



Laboratory of Biology of Cytoskeleton

Modulation of microtubule organization, microtubule proteins, signal transduction

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The long-term research programme of the laboratory has been focused on studying the structure-function relationships of microtubule [MT] proteins and their interactions with other cytoskeletal elements in cells under normal and pathological conditions. The organization of dynamic MT networks is controlled by microtubule organizing centres [MTOCs]. One of the key components of MTOCs is γ -tubulin, which is necessary for nucleation of MT. Current work focuses on the understanding of the modulation of MT properties by signal transduction molecules, the function of γ -tubulin forms, and molecular and functional characterization of regulators of microtubule nucleation. To address these questions, techniques of molecular biology, biochemistry and immunology are being used, as well as a variety of microscopic techniques, including live cell imaging using TIRF microscopy. Our results demonstrate that rearrangement of microtubules in activated mast cells depends on activity of the stromal interacting protein 1 [STIM1] and that Ca^{2+} plays an important role in the formation of microtubule protrusions in activated cells. We have also shown that non-receptor protein tyrosine kinases play an important role in MT nucleation. We have demonstrated that prediction of epitope exposure on microtubules by means of homology modelling combined with site-directed antibodies can contribute to better understanding of the interactions of microtubules

with associated proteins. Finally, we have shown that ectopic expression of tubulins may represent a novel marker in pathobiology of glioblastoma multiforme, the most common and deadliest form of primary brain cancers.

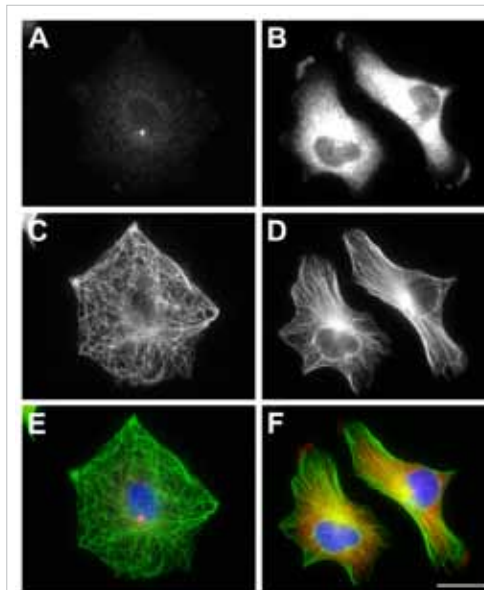


Fig. 1. Different subcellular distribution of γ -tubulin in fibroblasts and glioblastoma cells. Mouse fibroblasts 3T3 [A, C, E] or human glioblastoma cells T98G [B, D, F] stained with antibodies for γ -tubulin [A, B, red] and α -tubulin [C, D, green]. Superpositions of images are shown in E and F. Scale bar, 20 μ m.

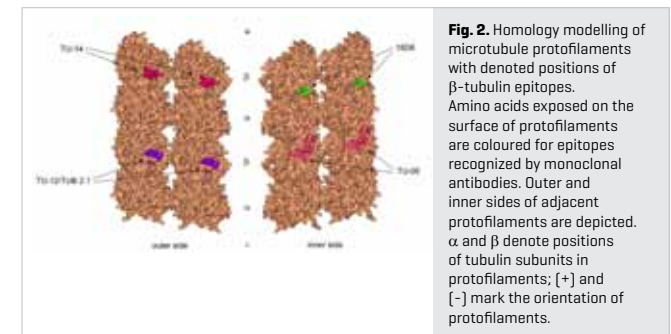


Fig. 2. Homology modelling of microtubule protofilaments with denoted positions of β -tubulin epitopes. Amino acids exposed on the surface of protofilaments are coloured for epitopes recognized by monoclonal antibodies. Outer and inner sides of adjacent protofilaments are depicted. α and β denote positions of tubulin subunits in protofilaments; (+) and (-) mark the orientation of protofilaments.

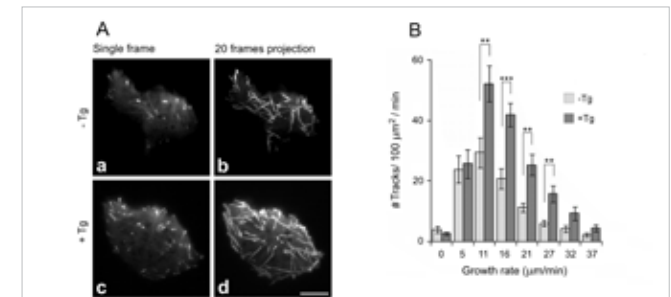


Fig. 3. Activation of mast cells increases the number of growing microtubules in cell periphery as determined by TIRFM time-lapse imaging. [A] Time-lapse imaging of resting [a-b] and thapsigargin [Tg]-activated [c-d] mast cells expressing EB1-GFP that marks growing microtubules. Still images of EB1 [a, c] and tracks of EB1 comets over 20 sec [b, d] in TIRFM. [B] Histogram of microtubule growth rates in cell periphery of resting [-Tg] and thapsigargin-activated [+Tg] cells.



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- GA CR, GAP302/10/1701 – Analysis of functional differences between γ -tubulins, 2010–2012, E. Dráberová
- Ministry of Education, Youth and Sports of the Czech Republic, LC545 – Centre of Functional Organization of the Cell, 2005–2011, Pavel Dráber
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