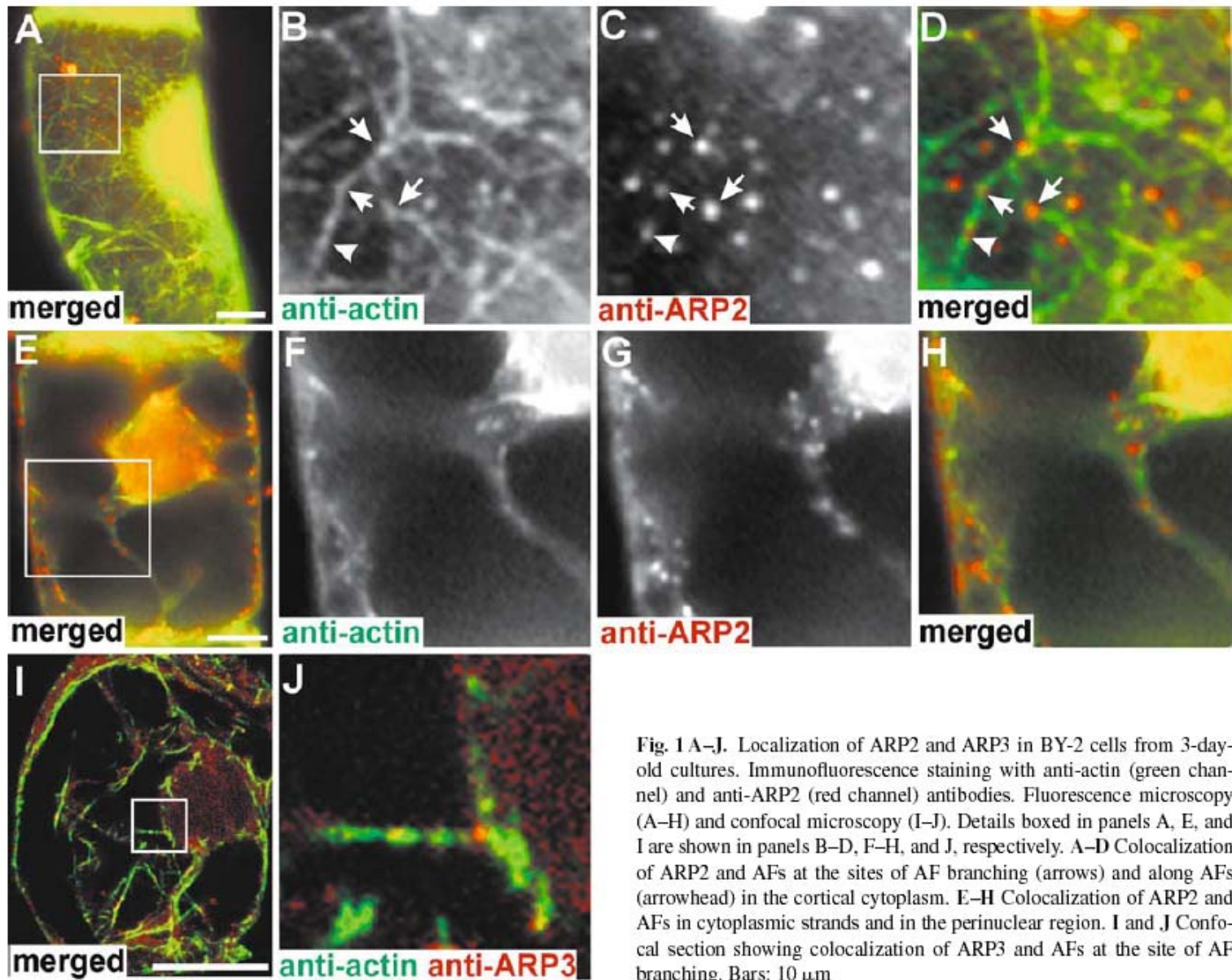
I and J Angles formed by two branches of newly formed AFs. I Branching of thin AFs with a dotted signal formed in the cortical cytoplasm. Rhodamine-phalloidin staining. J Frequency of branching angles of defined size categories. The angle of branching was measured for 156 cases in a total of 50 optical fields. Data are from one representative experiment. Bar: 10  $\mu\text{m}$

## ARP2 and ARP3 are localized to sites of actin filament nucleation in tobacco BY-2 cells

J. Fišerová<sup>1,\*</sup>, K. Schwarzerová<sup>1</sup>, J. Petrášek<sup>1,2</sup>, and Z. Opatrný<sup>1</sup>

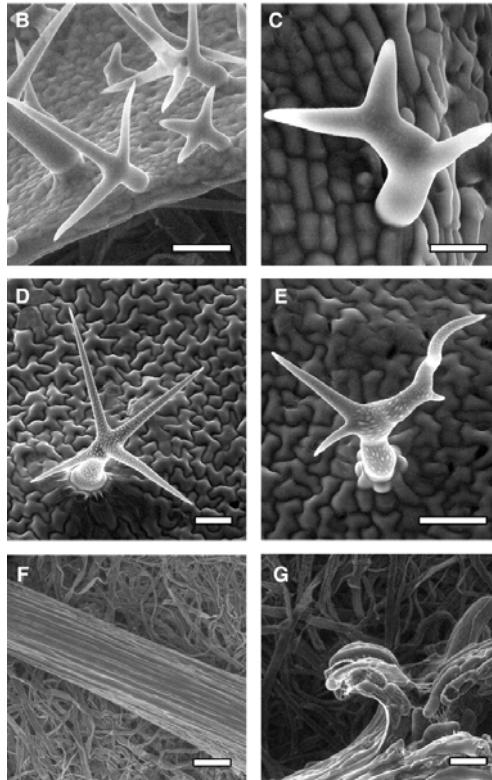
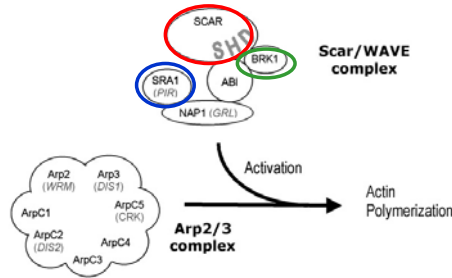
Protoplasma (2006) 227: 119–128  
DOI 10.1007/s00709-006-0146-6





**Fig. 1 A–J.** Localization of ARP2 and ARP3 in BY-2 cells from 3-day-old cultures. Immunofluorescence staining with anti-actin (green channel) and anti-ARP2 (red channel) antibodies. Fluorescence microscopy (A–H) and confocal microscopy (I–J). Details boxed in panels A, E, and I are shown in panels B–D, F–H, and J, respectively. A–D Colocalization of ARP2 and AFs at the sites of AF branching (arrows) and along AFs (arrowhead) in the cortical cytoplasm. E–H Colocalization of ARP2 and AFs in cytoplasmic strands and in the perinuclear region. I and J Confocal section showing colocalization of ARP3 and AFs at the site of AF branching. Bars: 10  $\mu$ m

# SCAR/WAVE: regulatory Arp2/3



WT

*itb1/scar2*

The evolutionarily conserved **Rac-WAVE-Arp2/3 pathway** links **actin filament nucleation** to **cell morphogenesis**. WAVE translates Rac-GTP signals into Arp2/3 activation by regulating the stability and/or localization of the **activator subunit Scar/WAVE**. The WAVE complex includes:

- 1) Sra1/PIR121/CYFIP1,
- 2) Nap1/NAP125,
- 3) Abi-1/Abi-2,
- 4) Brick1(Brk1)/HSPC300,
- 5) Scar/WAVE

**Arabidopsis BRICK1/HSPC300 Is an Essential WAVE-Complex Subunit that Selectively Stabilizes the Arp2/3 Activator SCAR2.**

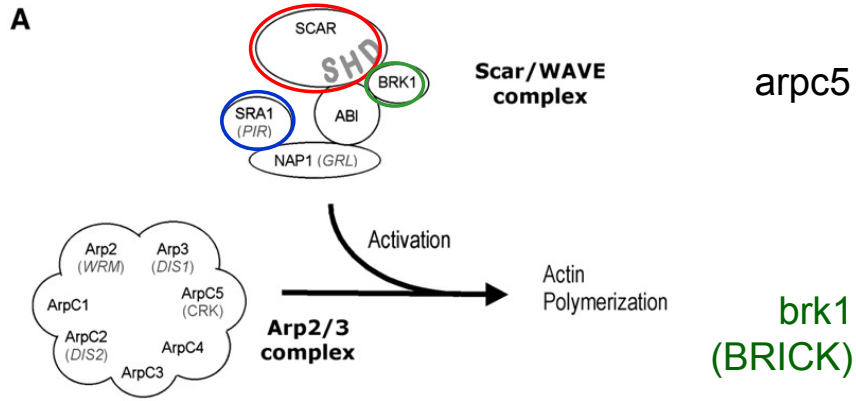
Current Biology, Volume 16, Issue 9, Pages 895-901

J. Le, E. Mallery, C. Zhang, S. Brankle, D. Szymanski

# BRICK1/HSPC300 functions with SCAR and the ARP2/3 complex to regulate epidermal cell shape in *Arabidopsis*

Stevan Djakovic\*, Julia Dyachok, Michael Burke†, Mary J. Frank‡ and Laurie G. Smith§

WAVE-complex stabilizes the Arp2/3 activator SCAR2.

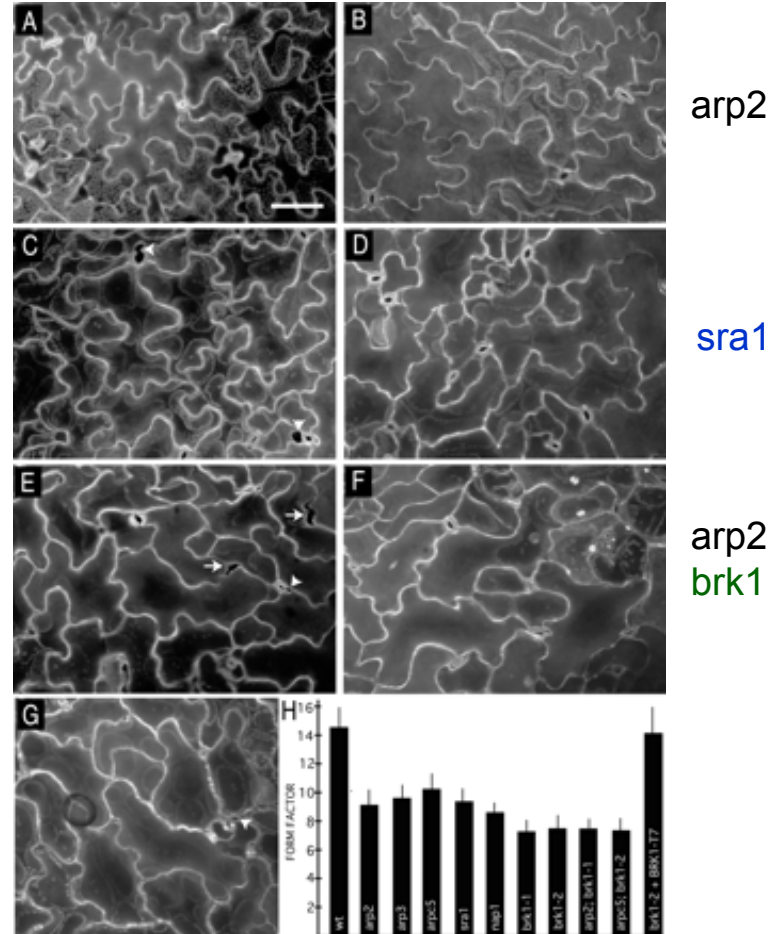


WT

arpc5

brk1 (BRICK)

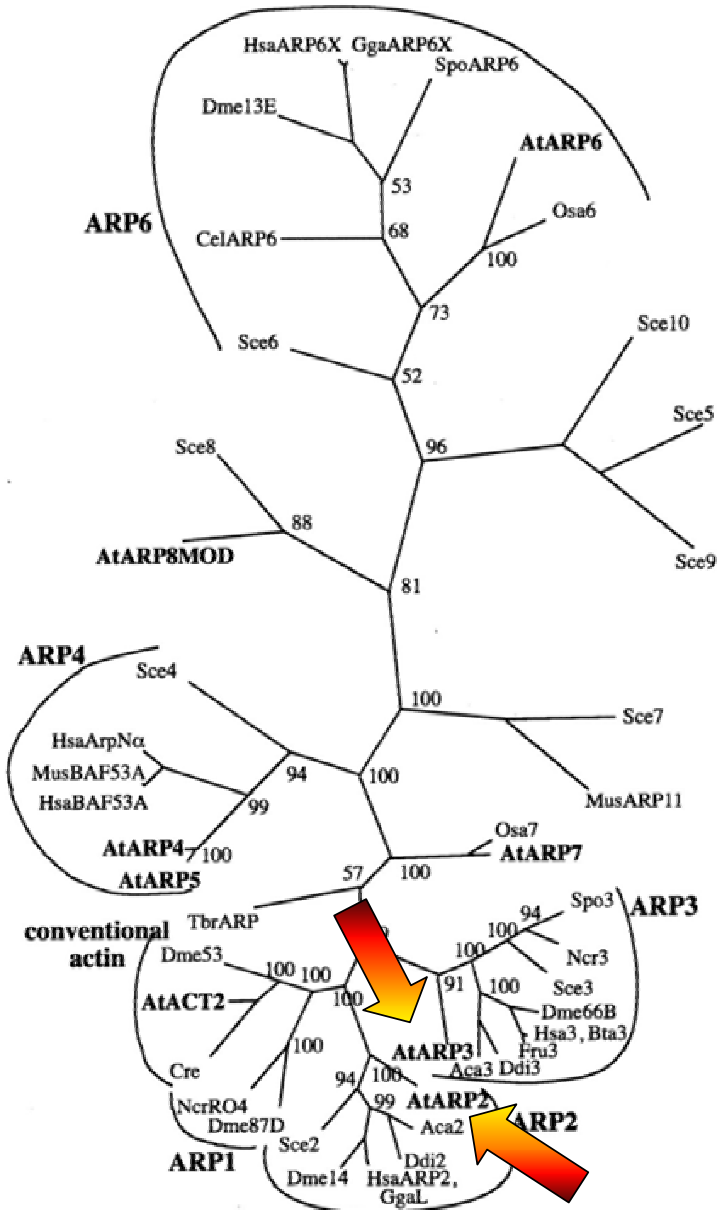
arpc5  
brk1



Podjednotky Arp2/3 komplexu jsou evolučně staré

Arabidopsis:

- ARPC1/Arc40p/p40 (2)
- ARPC2/Arc35p/p35 (2)
- ARPC3/Arc18p/p21 (1)
- ARPC4/Arc19p/p20 (1)
- ARPC5/Arc15p/p15 (1)



# Arabidopsis: mutanti třídy distorted1

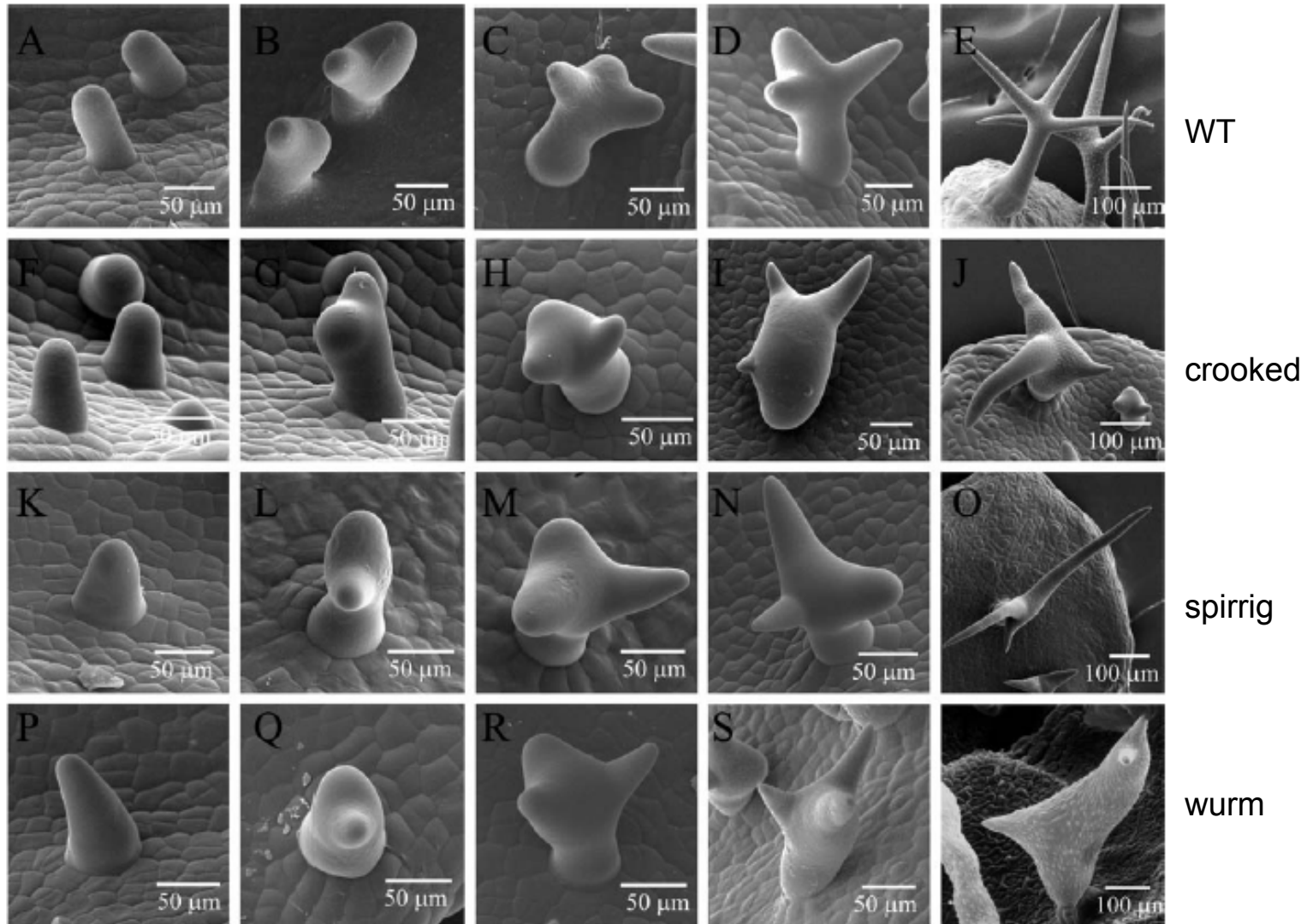
**Table 3.** The *DISTORTED* class of *Arabidopsis* genes

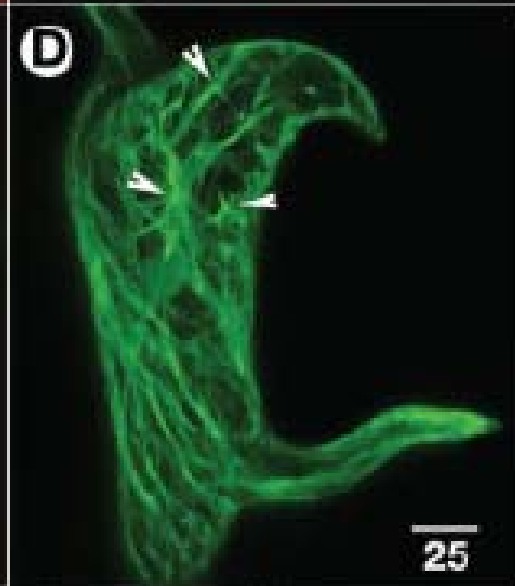
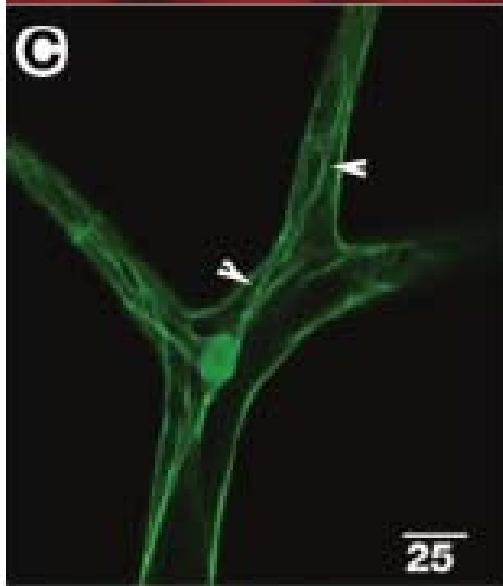
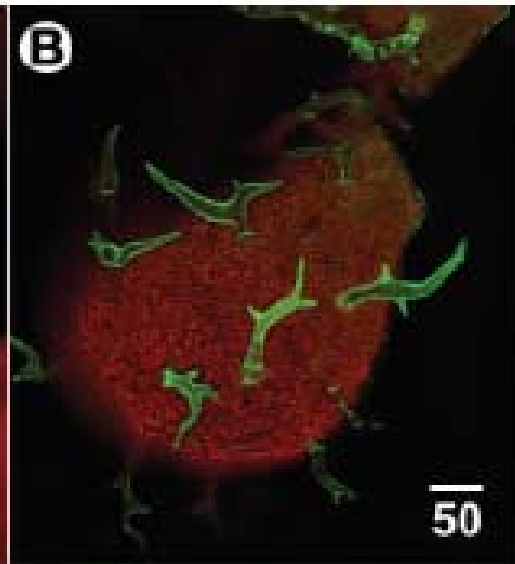
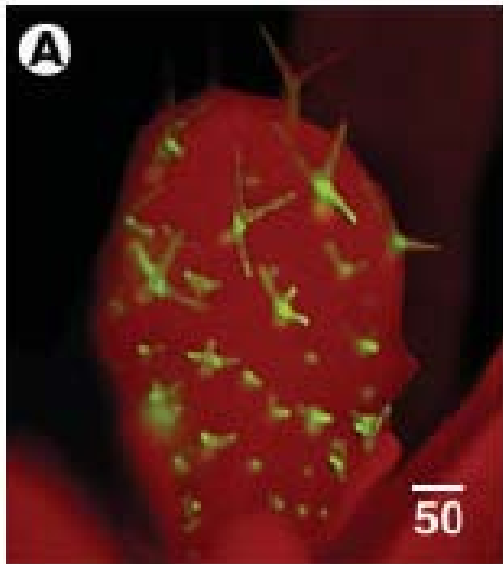
<b>Gene</b>	<b>Chr</b>	<b>AtDB Ac. No</b>	<b>Homolog for</b>
→ <i>ALIEN</i>	4	unknown	unknown
→ <i>CROOKED</i>	4	At4g01710	ARPC5
→ <i>DISTORTED1</i>	1	At1g13180	ARP3
→ <i>DISTORTED2</i>	1	At1g30825	ARPC2
<i>GNARLED</i>	2	At2g35110	NAP135
→ <i>KLUNKER*</i>	5	At5g18410	PIR121
→ <i>SPIRRIG</i>	1	Unknown	Unknown
<i>WURM</i>	3	At3g27000	ARP2

Podjednotky a regulátory Arp2/3 komplexu!

(Mathur, 2005)

# Arabidopsis: mutanti třídy *distorted*





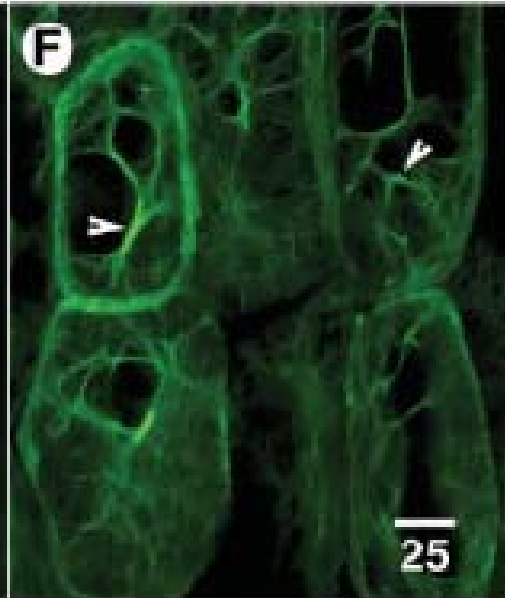
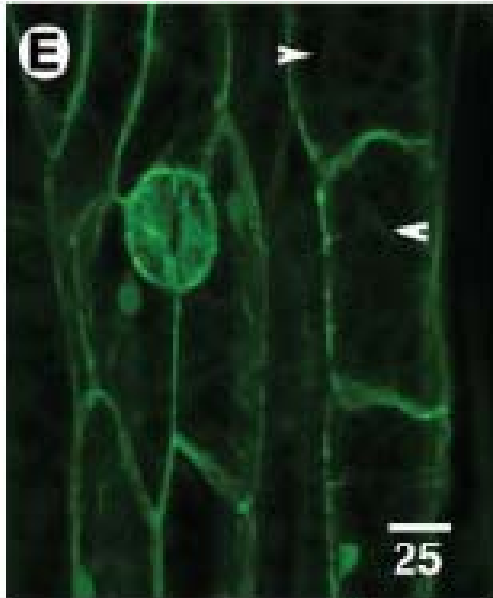
autofluorescence

GFP-mTalin

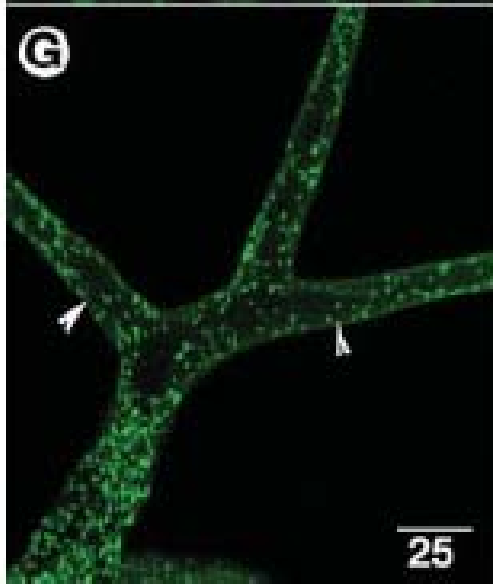
WT

*wurm*

hypokotyl  
GFP-mTalin



trichomy  
Golgi



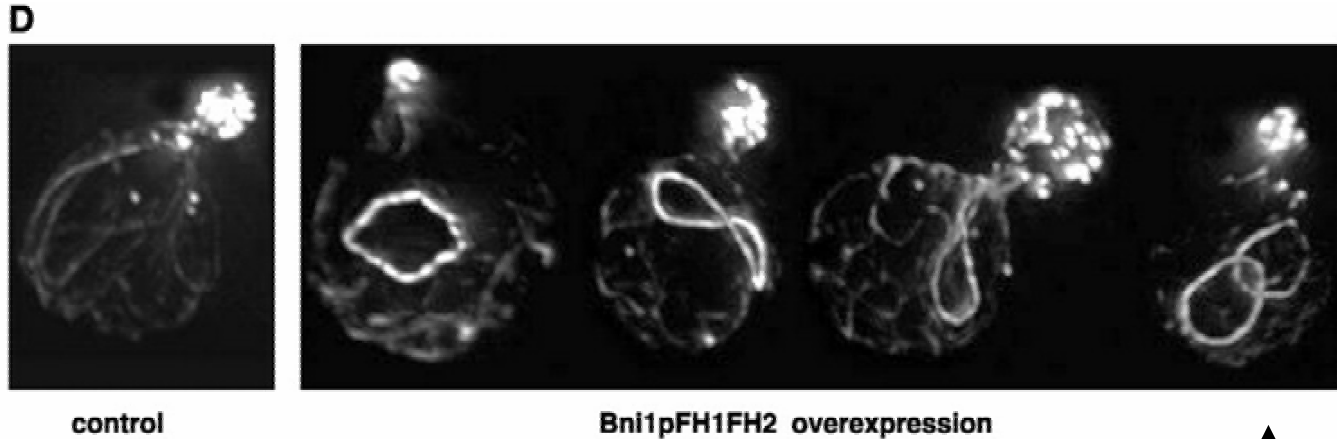
agregáty!

WT

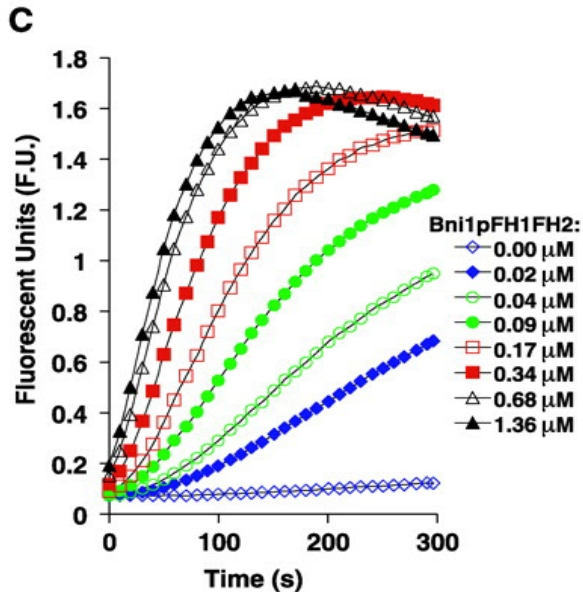
*crooked*



# Další způsoby nukleace: forminy (FH2 proteiny)

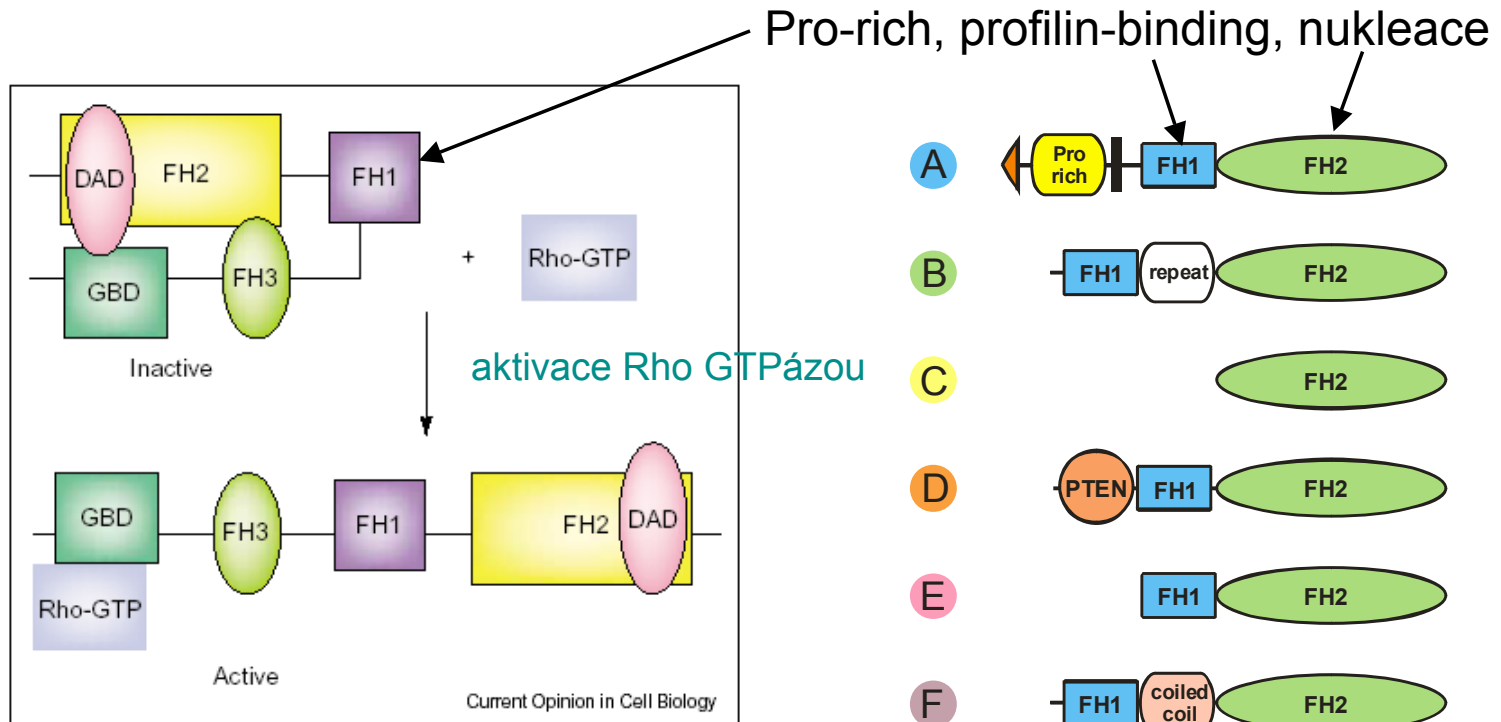


overexpression of a partial yeast formin causes formation of ectopic actin structures



overexpression of formins stimulates filament formation *in vitro*

# Doménová struktura forminů



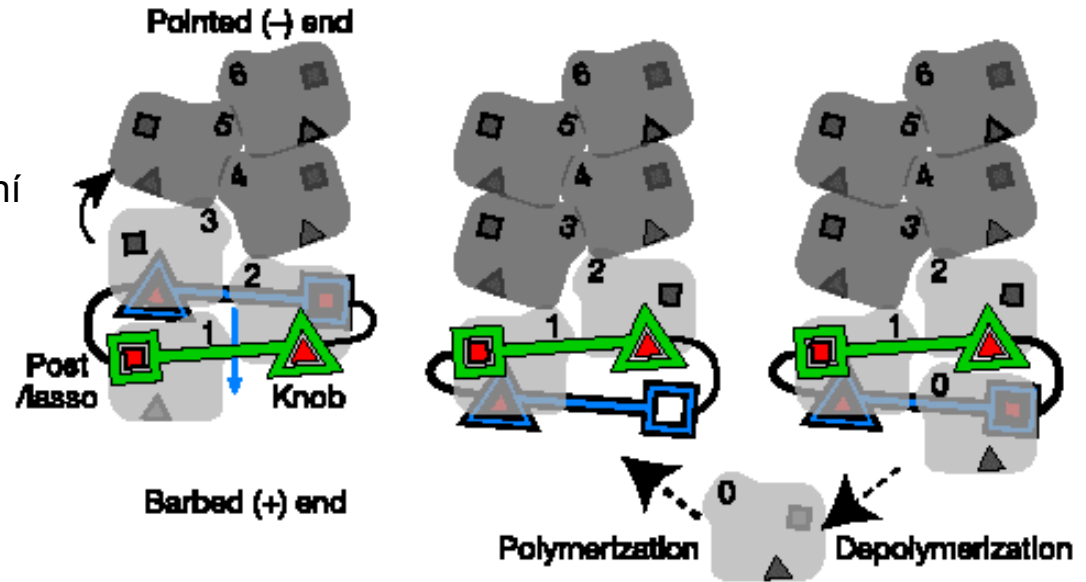
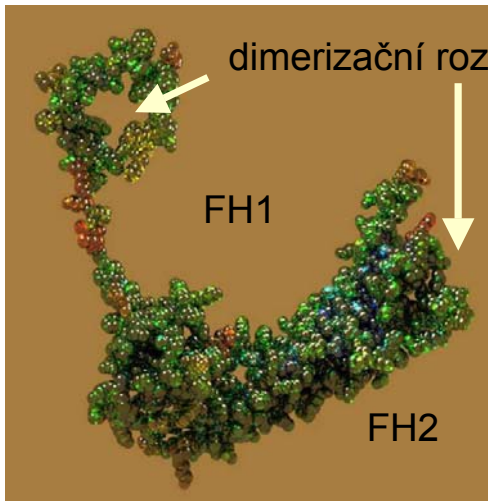
Activation of Diaphanous-related formin by Rho-GTPase. Diagram of domains present in Diaphanous-related proteins, illustrating the conformation change that occurs upon binding Rho-GTP.

Arabidopsis: 21 genů!

FH1-FH11 transmembránová doména (Class I)

FH12-FH21 bez tm. domény (Class II)

# „Kráčející nukleační komplex“

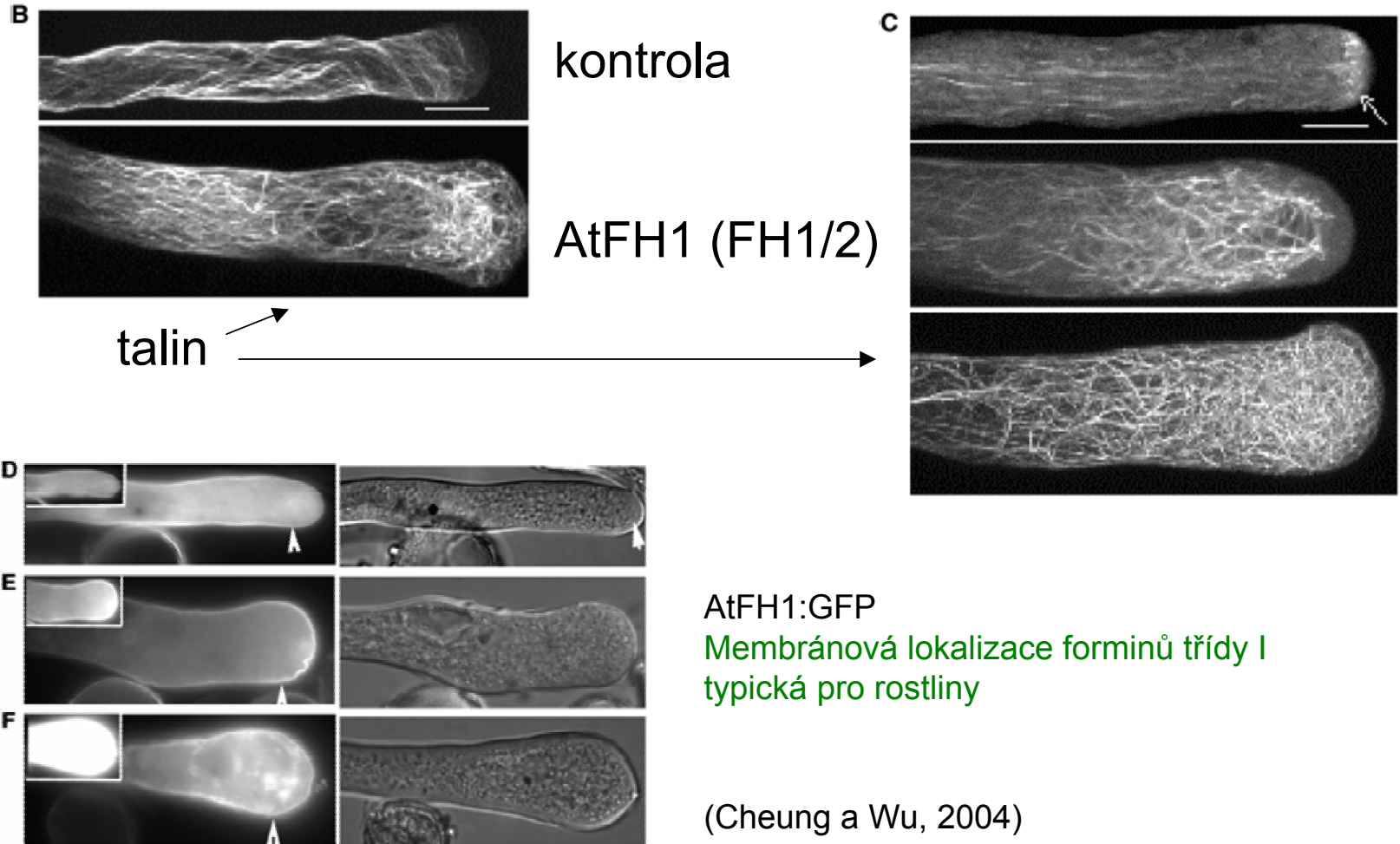


„leaky cap“ model - živočišný

Otomo et al. Nature 433:488-494, 2005

u rostlin?  
vazba na MT?

# Overexpresse AtFH1 v pylových láčkách

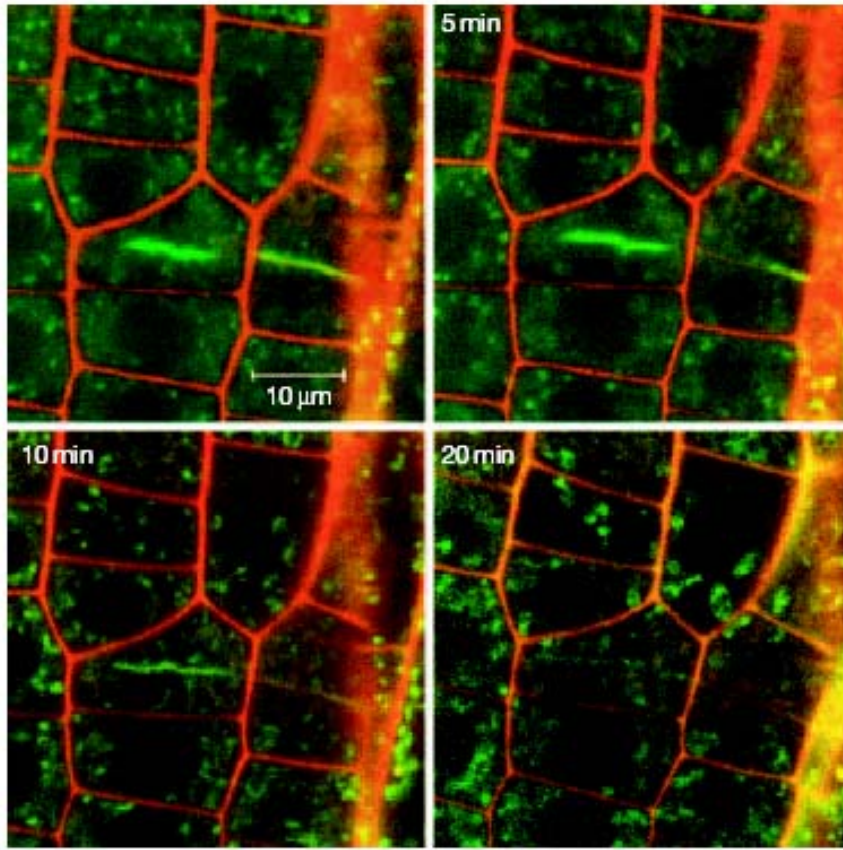


# Proč mají rostliny právě tyto geny v tolika kopiích?

- Kromě funkčního rozrůznění i „jemné ladění“ pro proměnlivé podmínky
  - v čase (rostlina je přisedlá)
  - v (mikro)prostoru buňky obklopené stěnou?
- Možnosti se nevylučují

(viz též Nasmyth, Dirick, Surana, Amon, Cvrčková, 1991)

# Specifické funkce isoformem forminů



- AtFH1: bundling
- AtFH5: cytokineze endospermu
- AtFH6: zvýšená exprese v hálkách indukovaných hlísticemi,
- suprimuje mutaci *bni1* u kvasinky
- AtFH4/AtFH8: kořenově specifické, slabý defekt na kořenových vláscích

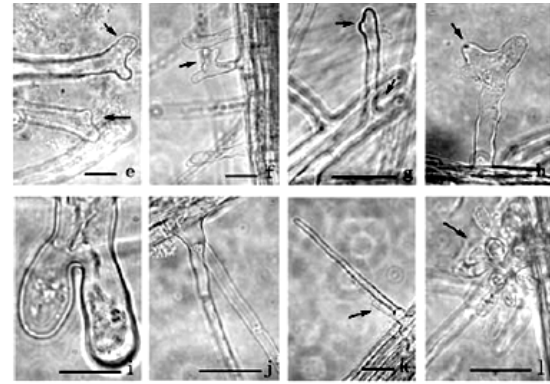


Figure 3 Formin AtFH5-GFP is targeted to the developing cell plate.

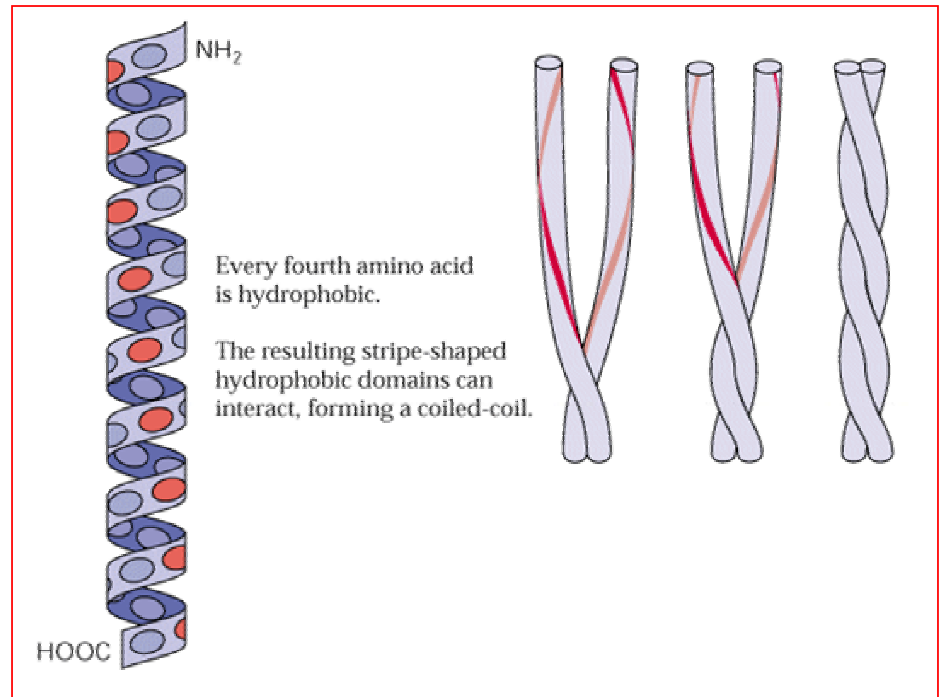
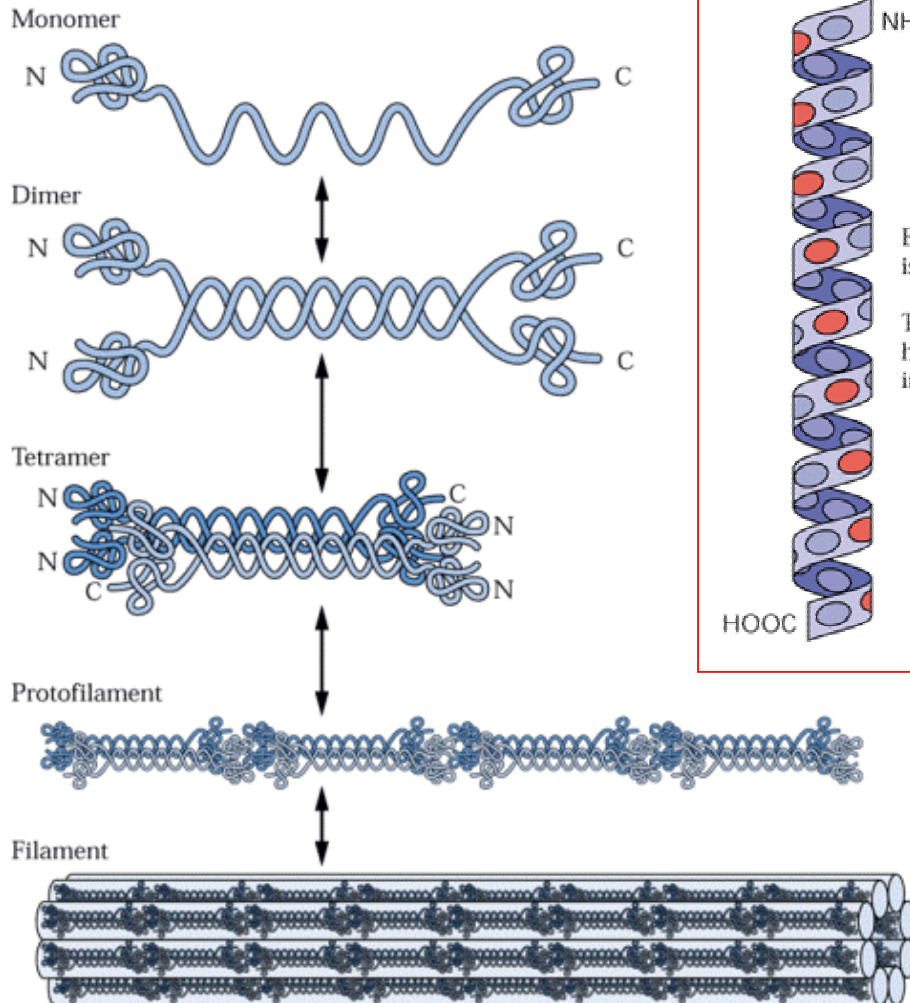
mutant *fh5* má zpožděnou celularizaci endospermu (jinde redundantní)

(Ingouff et al. 2005)

mutace ve forminech u různých organismů často vedou k defektům v cytokinezi

# Intermediální filamenta

# Intermediální filamenta: konvergence?

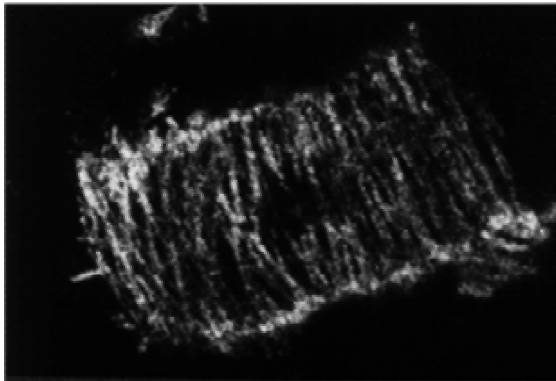


regulace např. fosforylací  
(CDK ... jaderná lamina)

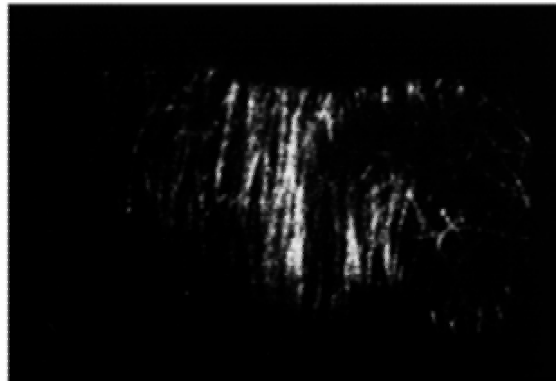
Výjimka z pravidla: rovnocenné konce!



(A)

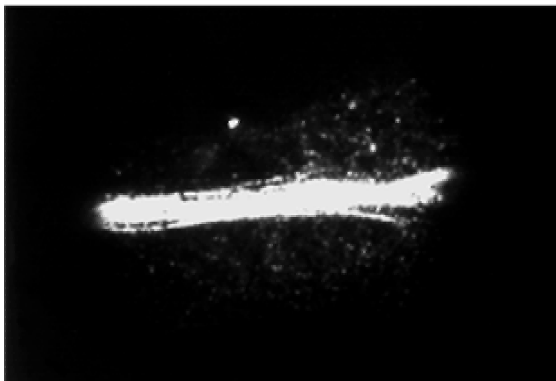


(B)

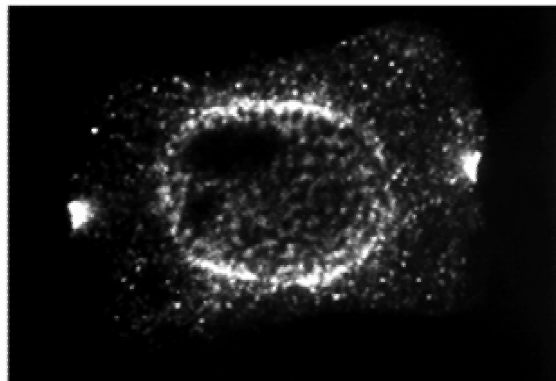


interfáze

(C)



(D)

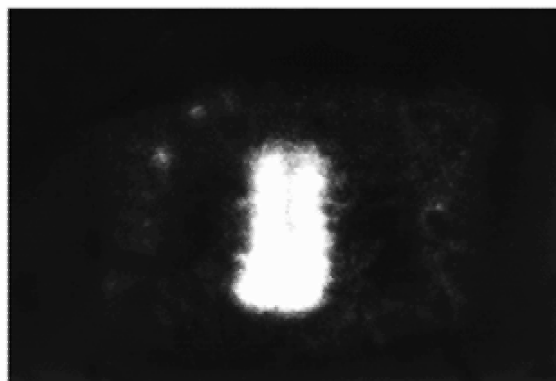


profáze

(E)



(F)

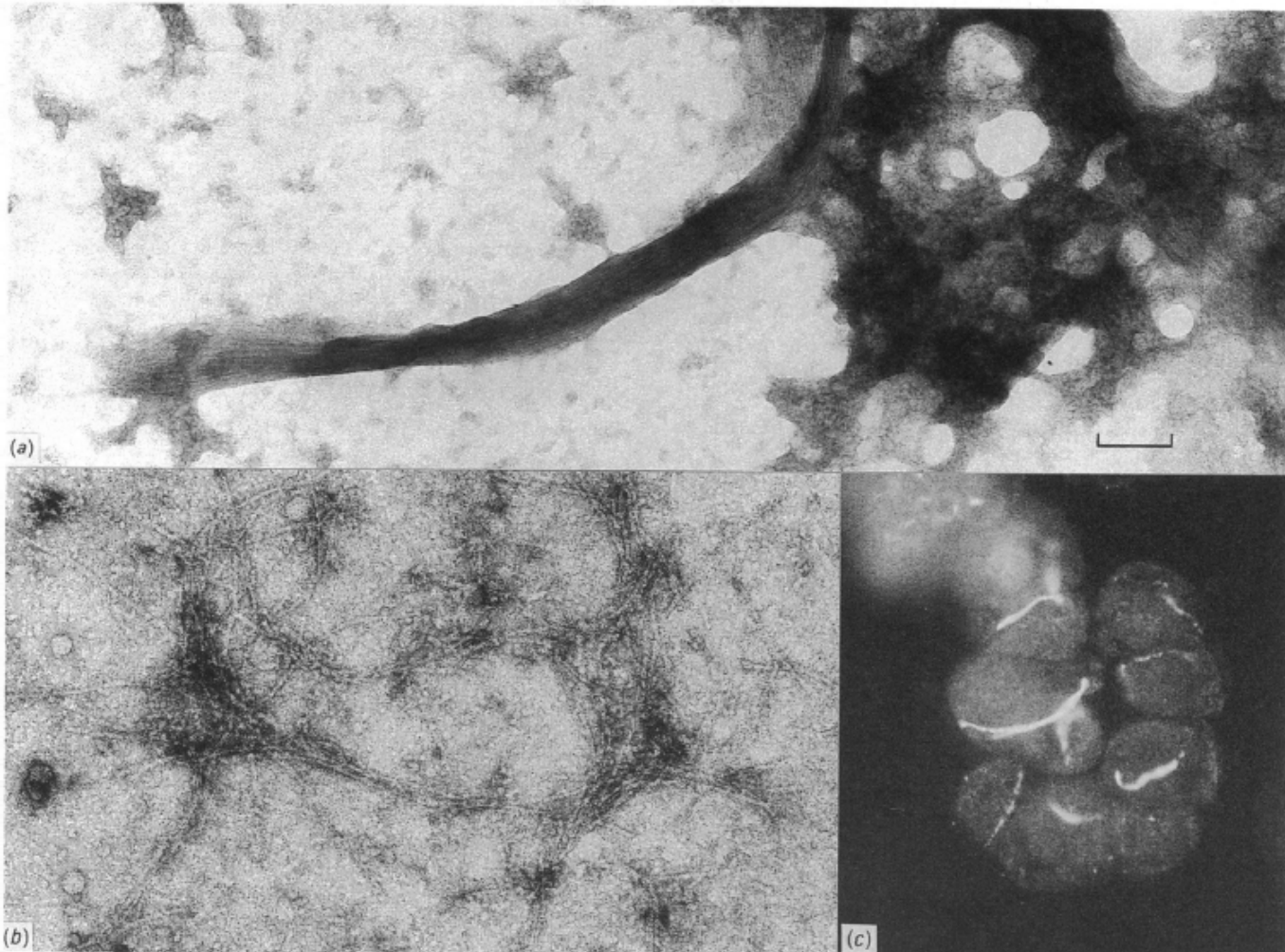


mitóza a cytokineze

# Epitopy příbuzné IF v rostlinách?

kolokalizace  
s MT?

wheat root tip,  
anti-keratin



**Fig. 1. Ultrastructural characterization of reconstituted 10 nm filaments and filament bundles**

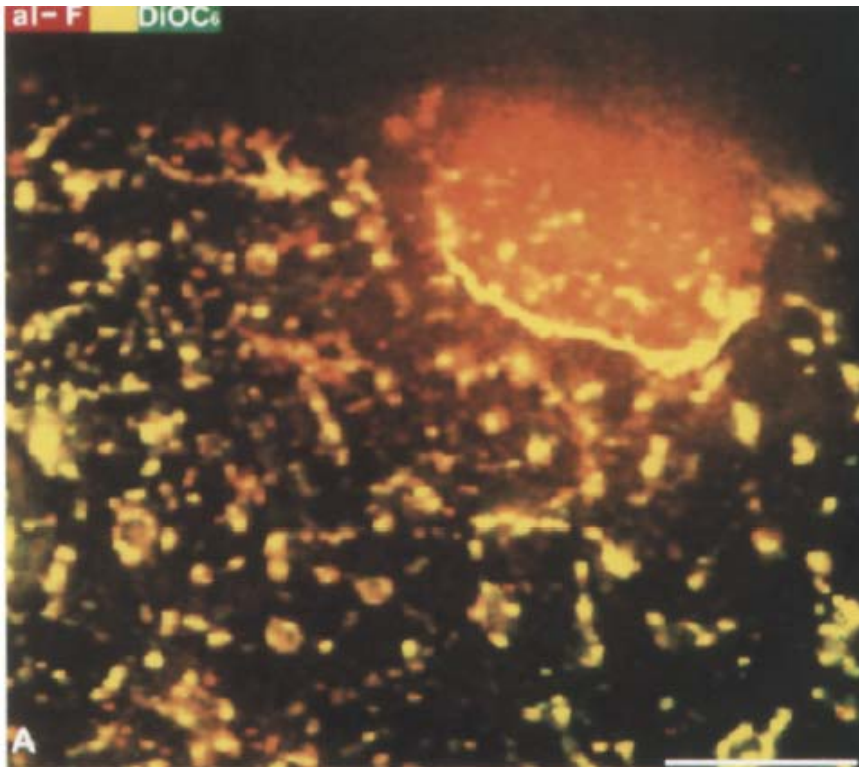
Shown are electron micrographs of a reconstituted filament bundle (a) and reconstituted 10 nm filaments (b), together with indirect immunofluorescence of carrot cells showing native fibrillar bundles stained with AFB followed by a fluorescein-conjugated goat anti-rat IgM (c). The bar (shown only in a) represents 100 nm in (a) and (b) and 10  $\mu$ m in (c).

# Mikroinjekce značených protilátek

**Covisualization in living onion cells of putative integrin, putative spectrin, actin, putative intermediate filaments, and other proteins at the cell membrane and in an endomembrane sheath**

Protoplasma (1997) 199: 173–197

Christophe Reuzeau\*\*, Keith W. Doolittle, James G. McNally, and Barbara G. Pickard\*

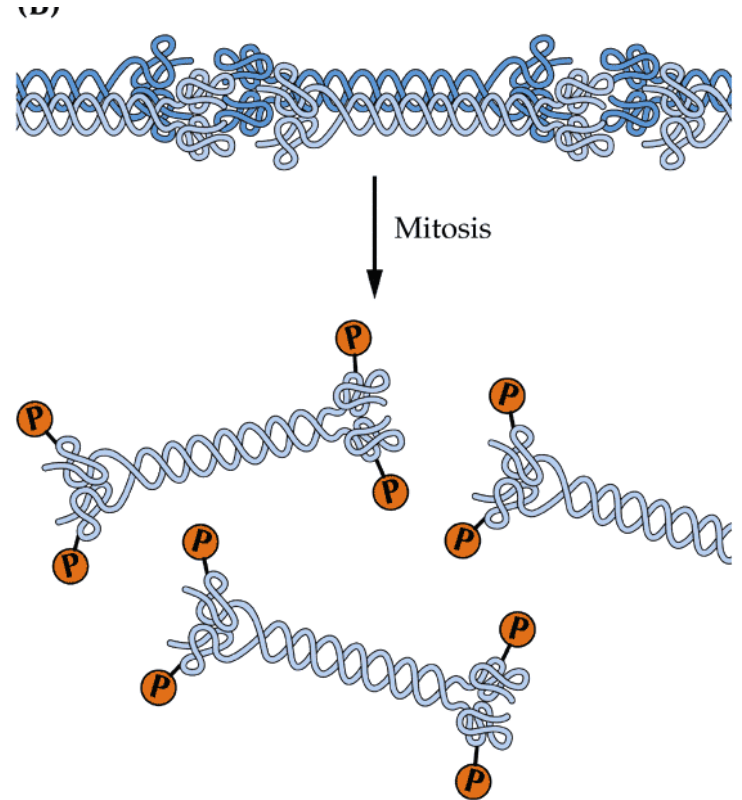
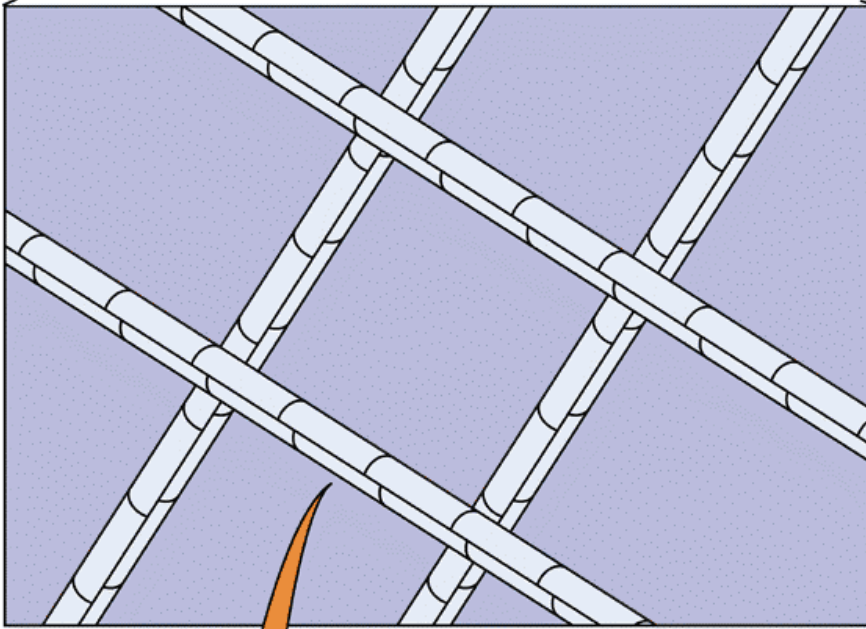
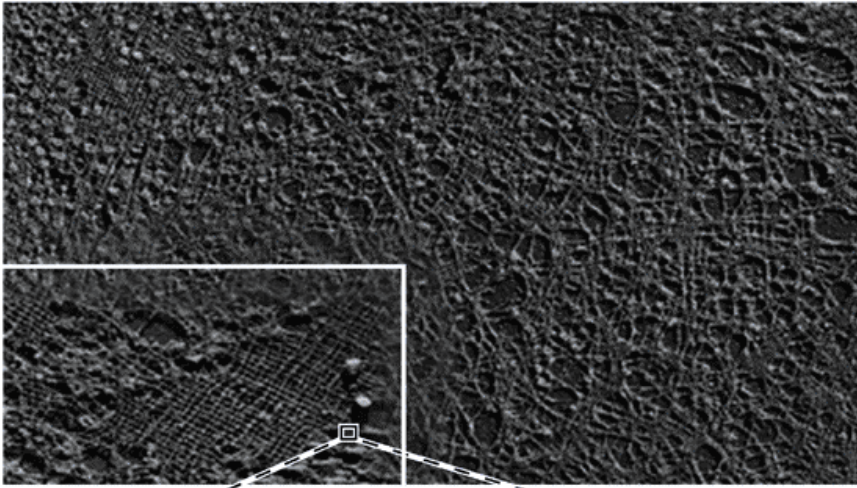


anti-IF

DIOC6 - membrány

# Jaderná IF - laminy

Asi jen u živočichů?



# Jak u rostlin?

- nacházíme „laminové“ epitopy, ale v genomu *A. th.* není ortolog
- rekonstrukce jader *Xenopus* v bezbuněčných extraktech z tabáku (Lu and Zhai 2001)
- savčí LBR se lokalizuje do jaderné membrány v tabáku (Irons et al. 2003)

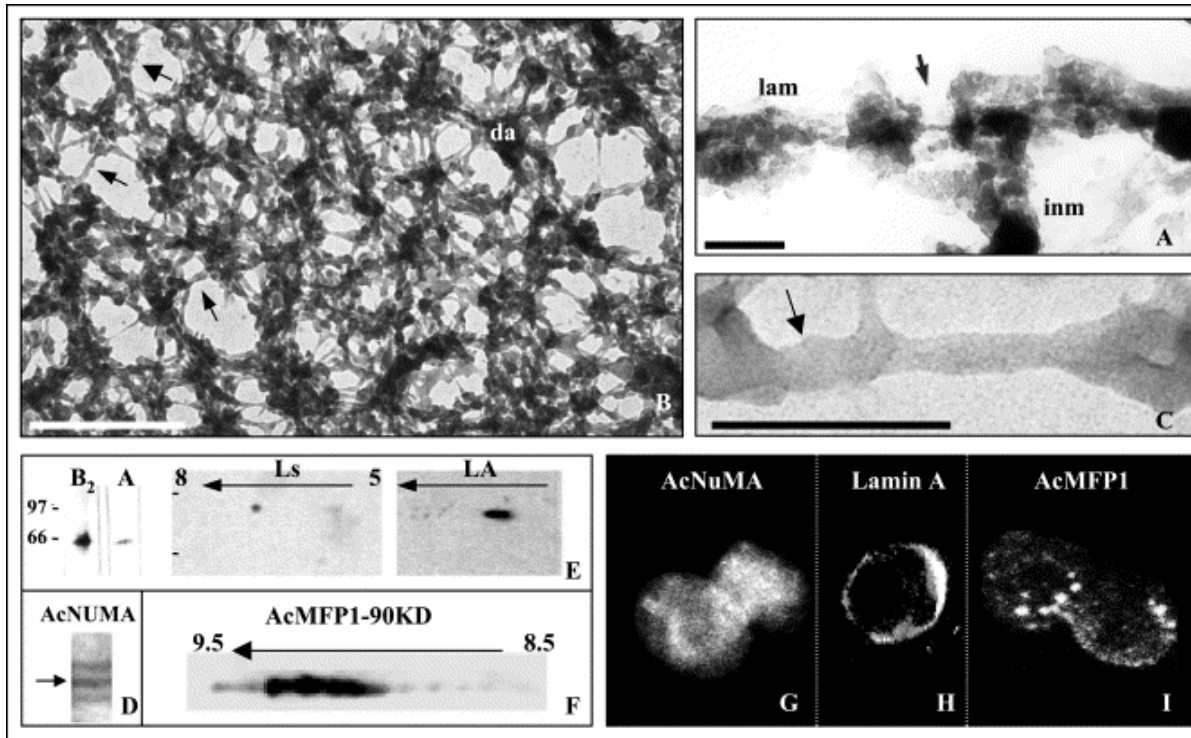
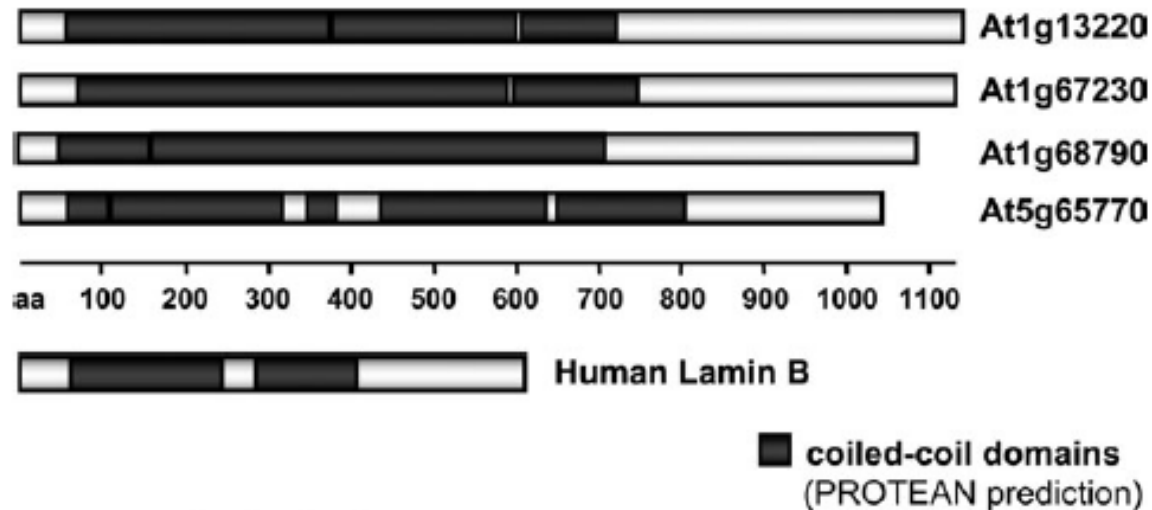


Fig. 1. Ultrastructural organisation of the plant NM in resinless sections. (A) Portion of the peripheral lamina with a complex structure (lam), connected with the internal matrix (inm). Individual filaments are observed (arrow). (B) A delicate anastomosed network of filaments forms the internal matrix: 15–25 nm knobbed filaments (arrows) and thicker ones, covered by globular structures (big arrow; dense aggregates (da)). (C) Higher magnification of a branched filament of the internal matrix, showing the junction of two filaments and a typical 25 nm knob. Bar in (A), 0.1  $\mu$ m; in (B), 0.5  $\mu$ m; and in (C), 0.1  $\mu$ m. Identification of the IF proteins by Western blot in the onion NM. (D) The anti-NuMA serum, S2, reveals three onion isoforms, the major one at 220 kD (arrow). (E) Antibodies against chicken B2 and A lamins, and a serum against the chicken lamina (Ls) recognise in all cases protein spots at 65 kD with pI varying from 5.65 to 6.8. (F) 2-D blot with serum 288 against LeMFP1. The 90 kD AcMFP1 shows up to 12 basic spots. In situ localisation of the IF proteins in the NM. (G) NuMA is associated with the internal matrix network and accumulates in small foci with a punctate pattern. (H) Lamin-like proteins accumulate not only at the peripheral lamina, but also in internal foci. (I) AcMFP1 accumulation in large replication foci from a late-S phase NM.

# Rostliny sice nemají laminy, ale ...

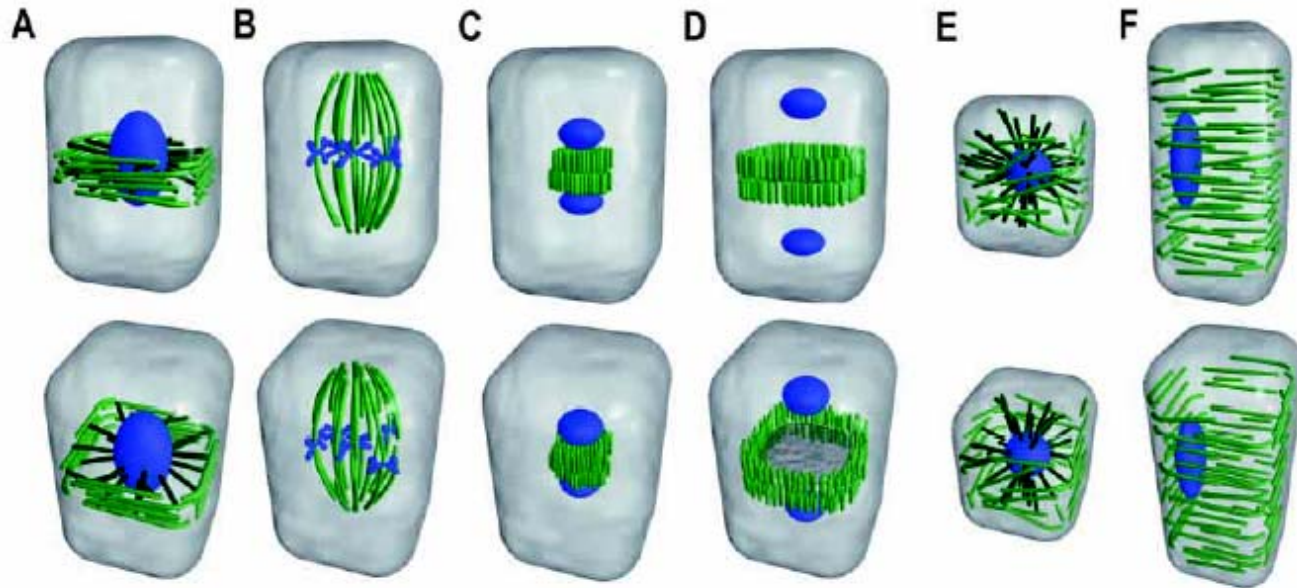


**Fig. 1.** Comparison of localization and structural organization of human lamin B and *Arabidopsis* nuclear matrix constituent proteins (NMCPs). While almost twice the size of lamins, NMCPs have a comparable domain organization of short head, coiled-coil centre, and longer tail domain. The gene identifiers of the four *Arabidopsis* homologues of DcNMCP1 are shown.

# Mikrotubulární cytoskelet

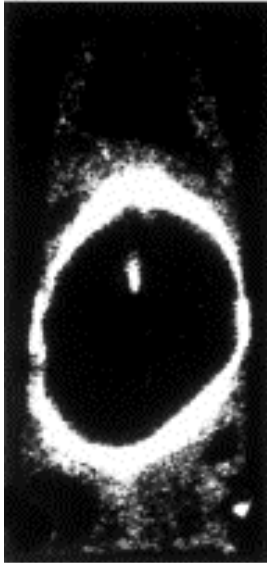


# Mikrotubulární systémy rostlinné buňky

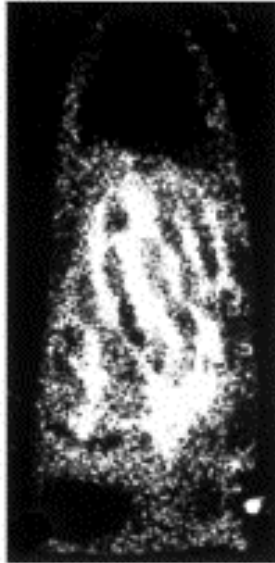


**Fig. 1.** These schematic illustrations, rendered in 3D at two aspects, show microtubule arrays through the plant cell cycle. (A) A preprophase band, linked to the nucleus by phragmosome microtubules, marks the future division site. (B) Metaphase spindle with a dispersed polar region. (C) In telophase, the phragmoplast forms as a concentrated cylinder of microtubules between daughter nuclei. (D) The cytokinetic phragmoplast expands centrifugally, leading the cell plate towards attachment sites previously established by the preprophase band. Microtubule plus ends meet at midplane. (E) Once cytokinesis is complete, microtubules extend from the nucleus toward the cell cortex and plasma membrane-associated microtubules appear. (F) Plant cells in interphase and those entering terminal differentiation often expand predominantly in one direction. During cell elongation, cortical microtubules are usually arranged in parallel arrays whose predominant orientation is at right angles to the axis of expansion.

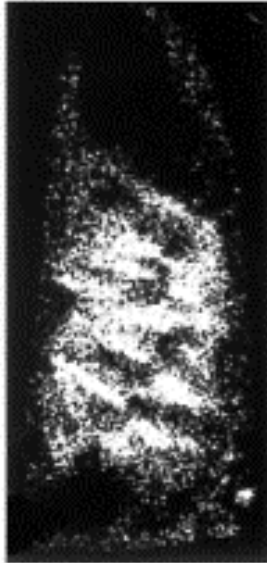
(A)



(B)



(C)



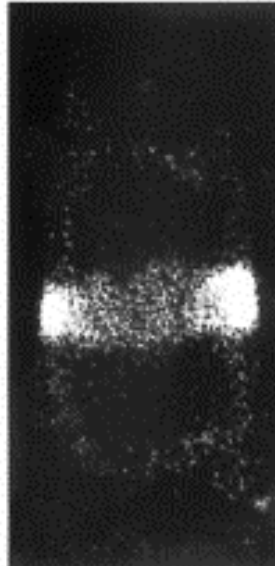
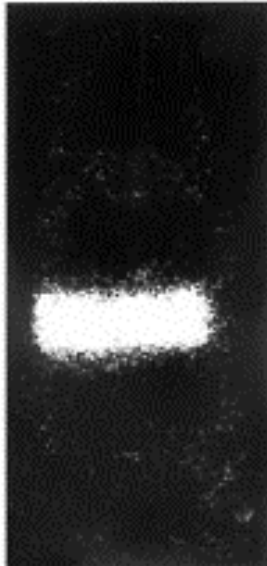
(D)



(E)



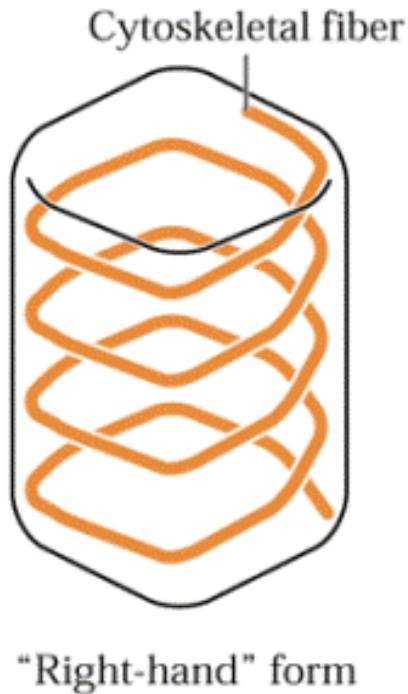
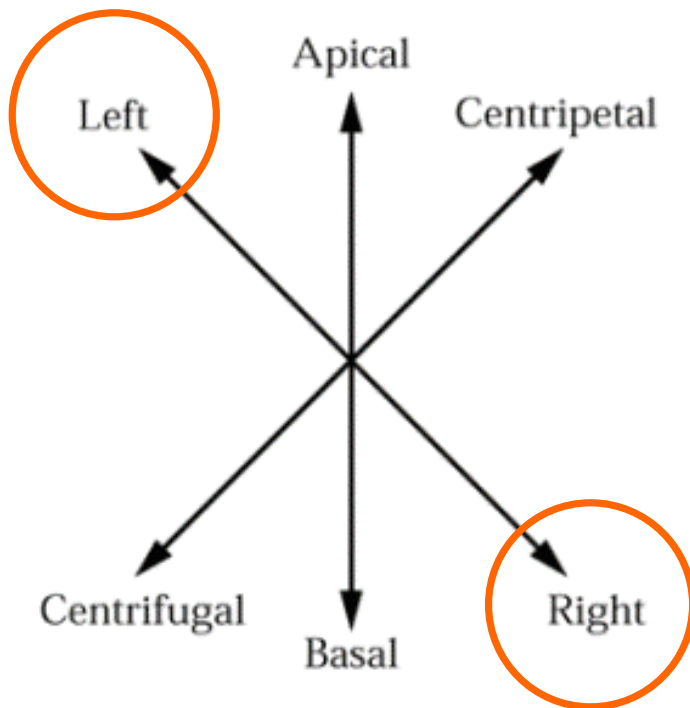
(F)



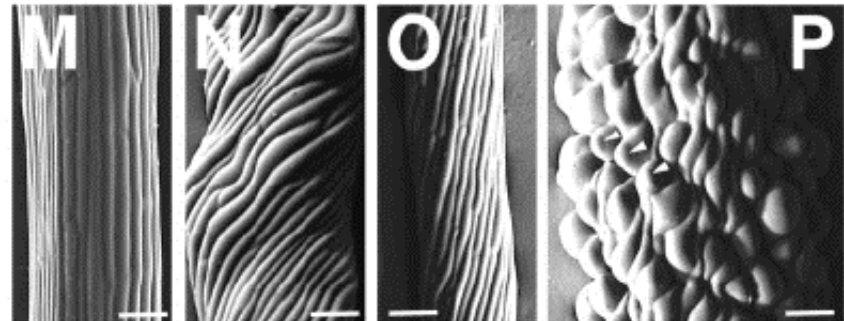
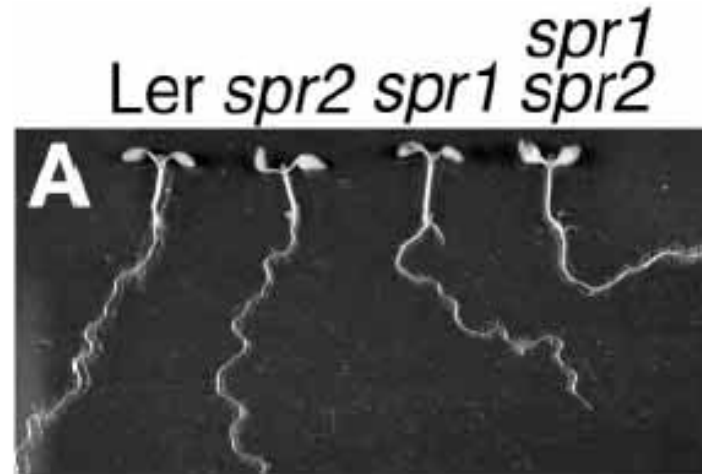
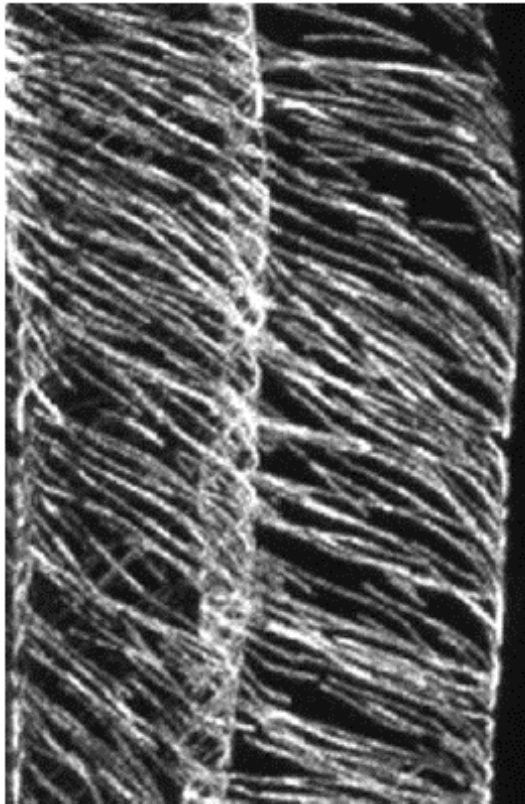
Tradescantia,  
injected  
bovine tubulin

# Helikální uspořádání MT

existuje preference směrů



# Helikální uspořádání MT

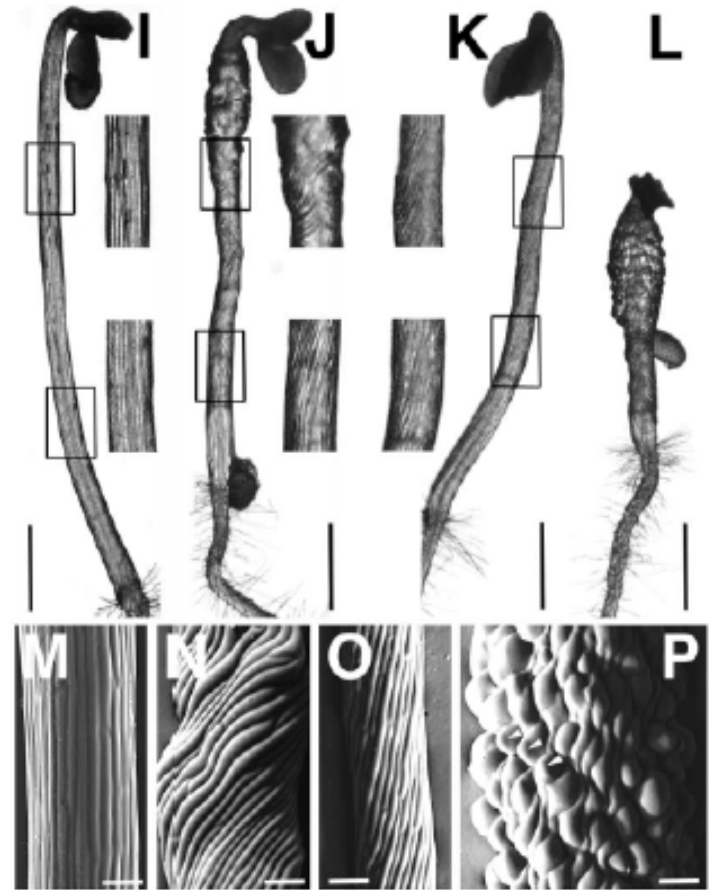
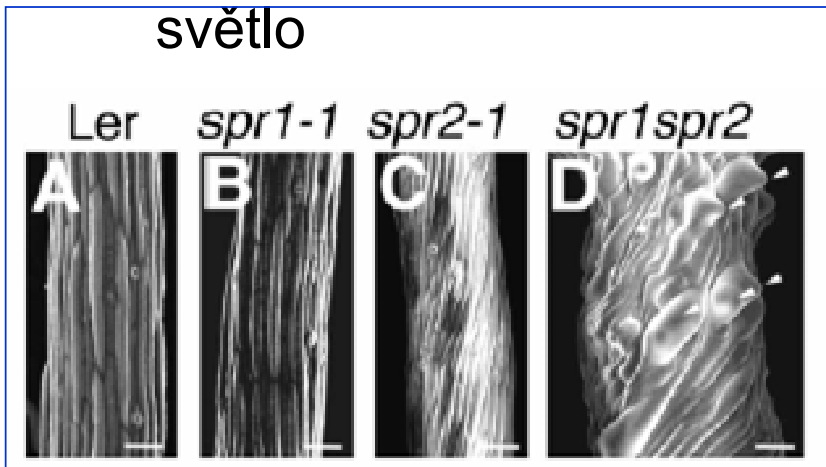


**Figure 1.** Microtubules in the *spiral* Mutant.

In the *spiral* mutant of *Arabidopsis*, the cortical microtubules of two epidermal cells wind around the cortex in left-handed S-shaped helices. Each single fluorescent strand is likely to represent a bundle of several microtubules. (Figure courtesy of Keiko Sugimoto.)

# SPR1, SPR2 (spiral)

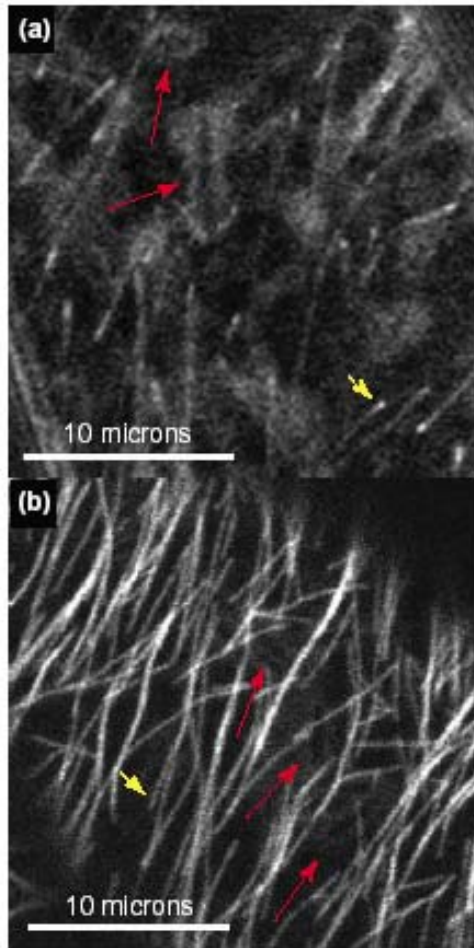
světlo



etiolované

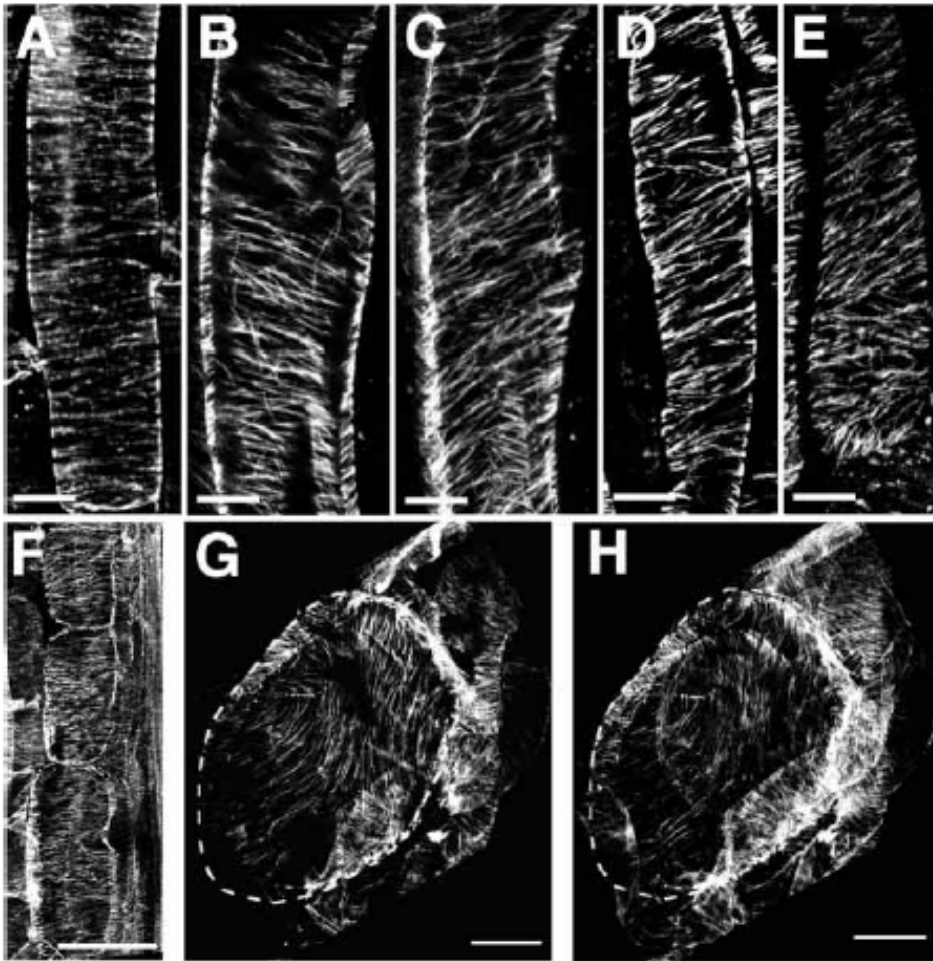
*Arabidopsis spiralis* (*spr1*) mutants show a right-handed helical growth in roots and etiolated hypocotyls due to impaired directional growth of rapidly expanding cells.

# SPR1: malý protein na +koncích MT

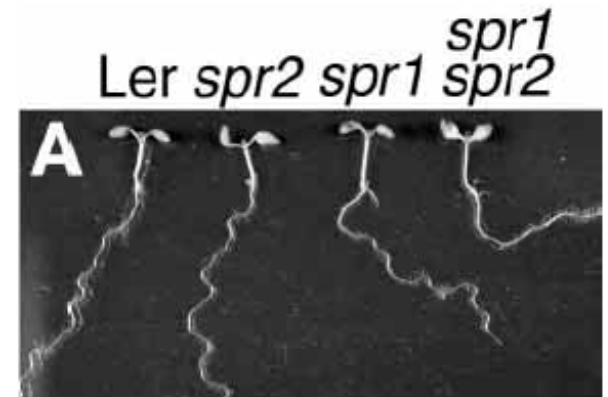


Localization of SPR-GFP and YFP-TUBULIN in the hypocotyl cells of transgenic *Arabidopsis* seedlings. GFP fluorescence from (a) an SPR1-GFP fusion protein and (b) an YFP-TUBULIN fusion protein (single confocal microscope slices). Note that the SPR1-GFP localizes preferentially to the MT plus ends (yellow arrowhead) as well as in the cytoplasm surrounding endomembranes (delineated by red arrows). By contrast, YFP-TUBULIN evenly labels MTs, yet is also found around endomembranes (delineated by red arrows). For a better view, please watch supplemental movies S1 and S3 in [35\*\*]. (b) is reproduced with permission from [35\*\*], copyright American Society of Plant Biologists (2004).

SPR1 encodes a small protein. Its localization to cortical microtubules (MTs) suggests that SPR1 maintains directional cell expansion by regulating cortical MT functions.



**Fig. 9.** Micrographs of cortical MTs in seedlings grown at 23°C. (A-E) Root epidermal cells at the basal elongation zone. (A) Ler, (B, C) *spr1-1*, (D) Ler grown on 1  $\mu$ M taxol and (E) Ler grown on 3  $\mu$ M propyzamide. Cortical MT arrays were located underneath the outer cell wall (A,B,D,E) or the inner cell wall (C). Panels B and C were from the same *spr1* cell. (F-H) Inner cortex cells at the upper region of 5-day-old etiolated hypocotyls. (F) Ler, and (G,H) *spr1-1*. G and H were from the same *spr1* hypocotyl optically sectioned at different focal planes. Broken line indicates shape of a cell in G and H. All images are positioned in their correct orientation relative to the long axis of the organ. Scale bars, 10  $\mu$ m in A-E; 50  $\mu$ m in F-H.



# „Točivost“ orgánů mutantů: směr nemusí souhlasit se směrem mt spirály

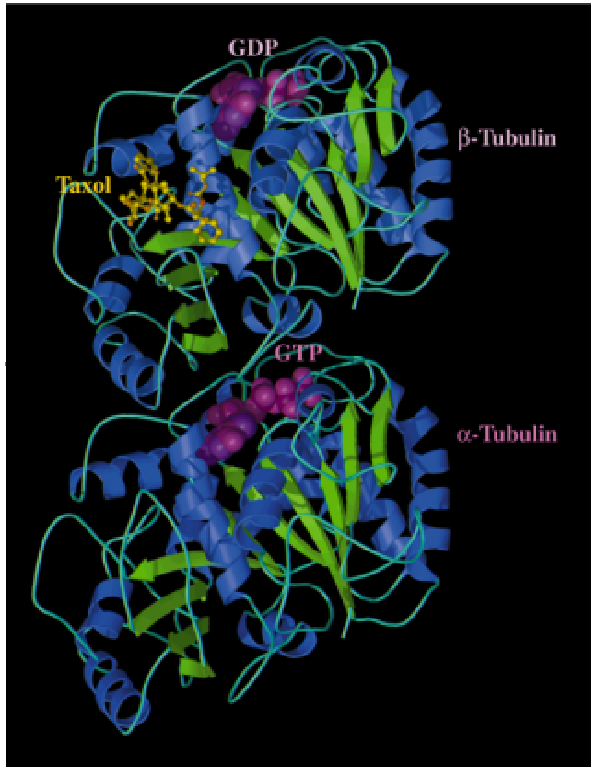
- Pravotočivé:
  - spr1
- Levotočivé:
  - inhibitory (taxol)
  - *mor1*
  - *lefty1, lefty2*: mutace tubulinu



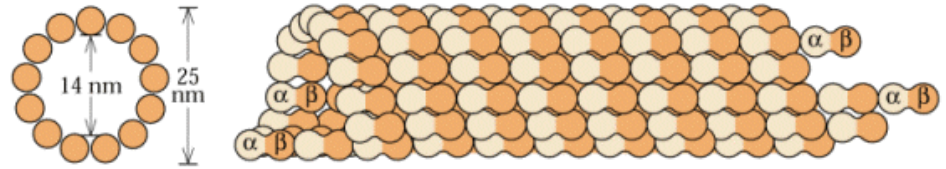
Točí se rostlina bez mikrotubulů doleva? A jak to dělá??



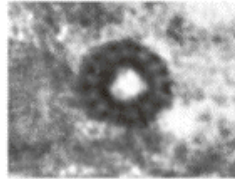
# Struktura mikrotubulů



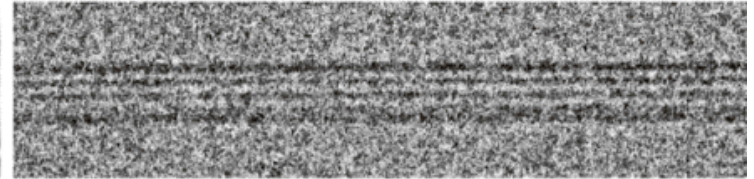
(B)



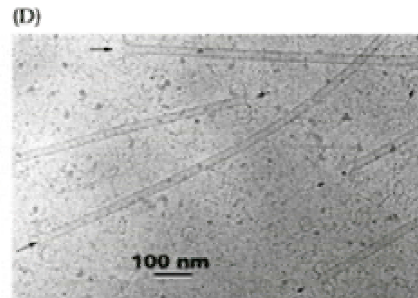
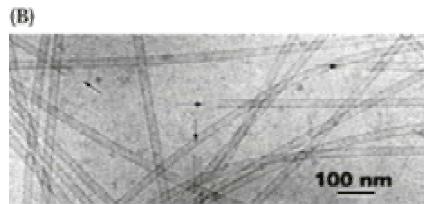
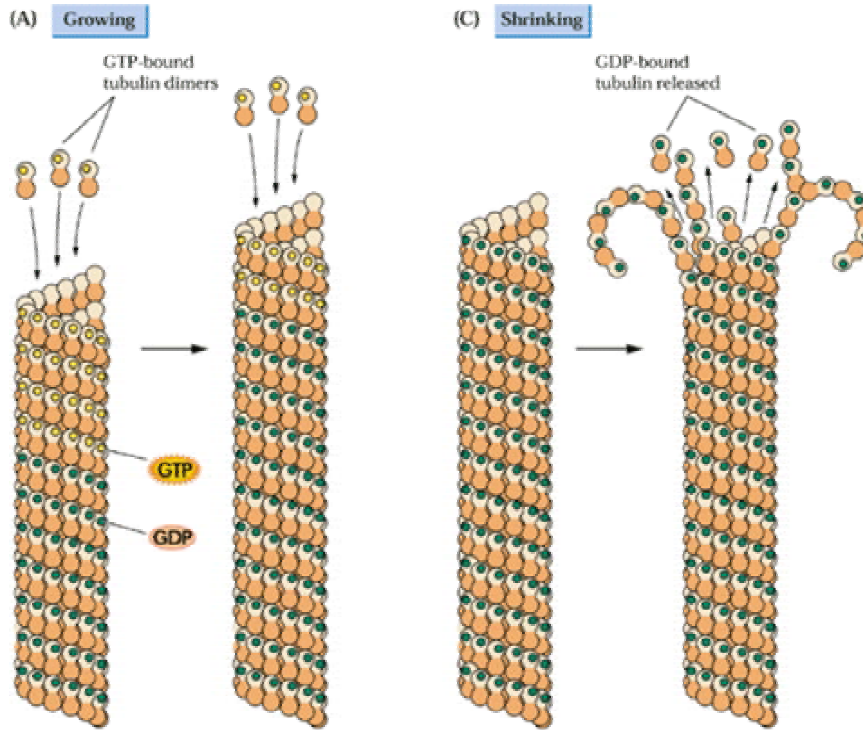
(C)



(D)

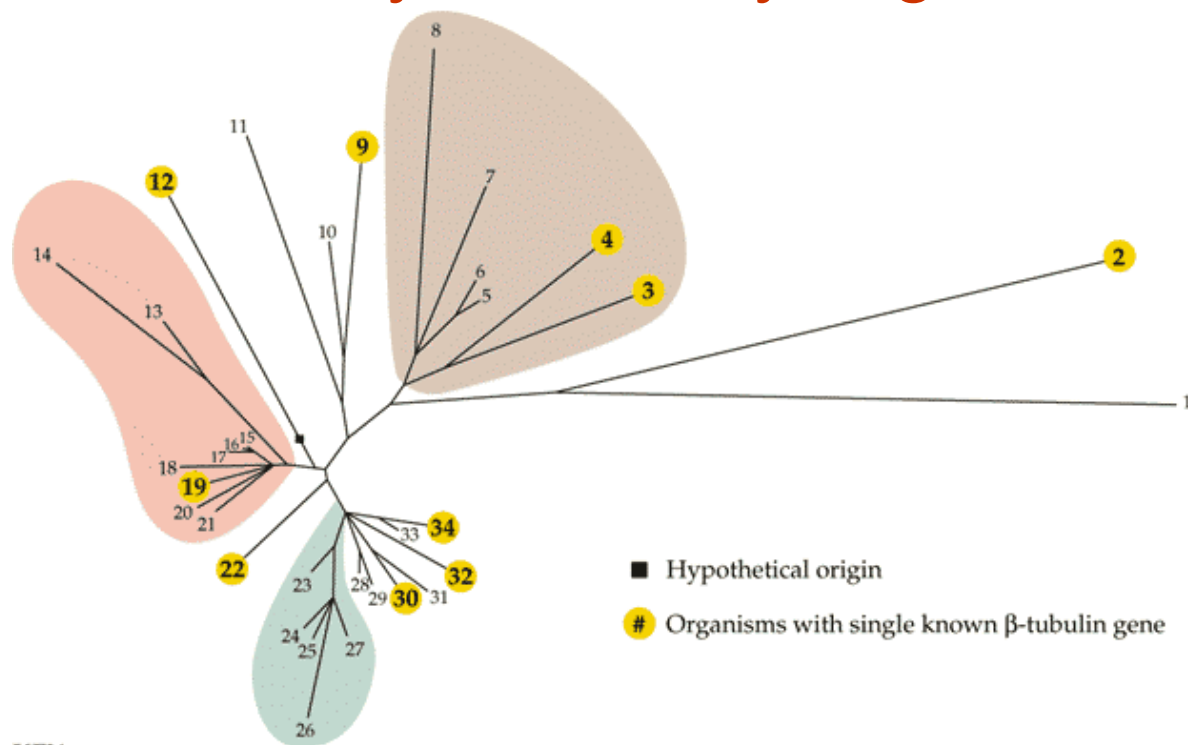


# Dynamika přestaveb mikrotubulů



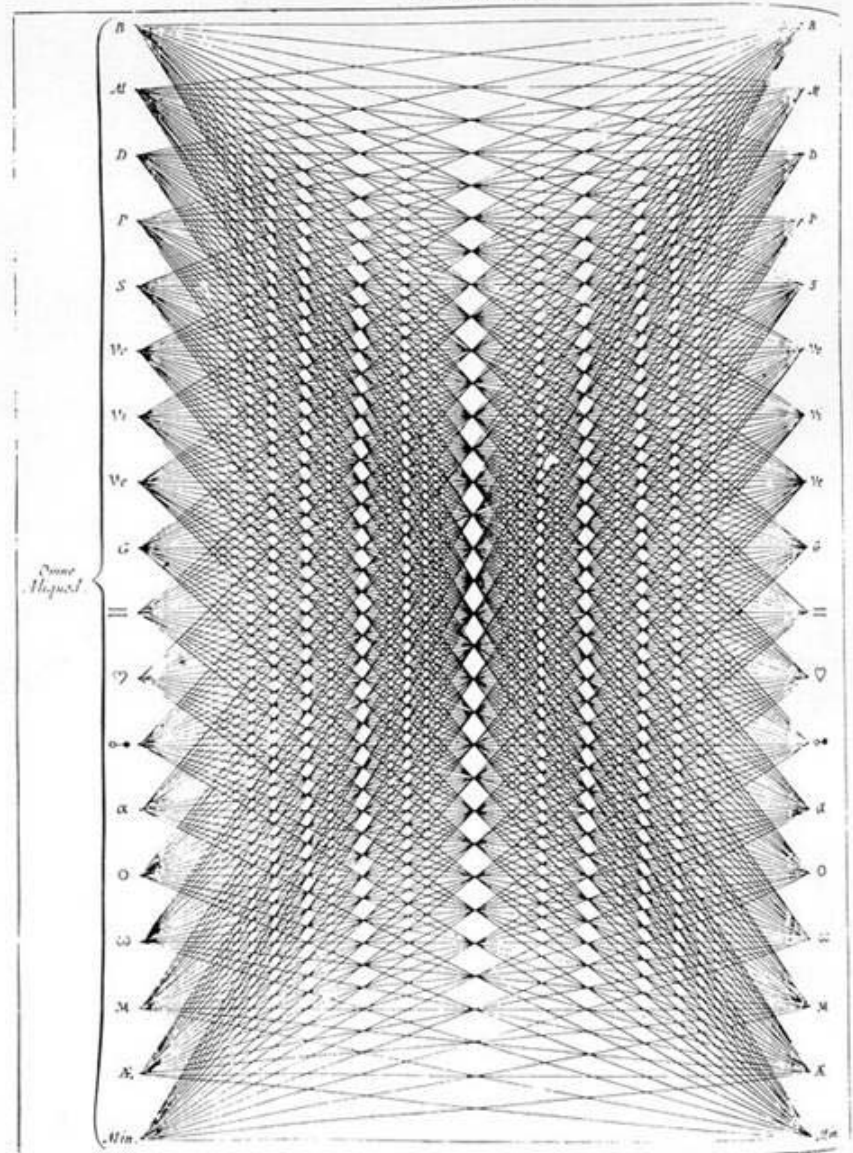
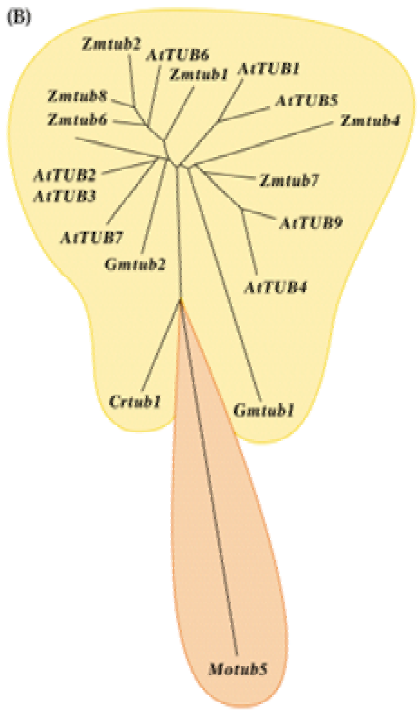
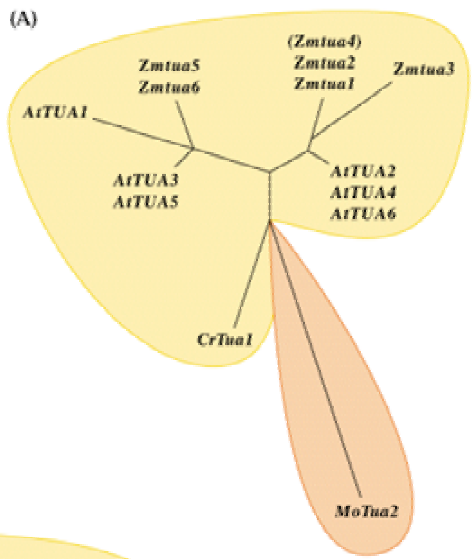
animace

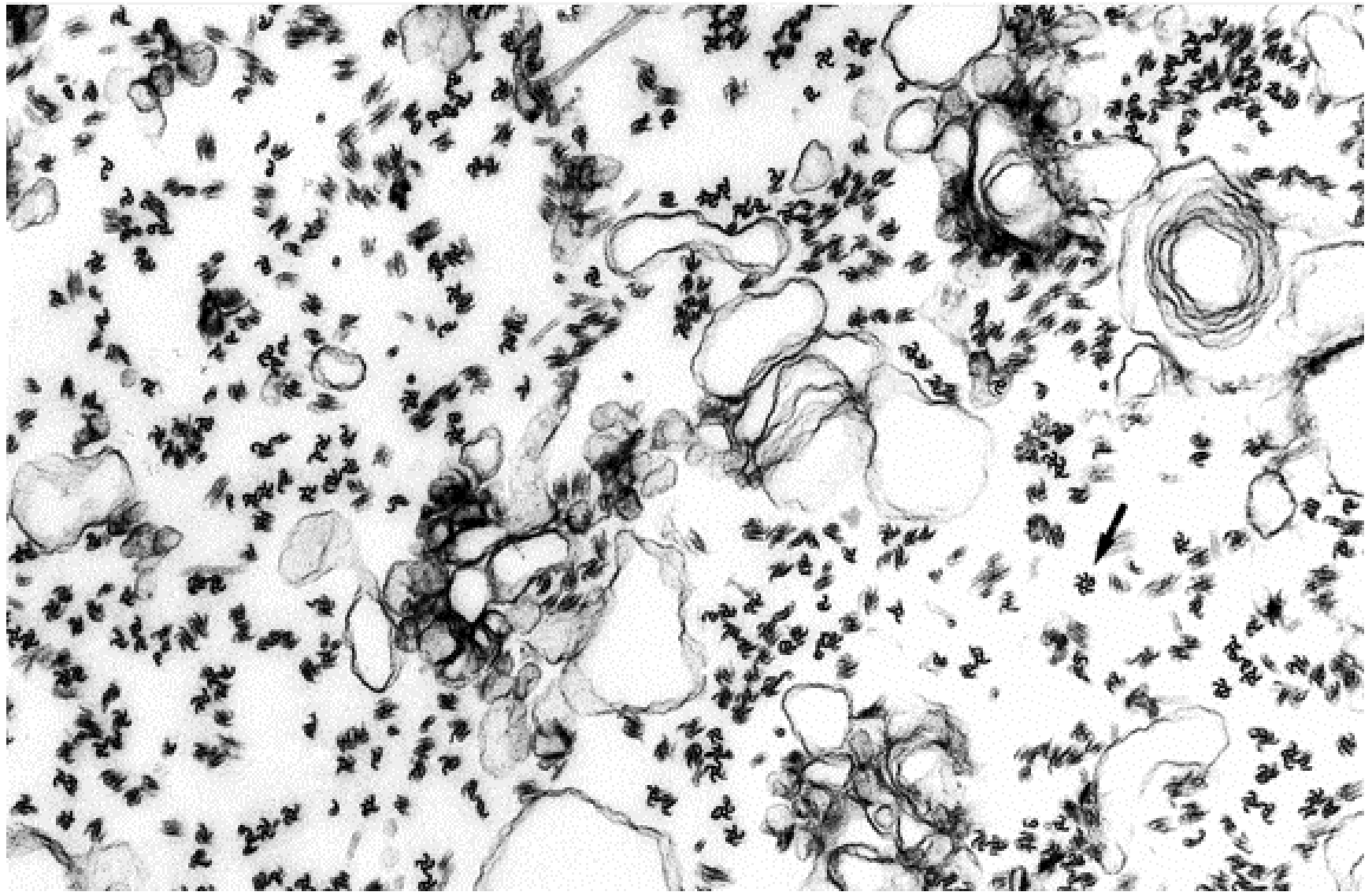
# Rodiny tubulinových genů



## KEY:

Fungi	Metazoa	Plants + green algae	Protozoa	Primitive eukaryotes
3: <i>S. pombe</i>	13: Chicken $\beta 6$	23: <i>Chlamydomonas</i> $\beta 1$	1: <i>Reticulomyxa</i> $\beta 2$	2: <i>Entamoeba</i>
4: <i>S. cerevisiae</i>	14: Mouse $\beta 1$	24: <i>Arabidopsis</i> $\beta 1$	9: <i>Dictyostelium</i>	12: <i>Trichomonas</i>
5: <i>Colletotrichum</i> $\beta 2$	15: <i>Xenopus</i> $\beta 2$	25: Maize $\beta 2$	10: <i>Physarum</i> $\beta 2$	22: <i>Giardia</i>
6: <i>Aspergillus</i> BenA	16: Human $\beta 5$	26: Soybean $\beta 1$	11: <i>Reticulomyxa</i> $\beta 1$	
7: <i>Aspergillus</i> TubC	17: Hamster $\beta 1$	27: Pea $\beta 2$	28: <i>Tetrahymena</i> $\beta 1$	
8: <i>Colletotrichum</i> $\beta 1$	18: <i>Haemonchus</i> $\beta 8$		29: <i>Plasmodium</i> $\beta A$	
	19: <i>Caenorhabditis</i>		30: <i>Achlya</i>	
	20: <i>Drosophila</i> $\beta 3$		31: <i>Ectocarpus</i> $\beta b$	
	21: Chicken $\beta 5$		32: <i>Trypanosoma</i>	
			33: <i>Physarum</i> $\beta 1$	
			34: <i>Euglena</i>	





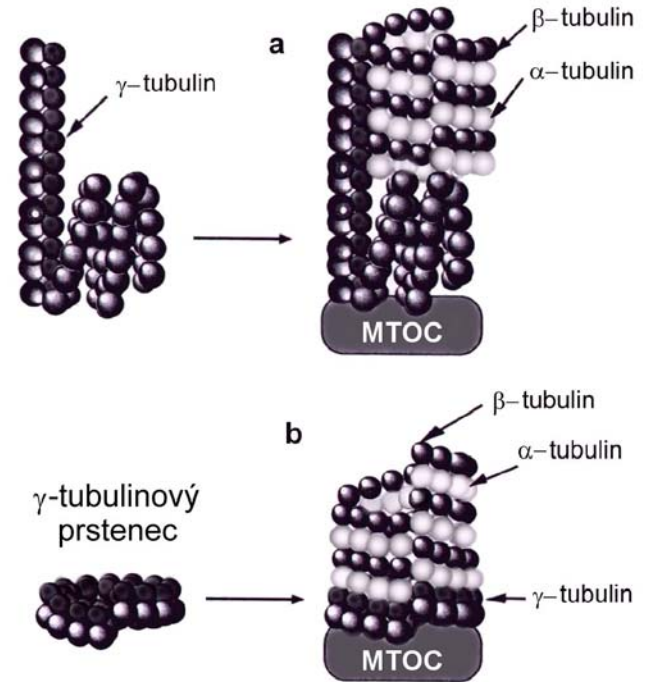
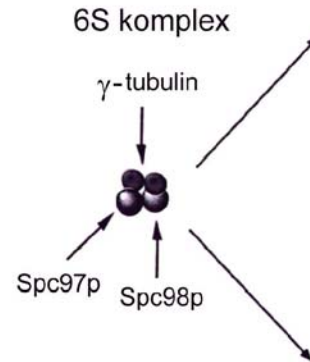
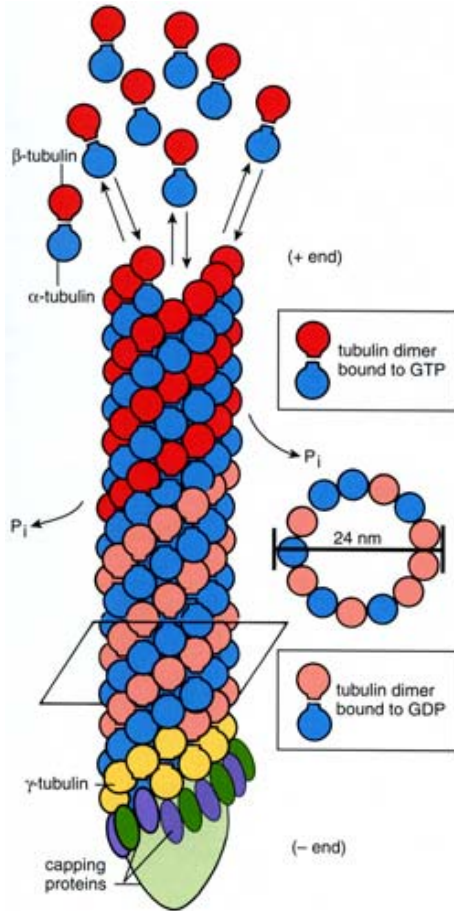
tubulinové „háčky“ v izol. fragmoplastech v nadbytku tubulinu

# Nukleace mikrotubulů

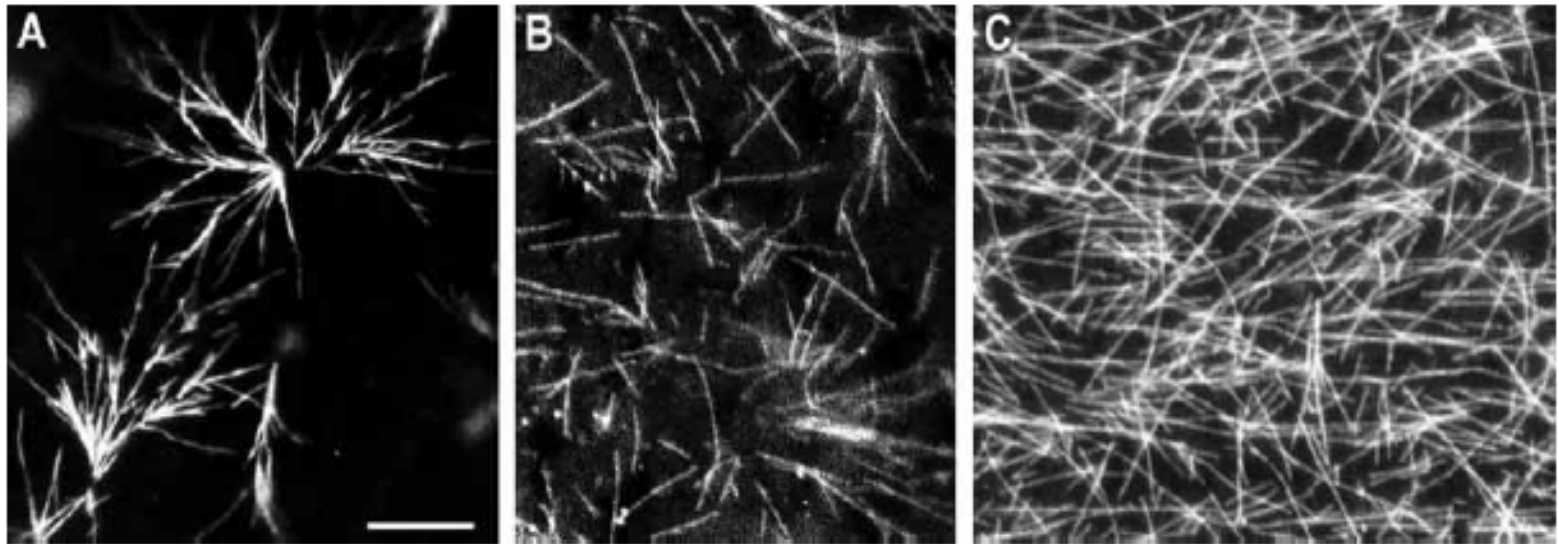
$\gamma$ -tubulin

mikrotubulární  
organizační centra  
(MTOC)

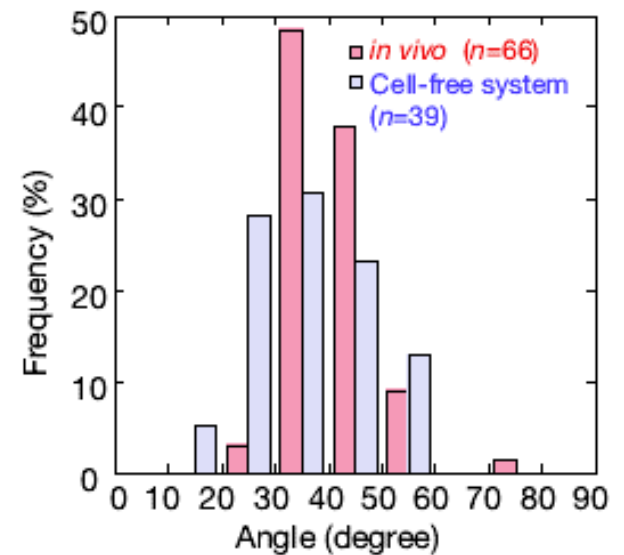
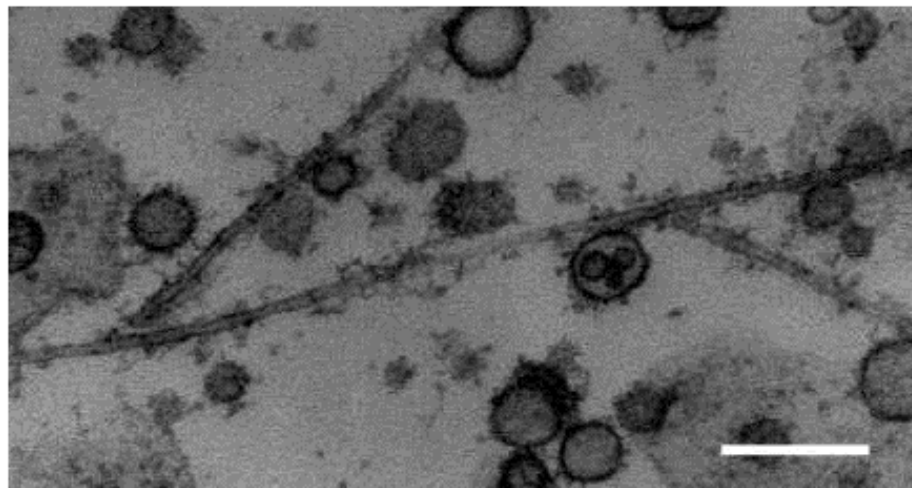
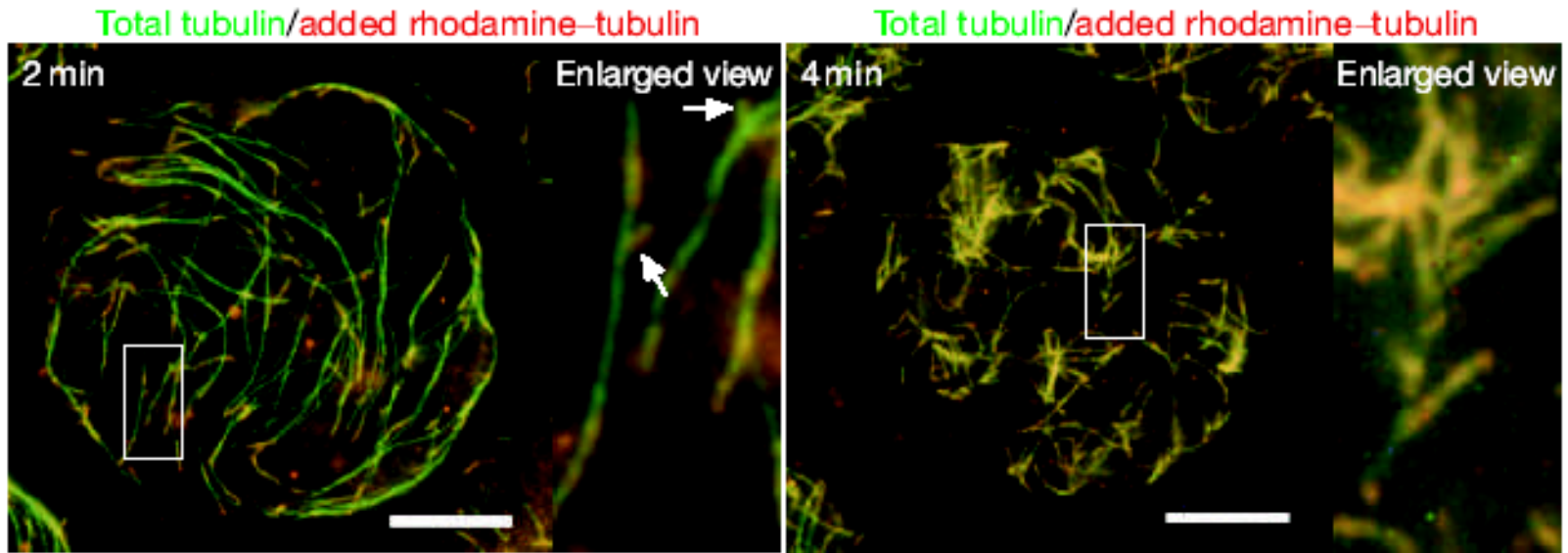
rostliny nemají žádná  
„definovaná“ MTOC



## MT recovery after drug-induced disassembly



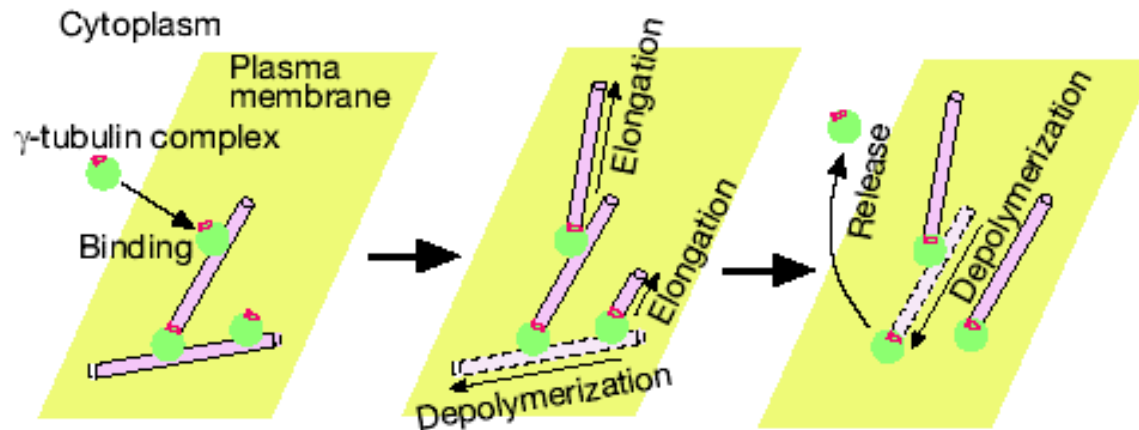
**Fig. 2.** Cortical microtubule recovery patterns after drug-induced microtubule disassembly. (A) Microtubules appear to diverge from the initial assembly site, forming fractal tree-shaped clusters, with microtubules diverging from each other at acute angles (figure adapted from Wasteney and Williamson, 1989b). (B) Clusters eventually break up. (C) Later in recovery, parallel microtubule order begins to consolidate but some branching configurations and discordant microtubules persist. Bar, 10  $\mu\text{m}$ .



Tobacco cell ghosts; Murata et al., Nat.Cell.Biol. Oct. 2005

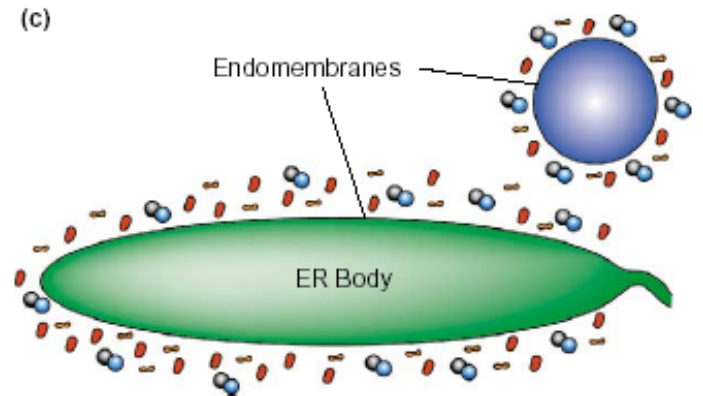
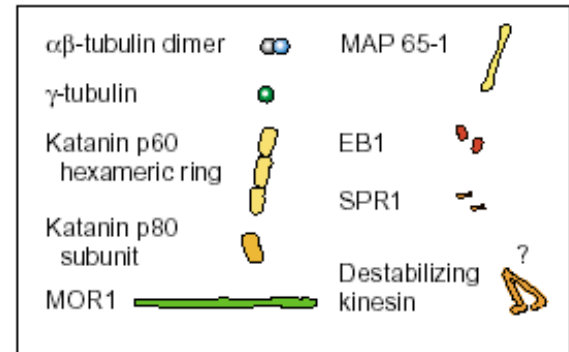
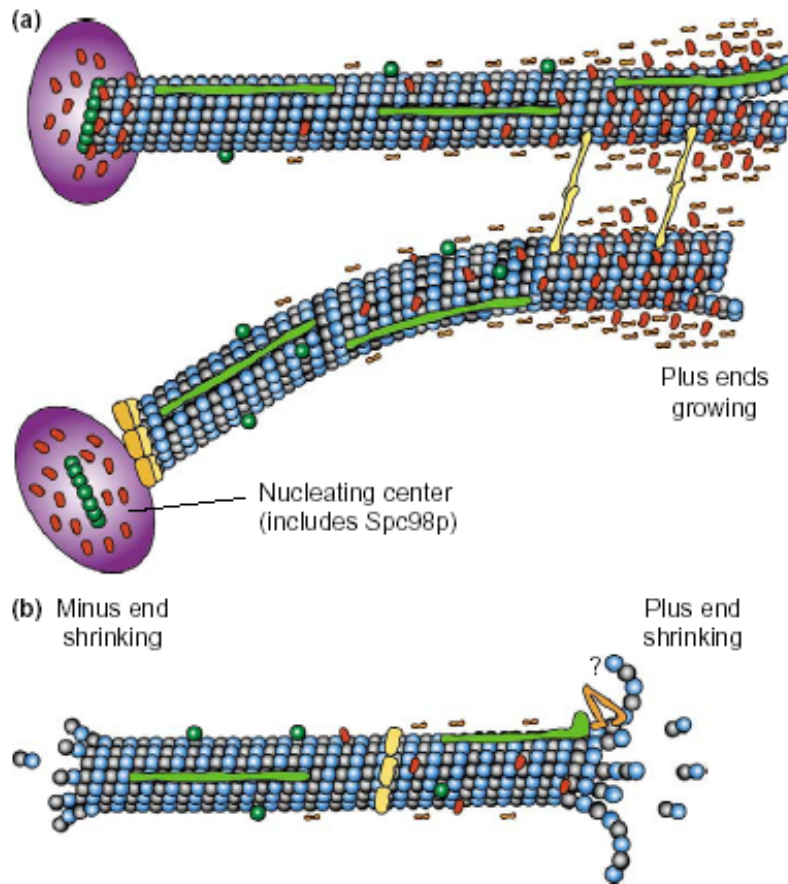


# Nukleace $\gamma$ -tubulinem - model



**Figure 5** Model for nucleation of cortical microtubules in plants. Cytosolic  $\gamma$ -tubulin complexes bind onto pre-existing cortical microtubules (left) and nucleate microtubules as branches (centre). The original microtubules frequently depolymerize (centre) but  $\gamma$ -tubulin complexes are not released until depolymerization of newly formed microtubules occurs (right).

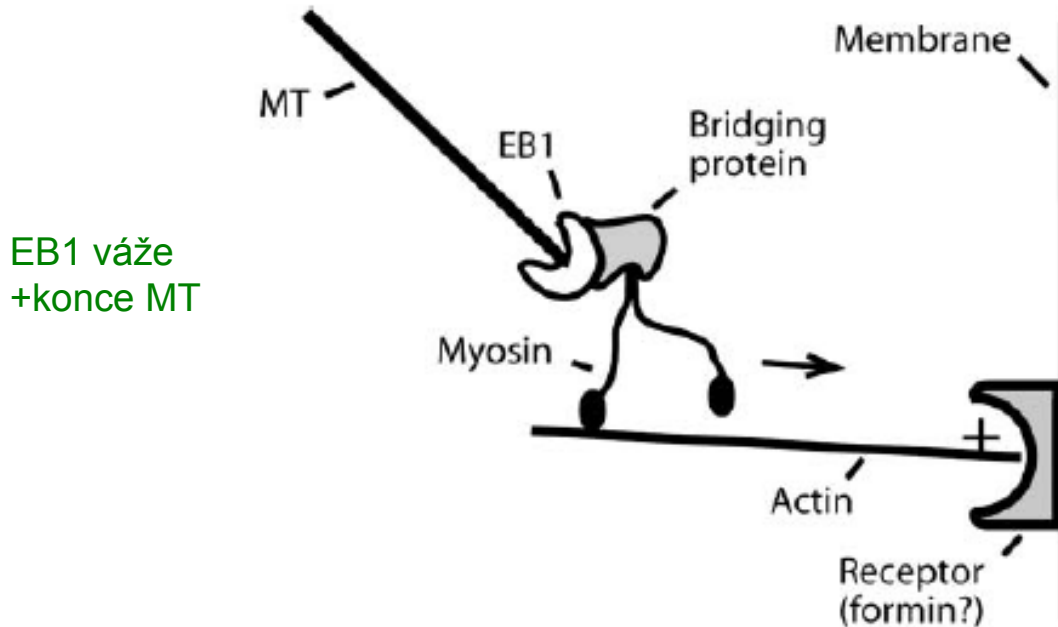
# Modulace dynamiky mikrotubulů - MAPs



**MAPS predicted to affect microtubule dynamics and organization.**

Predicted MAP (species)	Family size	Intracellular localization	Mutant name	Mutant phenotype	Predicted cellular function
AtEB1a, AtEB1b ( <i>Arabidopsis</i> )	3	Cortical MT plus ends, spindle poles, cortical MT nucleating sites, around endomembranes?	ND	ND	MT polymerization, recruiting MAPs to MTs
Katanin P60 subunit (AtKSS, AtKTN1) ( <i>Arabidopsis</i> )	1 <sup>a</sup>	Spindle poles; cortical MT nucleating sites, punctate along cortical MTs, perinuclear regions	<i>botero, fra2, frc2, erh3, lue1</i>	Disorganized cortical MTs, isotropic growth, cell wall composition/organization, ectopic root hairs, gibberellin signaling, trichome branching	ATP-dependent MT severing
Katanin P80 subunit ( <i>Arabidopsis</i> )	4	ND (presumed to colocalize with katanin P60)	ND	ND	Targeting and regulation of katanin P60?
MAP-65 (main isoform in carrot suspensions)	ND	Along MTs in interphase cells	ND	ND	Cortical MT bundling and organization in interphase cells
NtMAP65-1b (tobacco)	ND	ND (presumed to be along MTs)	ND	ND	MT bundling and organization, stabilize MTs against cold
AtMAP65-1 ( <i>Arabidopsis</i> )	9	Along MTs in preprophase band, at the phragmoplast midline, along a subset of cortical MTs	ND	ND	Crosslinking MTs
AtMAP65-3 (PLEIADE) ( <i>Arabidopsis</i> )	9	Along MTs in preprophase band, at phragmoplast midline where antiparallel MTs overlap	<i>ple</i>	Cytokinesis defect in roots, distorted phragmoplasts	Crosslinks MTs and/or promotes MT polymerization in phragmoplast

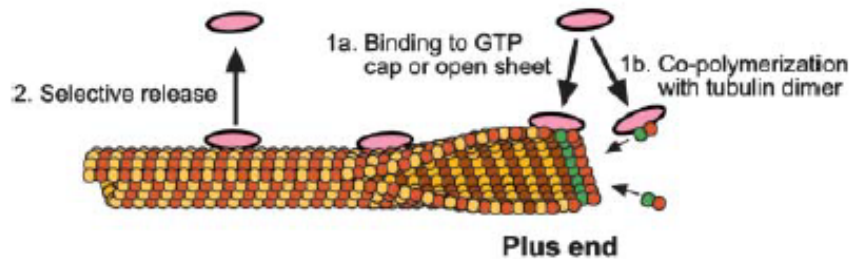
# End binding protein (EB1)



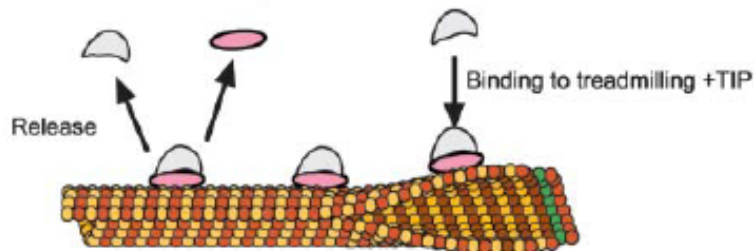
**Figure 2.** Hypothetical model for delivery of a plant MT to a cellular receptor at a specific site (e.g. PPB, phragmoplast, root hair tip) along actin filaments, based on models from yeast and fibroblasts (Gundersen et al., 2004). EB1 binds a bridging protein associated with myosin, which translocates toward the barbed (plus) end of the actin filament. Genome analysis identifies EB1, myosins, and formins in Arabidopsis.

# Jak se proteiny mohou dostat na +konec MT?

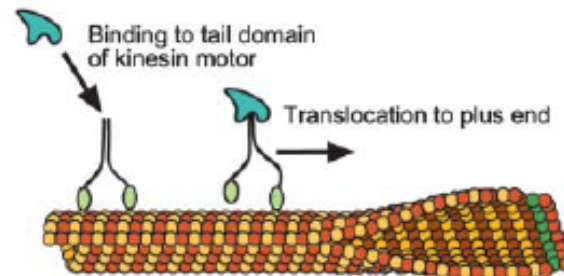
## A Treadmilling



## B Hitchhiking

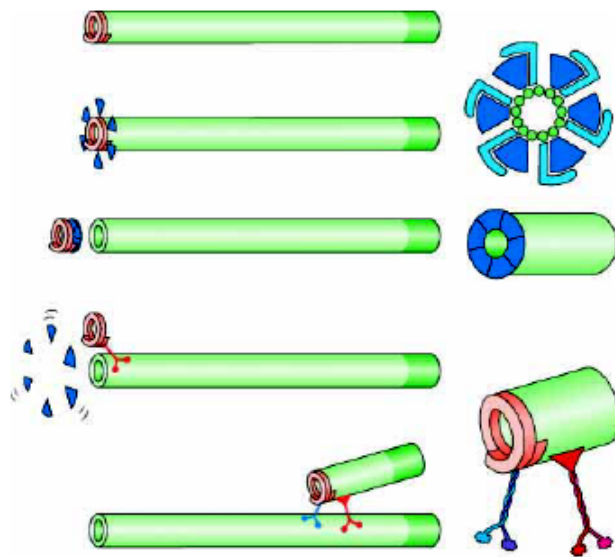


## C Motor-driven transport

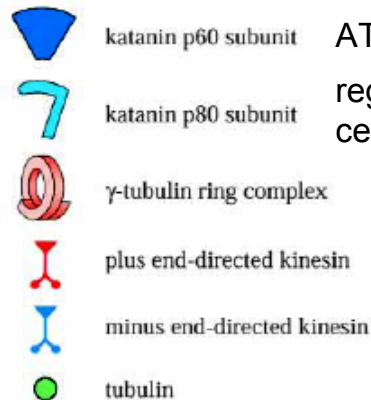


**Figure 1.** Proposed mechanisms for plus-end binding. Green circles at the plus end of the MT denote  $\alpha$ -tubulin bound to GTP. See text for details.

# Katanin - p60 a p80



**Fig. 3.** Model for microtubule assembly by severing and transport of nucleating templates. In this model, a  $\gamma$ -tubulin ring complex associates with the minus end of a microtubule, while the microtubule extends by the addition of tubulin subunits at the fast-growing, GTP-tubulin-containing plus end (dark green). Severing of the minus end is achieved by the formation of a hexamer of katanin p60 subunits, whose association with the microtubule wall is coordinated by the larger p80 subunit, which may transiently dimerize with the p60 subunits. Microtubule-mediated ATPase activity results in inward movement of the p60 subunits, an action that cleaves the ring complex from the microtubule minus end. Katanin subunits dissociate but the lock-washer-shaped ring complex is transported along the microtubule by a plus-end-directed kinesin. The extent of transport along the microtubule may be regulated by the relative activities of plus- and minus-end-directed kinesins. The ring complex serves as a template for the assembly of additional microtubules. Repeated generation, severing and transport of nucleating templates at the minus end of the original microtubule may explain how the fractal tree complexes shown in Fig. 2A develop.



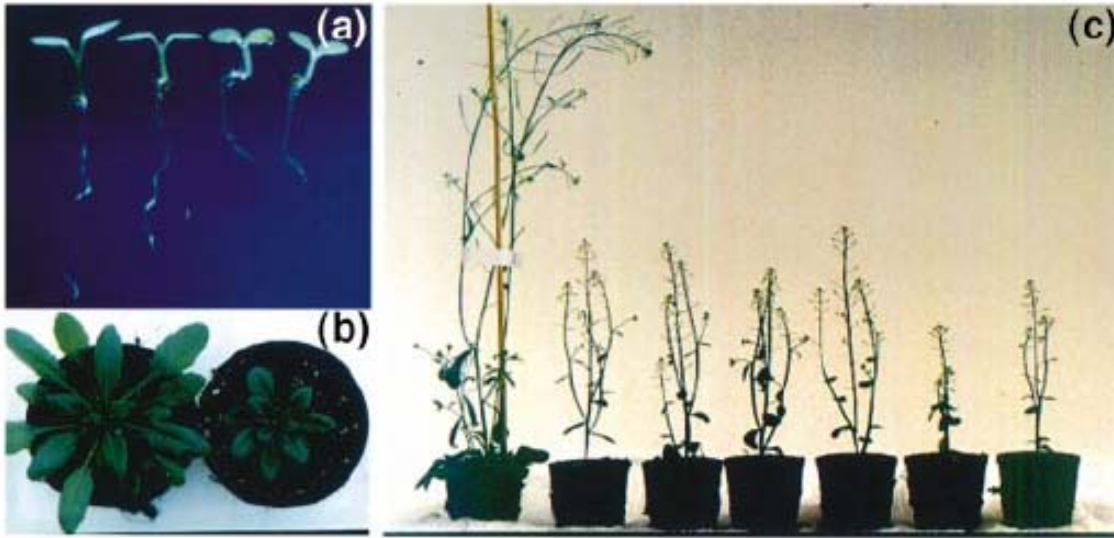
ATPase subunit, which functions to sever microtubules

regulates the activity of the ATPase and localizes the protein to the centrosomes, not present in plants

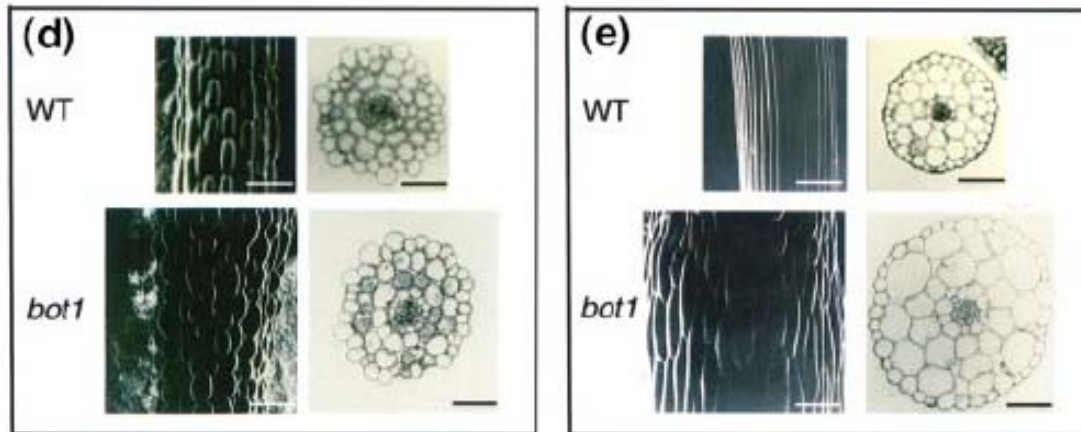
Mutace *botero*, *fragile fibre2*:

brittle swollen semi-dwarf, disordered cortical mt

# *botero1* mutant

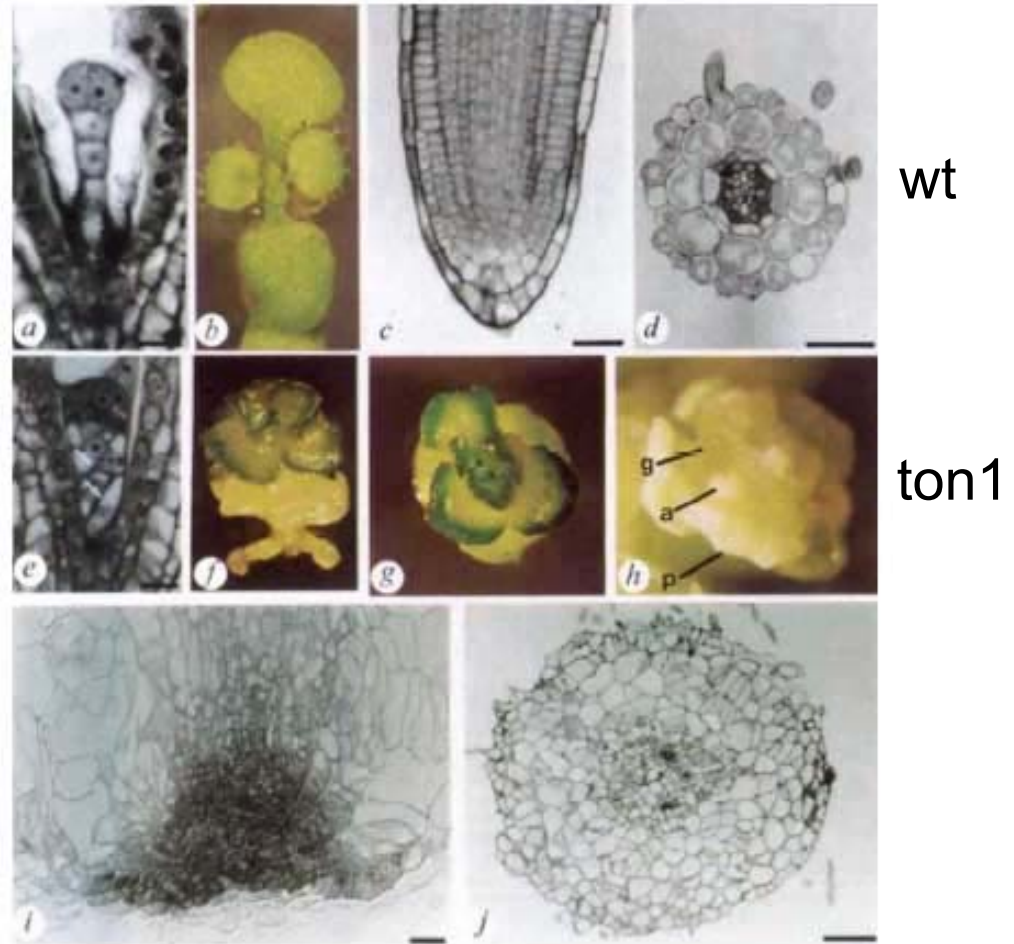


BOT1 is required for organizing cortical microtubules into transverse arrays in interphase cells, and that this organization is required for consolidating, rather than initiating, changes in the direction of cell expansion.



# TON1 (*tonneau*)

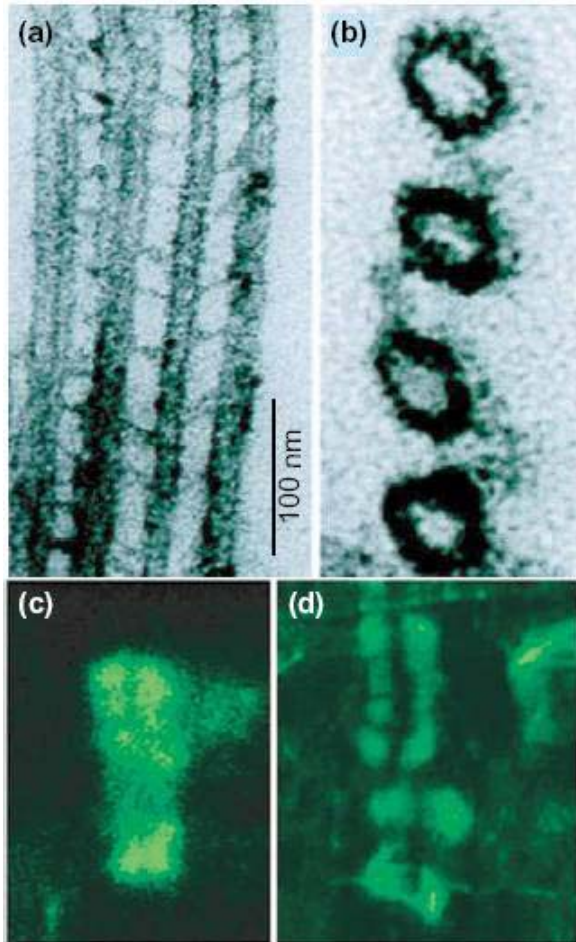
- defekt PPB a organizace MT
- *homolog LISSENCEPHALY1* (H.s. LIS1) - defekt migrace neuronů
- LIS1 váže +konce MT, interakce s dyneinem
- ale rostliny nemají dynein



(Traas et al. 1995)



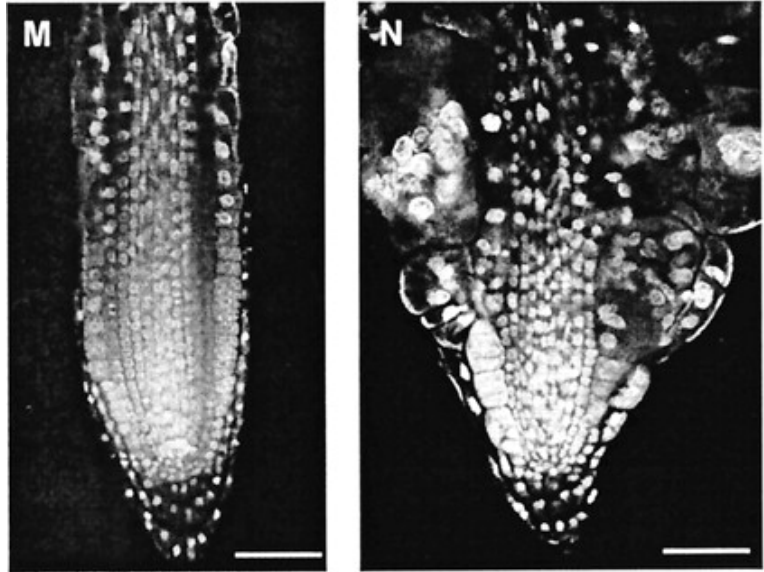
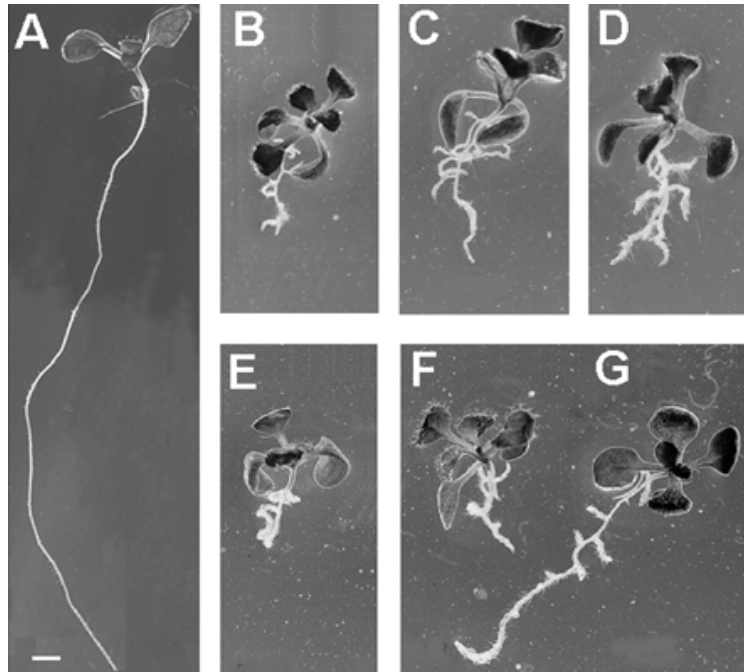
# MAP65: crosslinking („rostlinná specialita“)



MAP-65 family members that cross-link MTs. **(a)** Longitudinal and **(b)** transverse electron micrograph sections of MTs cross-linked *in vitro* by purified carrot MAP-65. MAP-65 can be seen as the evenly spaced filamentous cross-bridges (reproduced with permission from [46], copyright National Academy of Sciences, USA [1999]). **(c)** Wildtype and **(d)** *ple-6* phragmoplasts visualized with MAP4-GFP in *Arabidopsis* (confocal microscopy). Note the larger clear zone in *ple-6*, which is mutated in the *AtMAP65-3* gene (reproduced with permission from [51\*\*], copyright Cell Press [2004]).

Arabidopsis má 9 homologů MAP65

# *pleiade* – mutace v MAP65

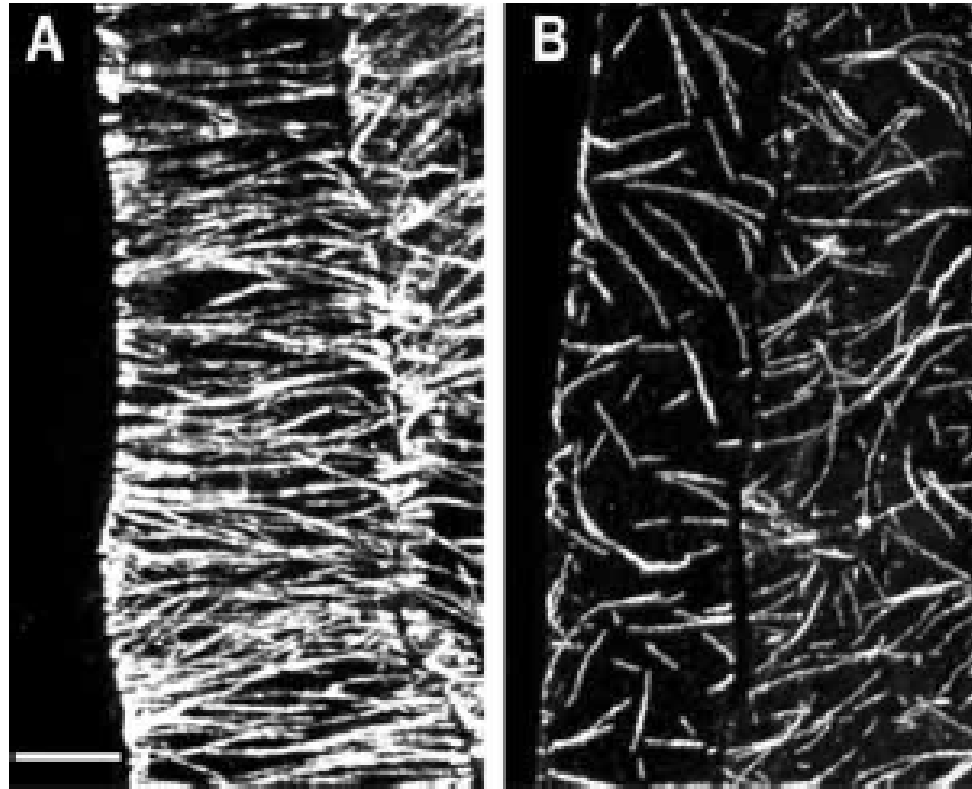


mnohojaderné buňky  
(kořenově specifické!)

(Muller et al. 2002)

MOR1 ( <i>Arabidopsis</i> )	1	Punctate along cortical MTs, at phragmoplast midline' spindle	<i>mor1, gem1</i>	<i>mor1</i> alleles are temperature sensitive with shortened/disorganized MTs and isotropic cell expansion, <i>gem1</i> alleles have cytokinesis defects and are homozygous lethal	Stabilize MTs, MT organization
TMBP200 (MOR1-like) (tobacco)	ND	ND	ND	ND	MT bundling?
p90 PHOSPHOLIPASE D (tobacco)	ND	Along cortical MTs and linking them to the plasma membrane	ND	ND	Conveys hormonal and environmental signals to cortical MTs?
PHOSPHOLIPASE D $\delta$ (PLD $\delta$ ) ( <i>Arabidopsis</i> )	12	Along cortical MTs and linking them to the plasma membrane?	PLD $\delta$ null	Increased sensitivity to H <sub>2</sub> O <sub>2</sub> -induced cell death	Oleate stimulated phospholipase activity linked to H <sub>2</sub> O <sub>2</sub> signaling
Spc98p ( <i>Arabidopsis</i> )	1 <sup>a</sup>	Punctate along nuclear surface and at cortex (presumably at MT nucleating centers)	ND	ND	MT nucleation?
SPIRAL1 (SPR1) ( <i>Arabidopsis</i> ) <sup>b</sup>	6	Cortical MT plus ends and uniformly covers MTs in other arrays, around endomembranes?	<i>spr1, sku6</i>	Organ axial twisting, root right skewing on agar surfaces, obliquely oriented cortical MTs	MT polymerization? Links proteins to MTs? Directional cell expansion
TANGLED1 (TAN1) (maize)	ND	Punctate throughout cytoplasm and along MTs in all four arrays	<i>tan1</i>	Abnormally oriented cell divisions	Orienting MT structures during cell division
WAVE-DAMPENED2 (WVD2) ( <i>Arabidopsis</i> )	8	Along MTs?	<i>wvd2</i>	Overexpression causes twisted, stockier organs, right-skewing of roots on agar surfaces and obliquely oriented cortical MTs	MT bundling?

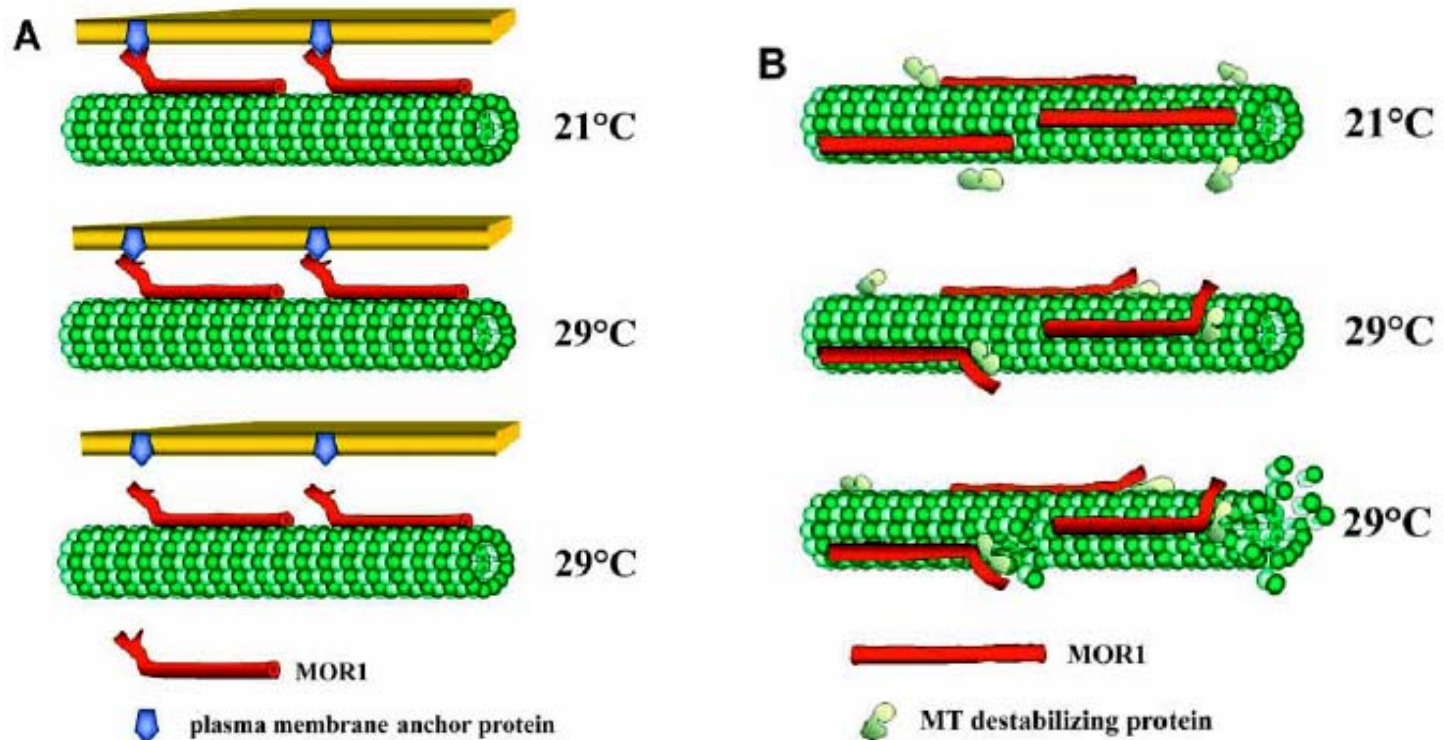
# MOR1 (microtubule organization)



(MAP215)

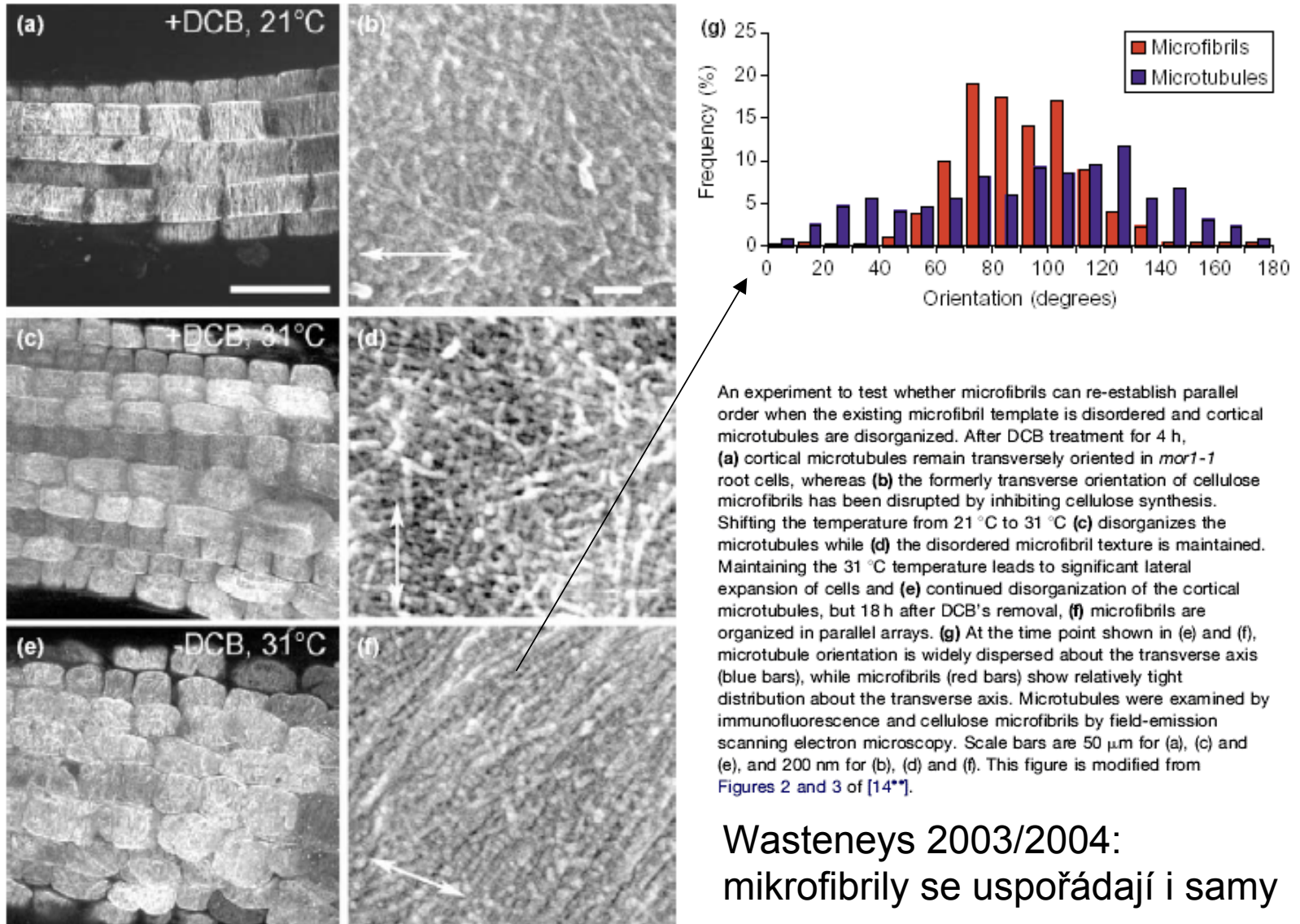
**Fig. 4.** Microtubule patterns in the epidermis of *Arabidopsis thaliana* cotyledons after 4 hours at 29°C. (A) Cortical microtubules are abundant and transversely oriented in wildtype. (B) In the *mor1* mutant, microtubules appear short and disoriented. Bar, 10  $\mu$ m.

# Modely funkce MOR1

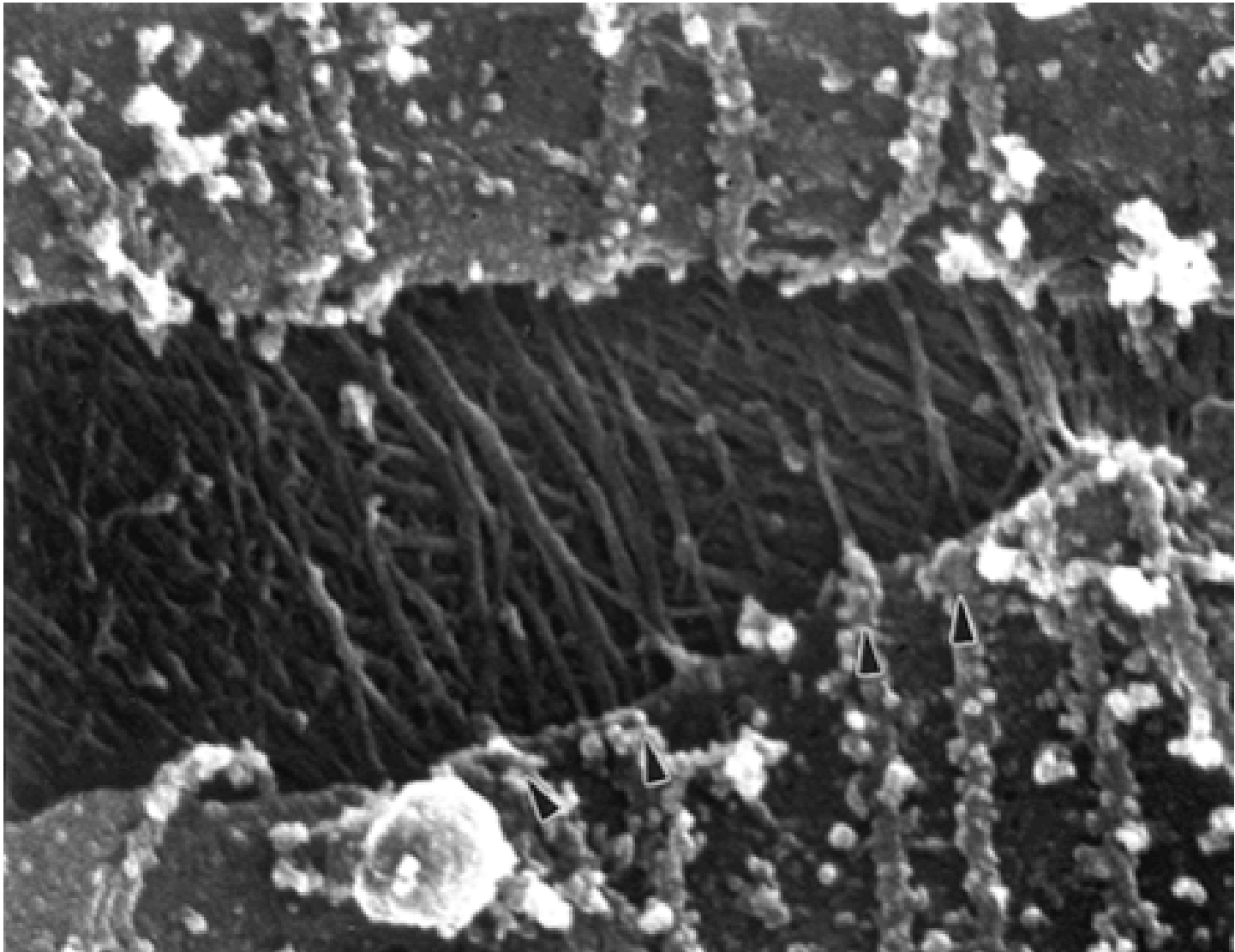


**Fig. 5.** Possible functions of the MOR1 HEAT repeat-1 (HR1) in microtubule stabilization. (A) HR1 links microtubules to the plasma membrane via a plasma-membrane-associated protein. At restrictive temperature, this loss of binding dissociates microtubules from the plasma membrane, promoting their destabilization. (B) HR1 competes with a destabilizing protein (probably a kin1-like kinesin) for binding. At permissive temperature, the high affinity of MOR1 for this site prevents destabilization. At 29°C, this affinity is lost, leading to kin1-dependent destabilization and microtubule shortening.

# MOR1

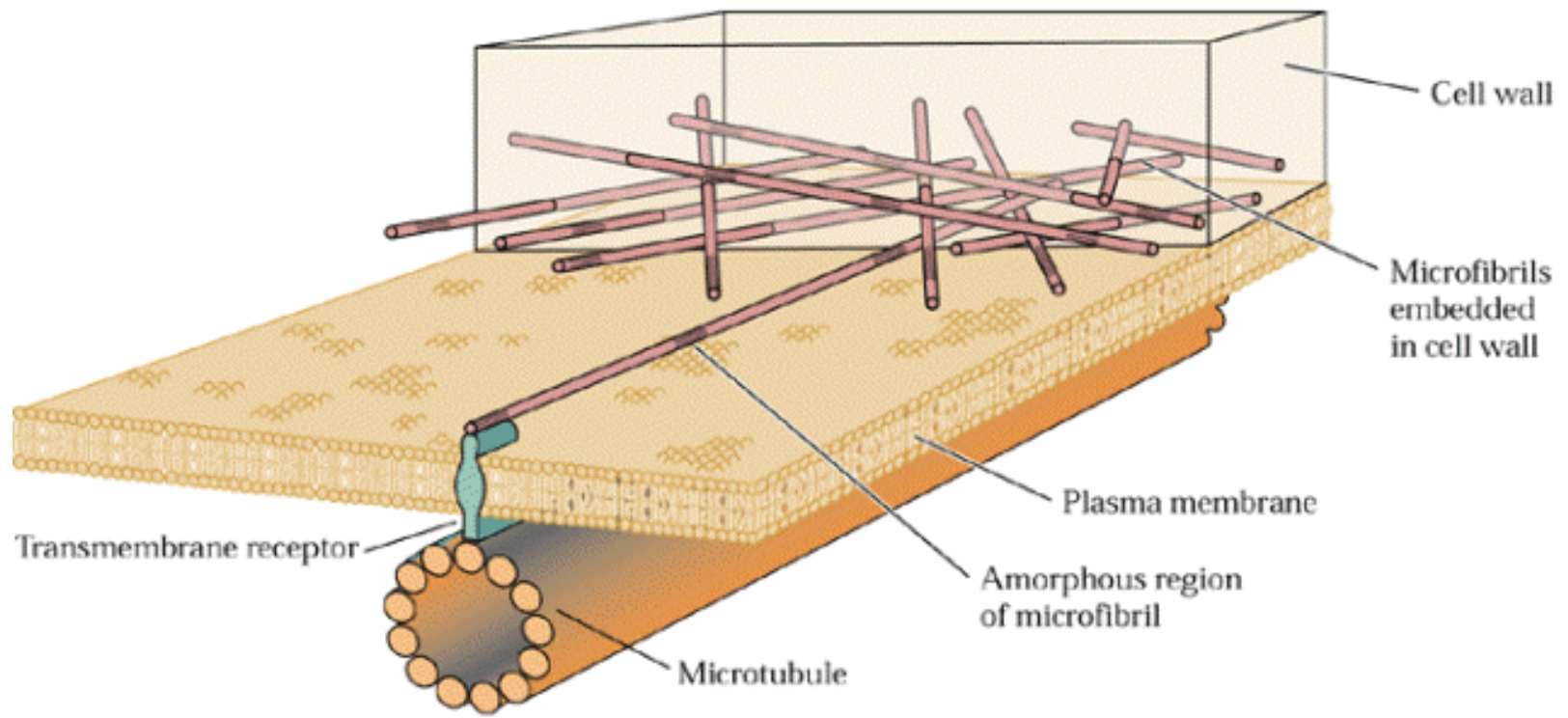


# Asociace mezi kortikálními mt a stěnou



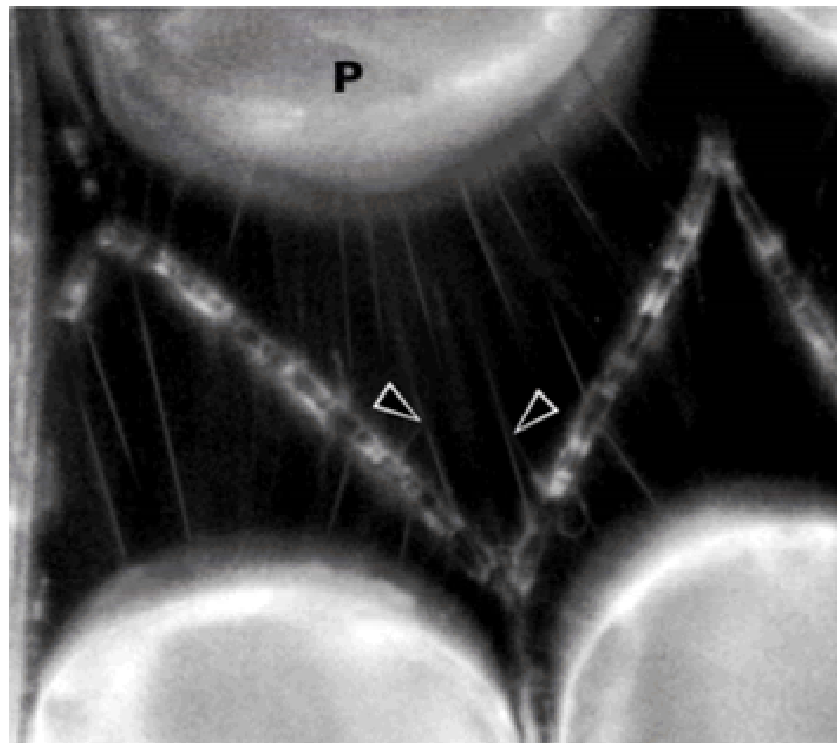
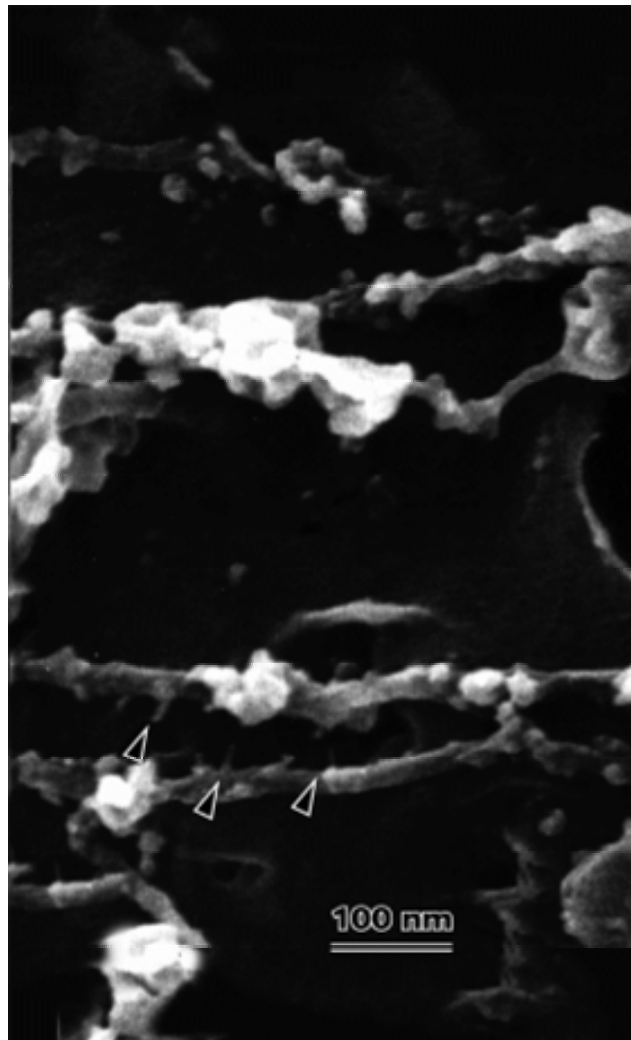
onion root

# Asociace mezi kortikálními MT a stěnou



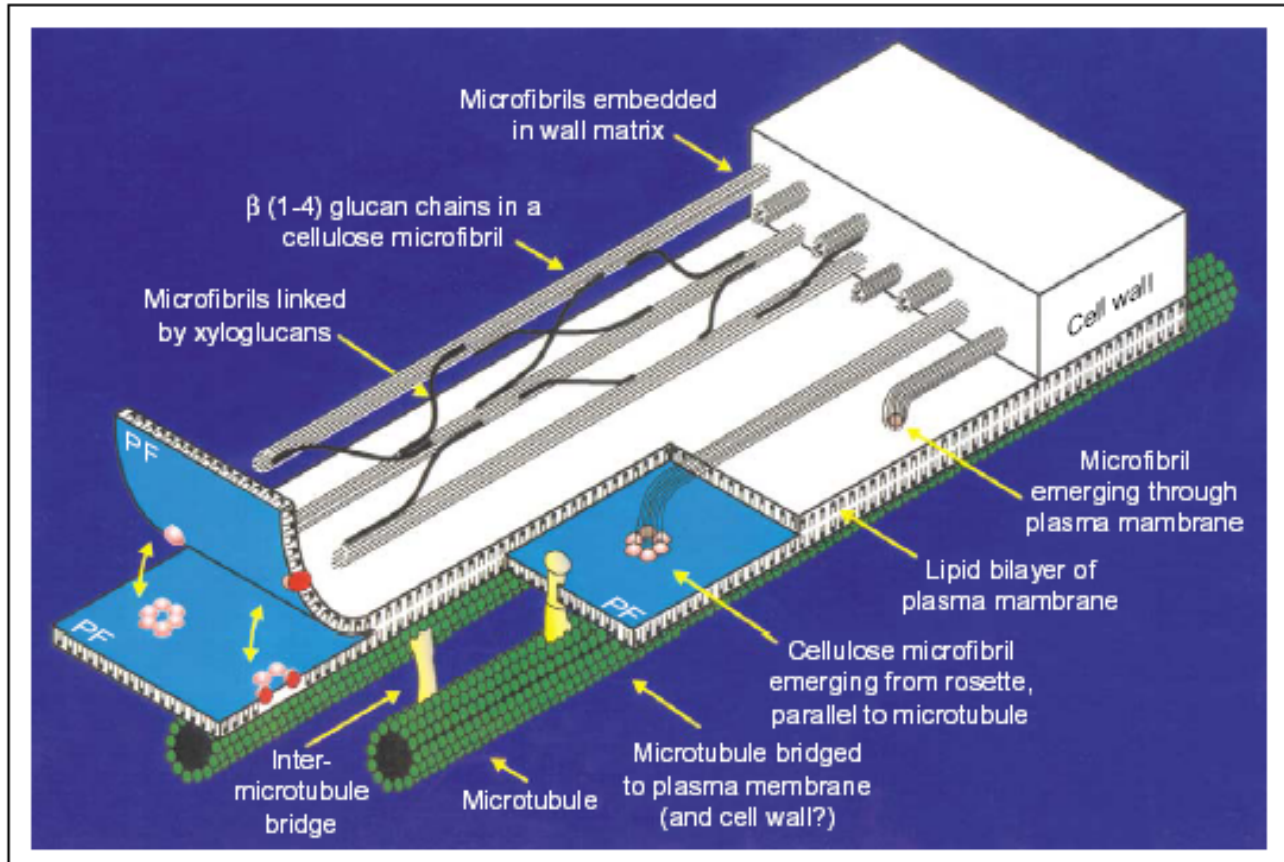


## Interakce MT a buněčné stěny



Hechtovy provazce

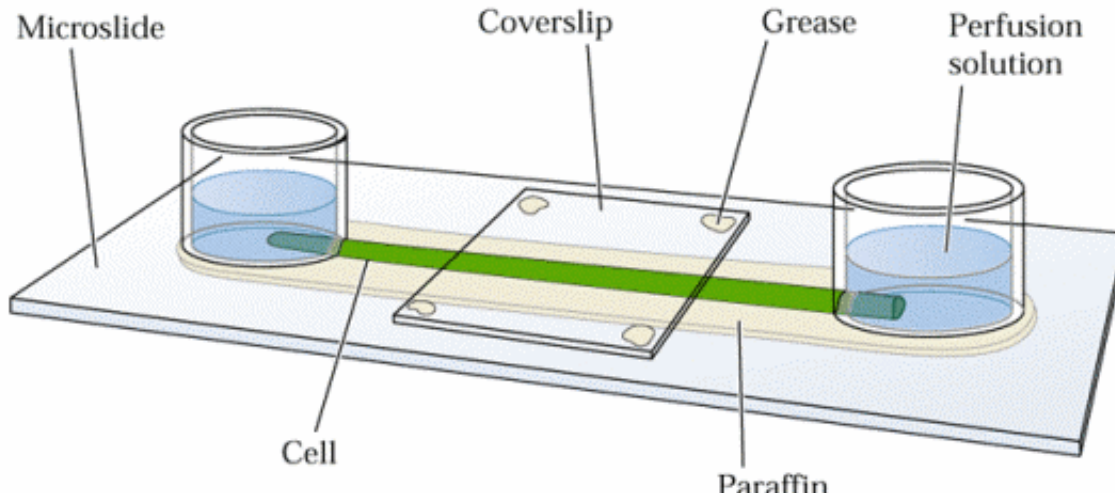
cibule



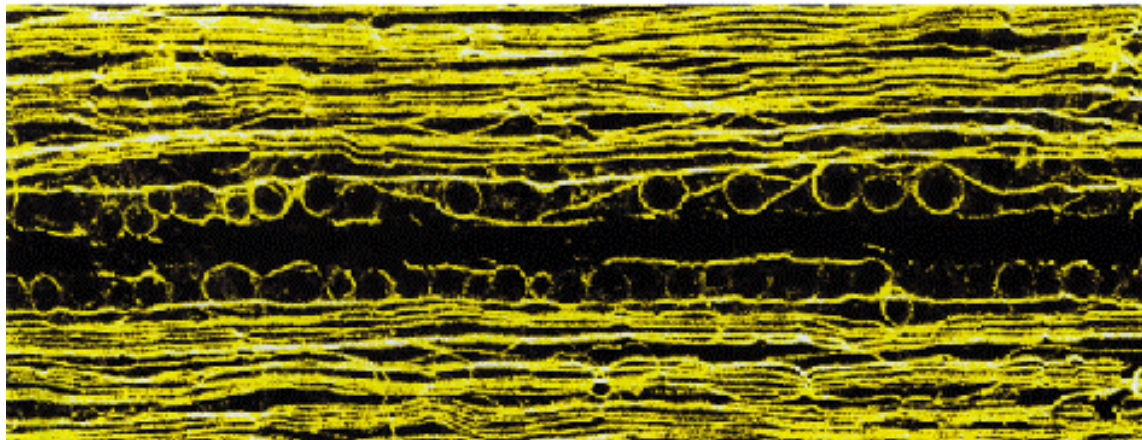
Relationships between cellulose-synthesizing complexes (rosette type), wall microfibrils, plasma membrane and microtubules. Diagram provided by Brian Gunning.

# Chara: jak se to měřilo

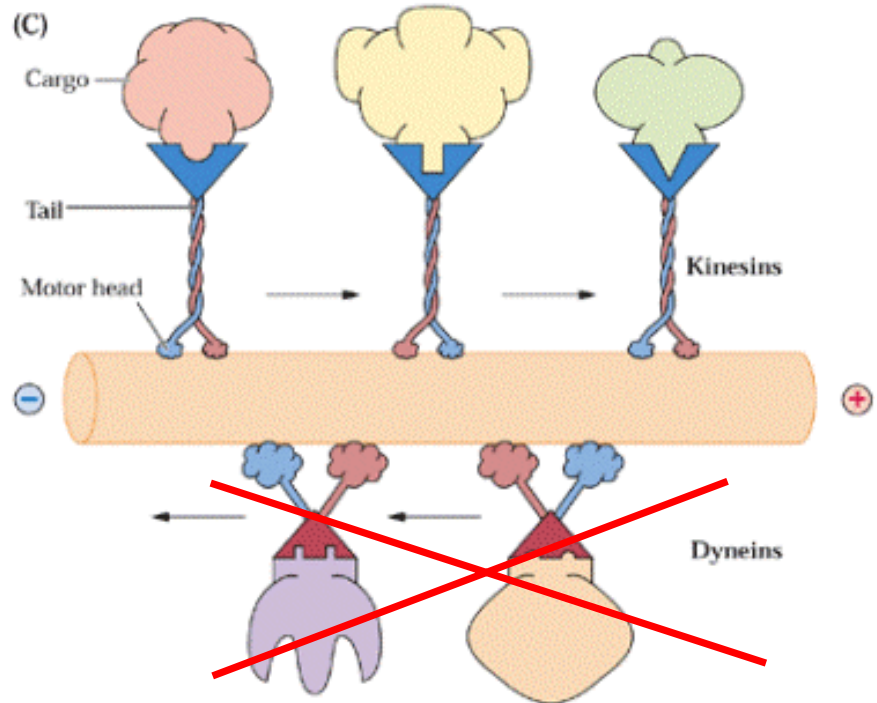
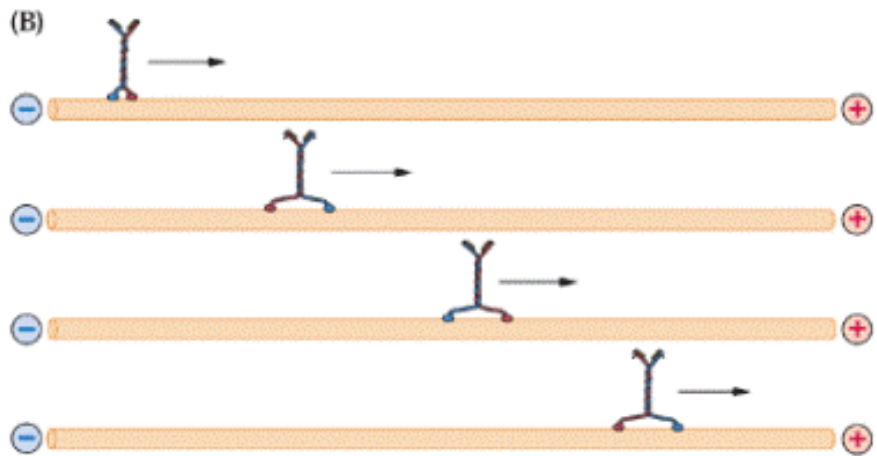
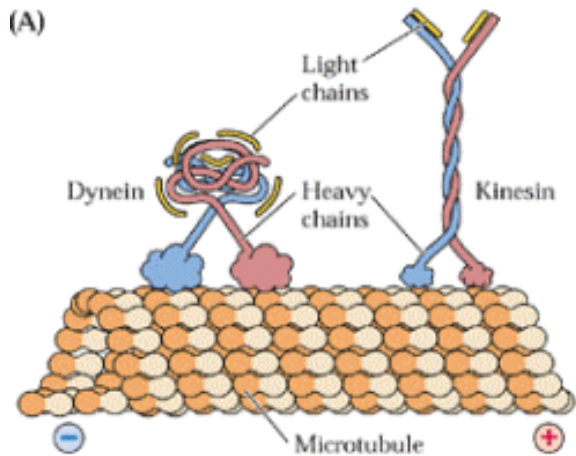
(A)



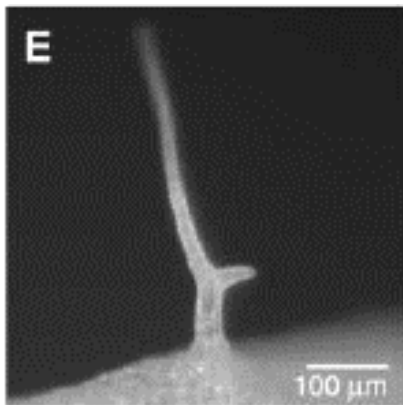
(D)



(aktin)

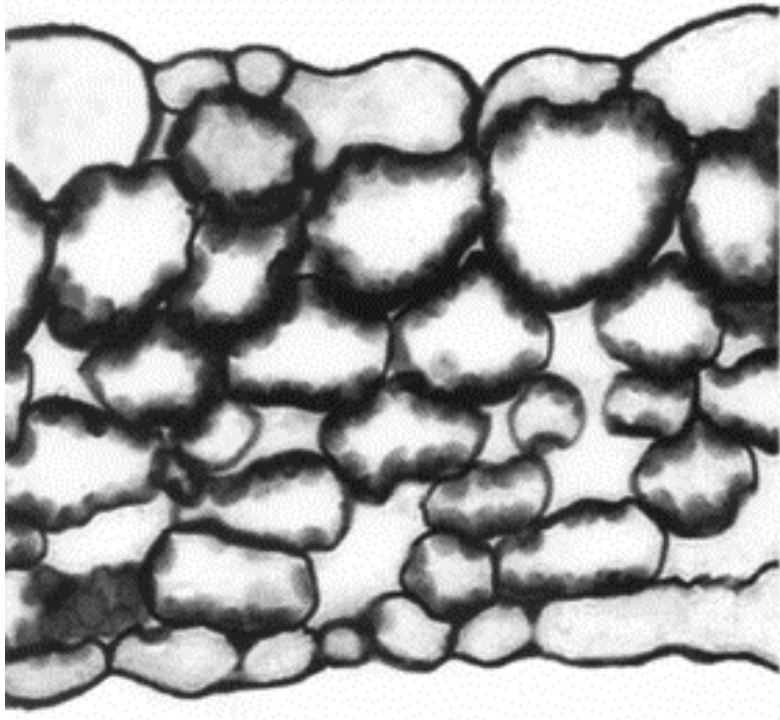


ZWICHEL kóduje kinesin!

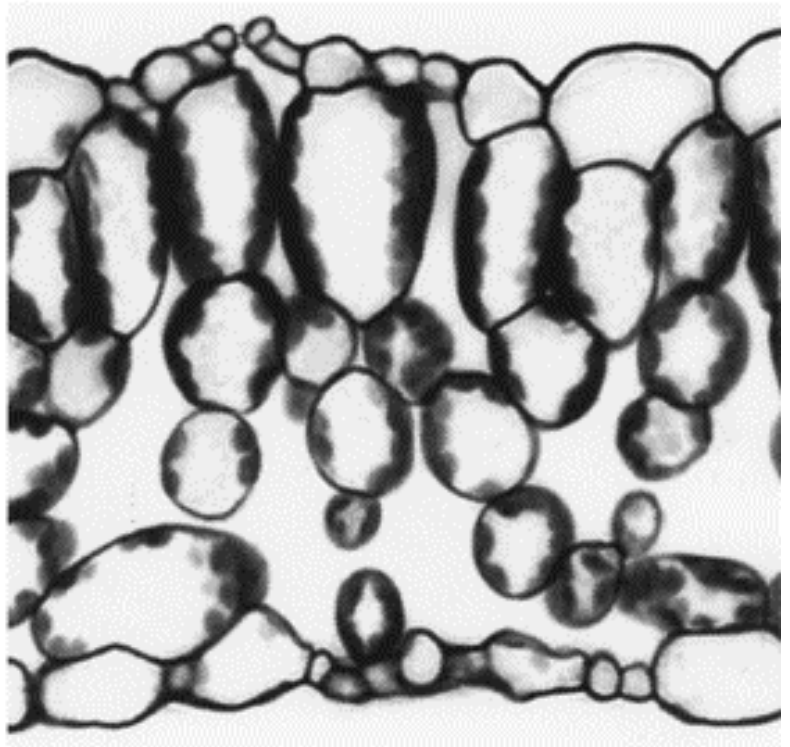


(jeden z mnoha!)

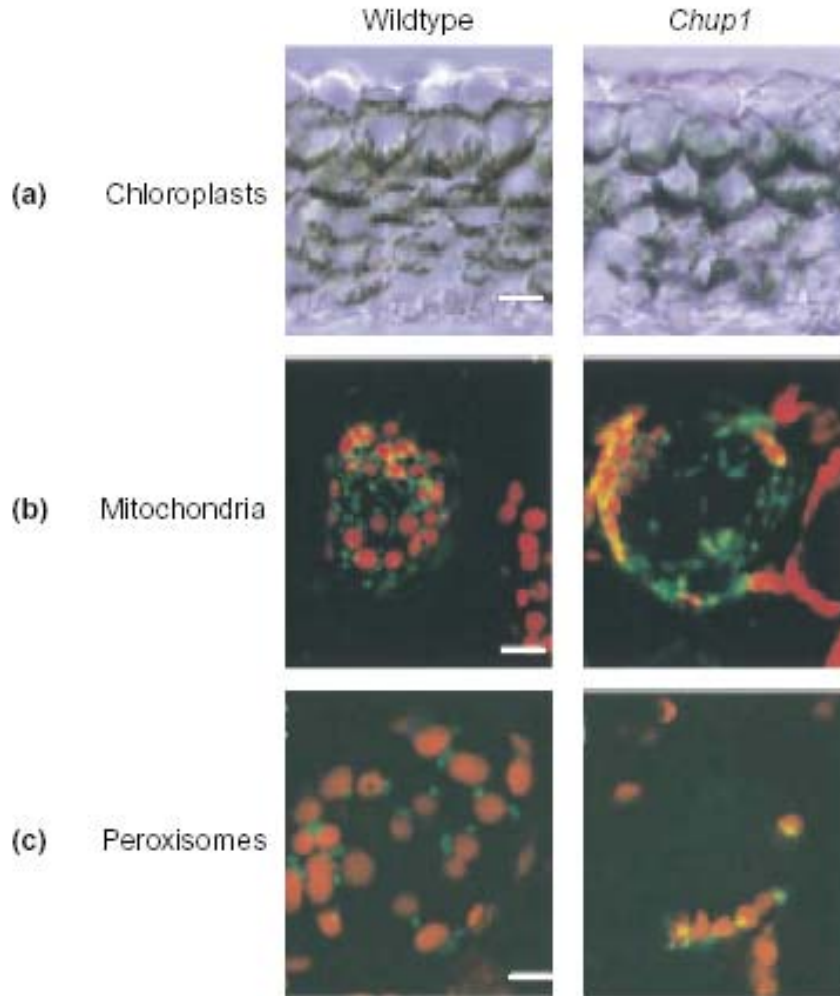
(A)



(B)

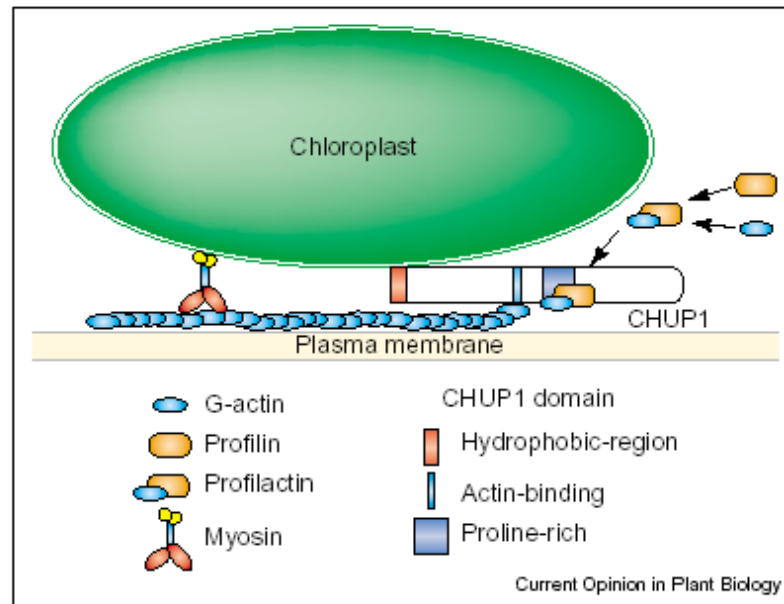


# Mutant *chup1* (chloroplast unusual positioning)



Distribution of chloroplasts, mitochondria and peroxisomes in wildtype and *chup1* mutants. **(a)** Cross-section of a light-adapted leaf. In the wildtype, chloroplasts are positioned on the upper and lower cell surfaces, whereas in the cells of *chup1* mutants, chloroplasts are aggregated on cell bottom. Bar represents 30  $\mu\text{m}$ . **(b)** Mitochondrial distribution viewed by transient expression of mitochondria-targeting GFP. The positioning of mitochondria is similar in wildtype and *chup1* mutants. Bar represents 10  $\mu\text{m}$ . **(c)** Peroxisomal positioning viewed by transient expression of peroxisome-targeting GFP. Peroxisomes are localized close to chloroplasts. In *chup1* mutant cells, peroxisomes are closely associated with aggregated chloroplasts. Bar represents 10  $\mu\text{m}$ .

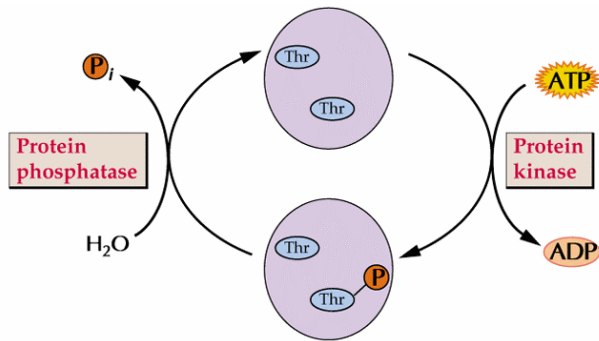
CHUP1: actin binding!



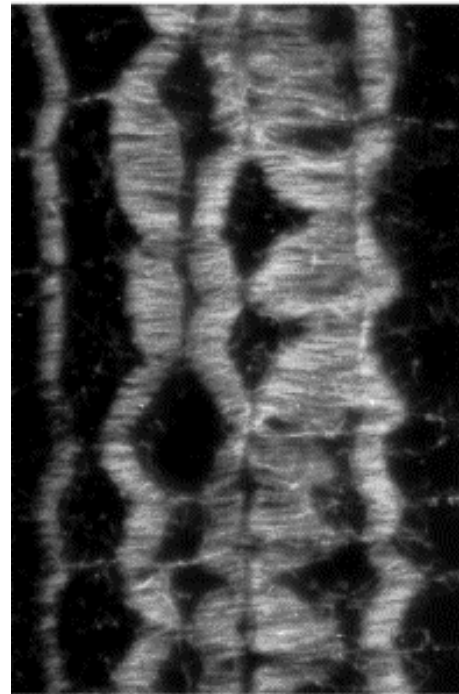
Working model of chloroplast positioning. The amino-terminal hydrophobic region of CHUP1 can target to the chloroplast outer membrane. The proline-rich motif of CHUP1 may serve in actin polymerization to recruit profilactin. CHUP1 binds polymerized F-actin through its actin-binding domain. These functions of CHUP1 may be important in anchoring chloroplasts to the plasma membrane. Myosin motor protein(s) may be necessary for chloroplast motility along actin filaments.

# Regulace struktury a funkce cytoskeletu

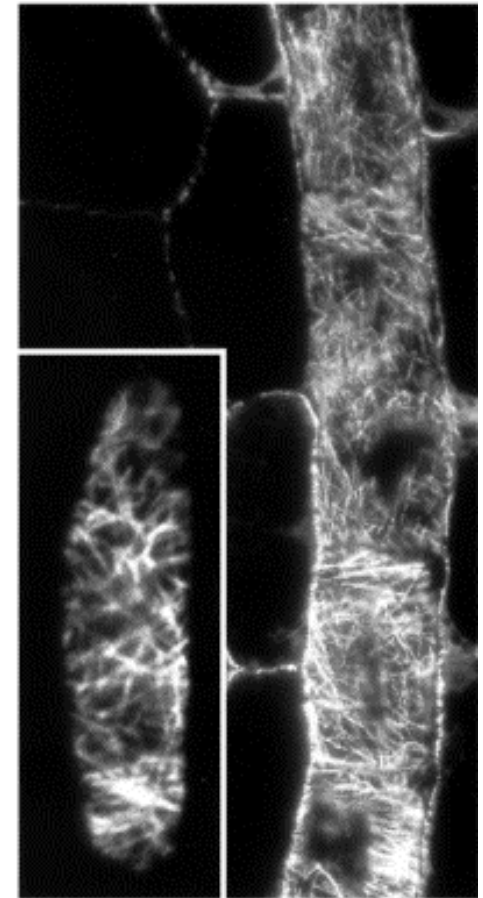
- Jako všude:  
proteinkinázy  
a fosfatázy



(A)



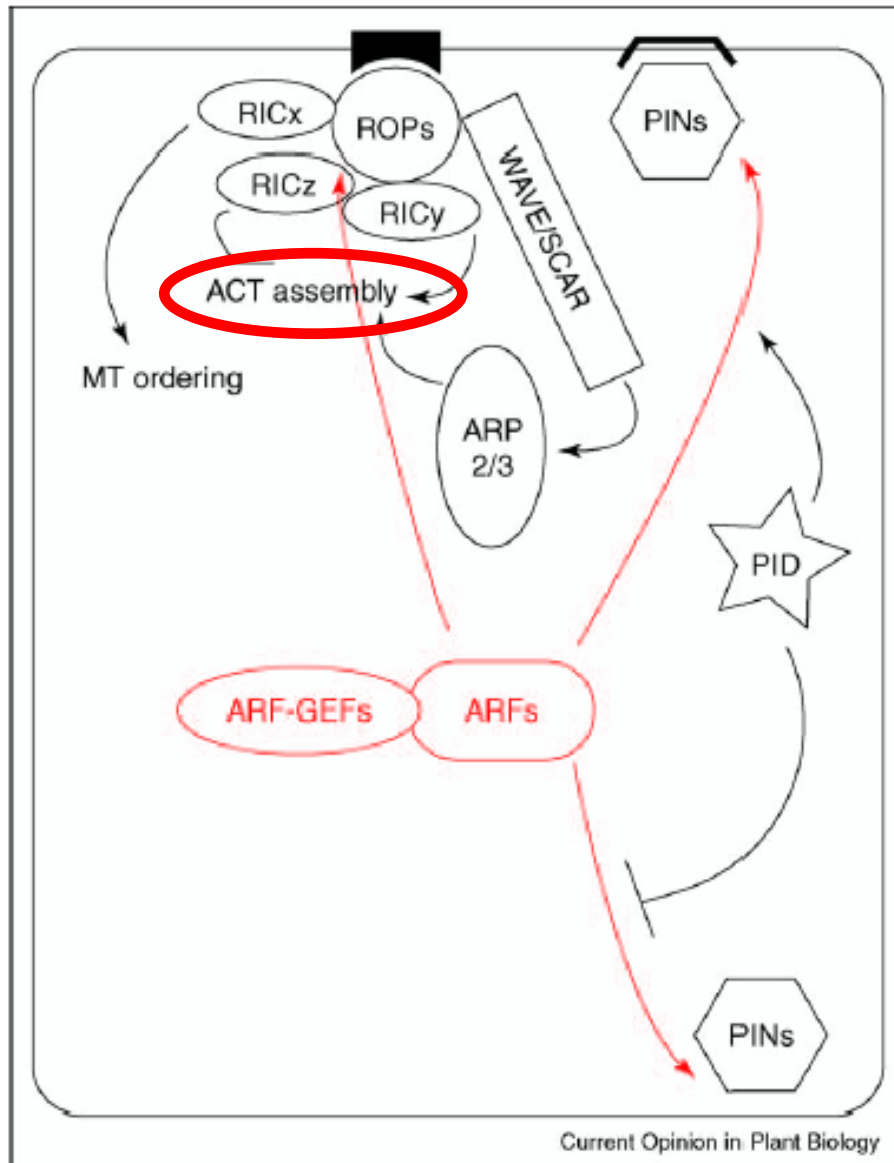
(B)



kořen A.th.  
po staurosporinu  
(mikrotubuly)



## Aktin: malé GTPázy (Rop, Arf)



Processes contributing to cell polarization. Vesicle trafficking that is mediated by Class 1 ARFs is required for the polar localization of ROP GTPases, which control actin (ACT) assembly through RICs and WAVE/SCAR-ARP2/3 pathways and microtubule (MT) bundling through other RICs. ARF-mediated vesicle trafficking and a specific ARF-GEF regulator of this process also control the localization of PIN proteins, and the polarity of this localization is controlled by the PID kinase, which functions as a binary switch. Polar localization cues for ROP localization or activation and for PIN localization (black bodies) remain unknown. Red arrows: vesicle trafficking control. Black arrows: protein activity control.

# Příklad: diferenciace epidermálních buněk

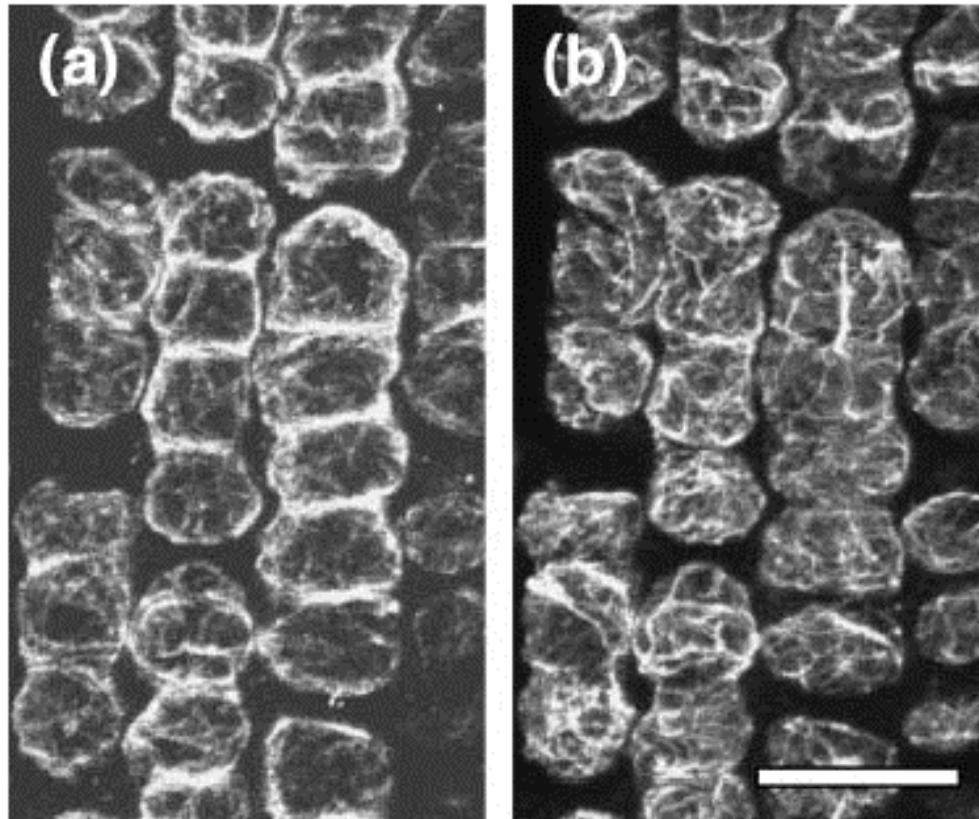
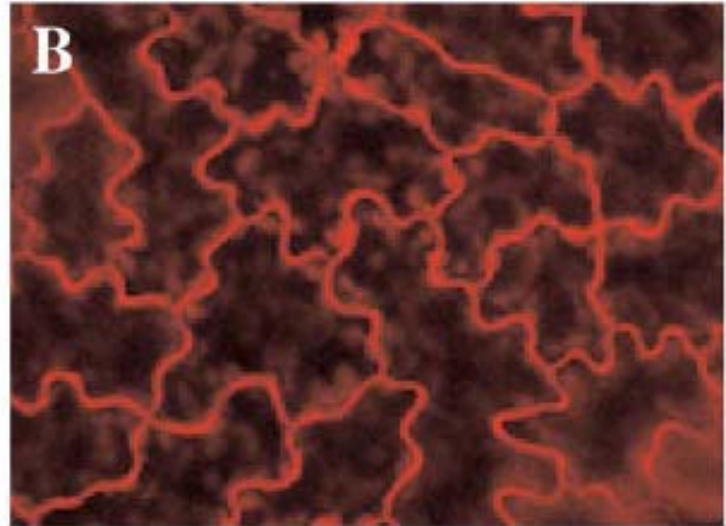
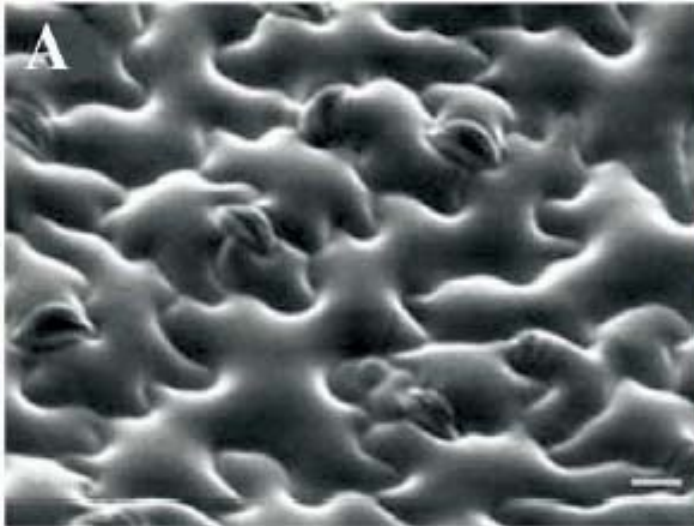


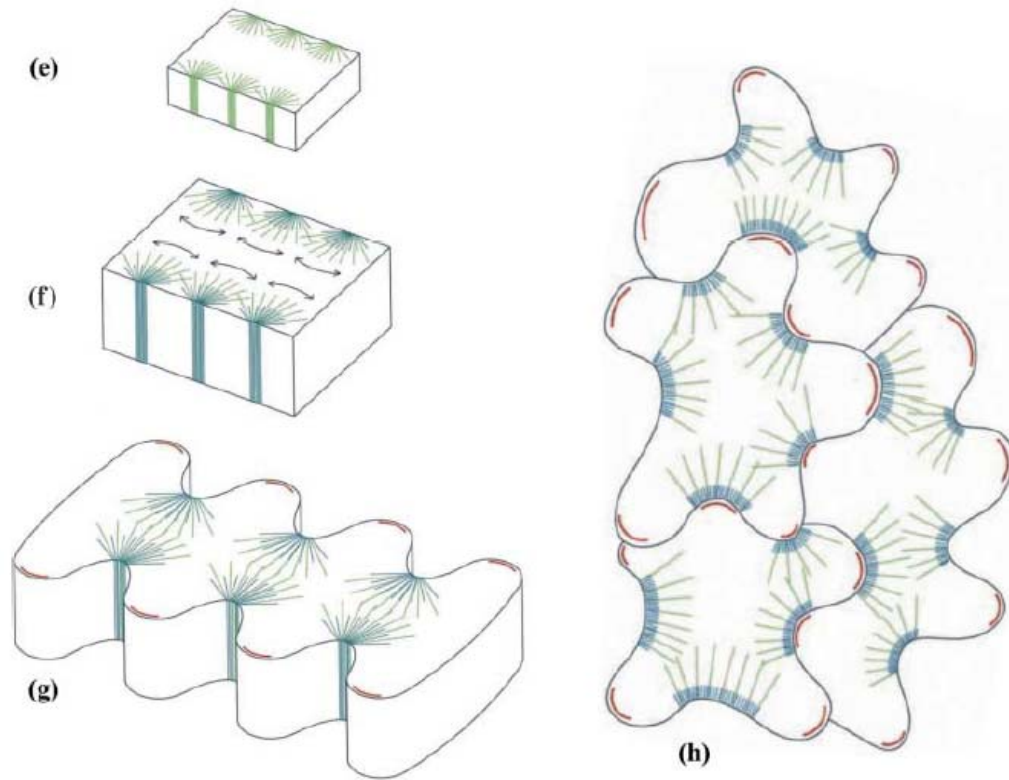
Fig. 1 Confocal-laser-scanning microscope (CLSM) images of lobed mesophyll cells of *Zea mays* after immunostaining of microtubules (MTs) (a) and visualization of actin filaments (AFs) with fluorescent phalloidin (b). Cortical MTs form ring-like bands (a), while AFs (b) exhibit diffuse arrangement. Bar, 20  $\mu$ m.

(Panteris a Galatis 2005)

# Regulační aspekty: ví se hlavně o aktinu

- *distorted* mutace (ARP2/3)
- *BRICK* (*Z. mays*)
- *SPIKE1* (multidomén. adaptor. protein)
- RICs/ROPs





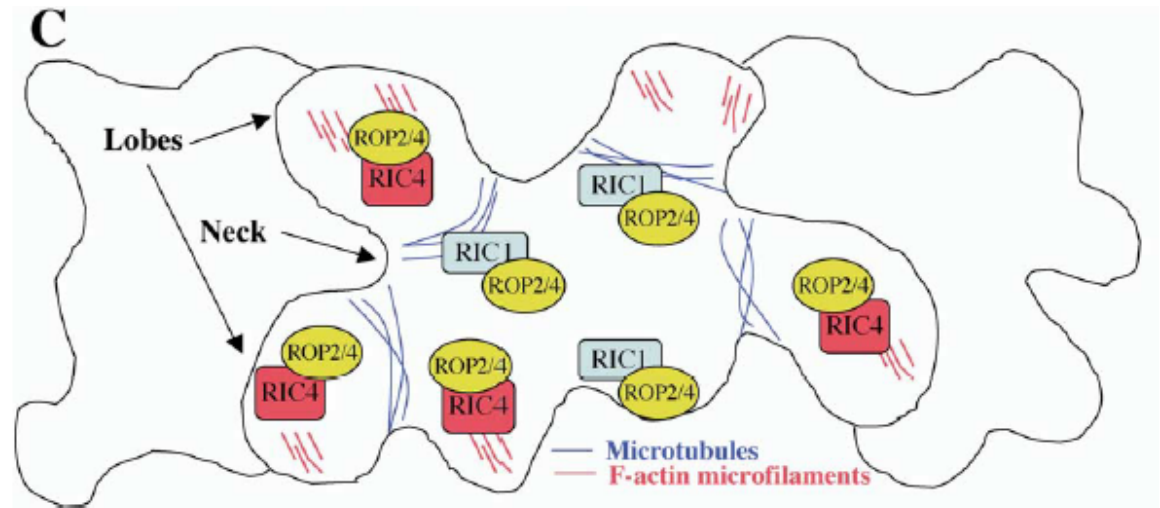
mikrotubuly  
 celulóz. mikrofibriily  
 aktin

(Panteris a Galatis 2005)

Fig. 3 Drawing presenting the successive stages of mesophyll (a–d) and pavement (e–h) cell morphogenesis. Microtubules (MTs) are shown in green, cellulose microfibrils (CMF) in blue and actin filament (AF) patches in red. In (a–d) cell wall thickenings, with CMFs parallel to MTs, are not shown. (a) Young mesophyll cell; the cortical MTs form interconnecting ring-like bundles. (b) As mesophyll cells grow, lobes and constrictions are created. New MT bundles also form. (c) Further mesophyll cell growth leads to formation of additional lobes. (d) Microtubule rings form at the base of growing mesophyll cell lobes, while AF patches (red lines) assemble at their domes. New axes of cell growth (arrows) are established. Apart from the AF patches, AFs exhibit diffuse pattern of organization and are not depicted. (e) Cortical MTs form bundles that line the anticlinal walls and radial MT arrays at the junctions of the external periclinal with the anticlinal walls in a young pavement cell. (f) Later morphogenetic stage of a pavement cell. Local cell wall thickenings with CMFs parallel to the underlying MTs are deposited. Arch-like tangential expansion (arrows) is imposed on the external periclinal cell face. (g) Lobes and constrictions are initiated in this morphogenetic stage of pavement cells, while AF patches (red lines) assemble at the domes of growing cell lobes. (h) Paradernal view of an epidermal area during cell lobe formation and expansion. Note that cortical MT arrays, wall thickenings with CMFs and AF patches are alternate between neighboring pavement cell resulting in a 'jigsaw-puzzle' top view.

# Regulace tvaru epidermálních buněk: RIC/ROP systém

- **Rop** Interacting **CRIBs**: efektory ROP, kontrolující aktin i tubulin!
- RIC1: MT bundling
- RIC4: actin polymerization
- ROP2/4:
  - inhibice RIC1
  - stimulace RIC4

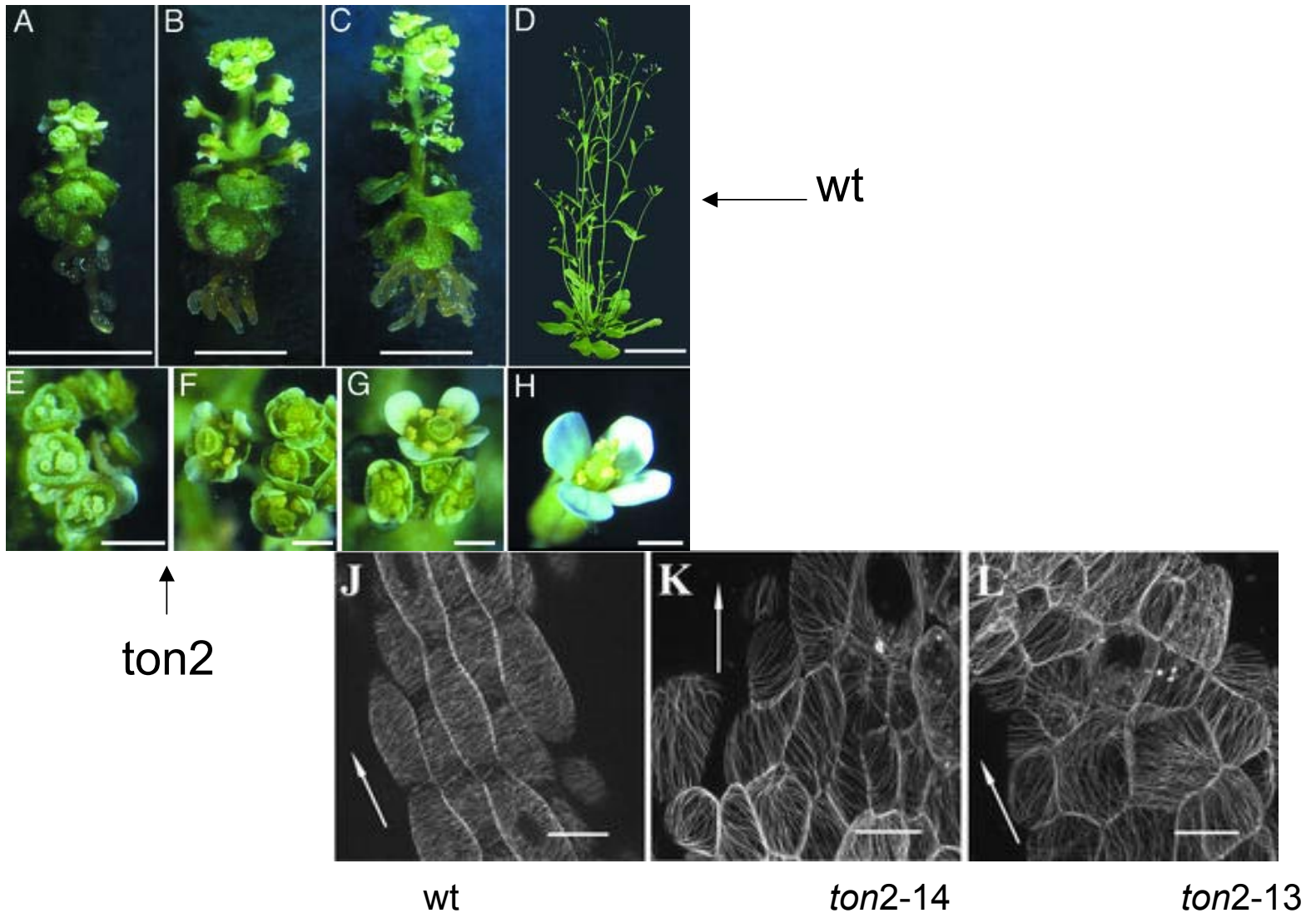


(C) Schematic representation of the model proposed by Fu et al. to explain the role of ROP GTPases and their effectors (RICs) in leaf morphogenesis. ROP2/4 GTPases, via activation of RIC4, promote actin microfilament formation in regions of growing lobes. At the same time, the ROPs, via RIC1 binding, promote microtubule bundling at neck regions to restrict widening and sequester RIC1 at the plasma membrane at sites of lobe initiation in order to prevent microtubules from organizing at those sites.

ROP GTPázy = Rho of plants (Rho GTPázy)

CRIB doména = Cdc42/Rac interactive binding domain

# TON2 kóduje protein fosfatázu A

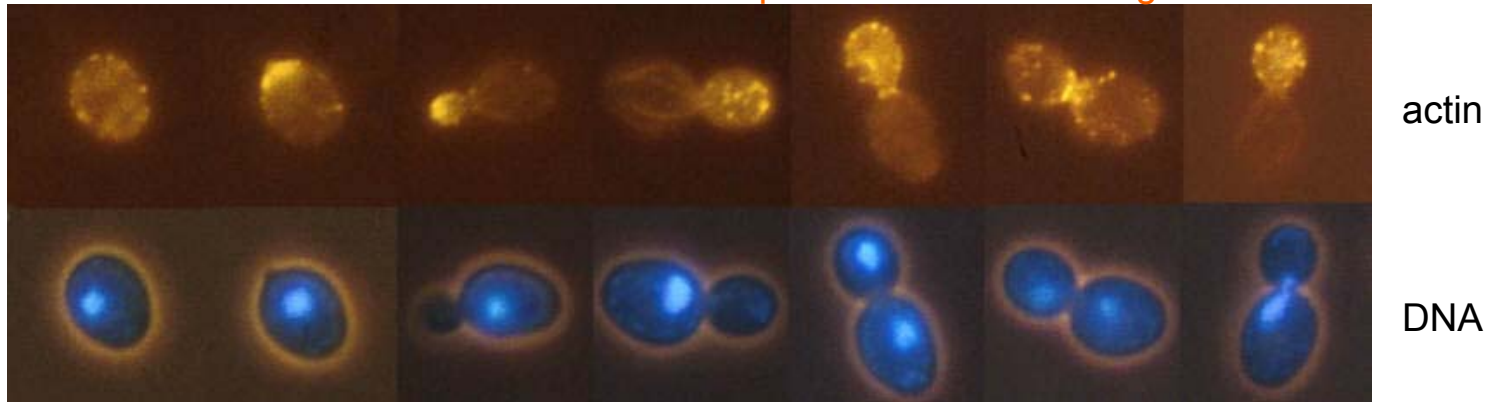


# Cytoskelet kvasinek

# Aktinový cytoskelet během buněčného cyklu

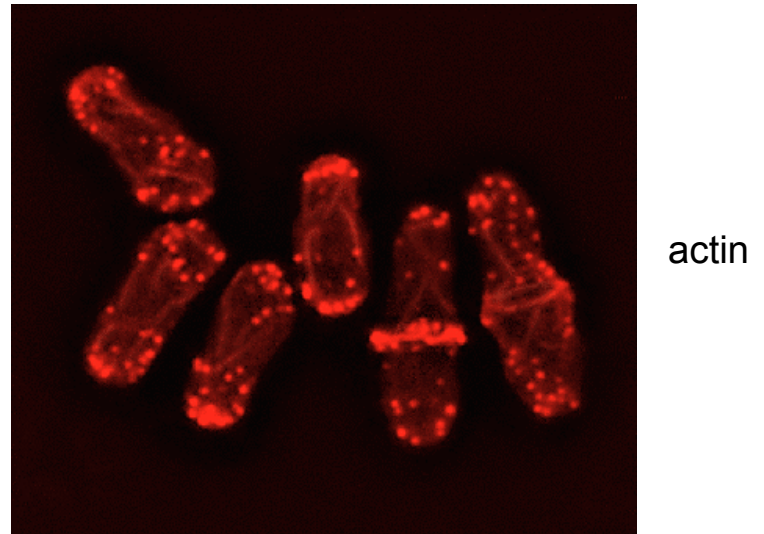
actin patches

actin ring



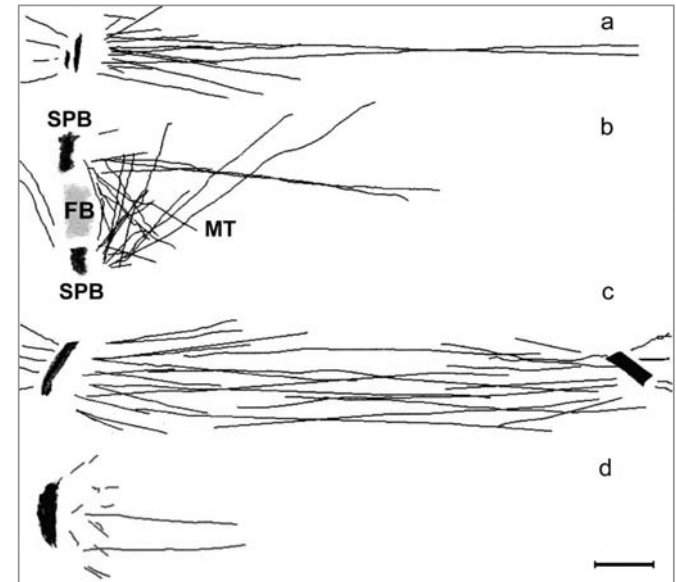
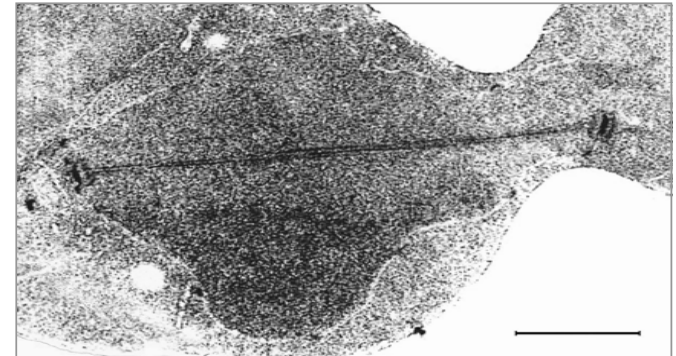
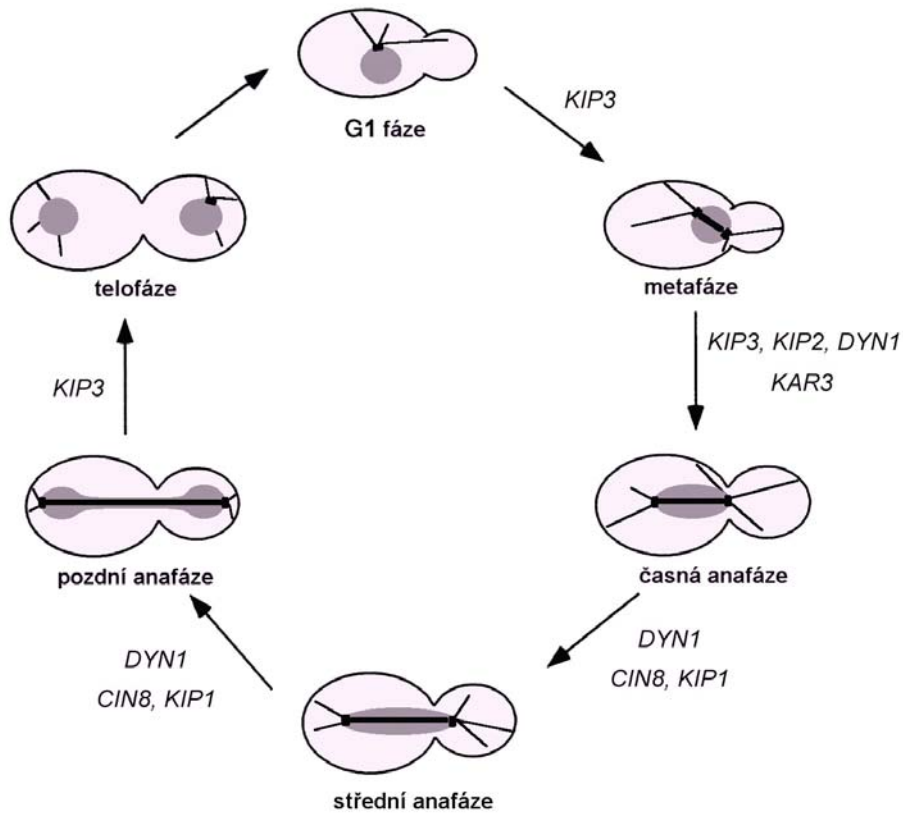
*Saccharomyces cerevisiae*

*Schizosaccharomyces pombe*

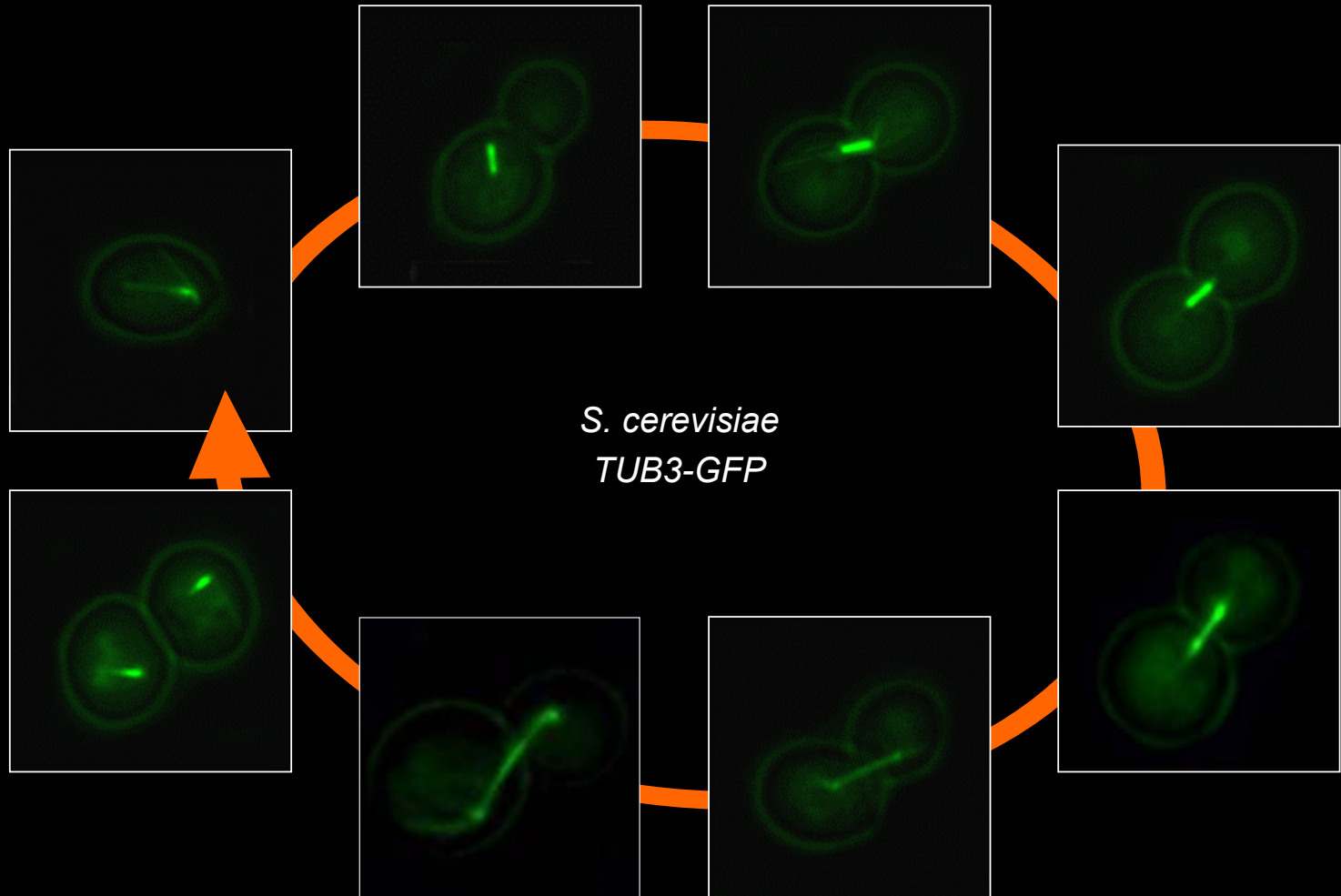




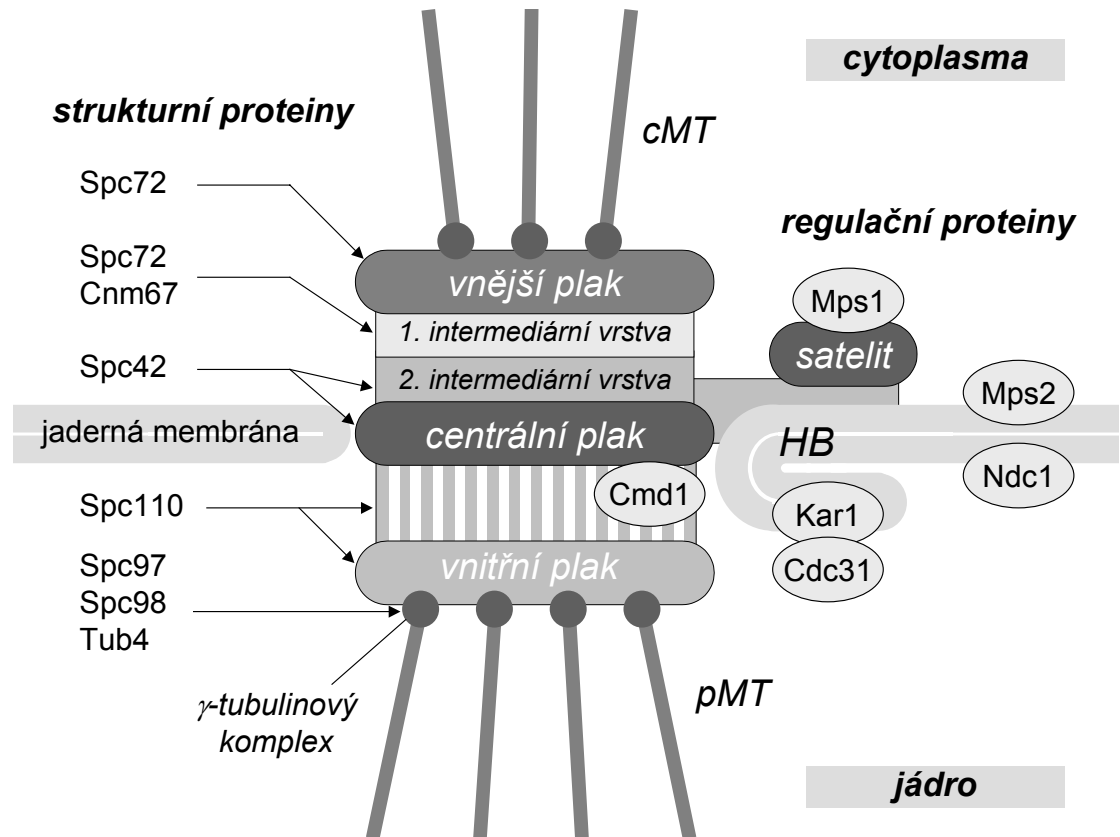
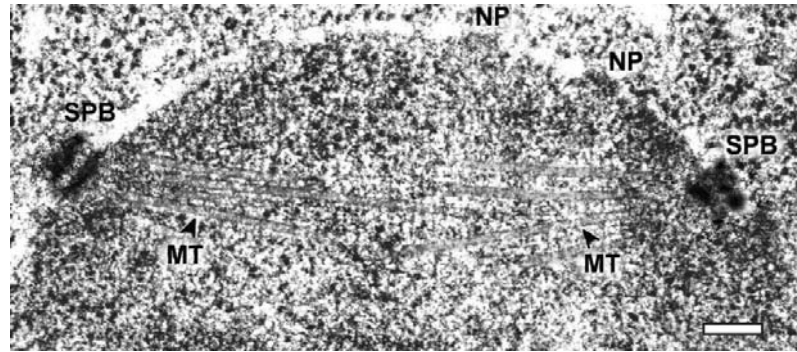
# Mitotické vřeténko během buněčného cyklu



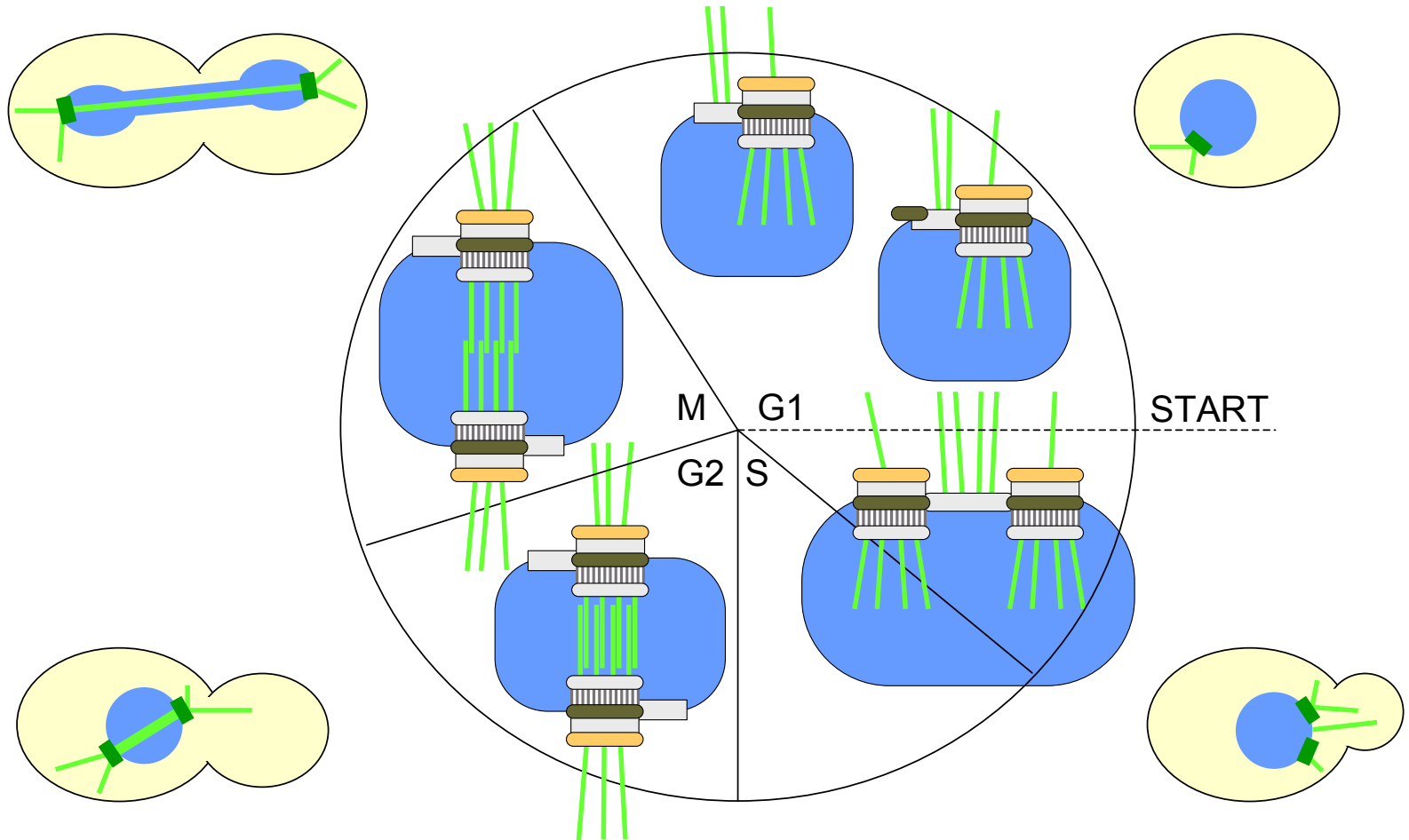
# Tubulinový cytoskelet během buněčného cyklu



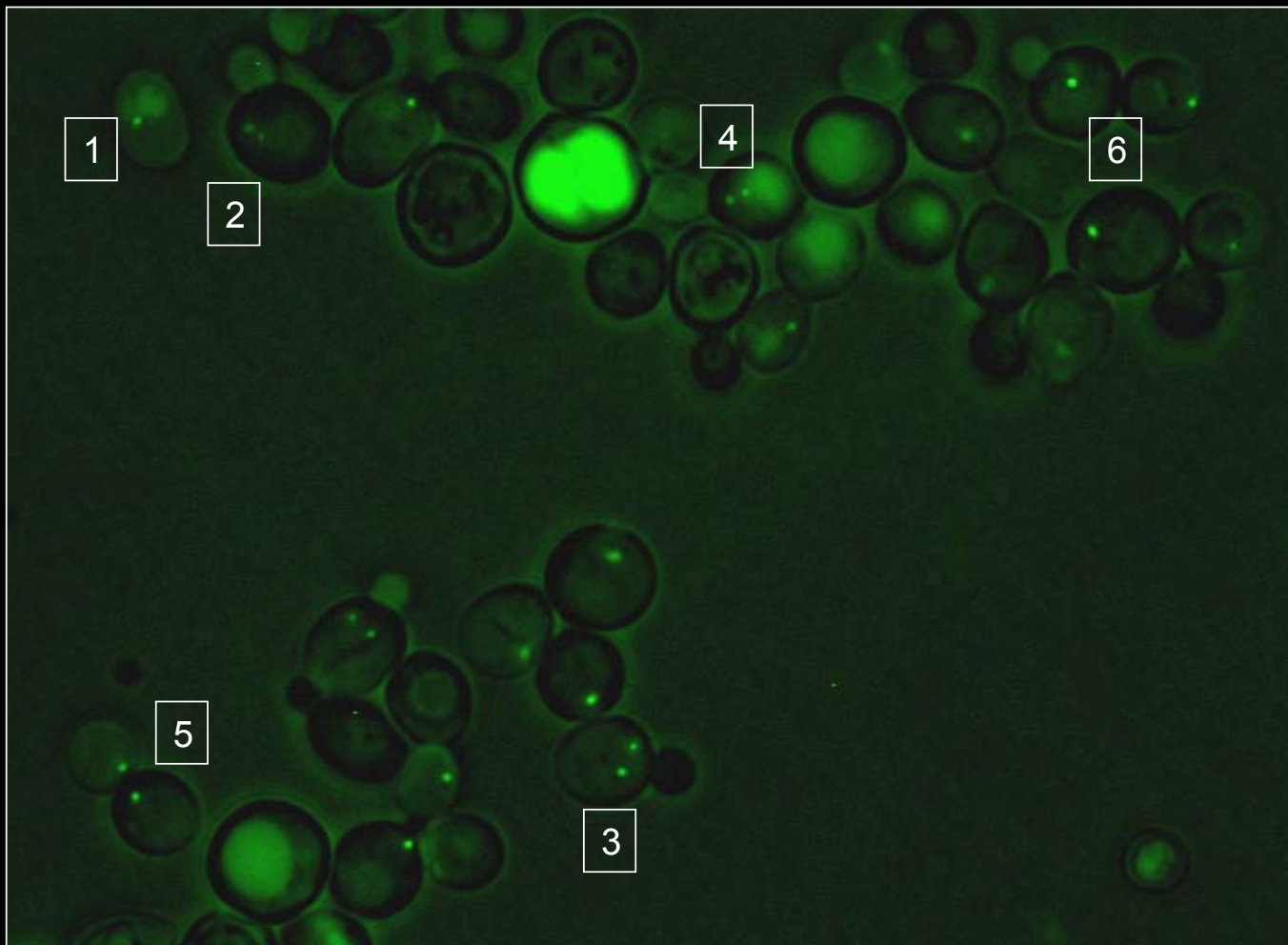
# Spindle pole body (SPB)



# SPB během buněčného cyklu



# SPB během buněčného cyklu



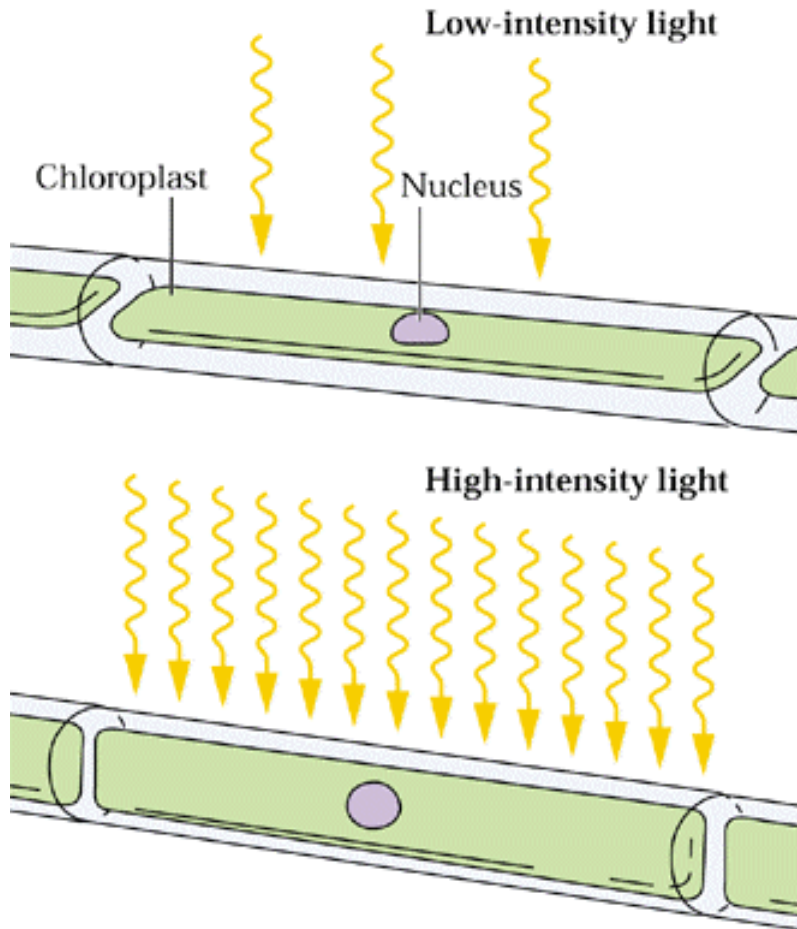
# Molekulové motory

# Molekulové motory

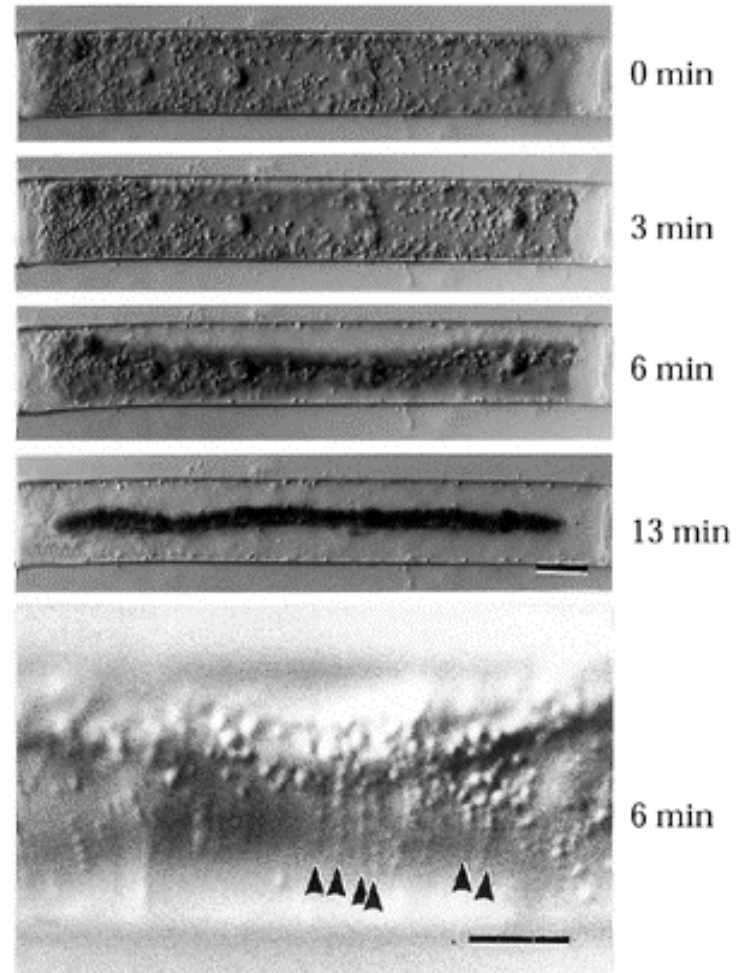
- Po **aktin. filamentech**:
  - až 100  $\mu\text{m/s}$  (*Chara*), typicky jednotky  $\mu\text{m/s}$
  - proudění cytoplazmy, pohyb organel
  - typický motor - **myosin**
- Po **mikrotubulech**:
  - cca 150 nm/s (řádově pomalejší)
  - pohyby chromosomů a mitotického vřeténka, membr. váčků, organizace mikrotubulů ...
  - **kinesiny** (typicky k +konci) a **dyneiny** (k -konci)

# Pohyby organel: plastidy jdou za světlem

(A)



(B)



*Mougeotia*



# MOLEKULOVÉ MOTORY

## DEFINICE

Proteiny, které jsou schopné posouvat se na úkor energie ATP po cytoskeletálních vláknech vždy jedním určitým směrem. Hydrolýza ATP vede k cyklickým konformačním změnám jejich molekul, což umožňuje jejich pohyb.

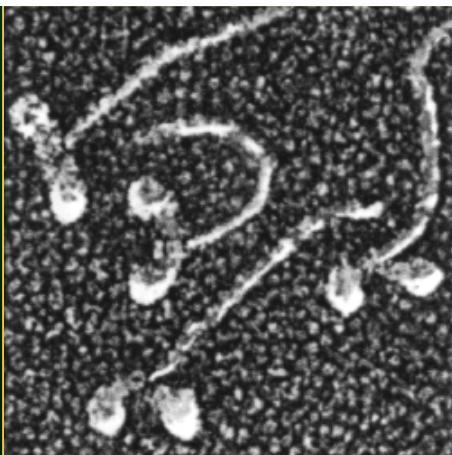
1965 - axonemální dynein (Gibbons)

1985 - kinezin (Brady, Vale, Scholey)

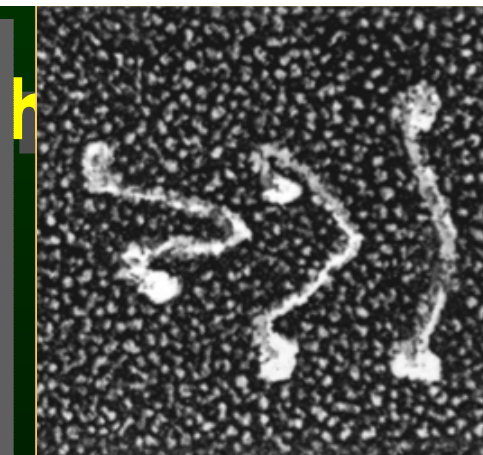
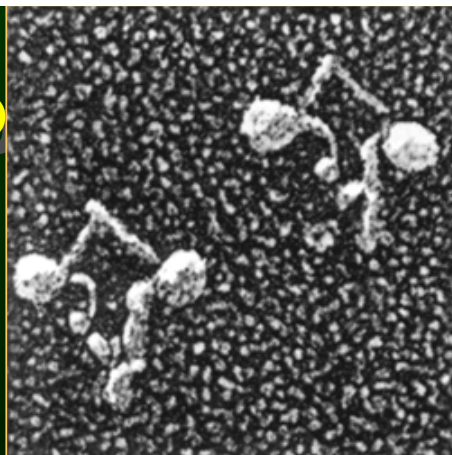
1987 - cytoplazmatický dynein (Lye, Scholey, McIntosh)

# FUNKCE MOLEKULOVÝCH MOTORŮ

- transport váčků  
(endocytóza, exocytóza, přenos signálů, atd.)
- transport molekul nebo komplexů (mRNA)
- pohyb organel
- proudění cytoplazmy
- segregace chromosomů v mitóze a meióze
- splývání jader (karyogamie)
- buněčná polarita, morfogeneze
- vliv na dynamiku mikrotubulů
- komunikace
- svalový stah



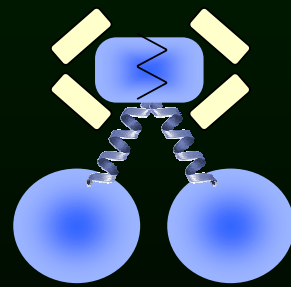
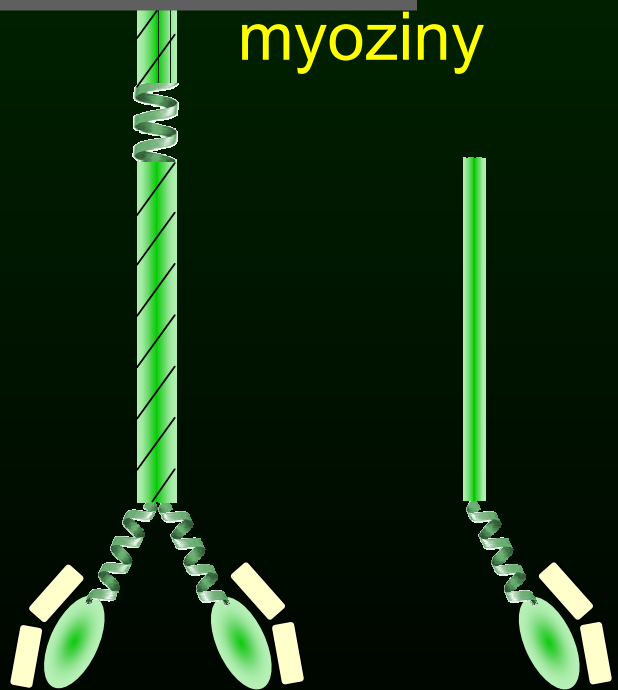
myo



myoziny

dyneiny

kineziny



ocas

tělo

krček

motorová doména

N-konec

mikrofilamenta

IF

mikrotubuly

# Nadrodiny molekulových motorů

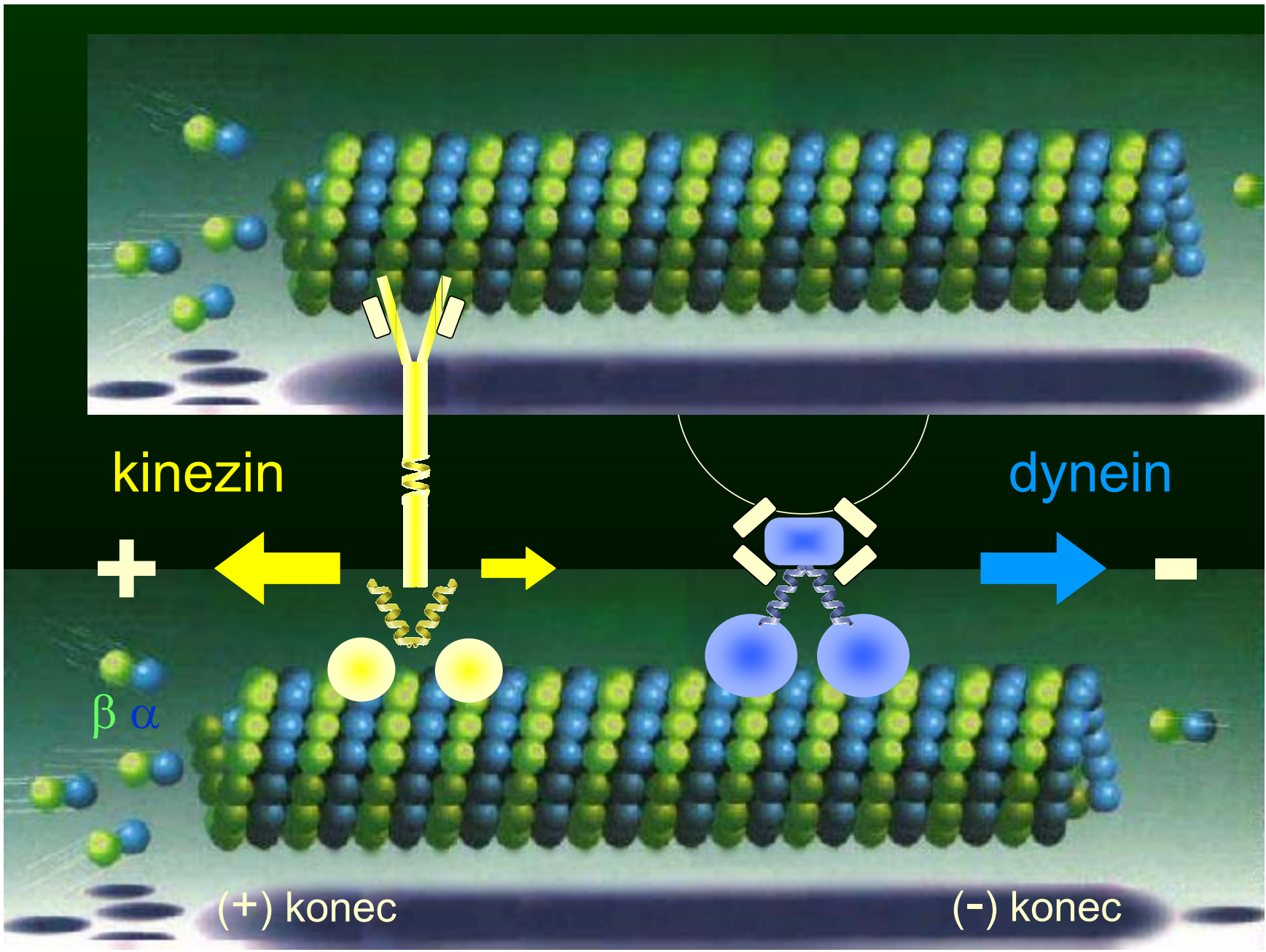
	<i>S. cer.</i>	<i>A. th.</i>	počet tříd
myoziny	5	17	15
kineziny	6	61	10 + nezař.
dyneiny	1	0	

- funkční redundance

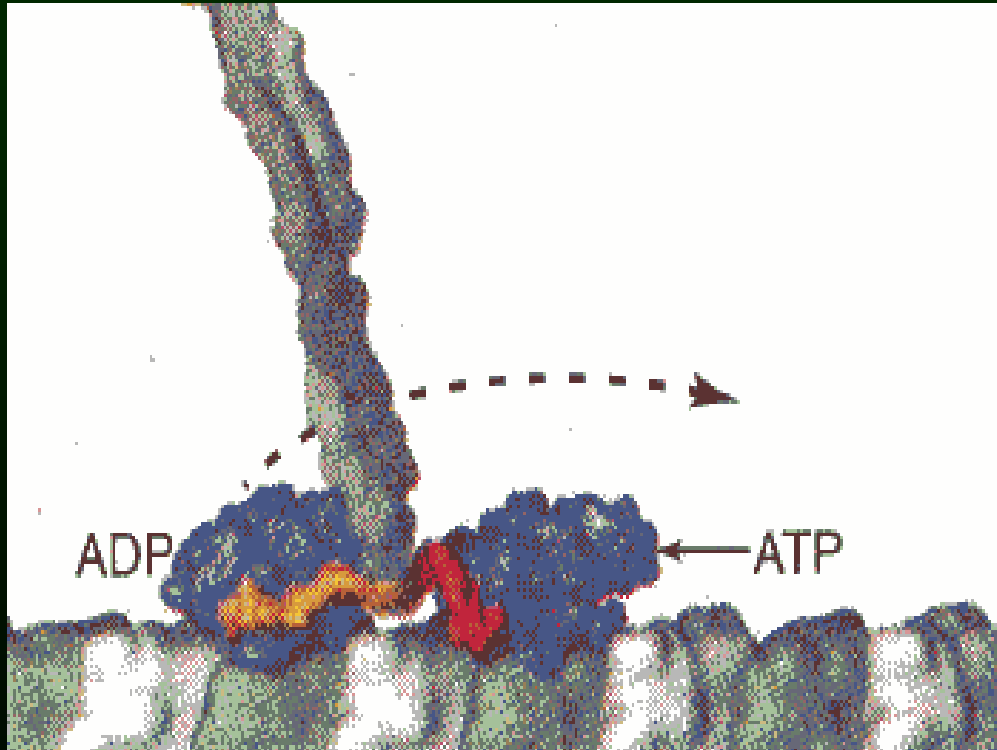
# Kineziны

## Arabidopsis má 61 kinezinů

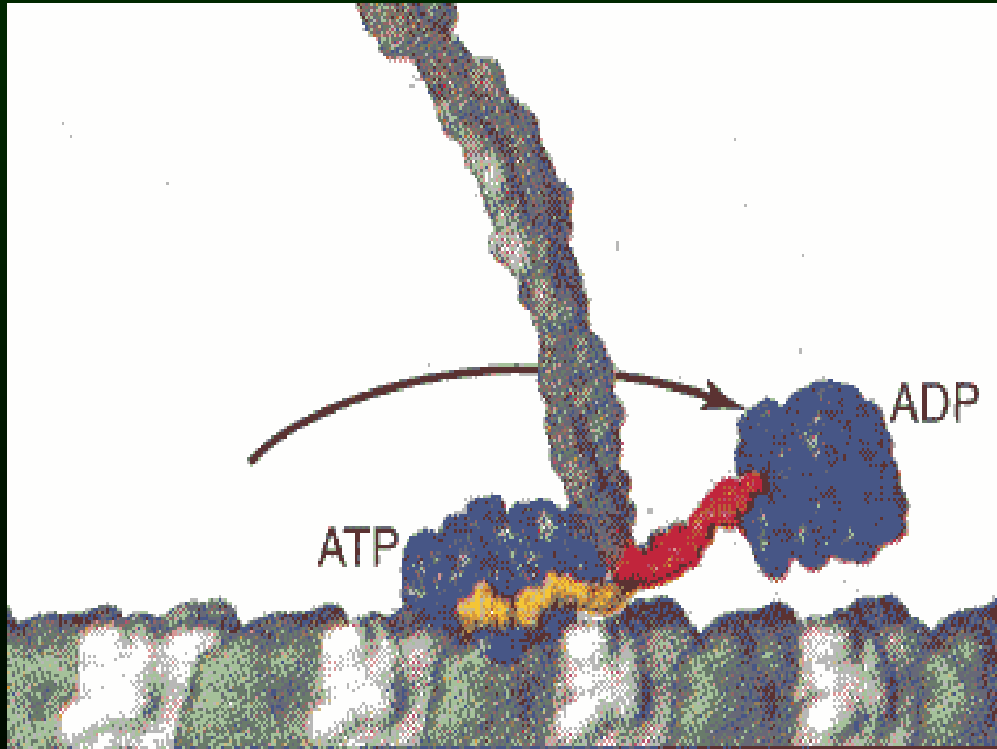
- pohyb po mikrotubulech oběma směry (zajišťován různými proteiny), směr určuje „krček“
- většina (živočišné a 40 u A.th.) + **end-directed**
  - často role v cytokinezi (PAKPs – phragmoplast-associated kinesin proteins)
- zbytek (21) – **end-directed**
  - ATK1 – funkce vřeténka
  - ZWI – regulace calmodulinem
  - KCH – calponin homology domain



# mechanismus pohybu kinezinu

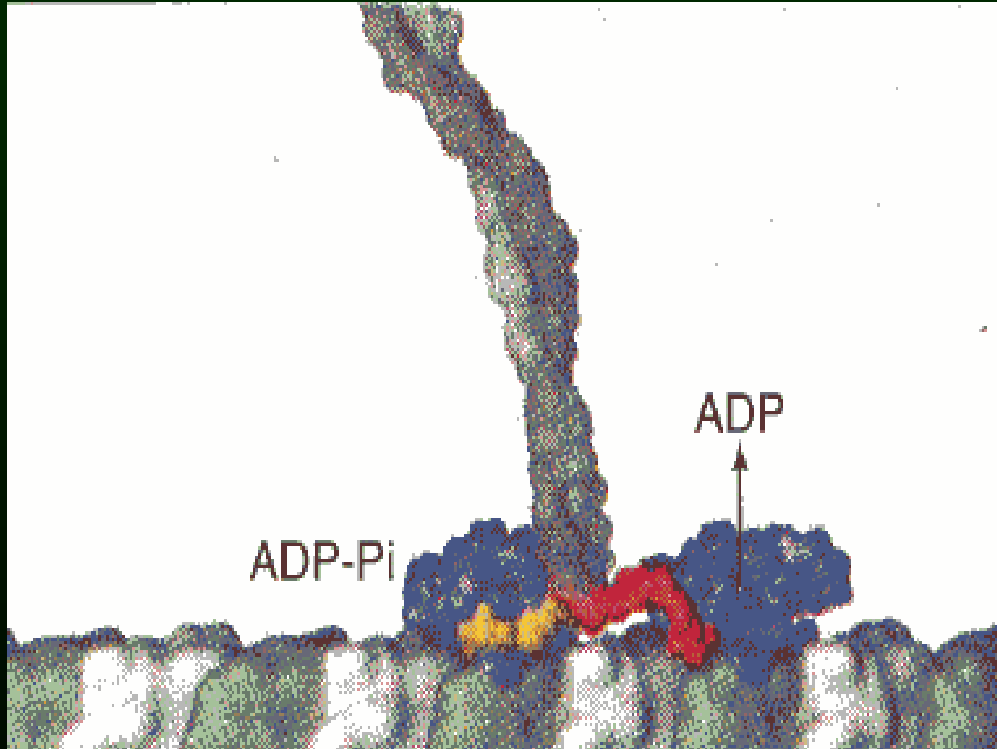


# mechanismus pohybu kinezinu

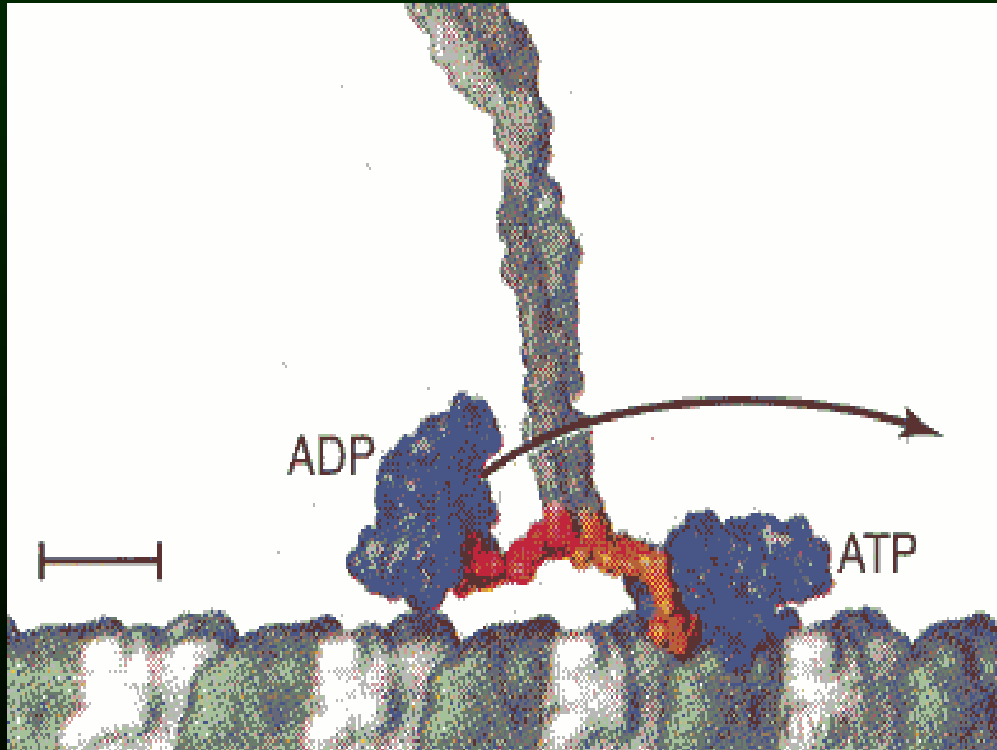




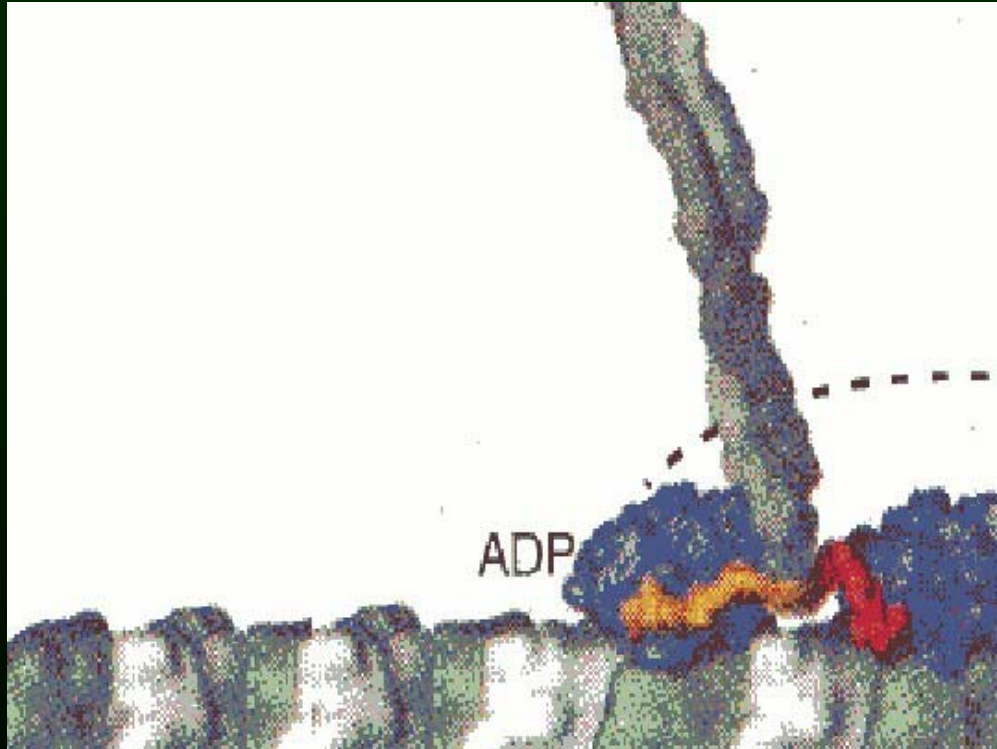
# mechanismus pohybu kinezinu



# mechanismus pohybu kinezinu

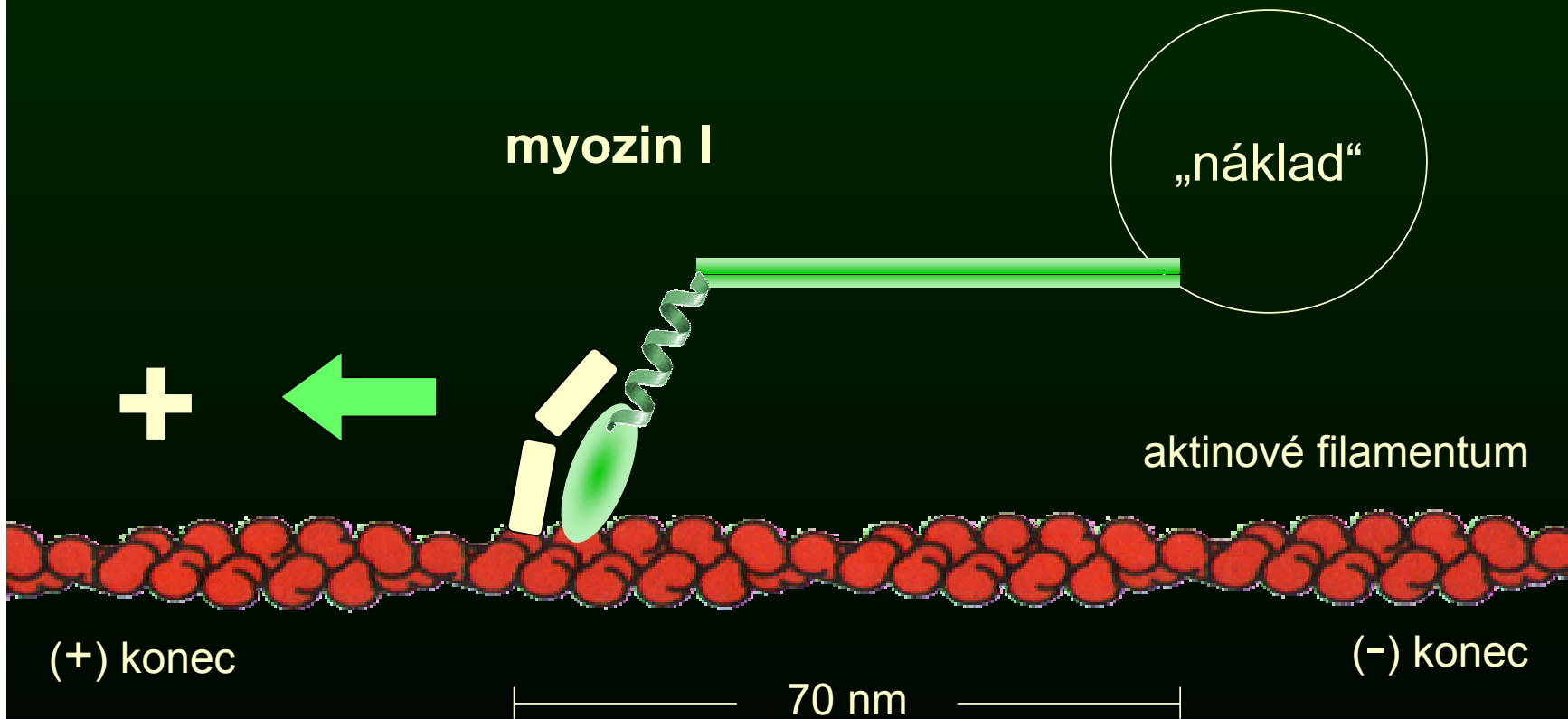


# mechanismus pohybu kinezinu

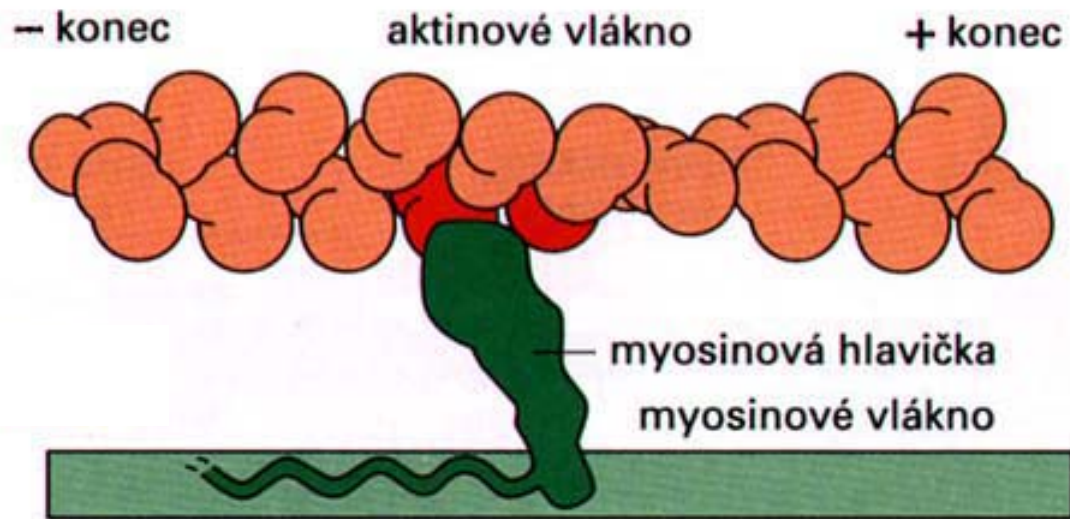


ANIMACE 1

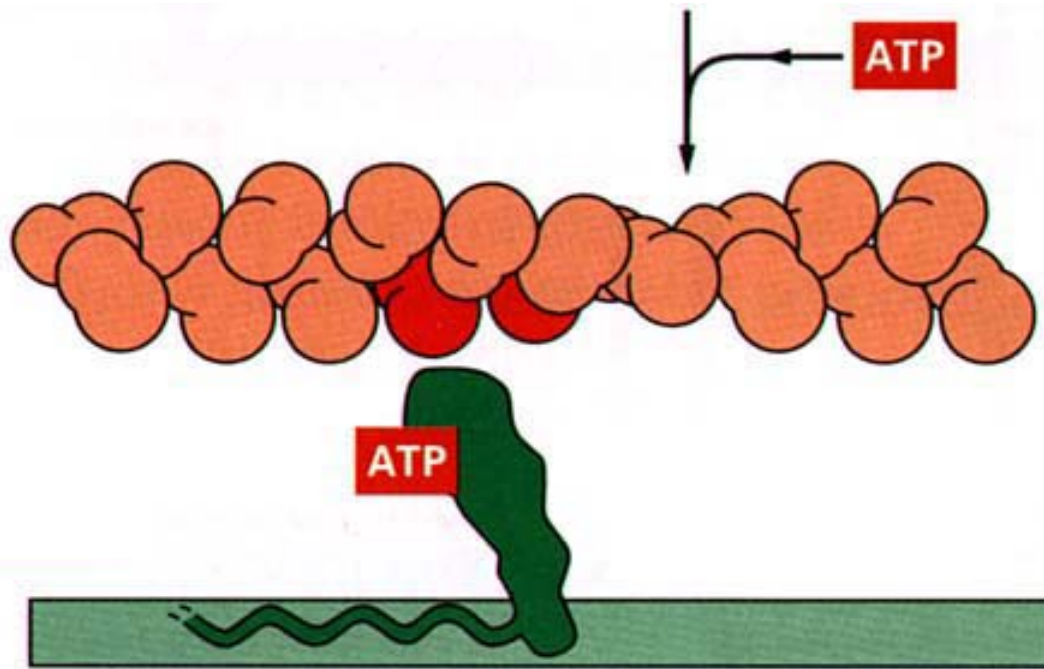
# Myoziny



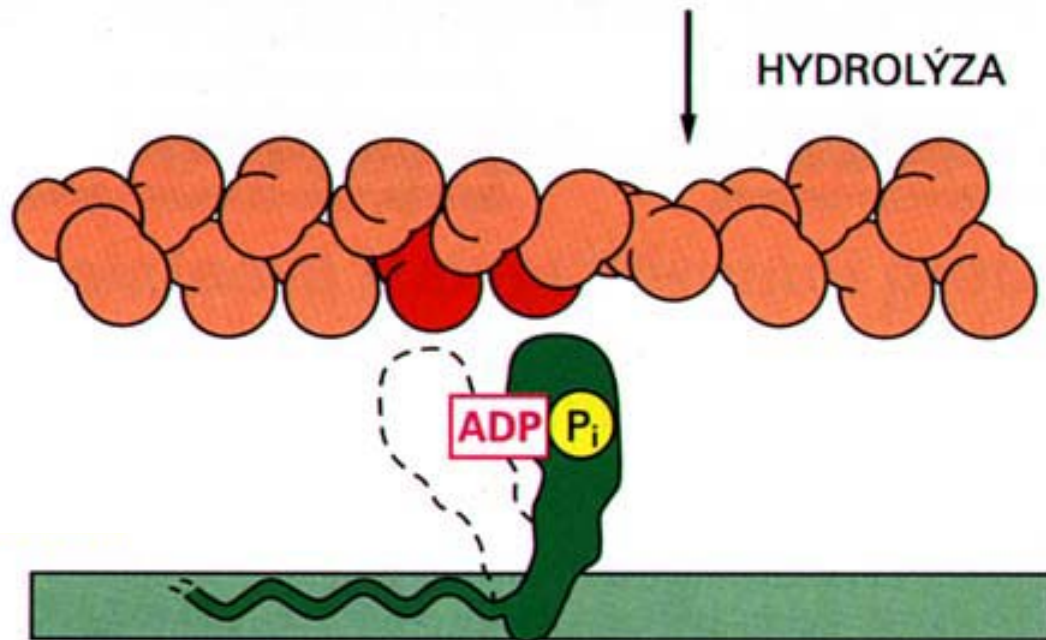
# mechanismus pohybu myozinu



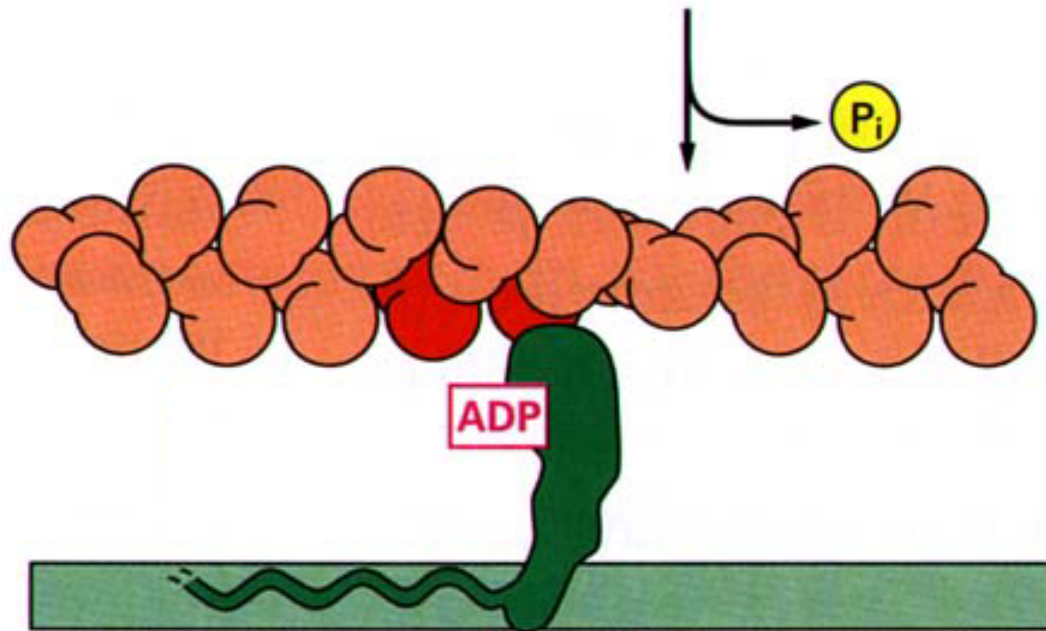
# mechanismus pohybu myozinu



# mechanismus pohybu myozinu

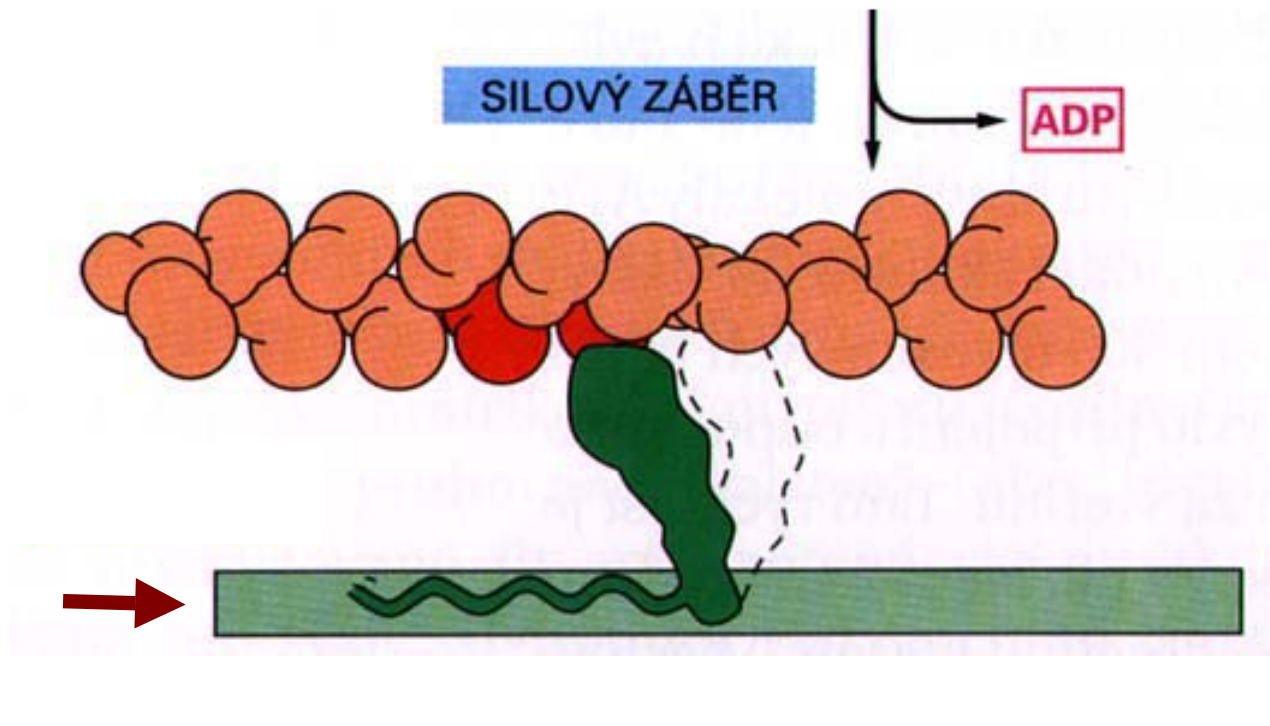


# mechanismus pohybu myozinu



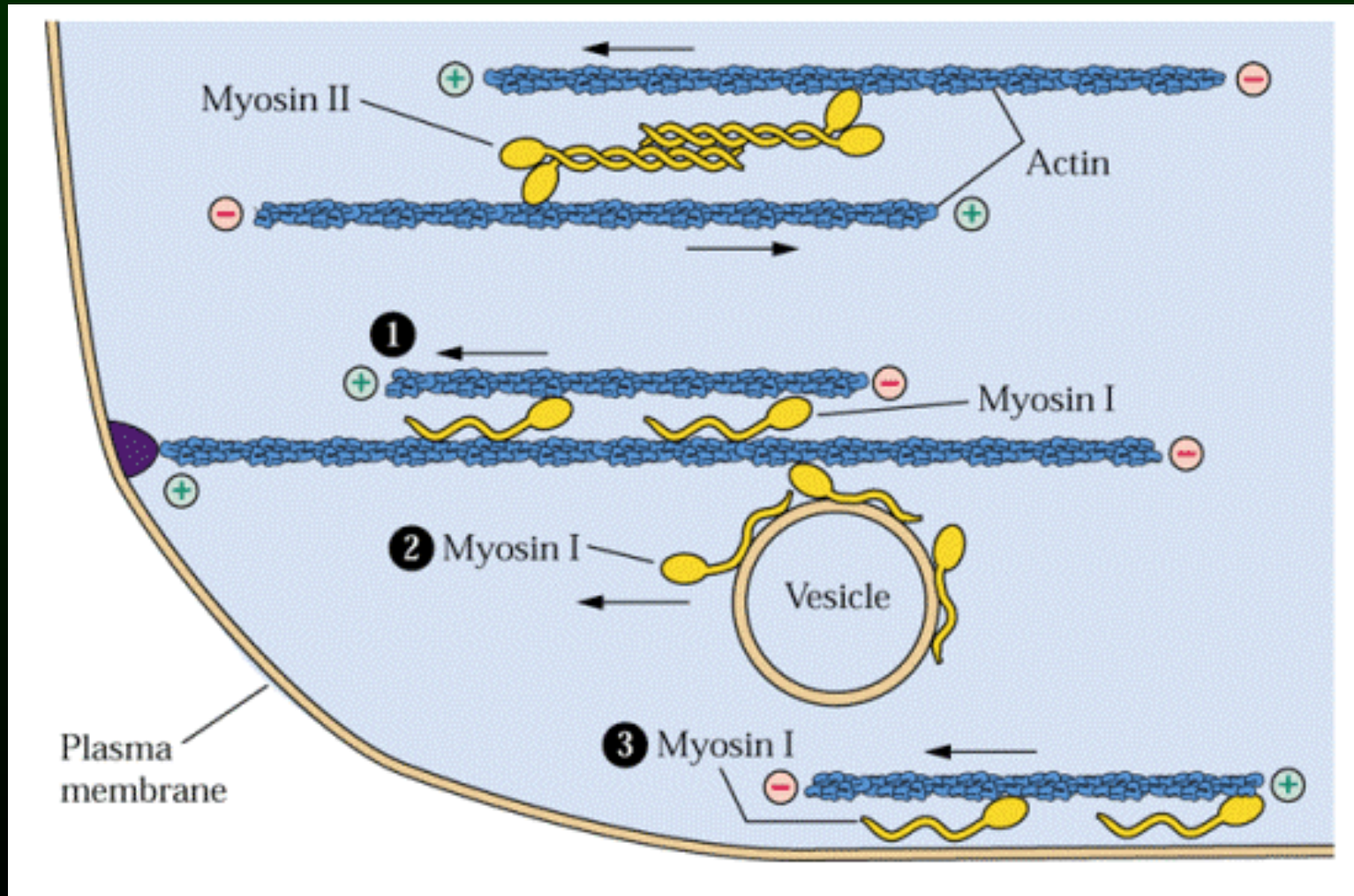


# mechanismus pohybu myozinu

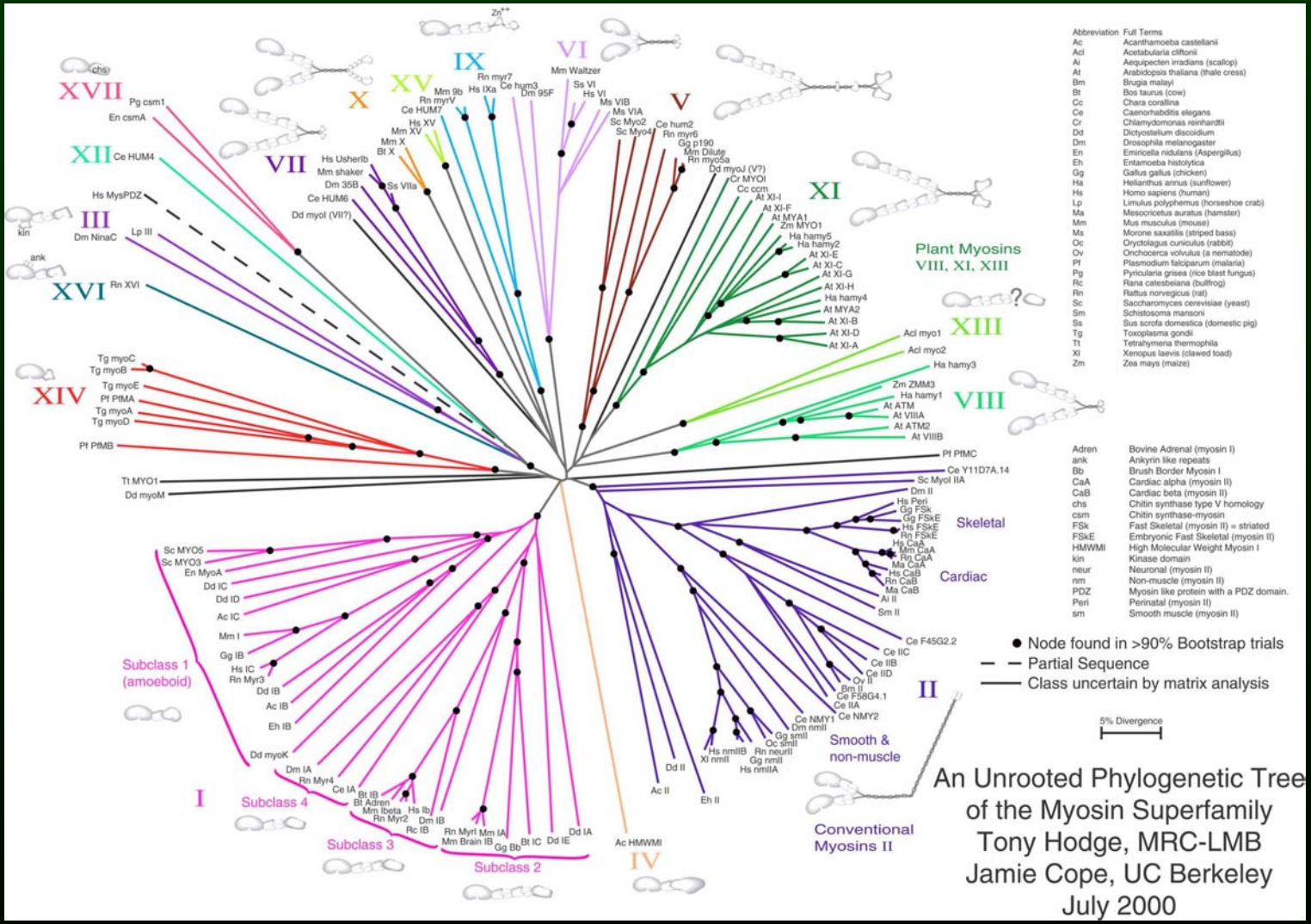


ANIMACE 2

# funkce myozinů



# Myosiny



An Unrooted Phylogenetic Tree of the Myosin Superfamily  
 Tony Hodge, MRC-LMB  
 Jamie Cope, UC Berkeley  
 July 2000

některé třídy specifické jen pro určité skupiny organismů

Arabidopsis má 17 myosinů (class VIII, XI, XIII)

# Regulace molekulových motorů

regulace přes  $\text{Ca}^{2+}$  a calmodulin

⇒ IQ motivy myozinů

⇒ KCBP kinezin

⇒ injekce  $\text{Ca}^{2+}$  urychluje pohyb chromosomů

fosforylace

# Interakce motorů operujících na mikrotubulech a mikrofinamentech

interakce obou skupin motorů → koordinace buněčného transportu (MT – long-distance, MF – short-distance)

- ⇒ SMY1 suprimuje mutaci v MYO2 (fyzická interakce jejich ocasových domén) – *S. cer.*
- ⇒ KCBP je z části kinezin, z části myozin - *A. th.*
- ⇒ přímá interakce MyoVa s KhcU - savci

# Cytoplasmic streaming

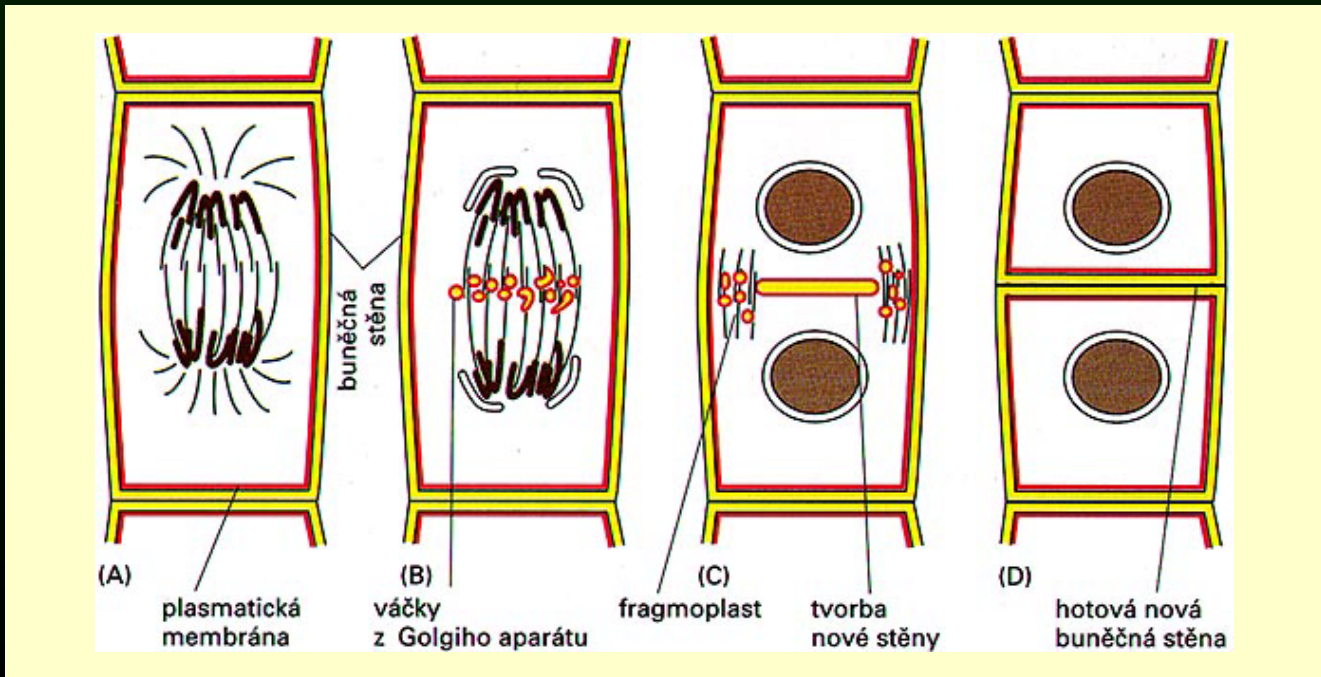
Pohyb organel, váčků a různého materiálu podél aktinového cytoskeletu ( $\sim 70 \mu\text{m/s}$ ), které provádějí motorové proteiny operující na aktinových filamentech.

- ⇒ kolokalizace mikrofilament a “proudů” cytoplasmy
- ⇒ depolymerizace mikrofilament inhibuje *streaming*
- ⇒ depolymerizace mikrotubulů neinhubuje *streaming*
- ⇒ kosedimentace myozinu s váčky
- ⇒ imunolokalizace myozinů:
  - myozin I kolem jádra, mitochondrií, chloroplastů
  - myozin II kolem mitochondrií, chloroplastů
  - myozin V kolem malých organel (váčky)

# Buněčné dělení

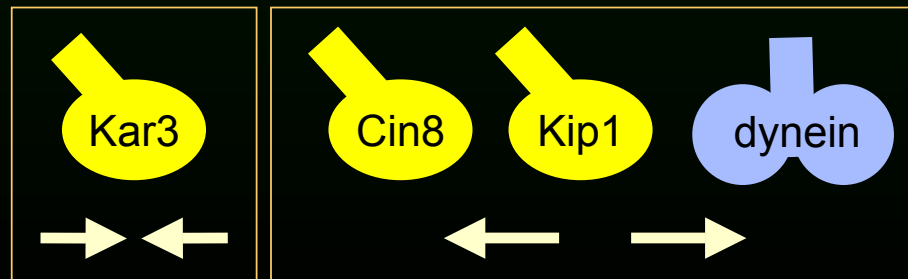
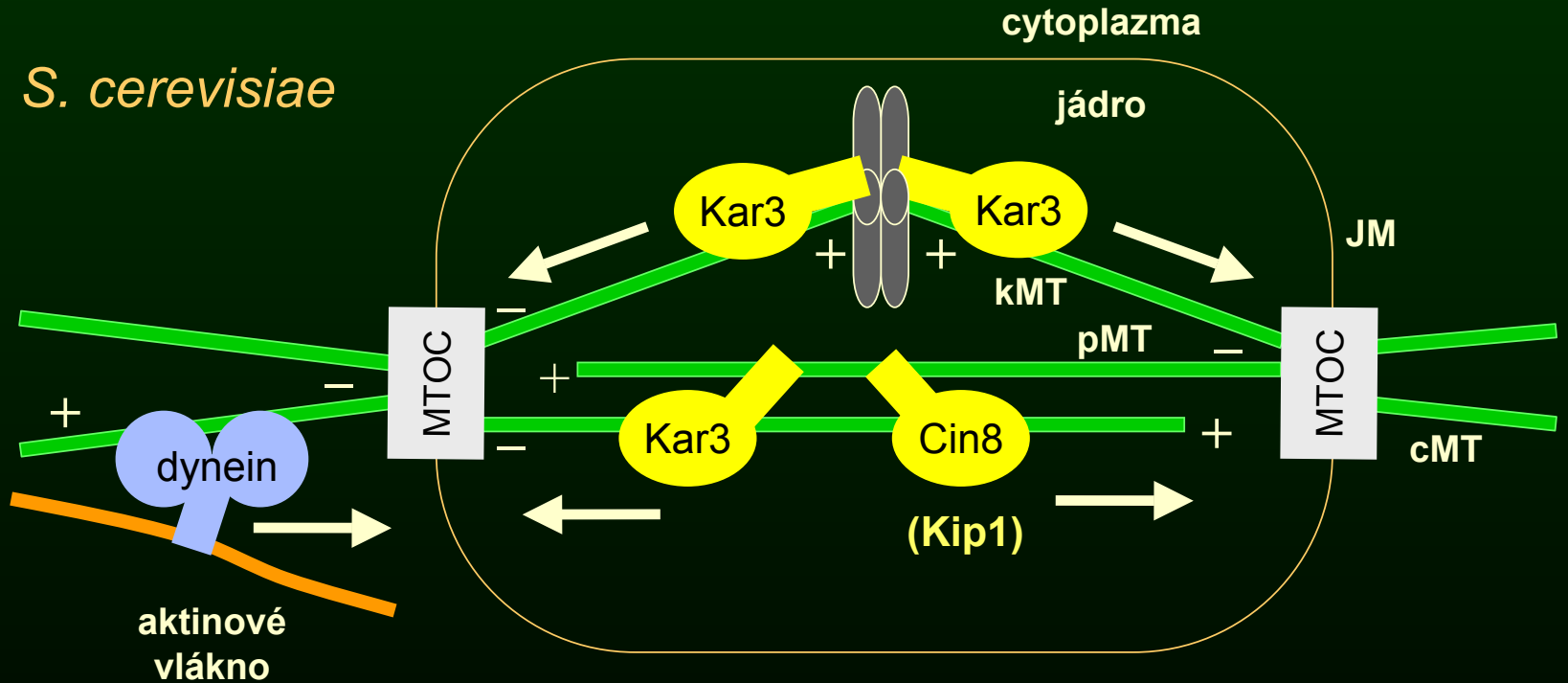
- kortikální mikrotubuly  $\Rightarrow$  preprofázový prstenec
- vytváření dělicího vřeténka
- segregace chromosomů
- tvorba plazmatické destičky

ANIMACE 3



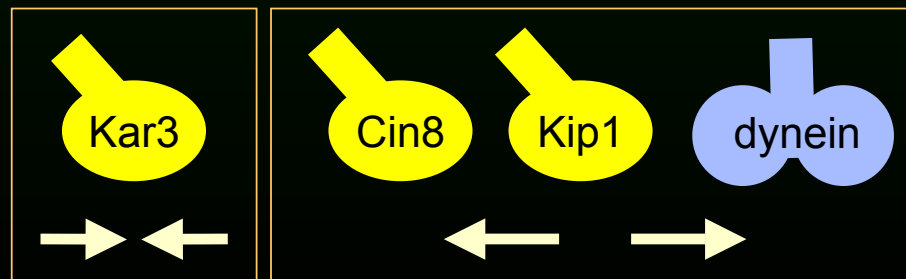
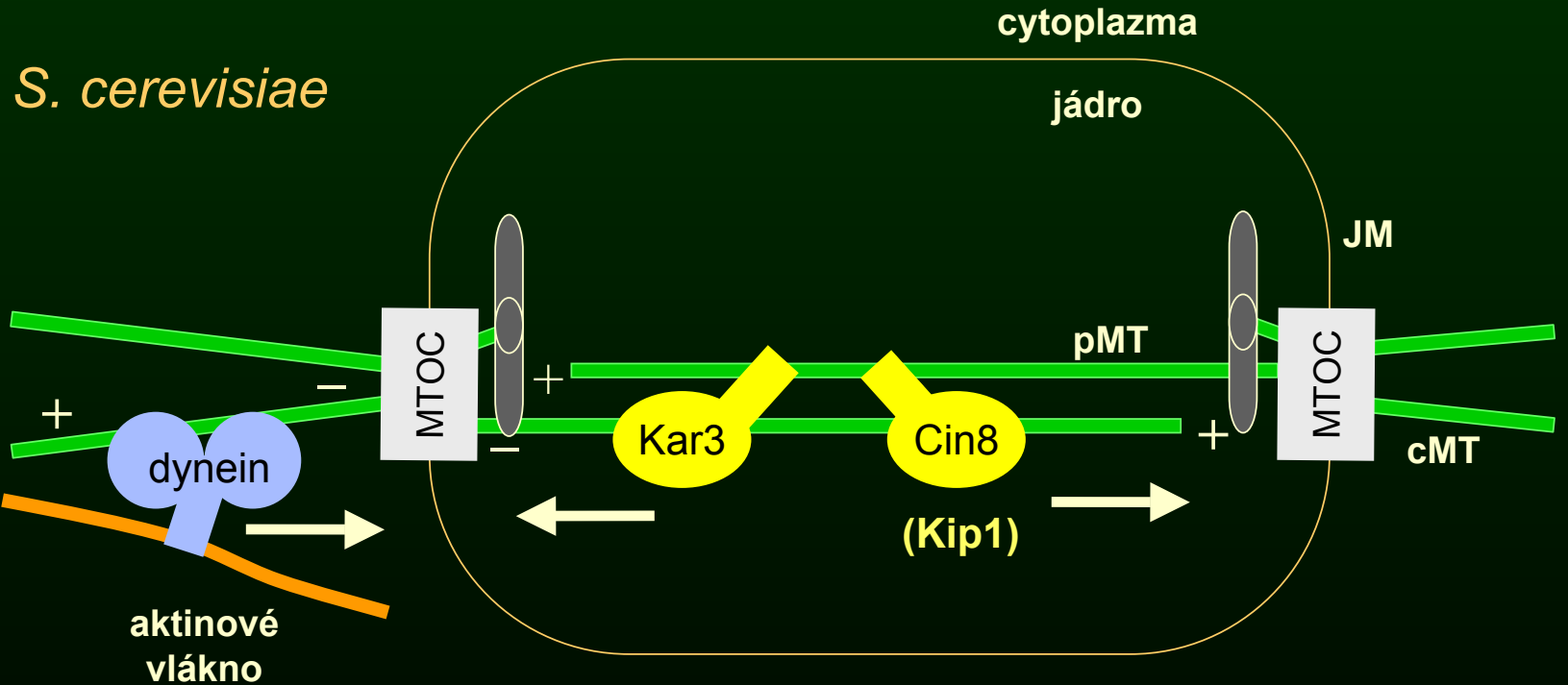
# segregace chromosomů

*S. cerevisiae*





# segregace chromosomů



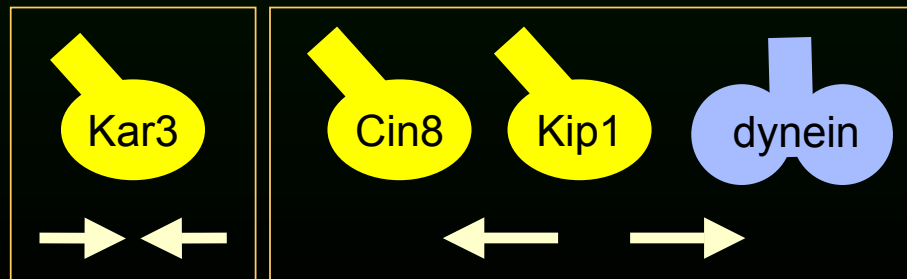
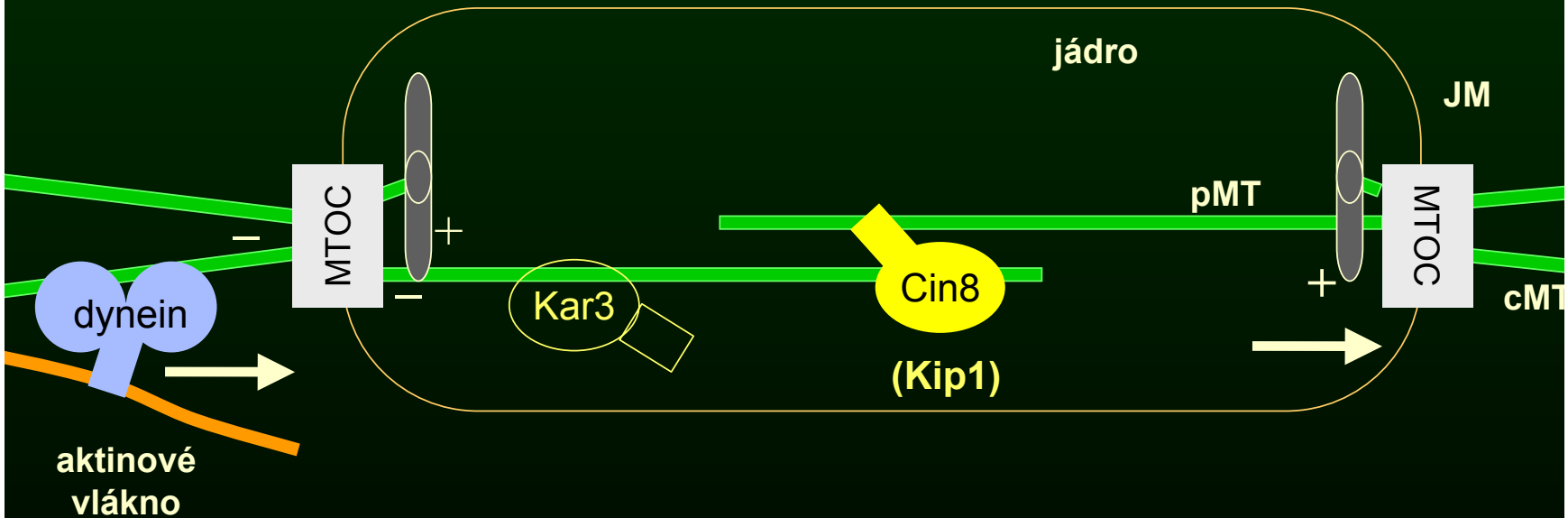
# segregace chromosomů

*S. cerevisiae*

cytoplazma

jádro

JM



# segregace chromosomů

rostliny

## motorové proteiny operující na mikrotubulech

- ⇒ zvýšená hladina exprese kinezinů během mitózy (KatB/C, KCBP, TKRP125)
- ⇒ lokalizace některých kinezinů podél mikrotubulů vřeténka (N- i C-koncové)
- ⇒ injekce KCBP-protilátky zablokuje mitózu v metafázi

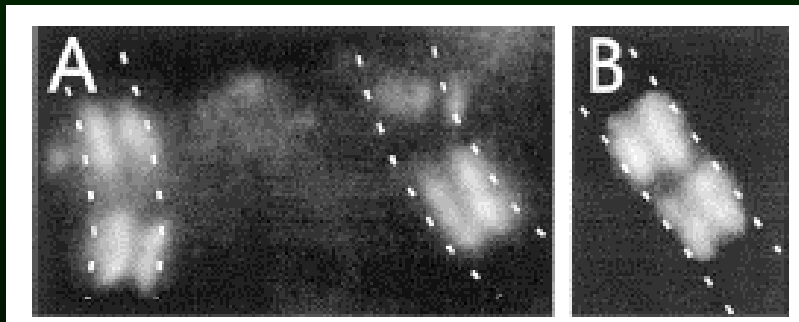
# tvorba plazmatické destičky

## motorové proteiny operující na mikrotubulech

- ⇒ asociace váčků s mikrotubuly fragmoplastu
- ⇒ depolymerizace mikrotubulů inhibuje tvorbu destičky
- ⇒ depolymerizace mikrofilament neinhibuje tvorbu destičky (vede pouze k abnormální pozici destičky)
- ⇒ lokalizace některých motorů u fragmoplastu; N-koncové kineziny (TKRP125) i C-koncové (KatA a KCBP)
- ⇒ injekce KCBP-protilátky po anafázi blokuje tvorbu destičky

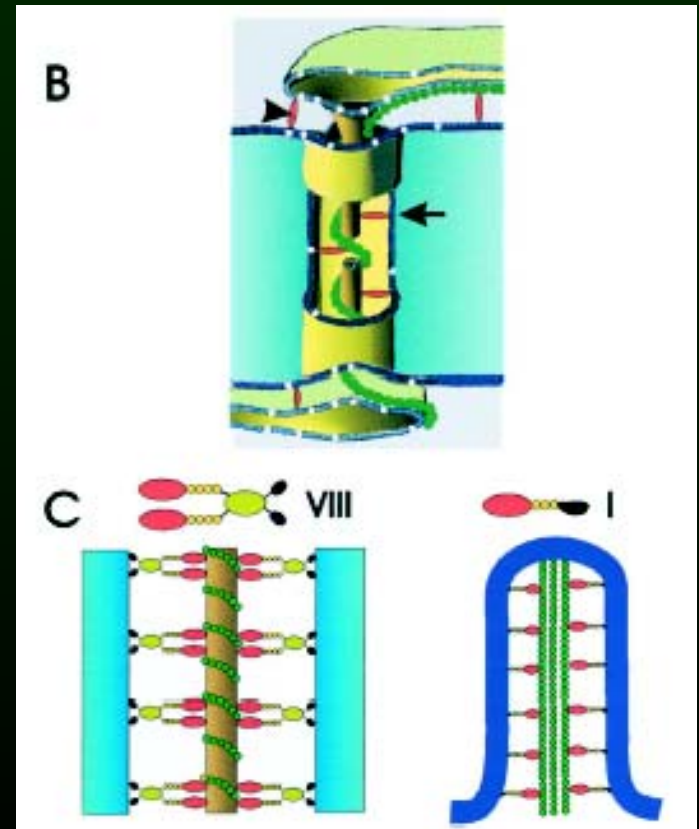
# buněčná komunikace

MyoVIII funguje v plasmodesmech



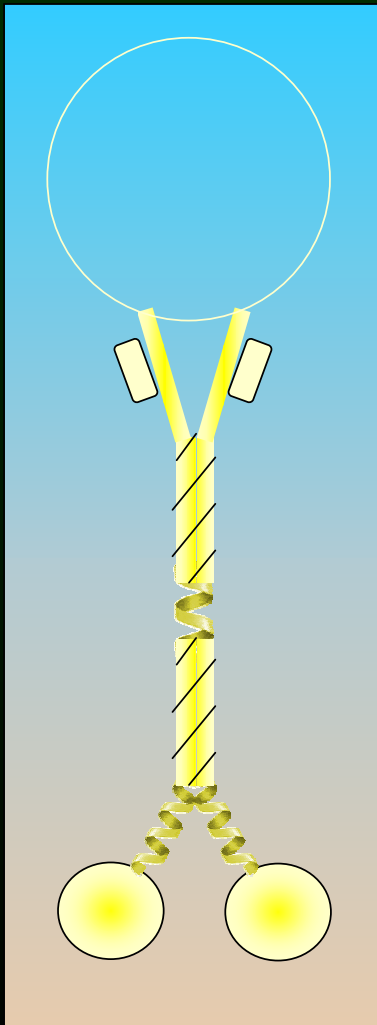
mikrotubuly a mikrofilamenta jsou nezbytné pro regulovaný transport makromolekul přes plasmodesmy

⇒ pravděpodobná účast molekulových motorů



„mikrovilus naruby“?

Baluška et al. 2001



náklad

tělo

motorová  
doména

