

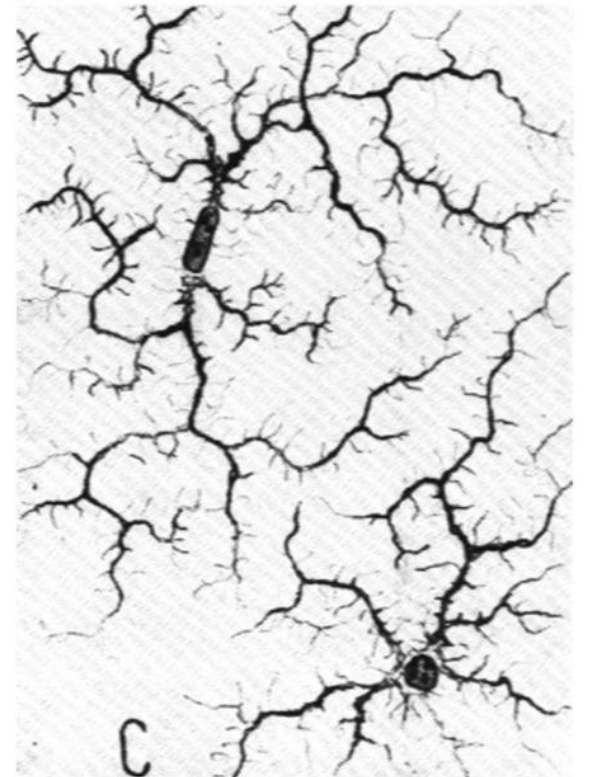
Microglia cytoskeleton rearrangement in homeostatic and activated states

Genetics and Molecular Biology
Molecular and Cellular Physiology
AA 2024/2025

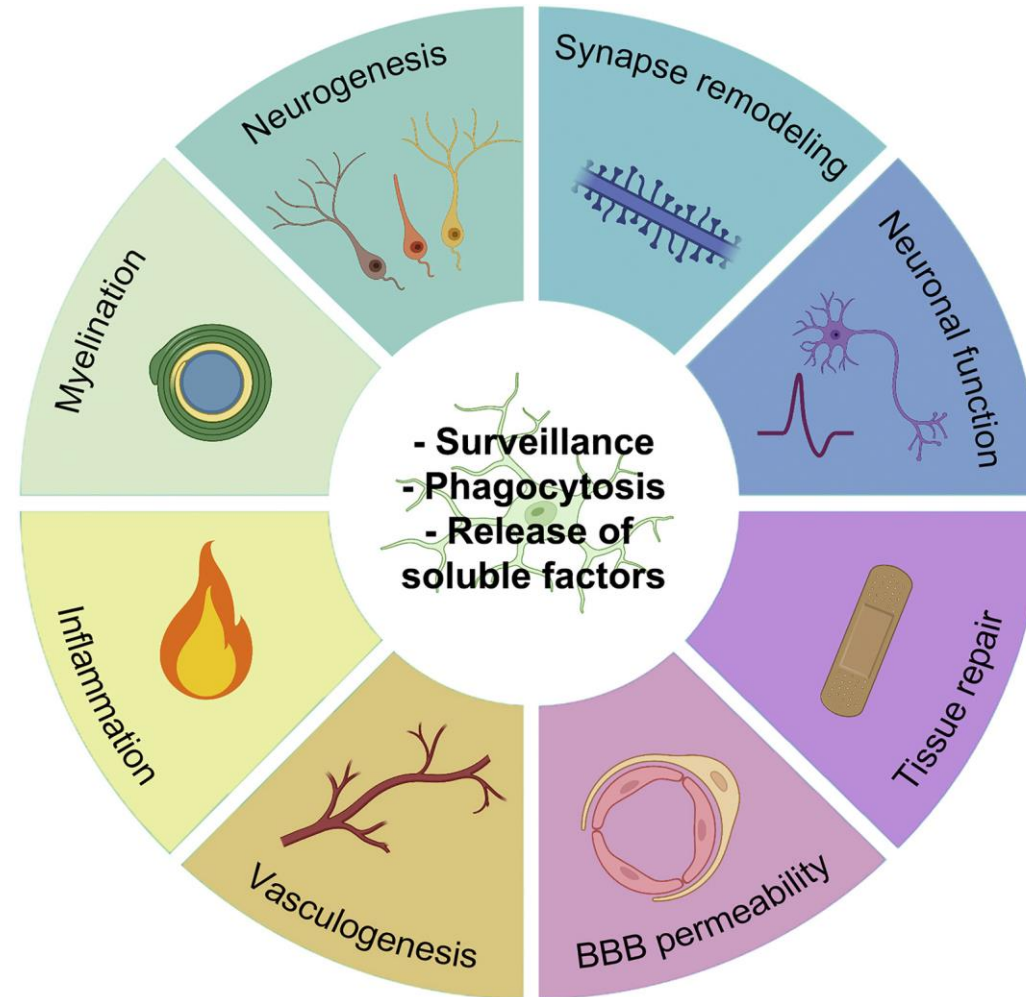
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Microglia origin and physiology

- Microglia are the primary **immune cells** of the brain
- Microglia originate from **myeloid** precursors and migrate early during development in the brain parenchyma
- Microglia cells in the healthy mature brain display a **ramified morphology**, with a small soma and fine cellular processes
- When subjected to external damaging stimuli, microglia undergo a process called «activation», characterized by profound changes in the microglial cell shape, gene expression and functional behavior

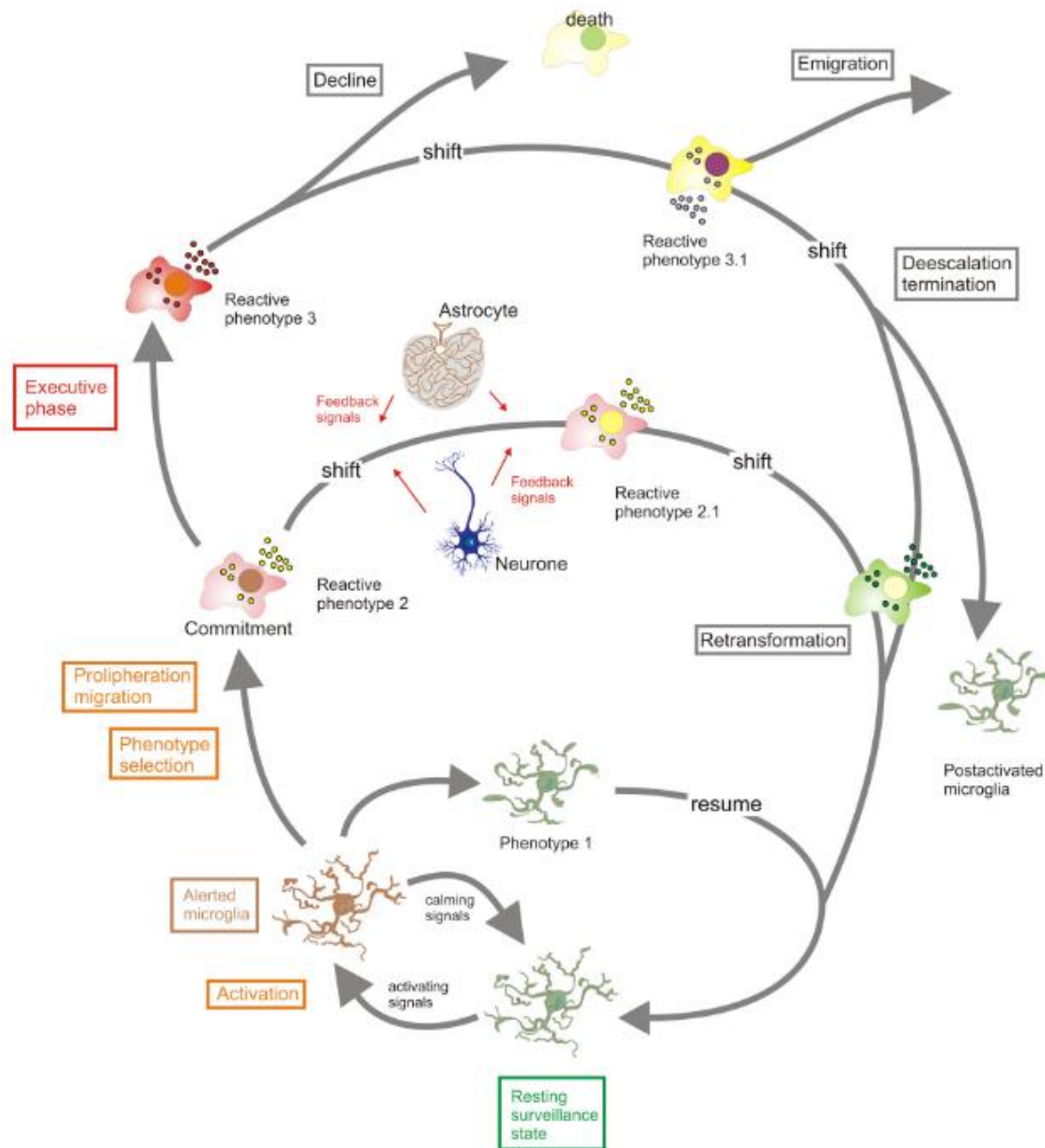


Microglial core properties and functions



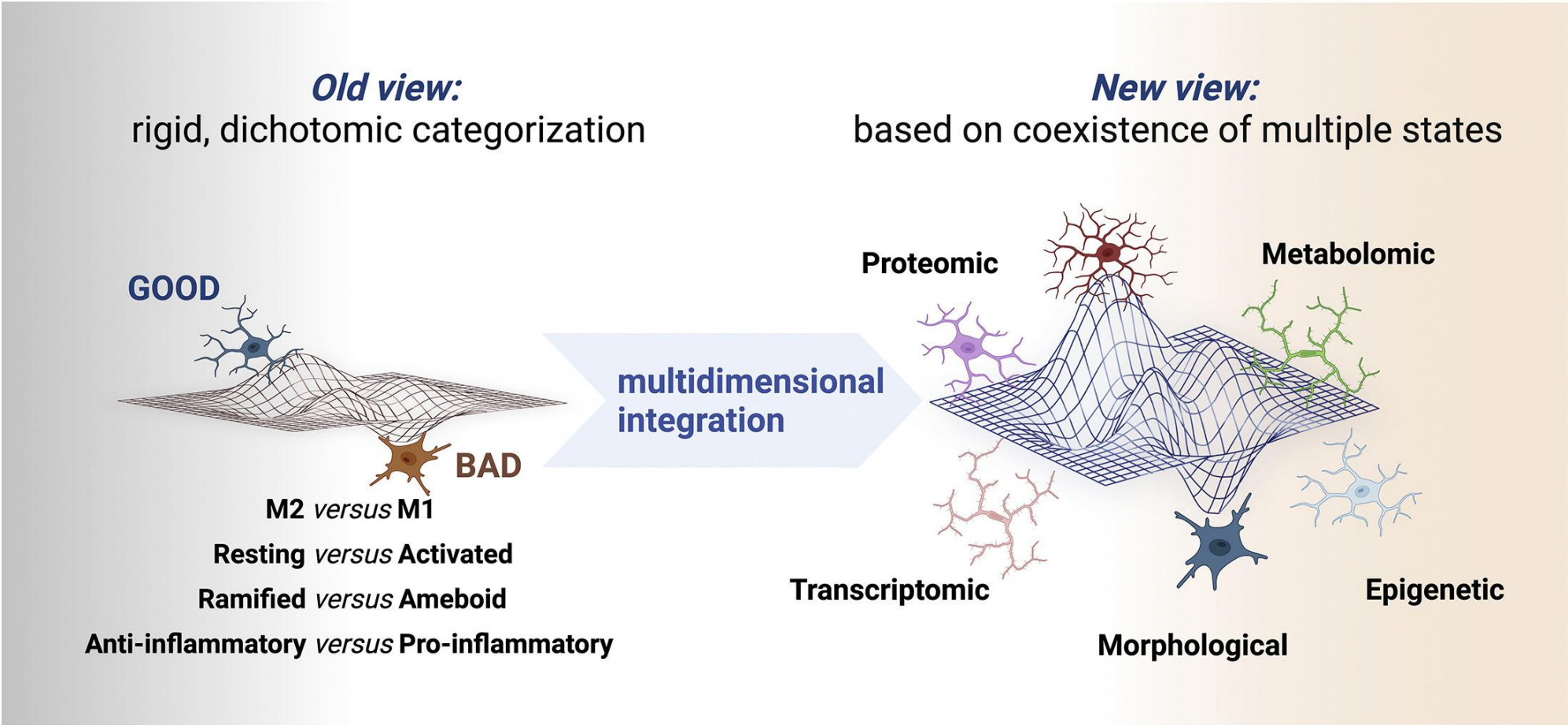
Phagocytosis, surveillance, and capacity for releasing soluble factors are core properties through which microglia contribute to key biological functions.

Microglia activation

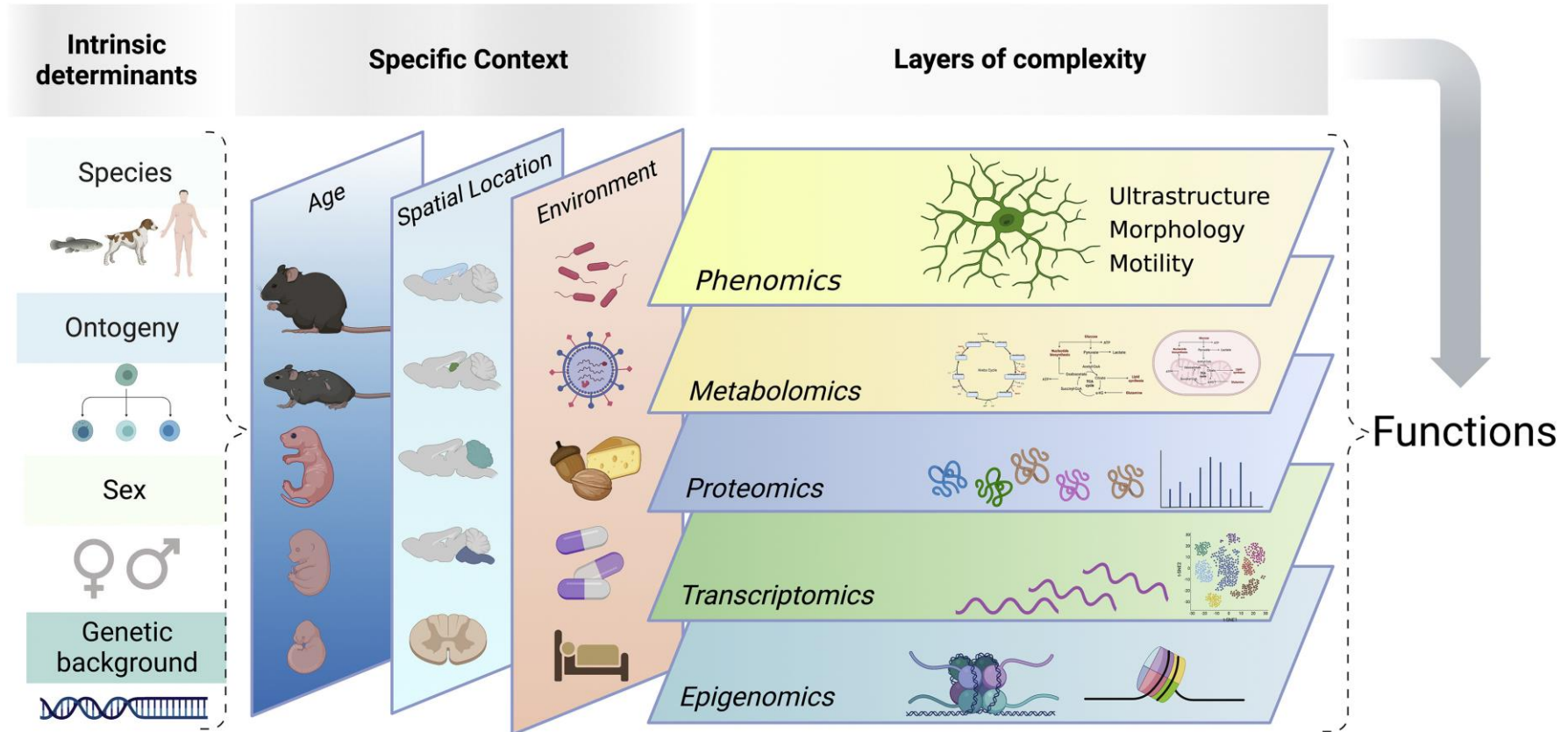


- Homeostatic microglia constantly and actively scan their environment for exogenous or endogenous signals indicating a threat to the homeostasis.
- Appearance of “activating” signals can then trigger transitions to alerted and activated states. Cells can commit to **distinct reactive phenotypes** depending on the challenging stimuli.
- Initial **reactive phenotypes** with **defense** orientation may **convert** to **repair-orientated** activity profiles.

Microglia have been traditionally framed into dichotomic categories, but the current integration of epigenetic, transcriptomic, metabolomic, and proteomic data favors a **multidimensional integration of coexisting states**.



Microglial states depend on **intrinsic determinants** (such as species, ontogeny, sex, or genetic background) as well as the **specific context** they inhabit, including age, spatial location, and environmental factors (such as nutrition, microbiota, pathogens, drugs, etc.). All together, these factors impinge on microglia at **multiple levels** (i.e., epigenomic, transcriptomic, proteomic, metabolomics, ultrastructural, and phenomic), which ultimately determine microglial functions

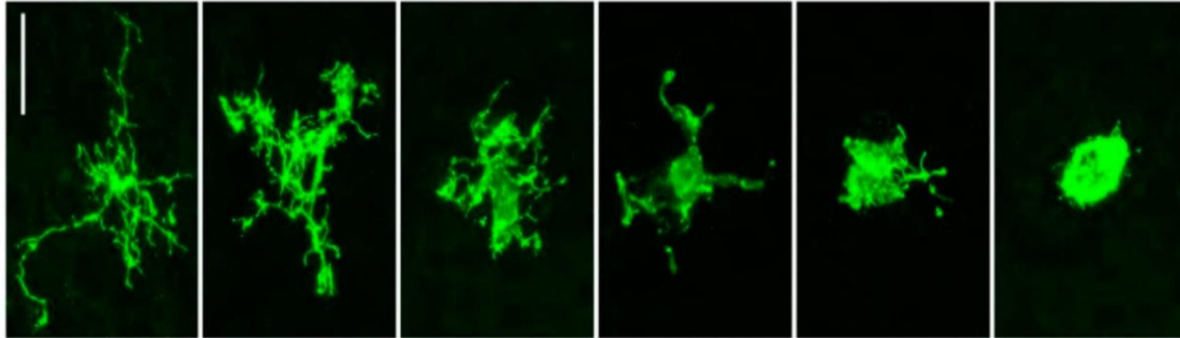


Microglia morphologies

In vivo

State of “Activating”

← Resting → “Activating” →



Rawlinson et al., 2020

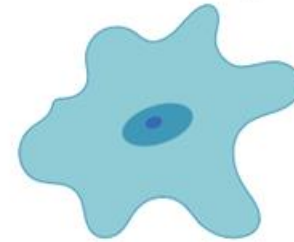
In vitro

Homeostatic



LPS-IFN γ

IL-4



Activated

Alternatively activated

“defensive”

“repairing”

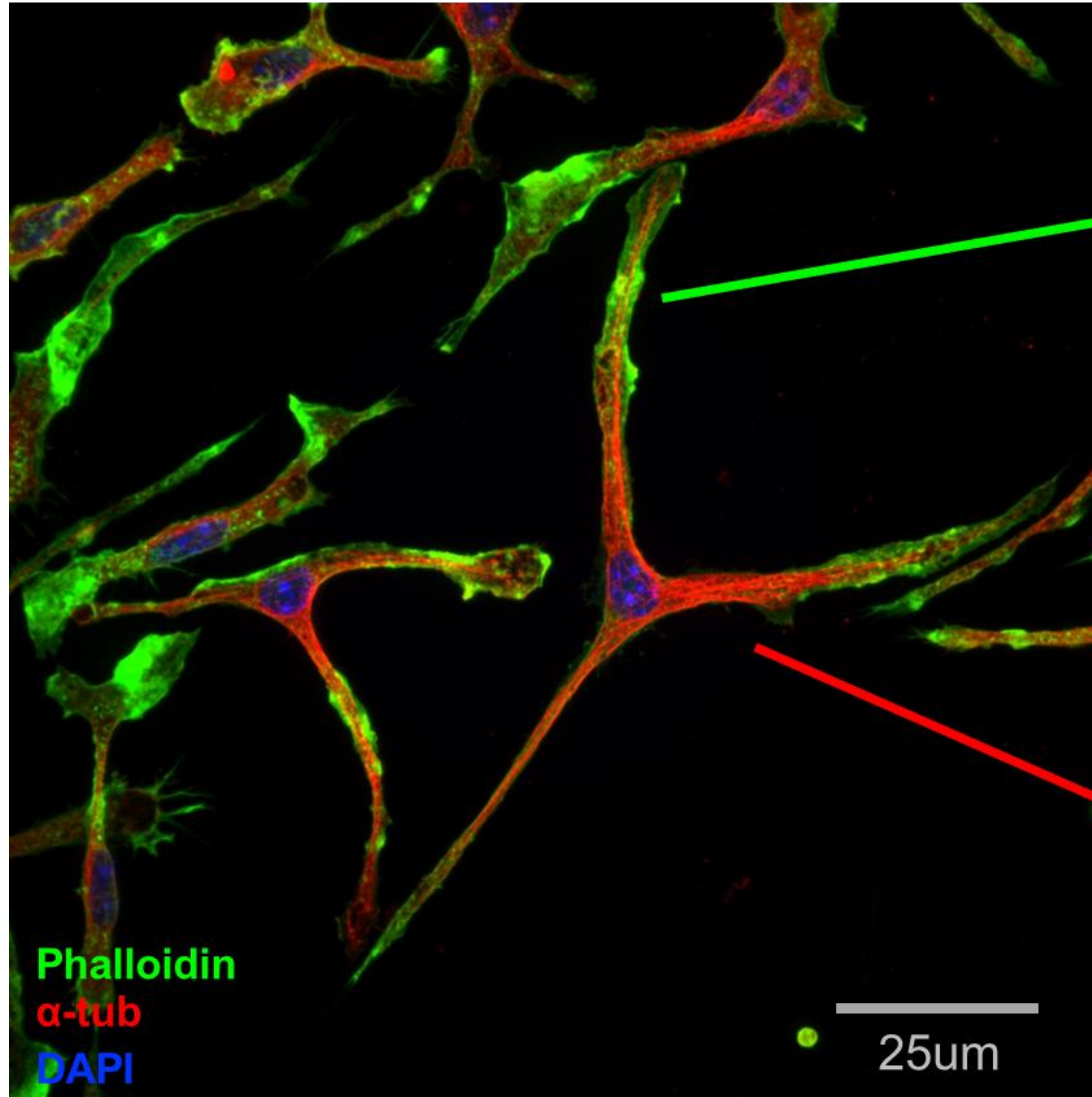
Why are we interested in microglia morphology?

- Microglia morphology remains a valid initial **proxy of microglia activation**, to be used as a **biomarker**
- Massive morphology changes are driven by **cytoskeletal rearrangements** → possibility to find **druggable targets**

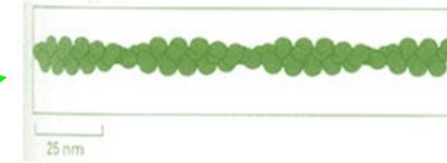
Although insufficient to define microglia state, morphology remains one of the first aspects to consider when approaching microglia functionality changes.

Microglia cytoskeleton

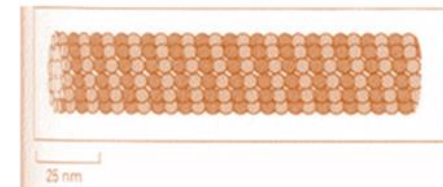
Primary murine microglia cells in culture



Actin



Microtubules



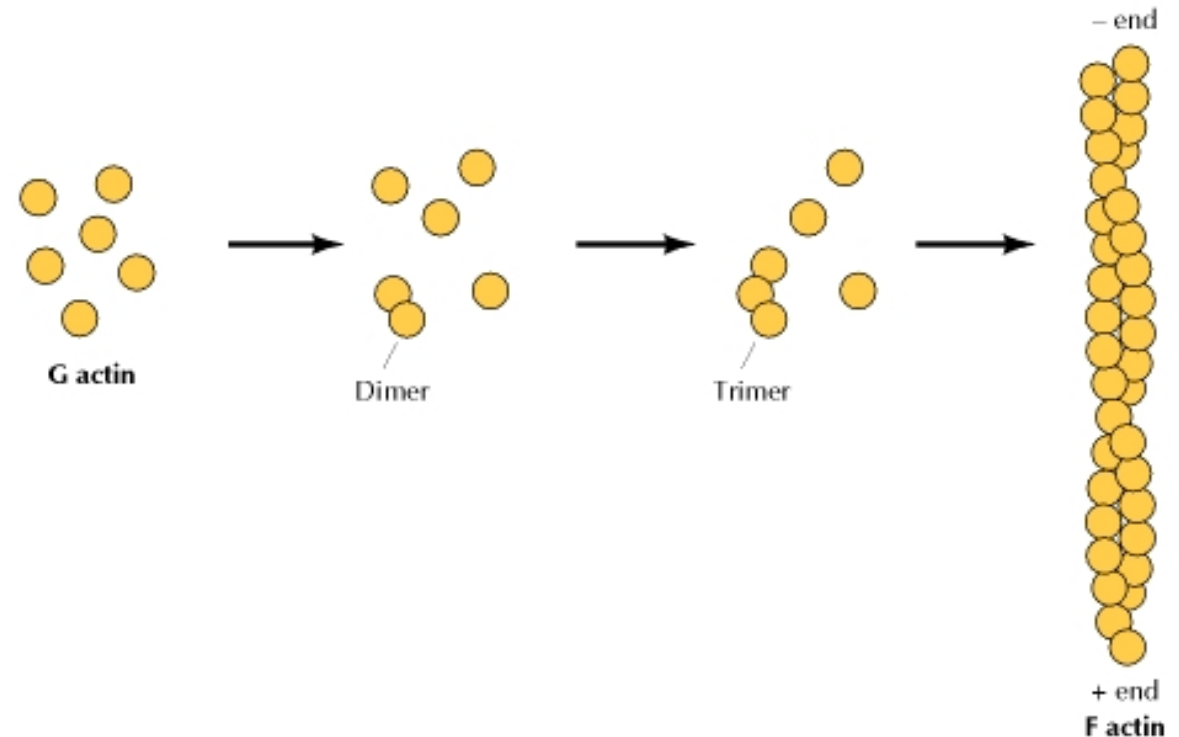
The actin cytoskeleton

Actin, a highly conserved protein found in all eukaryotic cells, plays a fundamental role in various cellular processes due to its dynamic structure and ability to form different types of filaments.

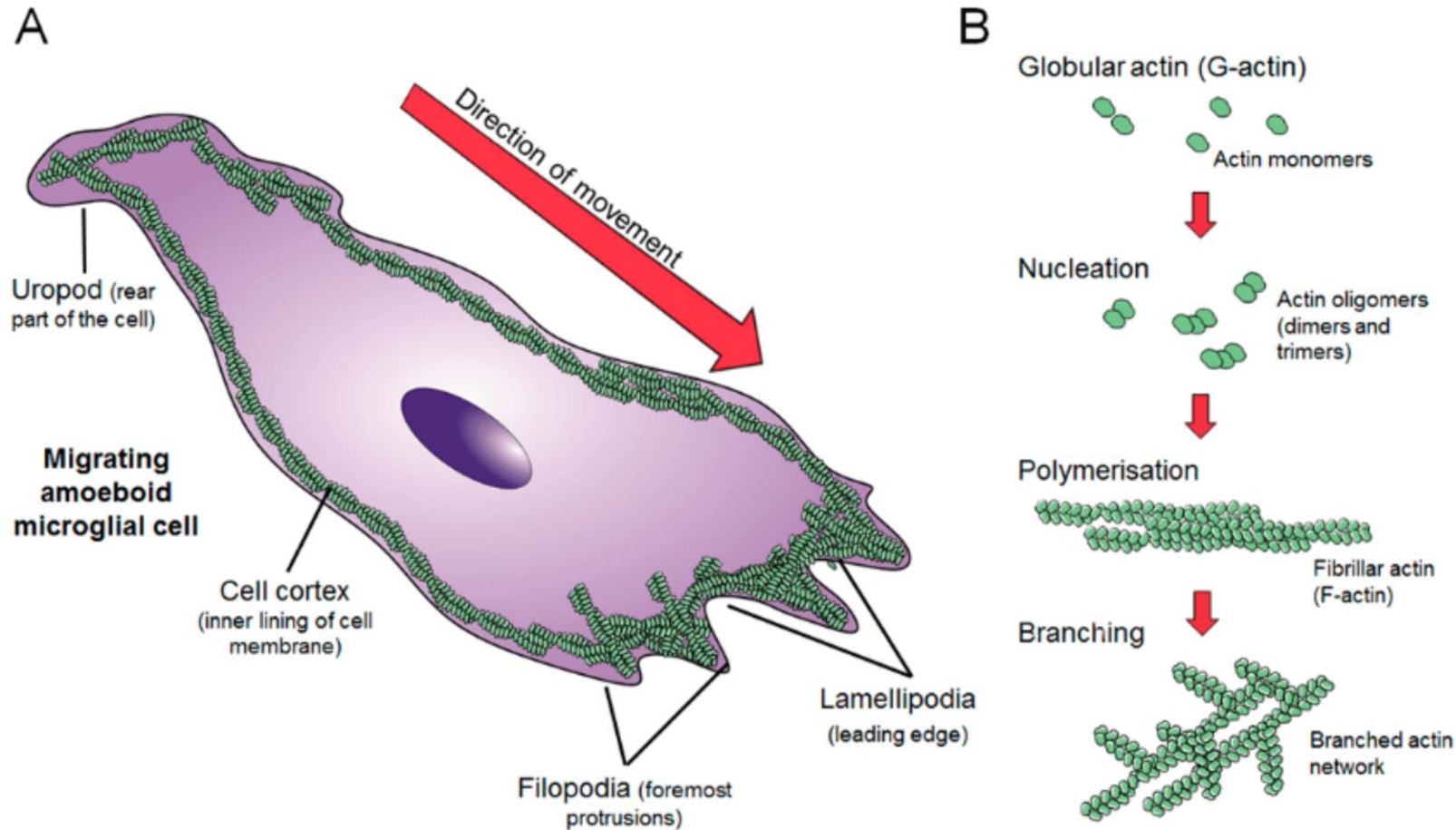
Actin monomeric form is known as **G-actin** (globular actin), and it has the potential to polymerize into **F-actin** (filamentous actin). The two forms of actin are maintained in a dynamic equilibrium.

This process is crucial for **cell motility, shape and division** and it is regulated by various actin-binding proteins.

Actin's versatility allows microglia to **rapidly remodel their cytoskeleton**, facilitating processes such as **migration, phagocytosis**, and the formation of cellular protrusions necessary for **nanoscale surveillance** of the brain environment



Actin and microglia migration



Different actin structures are present in microglia:

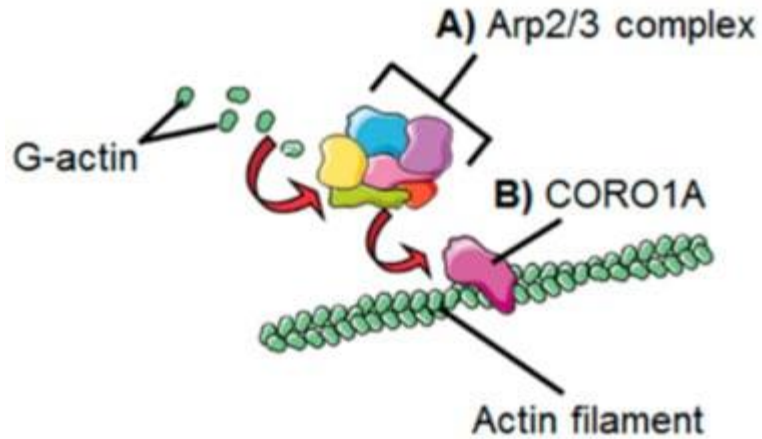
- **Cell cortex** (covering all the inner surface of the cell)
- **Filopodia** and **lamellipodia** (at the leading edge)
- **Uropod** (at the rear of the cell)

Mechanism of formation of the actin network includes:

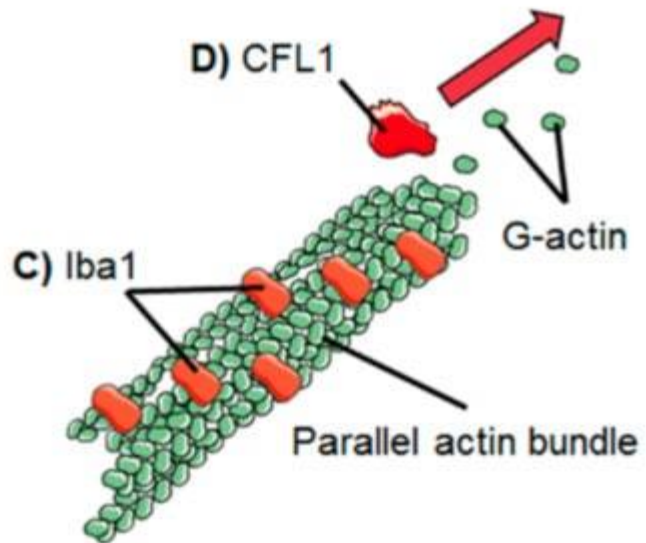
- Oligomerization of G-actin (nucleation)
- G-actin polymerization into F-actin
- Recruitment of additional globular actin to form **branches**

Actin filaments need to be branched to form the lamellipodia → branching is essential for **directed motility**

Branching



Cross-linking



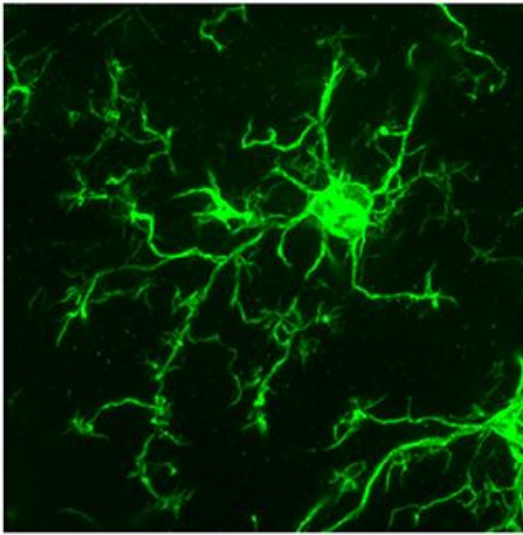
The process of **branching** at the lamellipodia and filopodia at the leading edge of moving cells is mainly controlled by the **Arp2/3** complex (actin-related proteins 2 and 3).

In the presence of ATP, the complex binds to the side of a filament and initiates a “subfilament” which stems from the “mother filament” at a characteristic angle of 70°. In microglia, this mechanism is regulated by **coronins**, such as coronin-1a (CORO1A).

To control the cell shape and movement, actin polymers are connected with each other by a process called **cross-linking**. Cross-linking allows the network to shape into more complex structures.

Iba1 promotes the formation of **parallel actin bundles**, scaffold-like structures that give shape to lamellipodia and filopodia.

Iba1

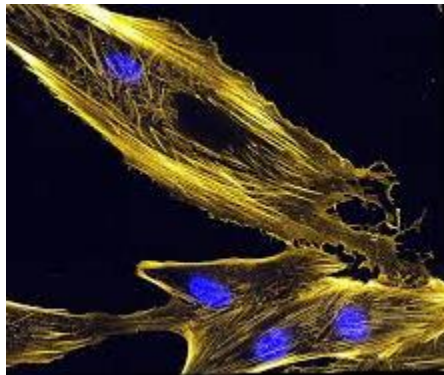
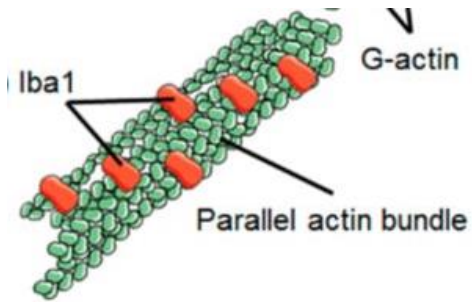


The ionized calcium-binding adapter molecule (Iba1) is also known as allograft inflammatory factor 1 (AIF1) is a widely used **microglia marker**

It is involved in **actin bundling** and **membrane ruffling**

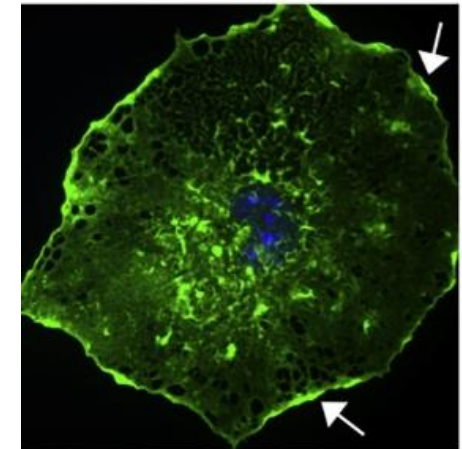
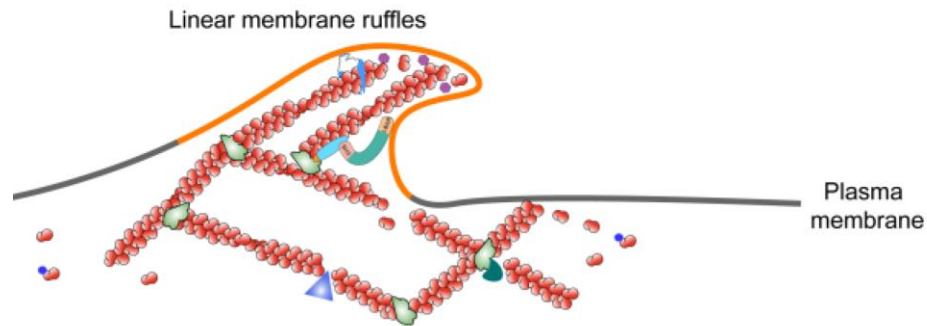
Actin bundles

Act like scaffolds, for lamellipodia and filopodia support



Membrane ruffling

Important for microglia migration and phagocytosis



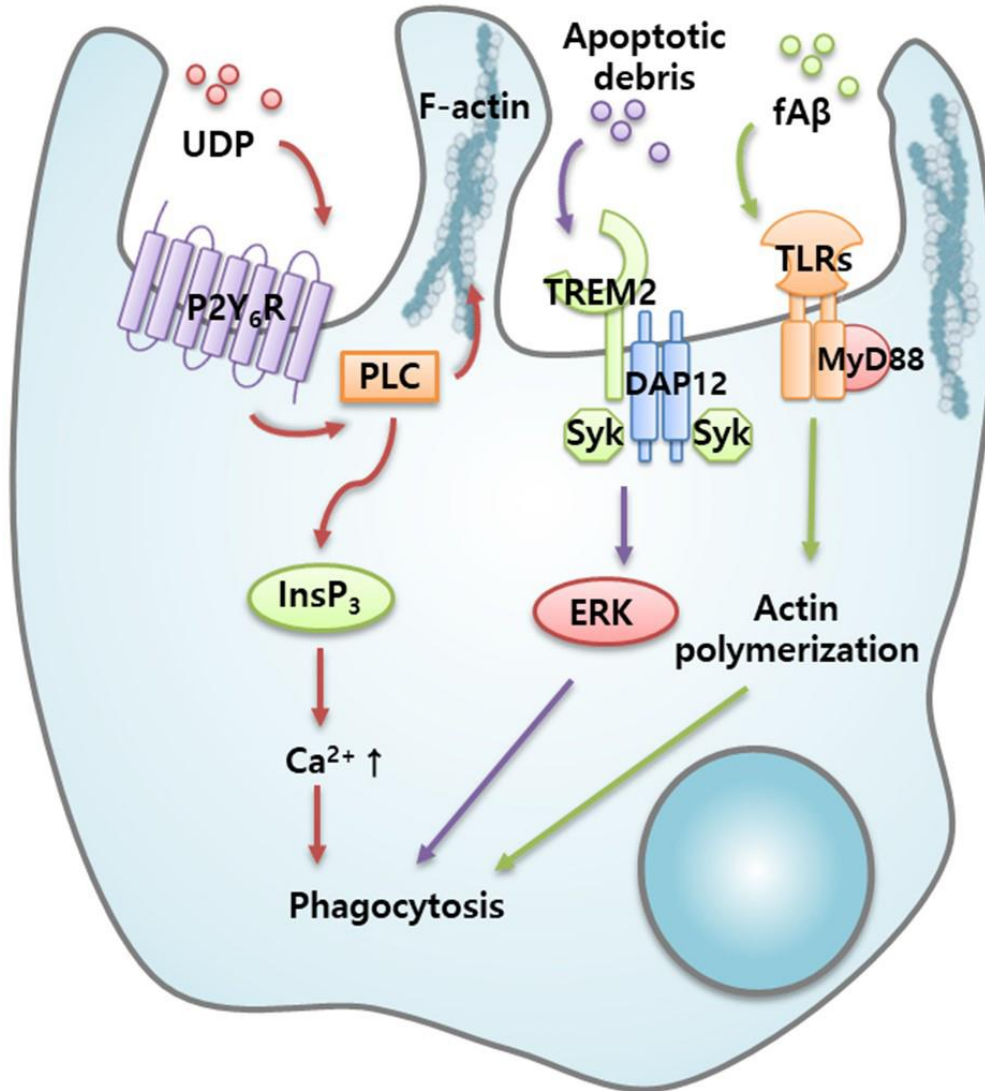
Actin and microglia phagocytosis

Specific **receptors** and signal transduction pathways that contribute to the reconstitution of actin proteins are utilized for microglial phagocytosis, such as:

- Toll like receptors (**TLRs**), high affinity receptors that bind to external microbial pathogens

- Trigger Receptors Expressed on Myeloid cells 2 (**TREM-2**), that recognize apoptotic cell substances, leading to the reconstitution of F-actin mediating the removal of apoptotic neurons

- Purinergic P2Y G-protein binding 6 receptor (**P2Y₆R**), that actively respond to uridine diphosphate UDP and activate phospholipase C to induce the synthesis of inositol 1,4,5-triphosphate and to release Ca²⁺ and promotes actin, mediating cytoskeletal polarization to form filopodia-like protrusions, thereby promoting cell phagocytosis



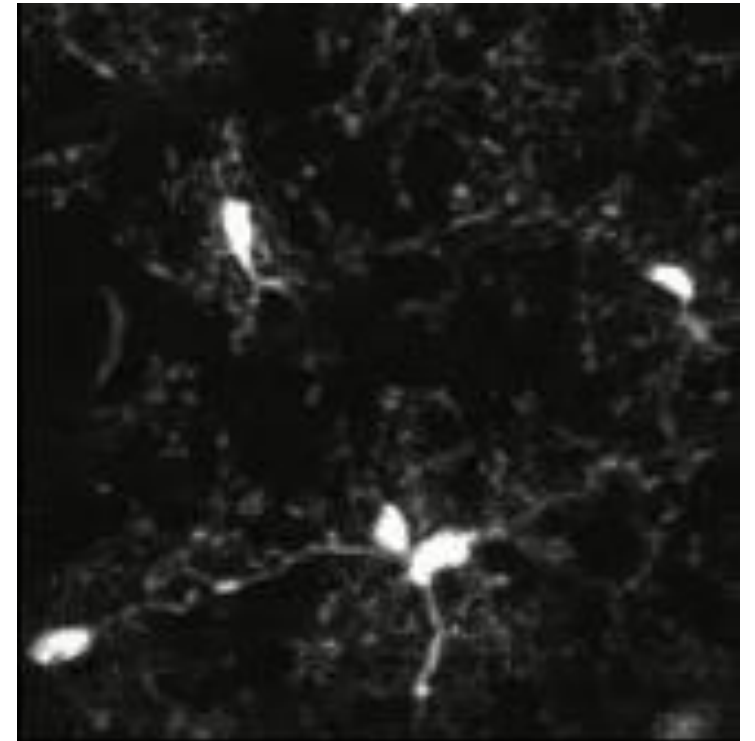
Actin and microglia surveillance

Video S1.

Filopodia motility at the tip of
large microglial processes

Duration: 12 minutes
45 seconds/frame

Microglia actively and continuously sense molecular cues within their local environment by using highly motile processes and ramified morphology

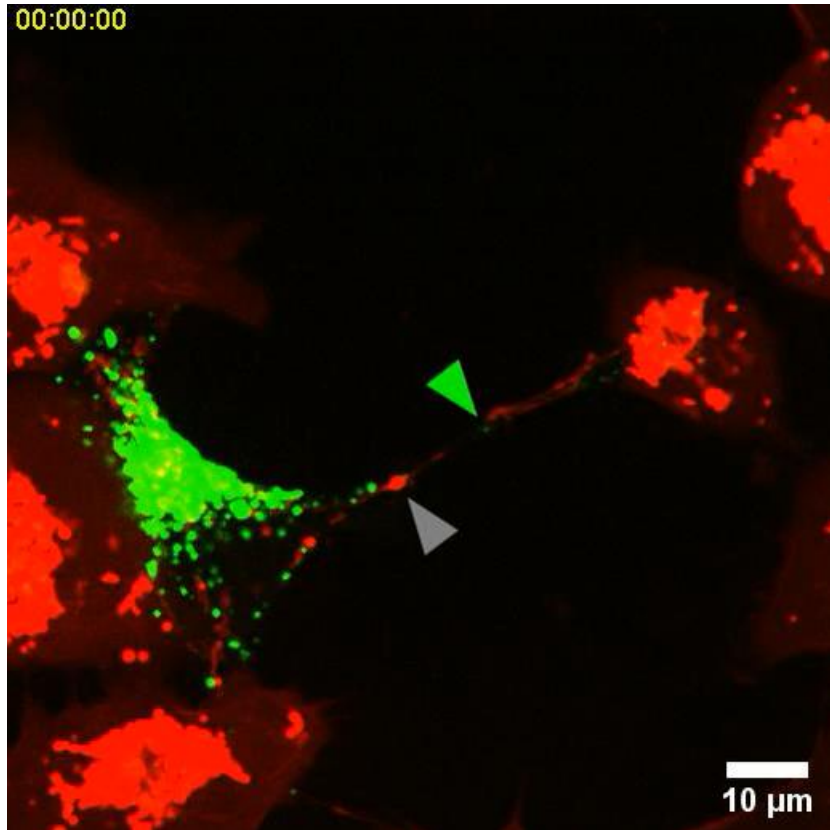


Bernier et al., 2019

Nimmerjahn et al., 2005

Microglia use **actin-dependent filopodia** to efficiently sample the brain parenchyma

Actin and microglia tunnelling nanotubes



Tunnelling Nanotubes (TNTs) facilitate **contact-mediated intercellular communication** over long distances.

TNTs are thin, membrane-enclosed, **F-Actin-rich protrusions** able to transfer cargoes of different kinds between the connected cells, like Ca^{2+} signals, messenger- and micro-RNAs, organelles such as lysosomes and mitochondria, pathogens, apoptotic signals, and protein aggregates

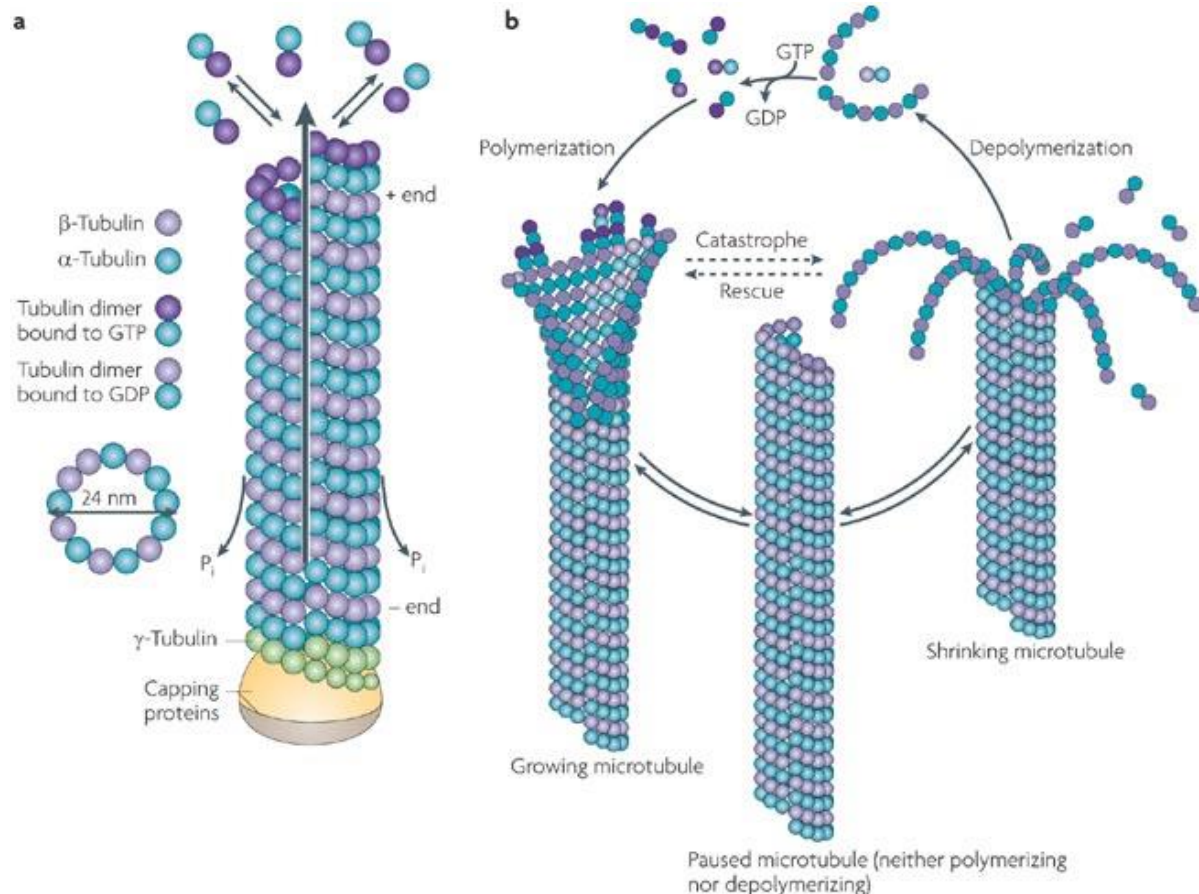
Bi-directional movement of α -Syn (green and yellow arrowheads) from neuronal cell to microglia and mitochondria (gray) in the opposite direction

Microglia actin cytoskeleton

- Actin is fundamental to drive rapid rearrangements of microglia cytoskeleton
- It is important for microglia **migration**, as it forms microglia lamellipodia and filopodia at the leading edge of migrating amoeboid microglia. The main involved processes are actin polymerization, branching and cross-linking that allow the formation of actin bundles and membrane ruffles.
- Many receptors on microglia cell surface trigger actin polymerization that finally allow **phagocytosis**
- Specialized microglia protrusions are formed mainly by actin, such as the filopodia that allow microglia **nanoscale surveillance**, and **tunnelling nanotubes** to drive contact-mediated cell communication.

The microtubule cytoskeleton

Microtubules (MTs) are dynamic structures composed of α/β -tubulin heterodimers, playing diverse roles in cell shape maintenance, intracellular transport and cell division



α/β -tubulin heterodimers string together to form the **protofilaments**.

Thirteen protofilaments form the microtubule

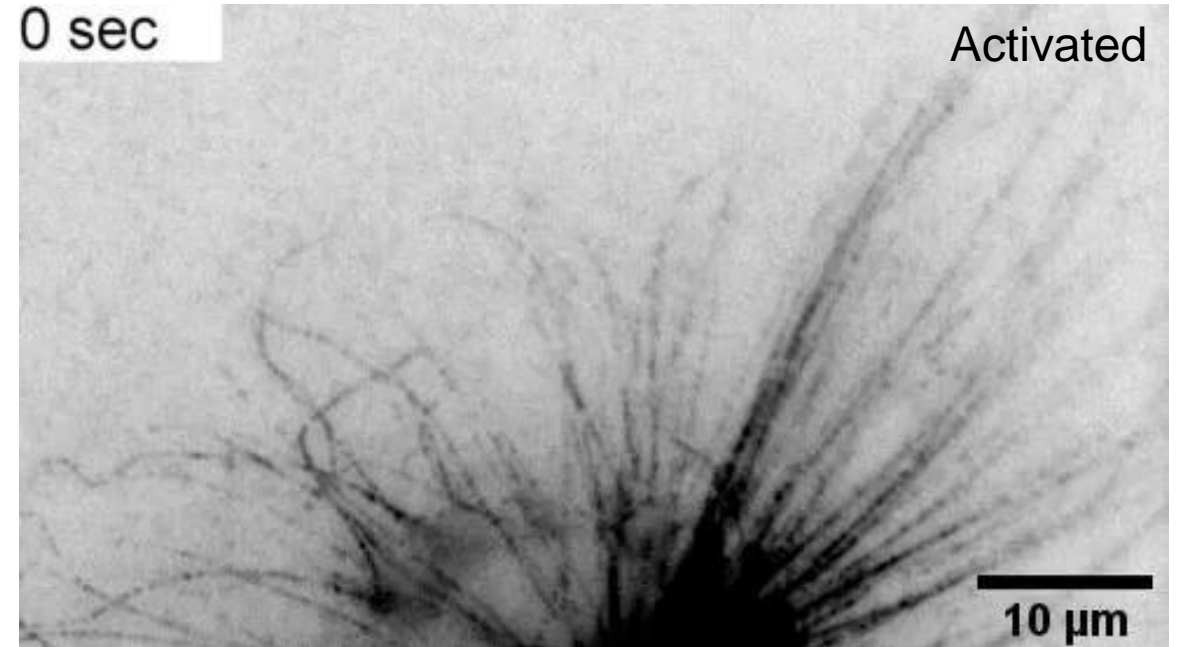
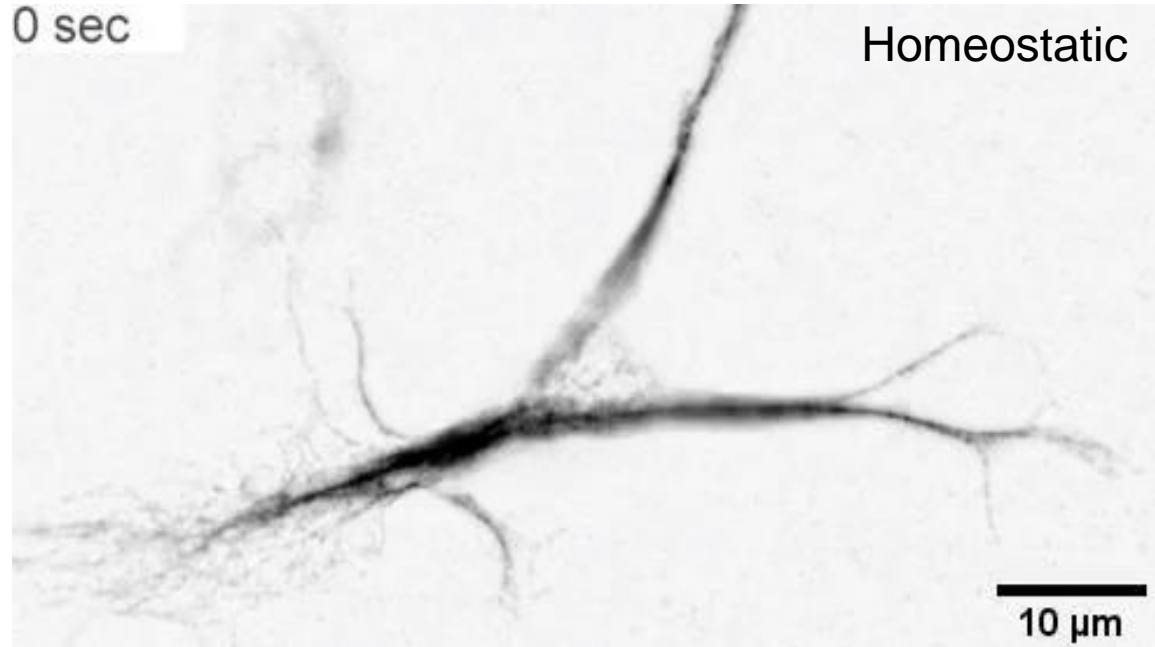
MTs are dynamics, with GTP-dependent reactions constantly adding and subtracting tubulin dimers at both ends of the filament.

One end grows more rapidly and is called the **plus end**, whereas the other end is known as the **minus end**.

The MT minus ends are anchored in structures called **microtubule organizing centers (MTOCs)**.

The primary MTOC in a cell is called the **centrosome**, and it is usually located adjacent to the nucleus.

Microtubule dynamics and microglia functions



MTs are dynamics, with GTP-dependent reactions constantly adding and subtracting tubulin dimers at both ends of the filament

MTs are usually associated with sustaining long-term cellular changes, as cell division and intracellular transport, but their dynamicity allows for cell motility as well

Microglia activation increases MT dynamicity

MT polarity and molecular transport

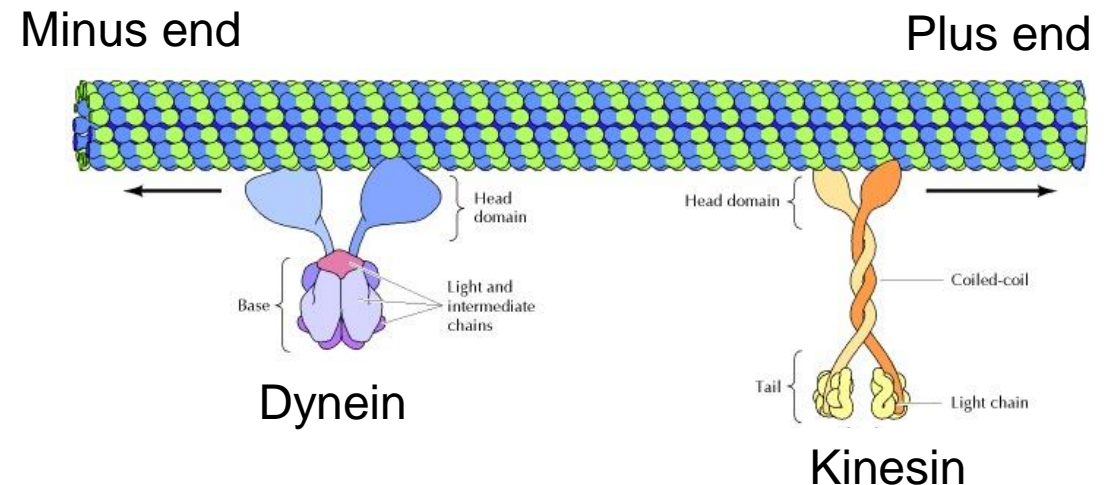
One MT end grows more rapidly and is called the **plus end**, whereas the other end is known as the **minus end**

The plus end can be visualized with end-binding proteins (EB)



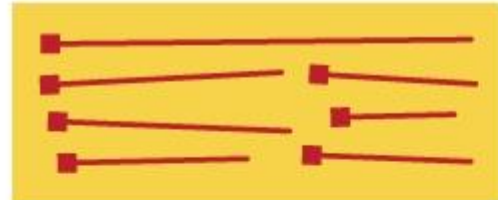
Movement along microtubules is based on the action of motor proteins that utilize energy derived from ATP hydrolysis to produce force and movement. Members of two large families of motor proteins—the **kinesins** and the **dyneins**— are responsible for powering the variety of movements in which microtubules participate.

Kinesin and dynein move in opposite directions along MTs, toward the plus and minus ends, respectively

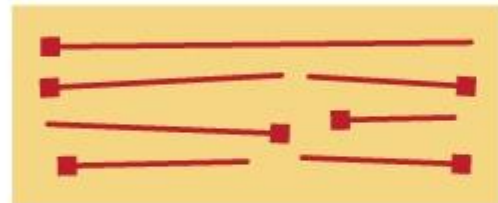


Microtubule polarity in neurons

Microtubule Polarity



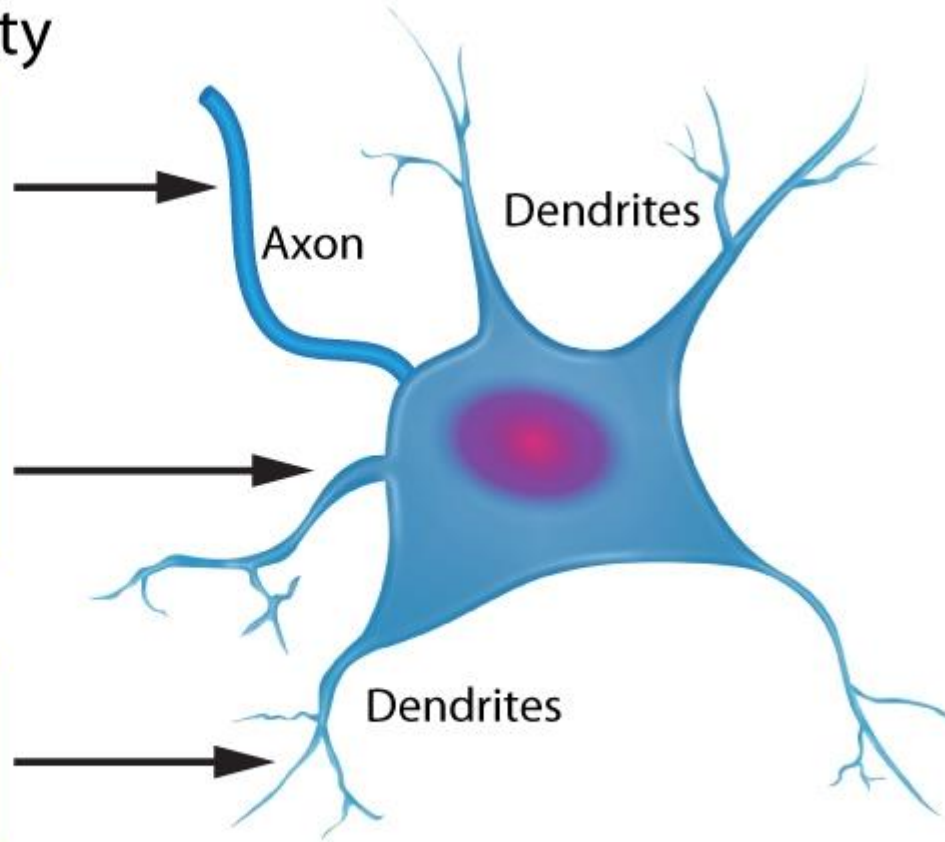
Axons: > 95% plus-ends out



Proximal dendrites: mixed orientations

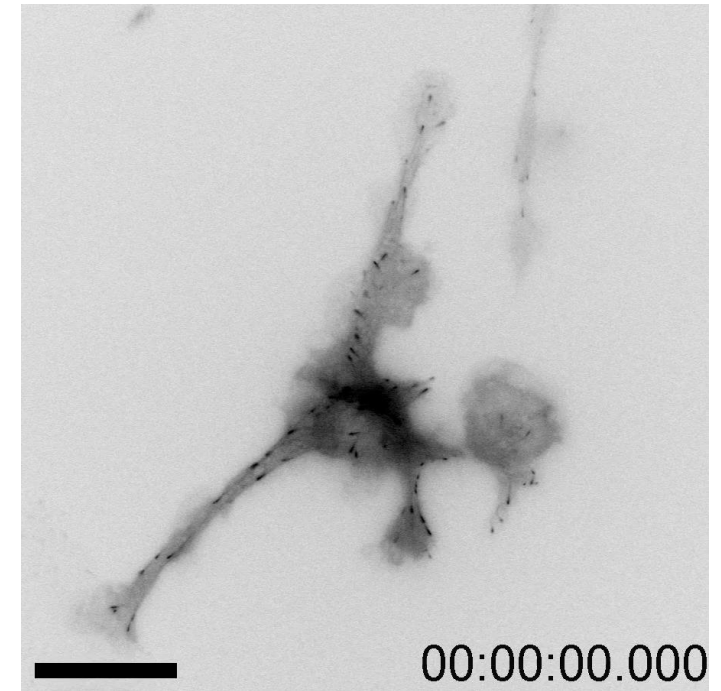
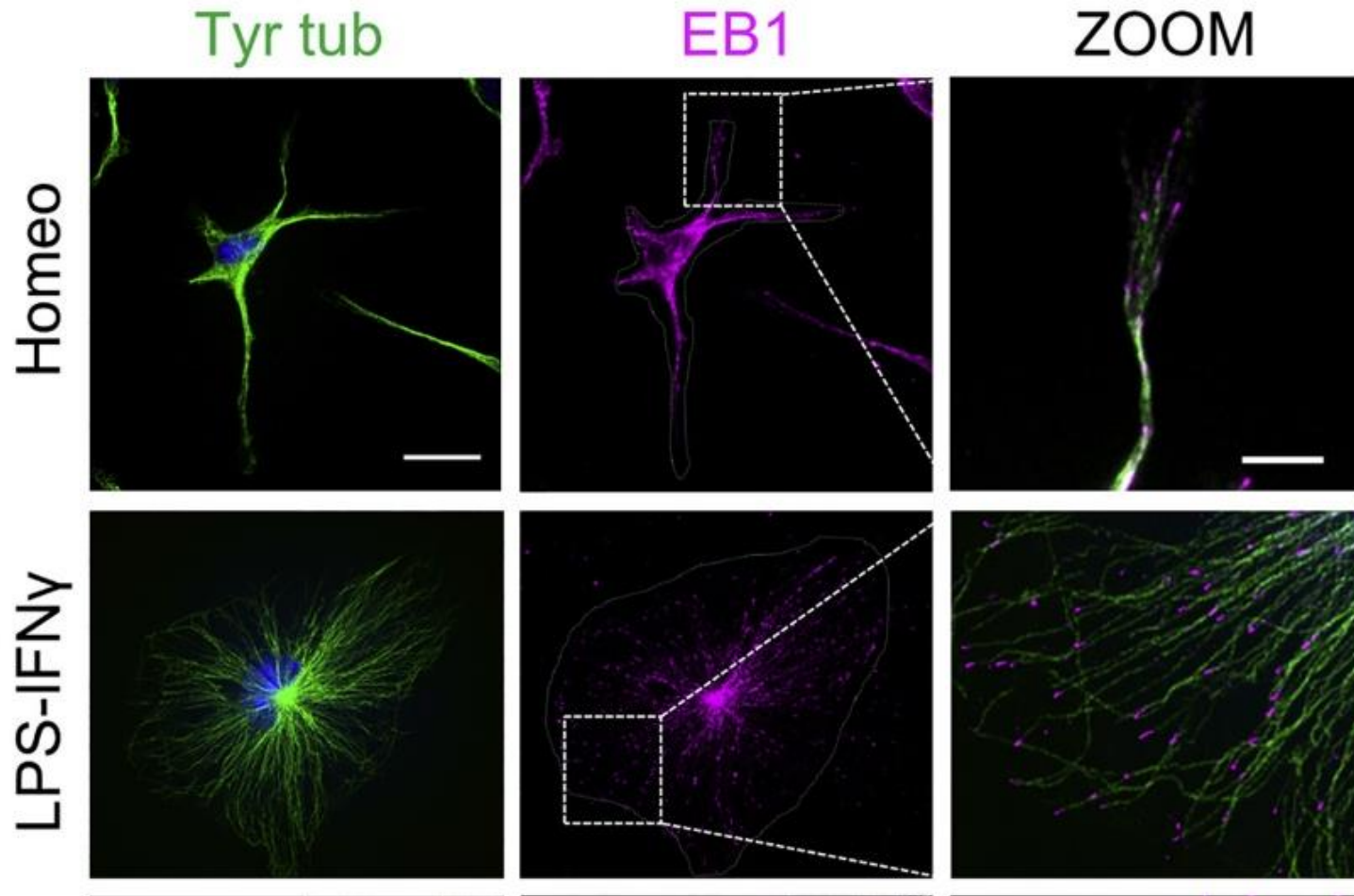


Distal dendrites: majority plus-ends out



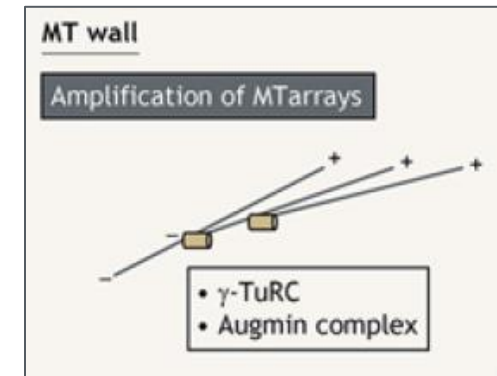
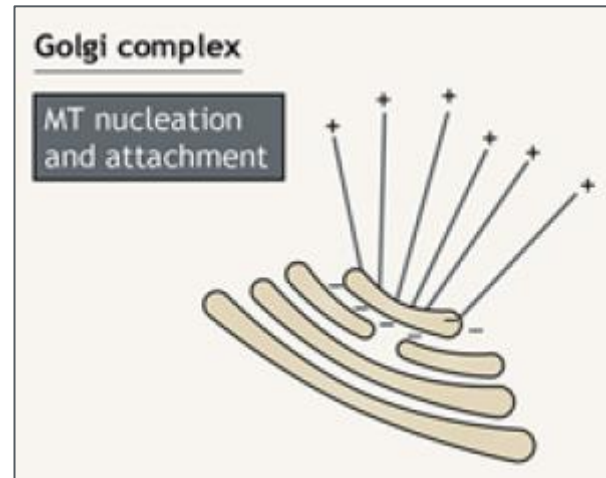
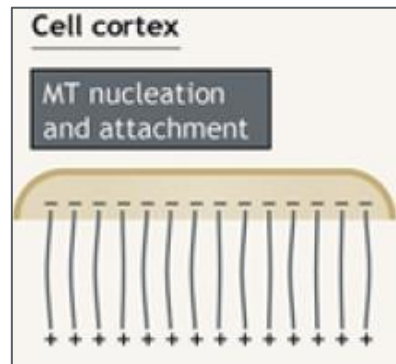
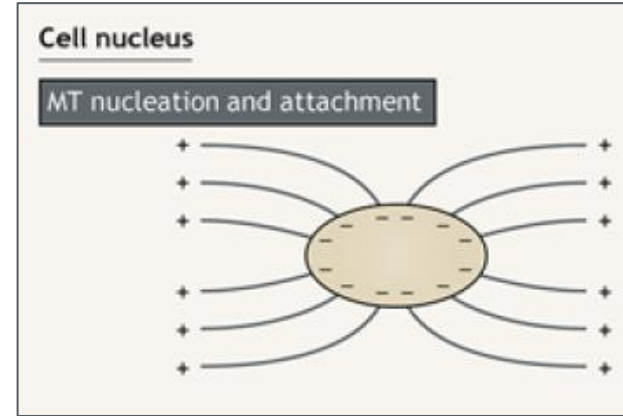
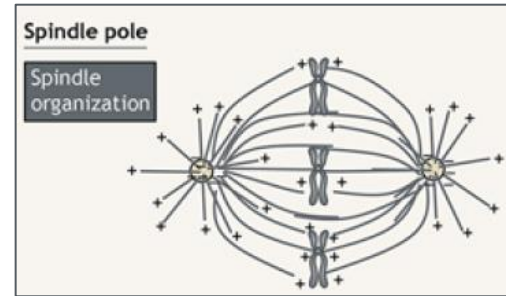
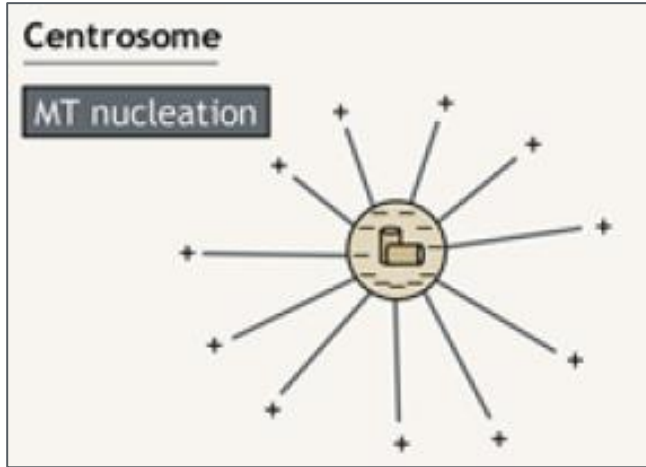
MT polarity is **uniform in the axon, mixed in the dendrites**. This allows for the control of the direction of the molecular cargos along the MTs.

This peculiar pattern allows for homogeneous transport along the axon (to send signals), and more flexible and heterogeneous transport in the dendrites (to receive signals)

B

Microglia MTs display mixed polarity when ramified, uniform polarity when amoeboid

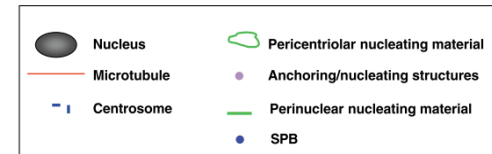
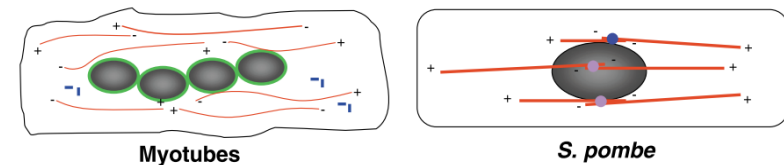
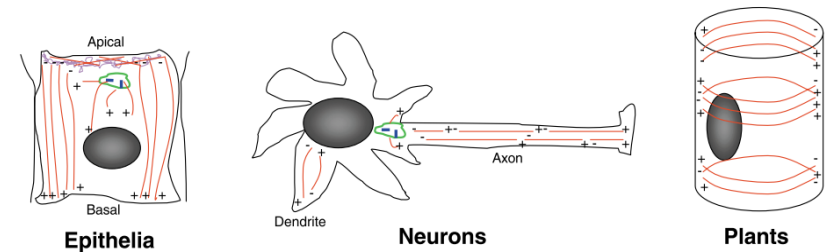
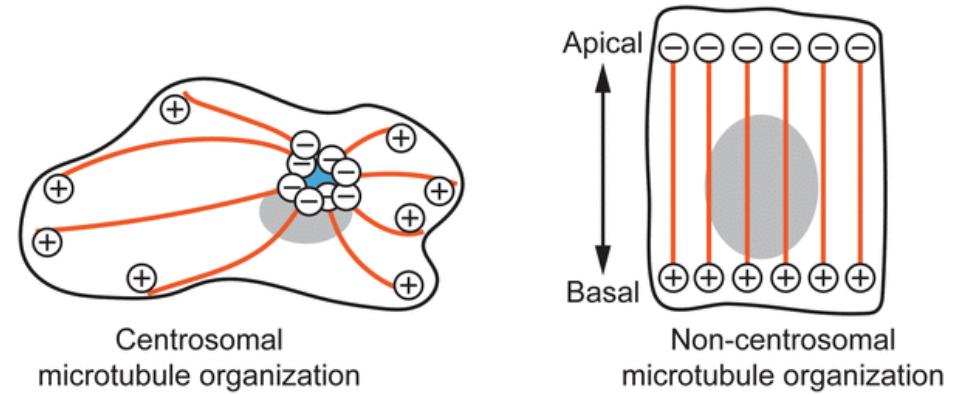
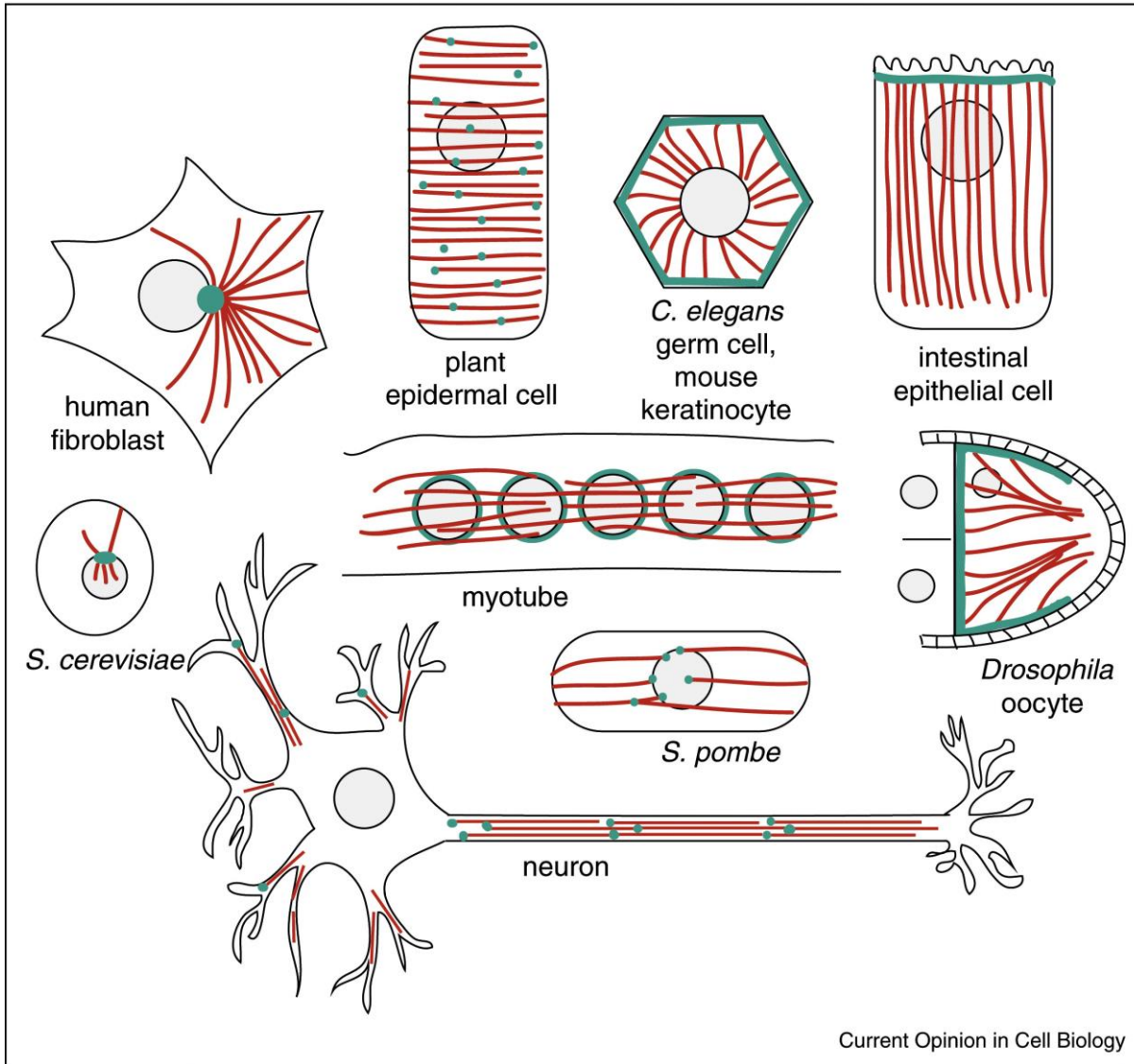
Microtubule organizing centres



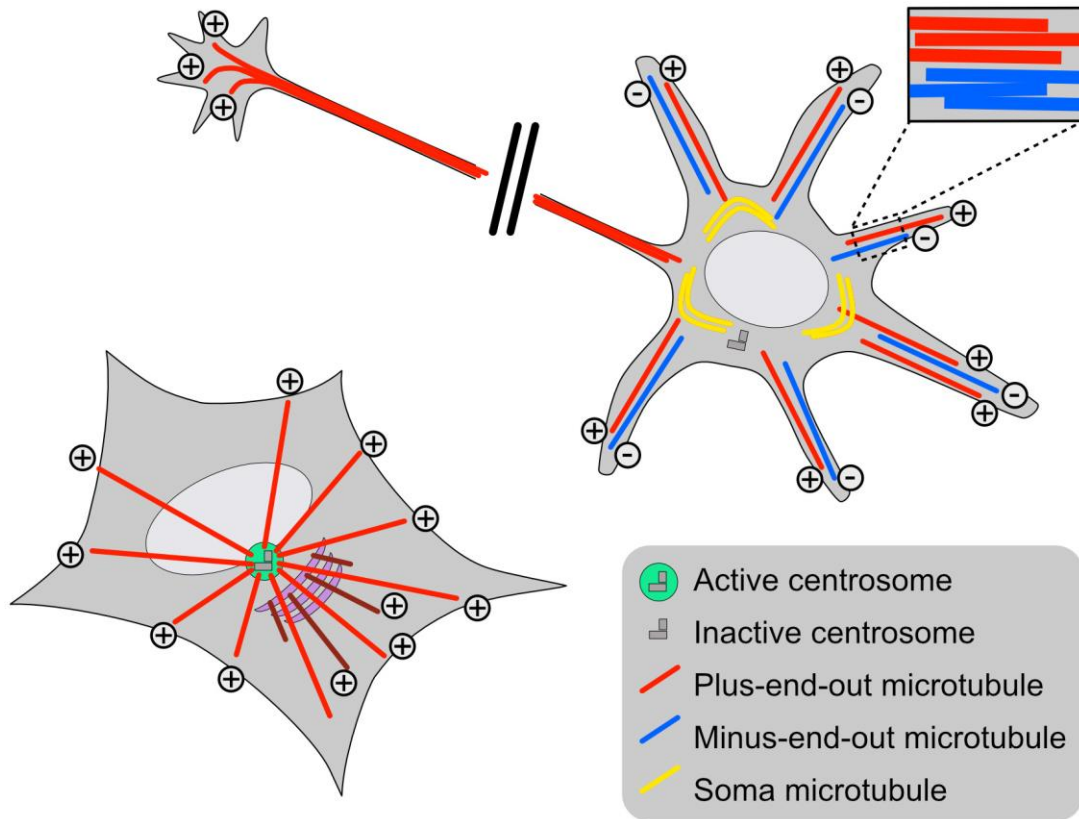
The main MTOC is the centrosome but there are many other structures that can anchor MTs and act as MTOCs

MTs are therefore distinguished between **centrosomal** and **non-centrosomal** MTs.

Differentiated animal cells often establish non-centrosomal MTOCs



Centrosomal and non-centrosomal microtubules



In differentiating, cycling cells, MTs are radially organized around the centrosome, the main MTOC. The centrosome is active.

In neurons, which are post-mitotic cells, centrosome is present but has lost MTOC activity.

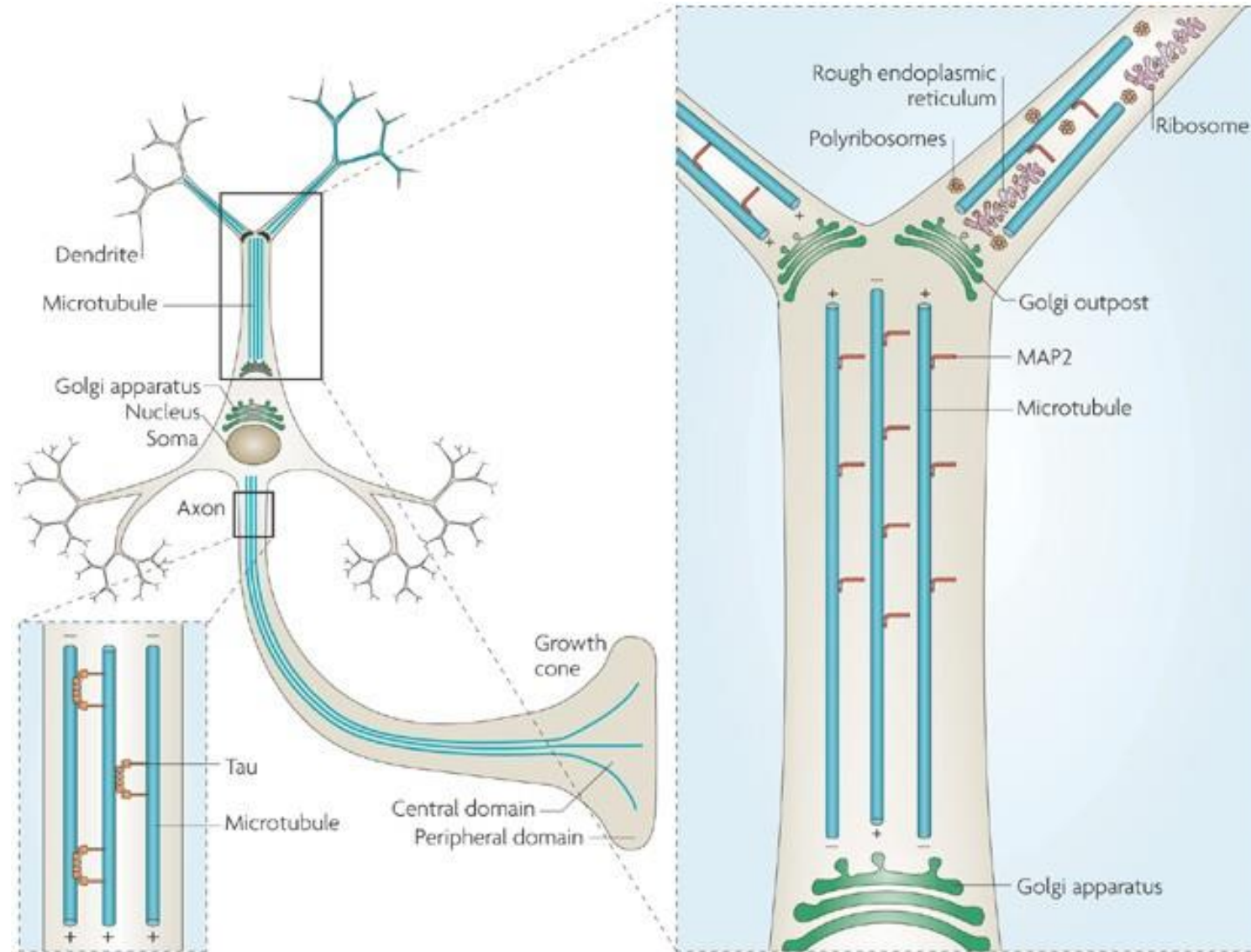
Neurons rely on local MT nucleation and stabilization to accommodate their extended morphology, such as axons and dendrites. This organization allows:

1.Support for Neurite Growth: Non-centrosomal microtubules enable the elongation of axons and dendrites.

2.Efficient Intracellular Transport: They ensure proper transport of organelles and molecules along long distances.

3.Structural Plasticity: They adapt to changes in neuronal architecture during development and synaptic plasticity.

The alternative MTOCs and the Golgi outposts

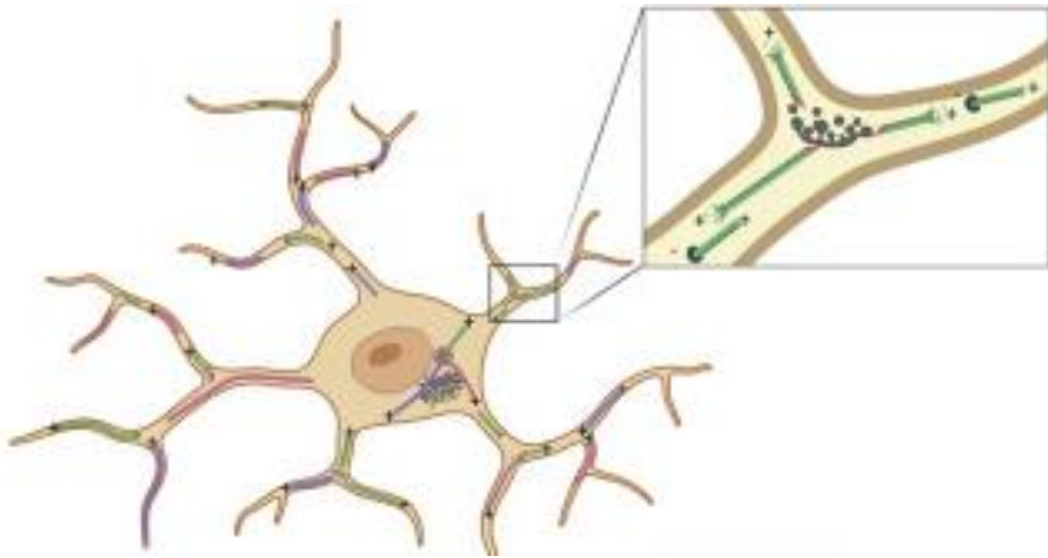


Non-centrosomal microtubules can be locally nucleated at alternative MTOCs, such as structures called **Golgi outposts**

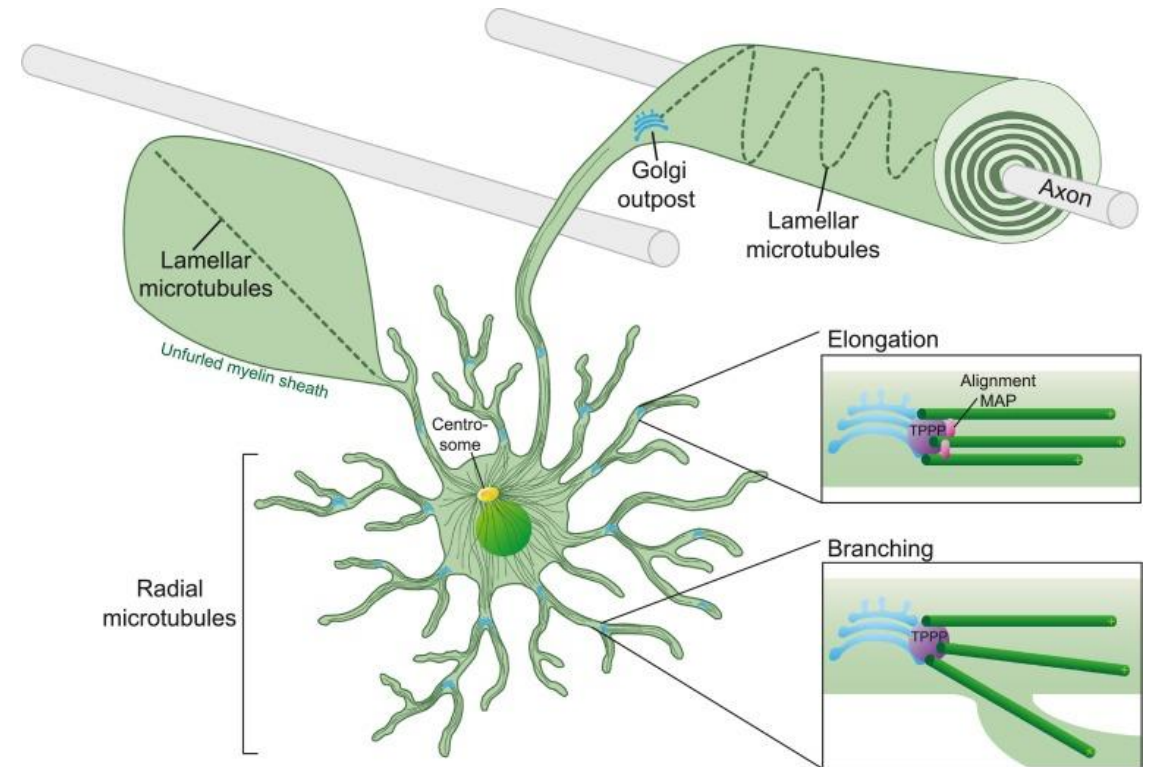
Glial Golgi outposts

Golgi outposts are present in glial cells such as oligodendrocytes and microglia

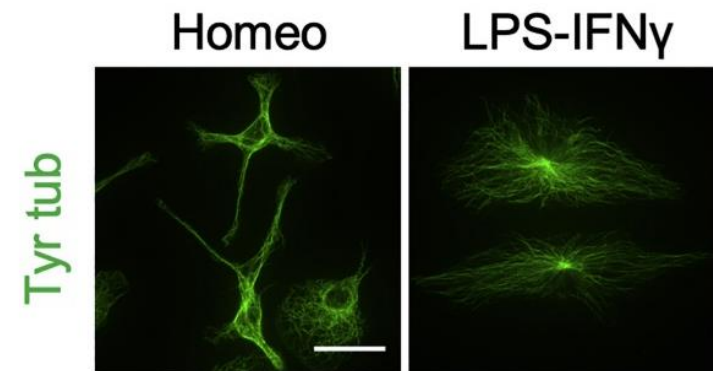
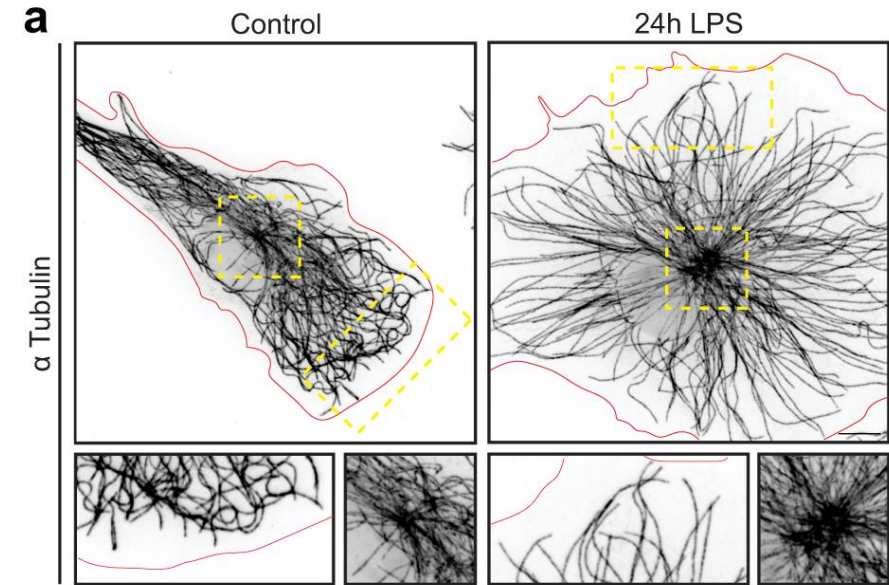
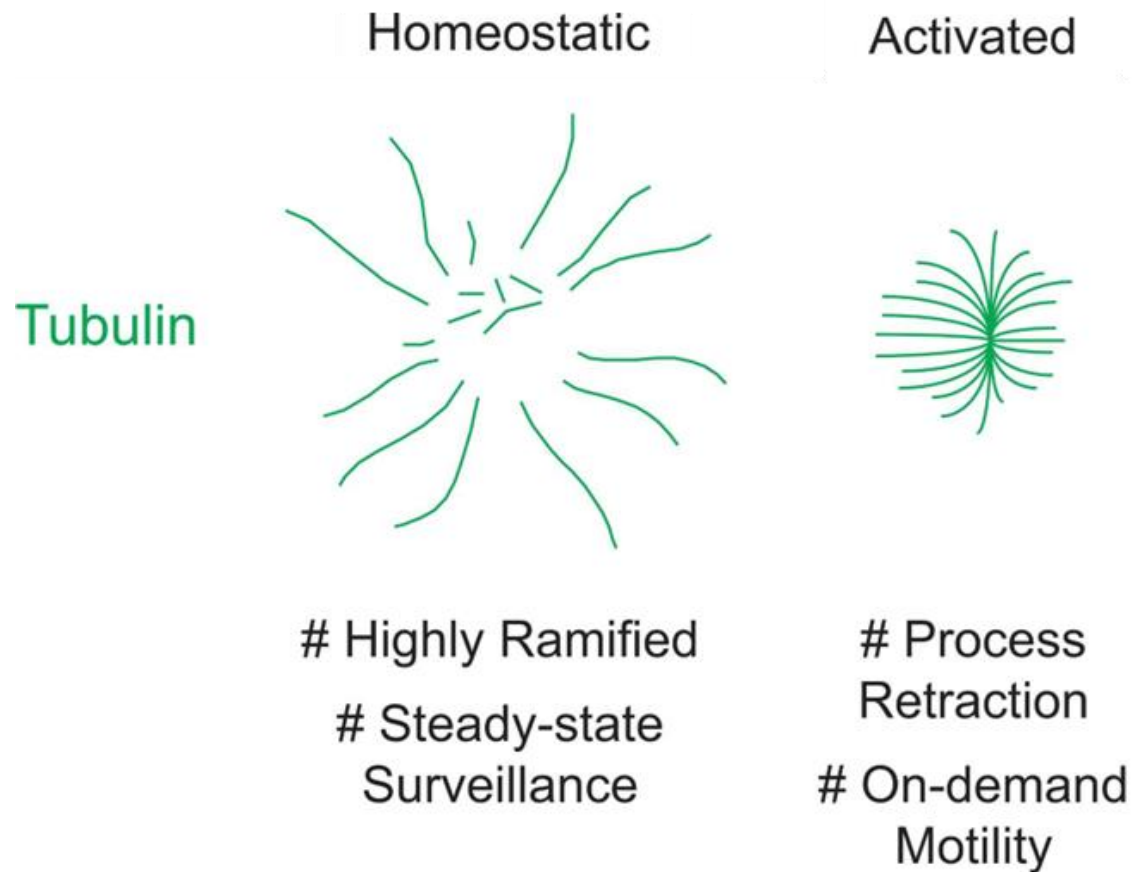
In microglia, they are important for MT elongation and branching in highly ramified morphologies



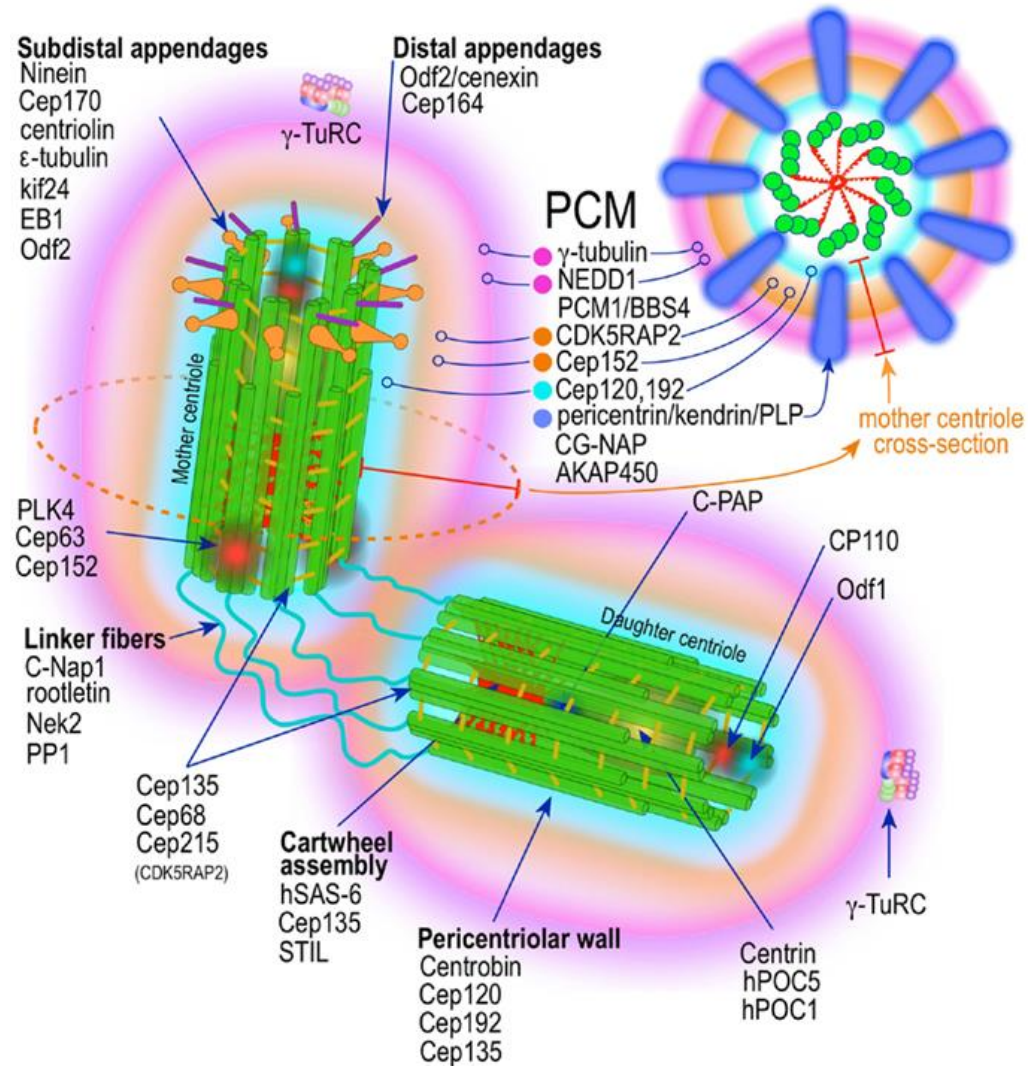
In oligodendrocytes, they are functional for MT elongation, branching and for the formation of myelin sheath



Microglia display different morphologies associated with different activation states, with **non-centrosomal** microtubules present in **ramified** morphologies, and **centrosomal** MT organization in **amoeboid** morphologies



The centrosome



The centrosome is the microtubule organizing center of animal cells, whereas the equivalent in yeast is called the 'spindle pole body'. The centrosome and the spindle pole body are the best-characterized microtubule organizing centers.

The centrosome has three important activities:

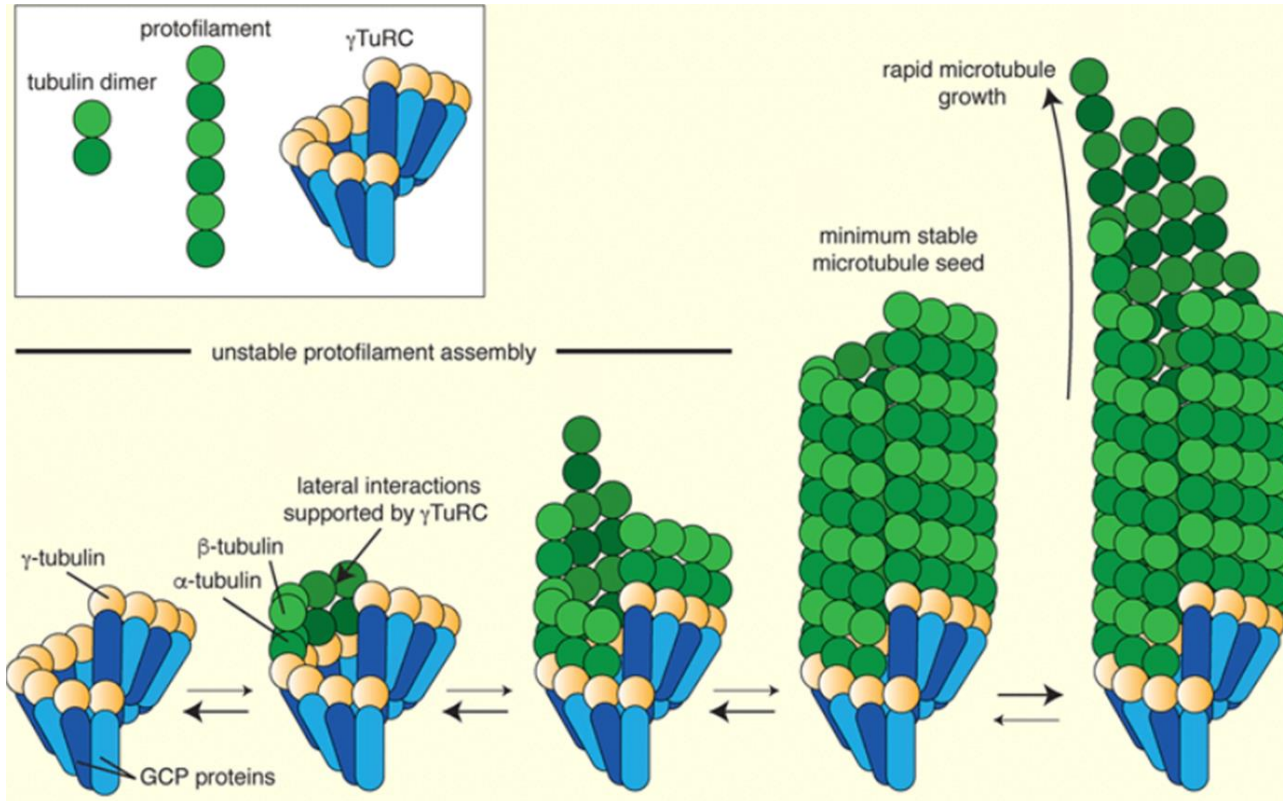
- it **nucleates** the polymerization of tubulin subunits into the long polymers that are **microtubules**;
- it **organizes** the nucleated **microtubules** into useful arrays;
- it **duplicates** once every cell cycle

The centrosome consists of two short cylindrical centrioles, usually oriented perpendicular to each other, surrounded by pericentriolar material (PCM)

The PCM contains elements that nucleate and organize microtubules, such as γ-tubulin

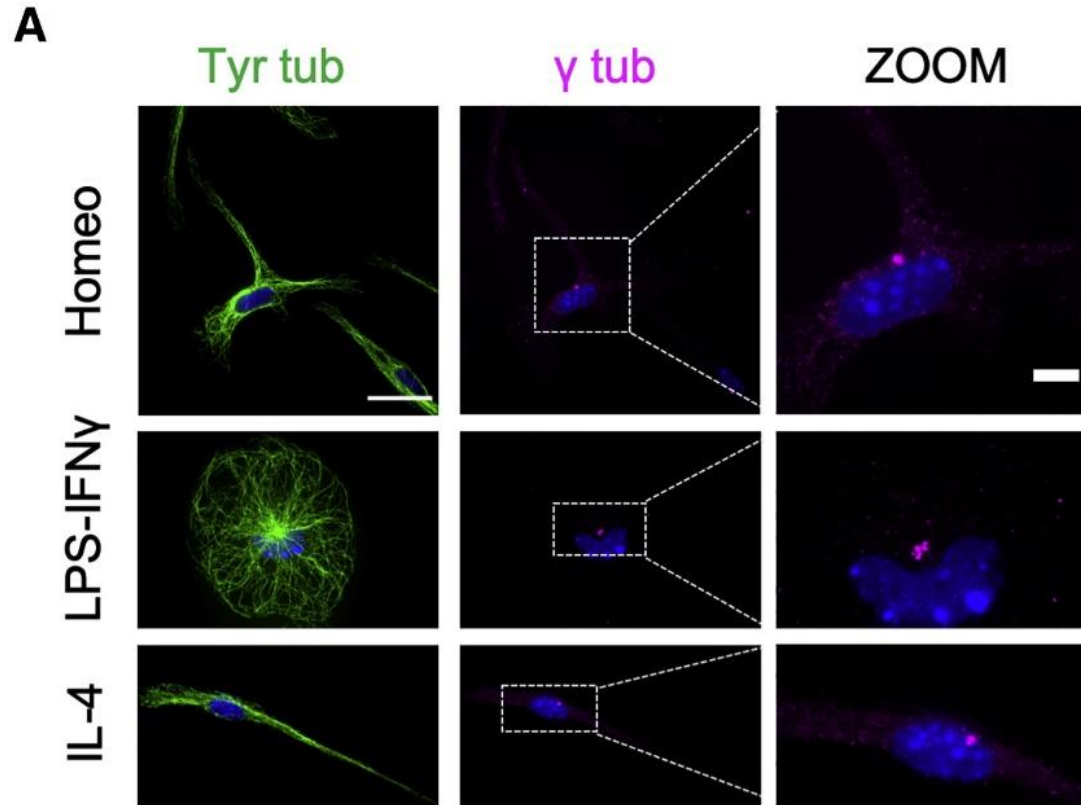
Microtubule nucleation

γ -tubulin is a conserved component of all microtubule organizing centers, localized to the pericentriolar material of the centrosome



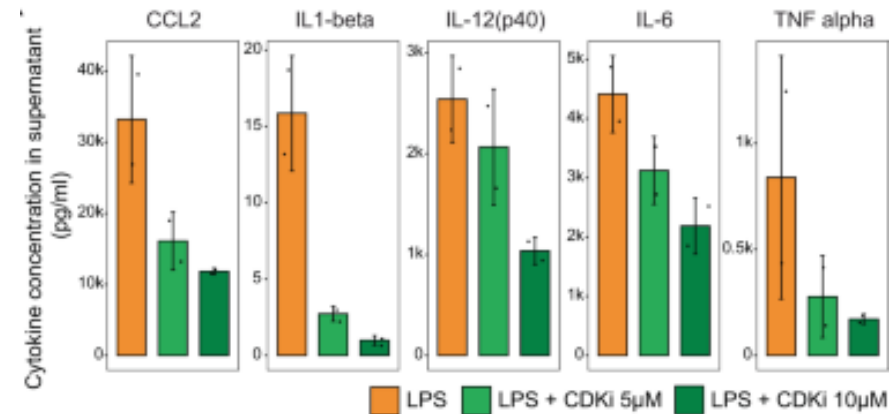
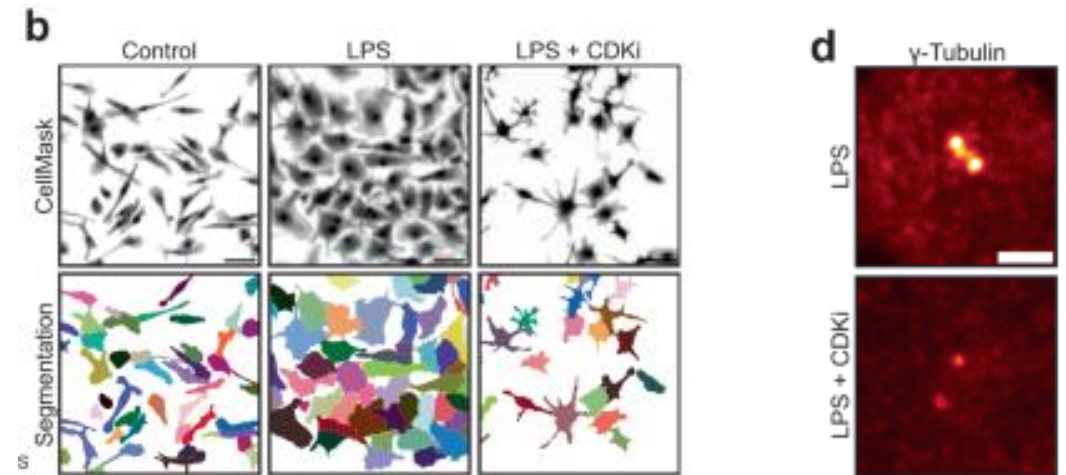
Microtubule nucleation occurs in the pericentriolar material, and γ -tubulin is required for this nucleation activity. The γ -tubulin is part of a large protein complex that is organized into an open ring of roughly 25 nm diameter, the same as the diameter of a microtubule

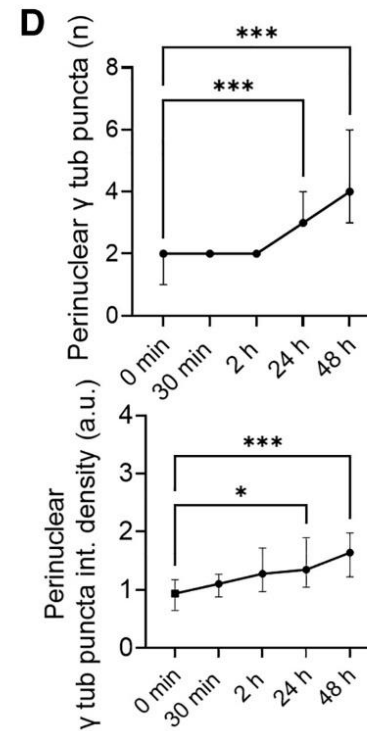
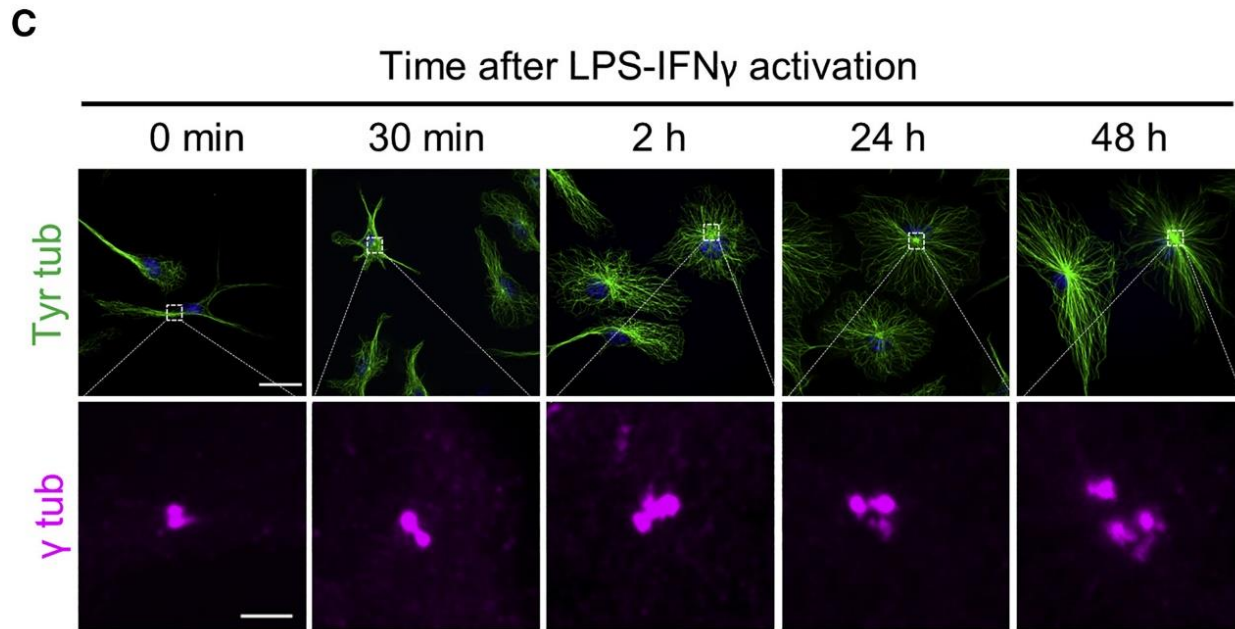
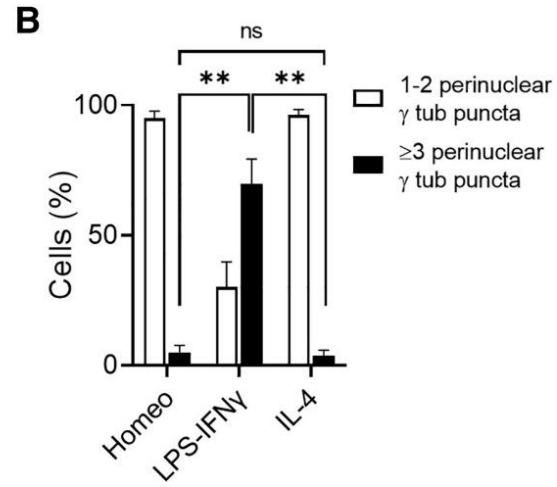
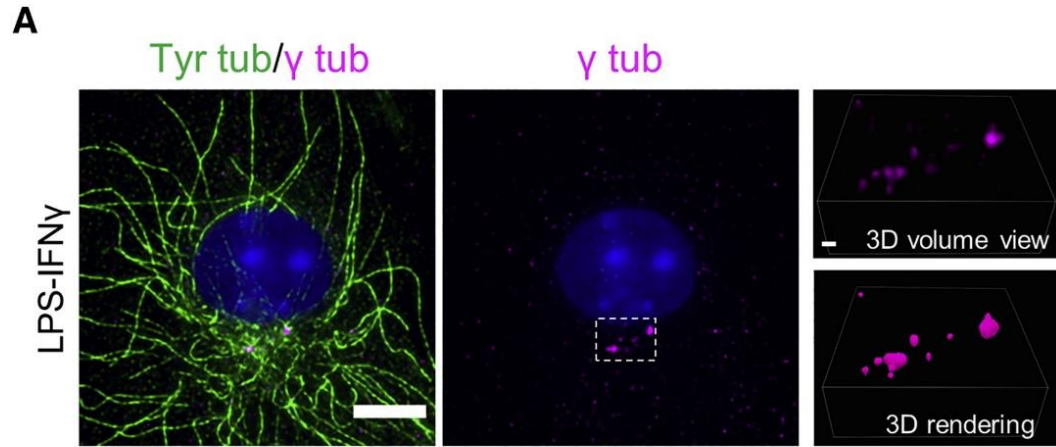
In microglia, γ -tubulin redistributes from a diffuse pattern in homeostatic cells to a concentrated pericentrosomal area in activated microglia



This redistribution is important for cytokine release

Cyclin-dependent kinase 1 (Cdk1) is an upstream regulator of microtubule remodeling and morphological changes
Cdk1 activation is essential for MT remodeling and functional cytokine release



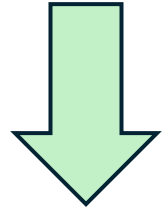


Microglia activation with LPS-IFN γ drives the formation of multiple pericentrosomal γ -tubulin positive structures

Microglia microtubule cytoskeleton

- MTs are fundamental for cell division, intracellular transport, to maintain cell shape and drive motility
- In microglia, MTs dynamicity increases with activation
- MT polarity, fundamental to establish intracellular transport direction, is mixed in ramified microglia and uniform in amoeboid microglia
- Ramified microglia display non-centrosomal MTs, amoeboid microglia rearrange MTs in centrosomal organization
- Ramified microglia display Golgi outposts as alternative MTOCs
- Microglia reorganize γ -tubulin during activation, and this drives cytokine release

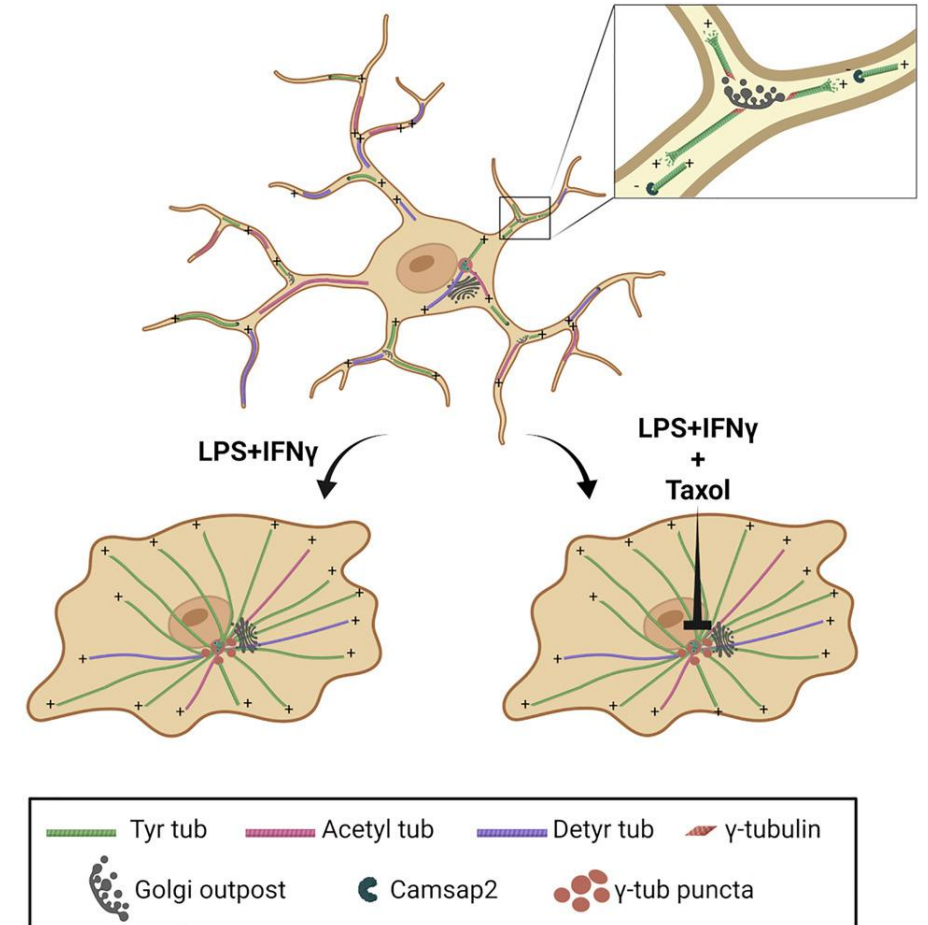
Why is microglia microtubule cytoskeleton interesting?



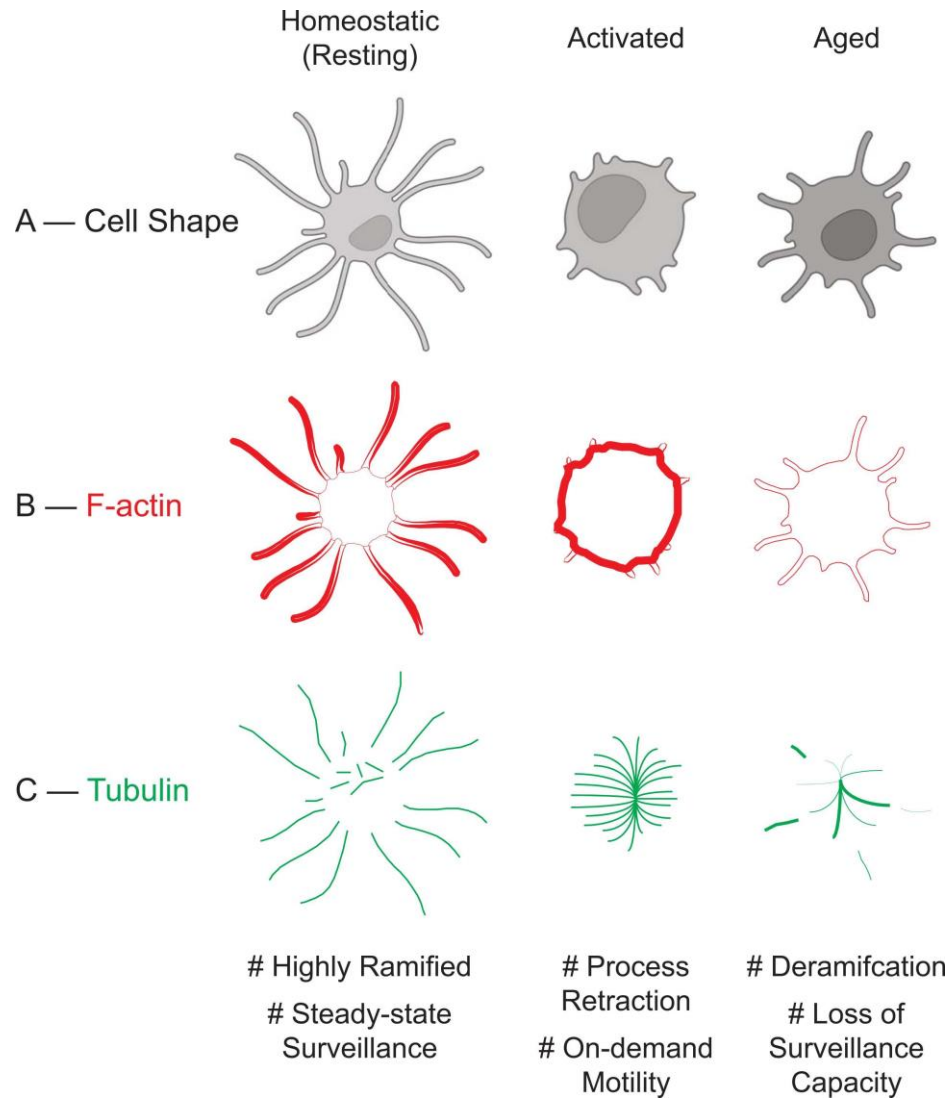
To find novel target to modulate neuroinflammation

The huge morphological changes in microglia morphologies rely on the MT cytoskeleton rearrangements, therefore MT cytoskeleton components can become:

- Proxies for neuroinflammation and aging
- Target for neuroinflammation modulation



A cytoskeleton symphony: Actin and microtubules in microglia dynamics and aging



Actin and microtubule cooperatively shape the microglial cytoskeleton behavior.

- Actin and microtubule dynamics underpin microglia homeostatic functions.

- Several MT reorganizing molecules can bridge microtubule dynamics to microglia reactivity.

- Actin polymerization coordinates microglial motility across diverse functional states.

- Disorganization of the actin-microtubule network is a hallmark of aged microglia.

- Aging in microglia is associated with notable changes in MT stability, leading to a disrupted MT network.

Useful links

[10.1016/j.neuron.2022.10.020](https://doi.org/10.1016/j.neuron.2022.10.020) microglia states and nomenclature

<https://doi.org/10.1038/s41419-023-05835-8> tunneling nanotubes

[https://www.cell.com/current-biology/fulltext/S0960-9822\(99\)80201-2](https://www.cell.com/current-biology/fulltext/S0960-9822(99)80201-2) the centrosome

[10.1016/j.pneurobio.2024.102586](https://doi.org/10.1016/j.pneurobio.2024.102586) actin and MTs in microglia

[10.3390/cells8060639](https://doi.org/10.3390/cells8060639) microglia actin

<https://doi.org/10.1016/j.celrep.2023.112104> microglia MTs

<https://doi.org/10.1038/s41467-023-41891-6> microglia MTs

<https://doi.org/10.1242/jcs.03227> generation of non-centrosomal arrays