

Lesson (11)

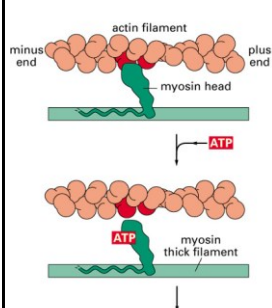
Inside the neuron – Molecular Neurobiology I:

Cellular transport and the molecular motors

MOLECULAR MOTORS

- **There Are Two Types of Microtubule Motor Proteins: Kinesins and Dyneins**
- **One Type of Actin-based Motor Proteins : Members of the Myosin Superfamily**
- **Motor Proteins Generate Force by Coupling ATP Hydrolysis to Conformational Changes**
- **Cilia and Flagella Are Motile Structures Built from Microtubules and Dyneins**

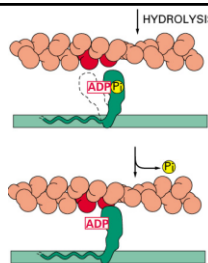
Motor Proteins Generate Force by Coupling ATP Hydrolysis to Conformational Changes



ATTACHED At the start of the cycle shown in this figure, a myosin head lacking a bound nucleotide is locked tightly onto an actin filament in a *rigor* configuration (so named because it is responsible for *rigor mortis*, the rigidity of death). In an actively contracting muscle, this state is very short-lived, being rapidly terminated by the binding of a molecule of ATP.

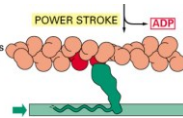
RELEASED A molecule of ATP binds to the large cleft on the "back" of the head (that is, on the side furthest from the actin filament) and immediately causes a slight change in the conformation of the domains that make up the actin-binding site. This reduces the affinity of the head for actin and allows it to move along the filament. (The space drawn here between the head and actin emphasizes this change, although in reality the head probably remains very close to the actin.)

Figure 16-58 part 1 of 3. Molecular Biology of the Cell, 4th Edition.

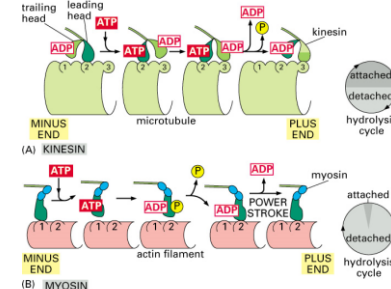


COCKED The cleft closes like a clam shell around the ATP molecule, triggering a large shape change that causes the head to be displaced along the filament by a distance of about 5 nm. Hydrolysis of ATP occurs, but the ADP and inorganic phosphate (Pi) produced remain tightly bound to the protein.

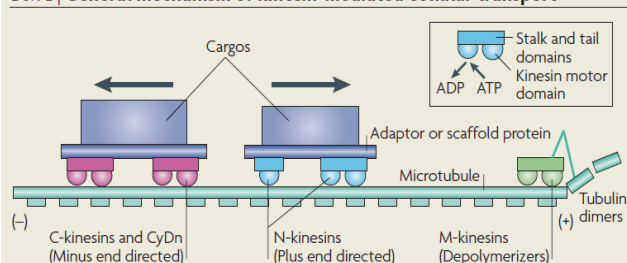
FORCE-GENERATING A weak binding of the myosin head to a new site on the actin filament causes release of the inorganic phosphate produced by ATP hydrolysis, concomitantly with the tight binding of the head to actin. This release triggers the power stroke—the force-generating change in shape during which the head regains its original conformation. In the course of the power stroke, the head loses its bound ADP, thereby returning to the start of a new cycle.



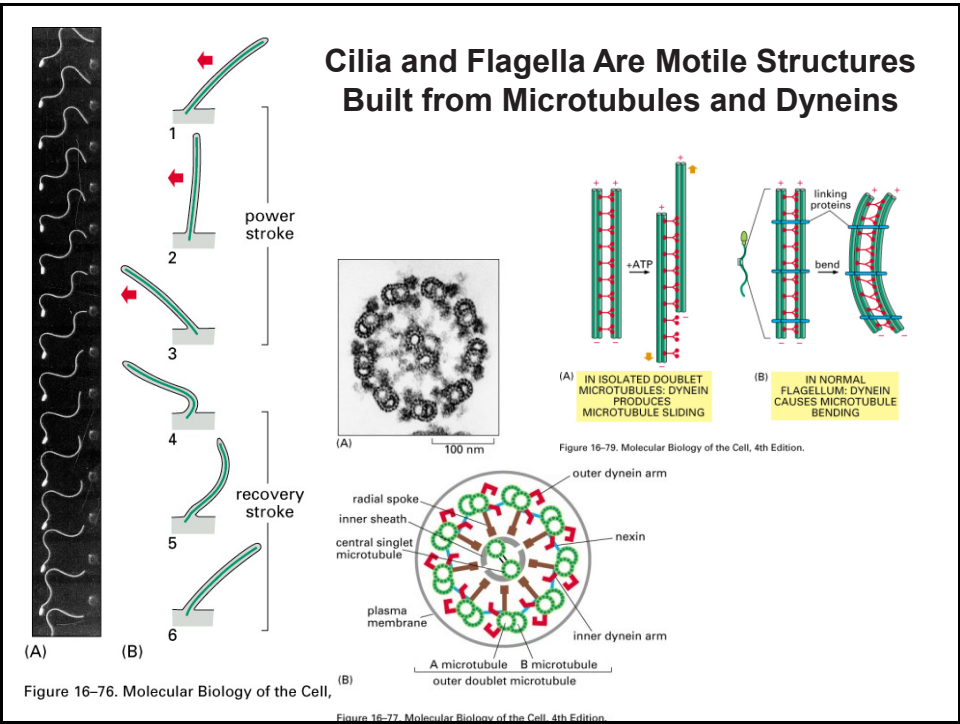
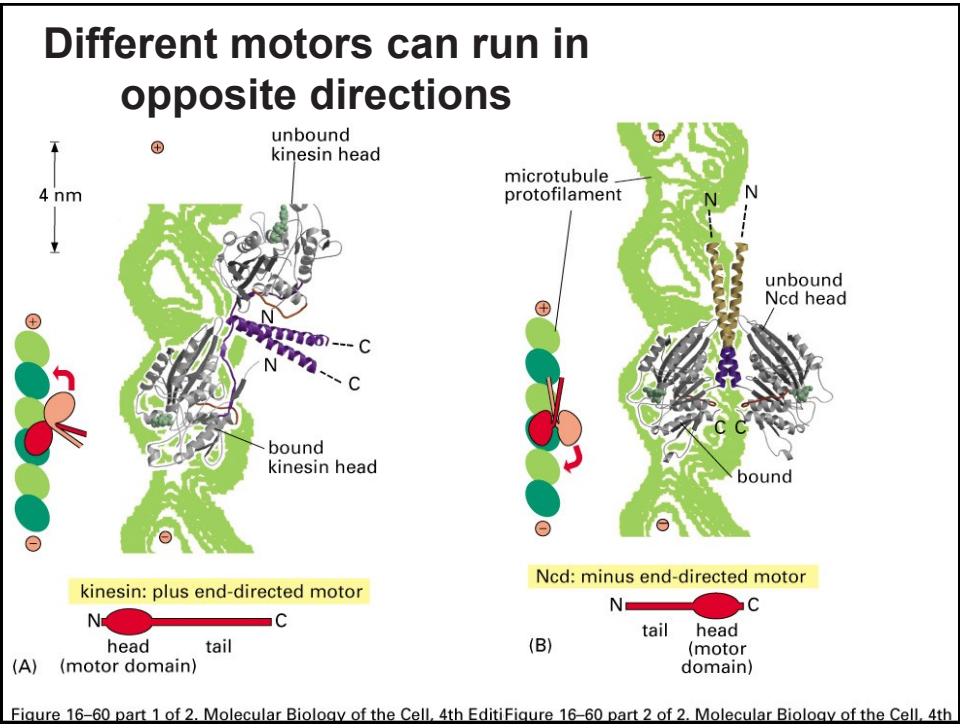
ATTACHED At the end of the cycle, the myosin head is again locked tightly to the actin filament in a *rigor* configuration. Note that the head has moved to a new position on the actin filament.



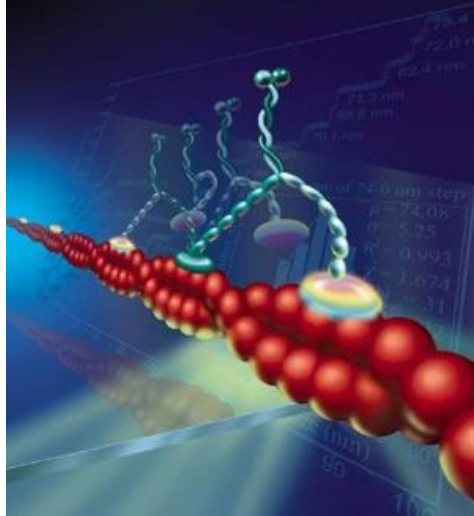
Box 1 | General mechanism of kinesin-mediated cellular transport



Kinesins transport cargos along microtubules by a standard mechanism (depicted here using a freight train model; see the figure). A kinesin motor generally consists of a kinesin motor domain, which is conserved among kinesin superfamily proteins (KIFs), and unique stalk and tail domains that are used for kinesin dimerization and/or kinesin binding to cargos, adaptors or scaffold proteins. The kinesin motor domain generates force by hydrolysing ATP. Kinesins are largely classified as N-kinesins, M-kinesins or C-kinesins, which contain their motor domain at the amino terminus, in the middle or at the carboxyl terminus, respectively. N-kinesins generally provide plus end-directed motility that is anterograde towards the cell periphery or axon terminals in neurons. Some N-kinesins act as monomers and others act as dimers. C-kinesins, together with cytoplasmic dynein (CyDn), provide minus end-directed motility that is generally retrograde towards the cell centre. M-kinesins depolymerize microtubules. In some cases, adaptors and scaffolds provide a mechanistic link between kinesins and cargos, and they might also have regulatory roles in kinesin-driven intracellular transport, namely in the recognition of specific cargos and the regulation of cargo loading and unloading.



Motors Walk Along Filaments

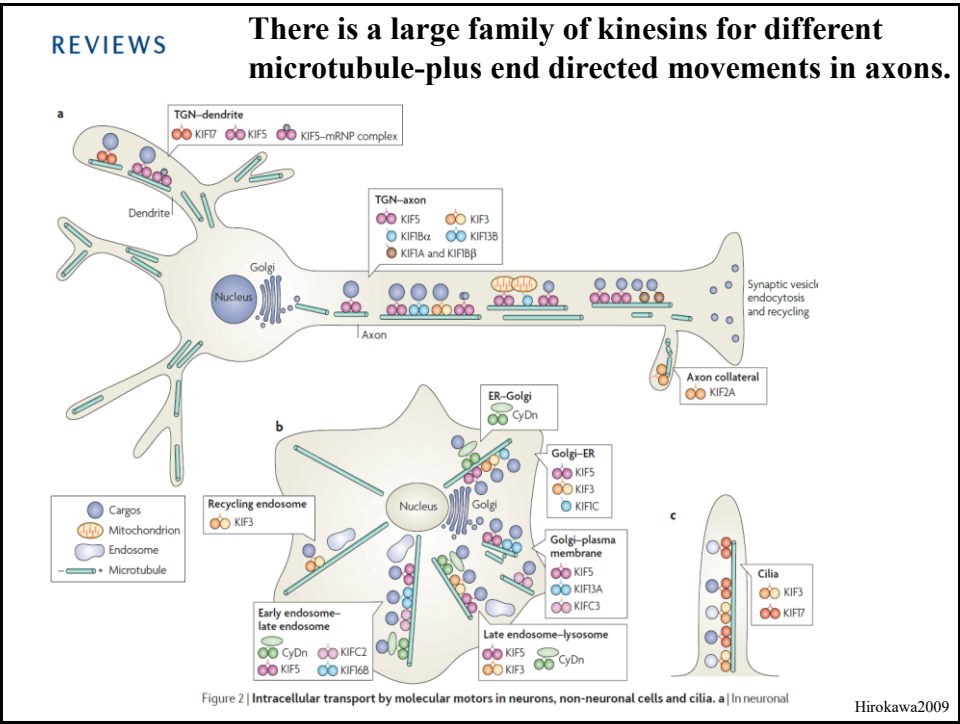
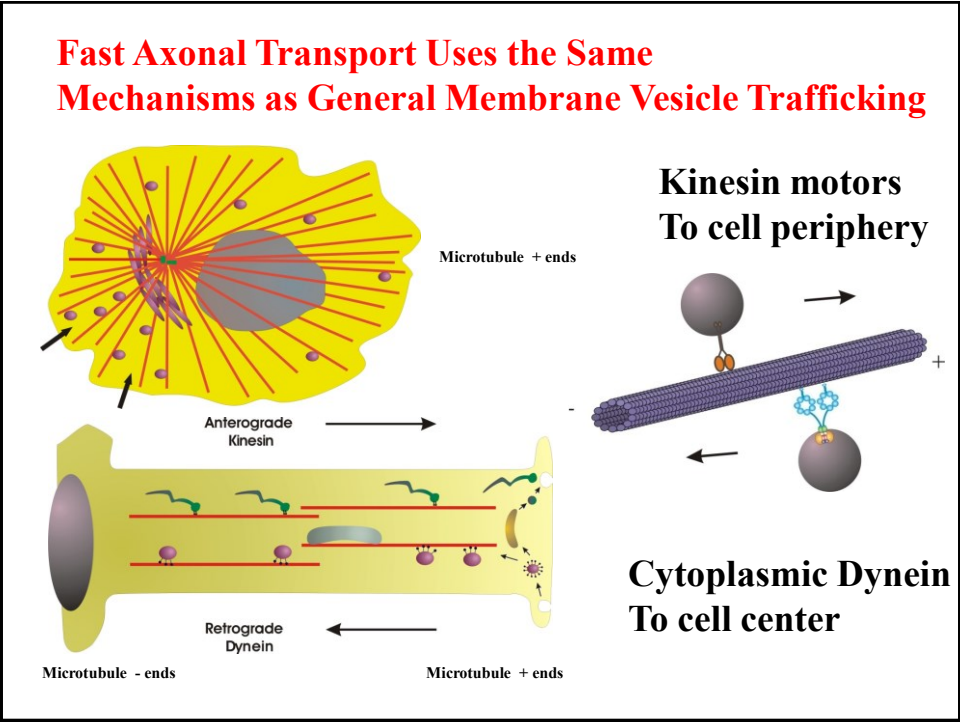


https://www.youtube.com/watch?v=B_zD3NxSsD8

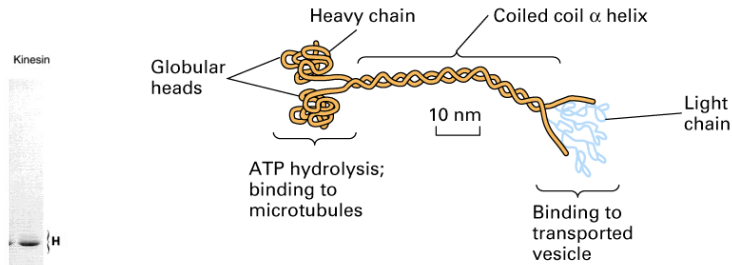
Yildiz *et al.*
Science 2003.

Motor proteins

- Move unidirectionally
- Stepwise
- Series of conformational changes
 - A *mechanical cycle*
 - Coupled to chemical cycle – Energy
 - Steps –
 - » ATP binding to motor
 - » Hydrolysis of ATP
 - » Release of ADP and P_i
 - » Binding of new ATP



Kinesin: Polypeptides and Structure



Tetramer

2 identical heavy and 2 identical light chains

Functional domains

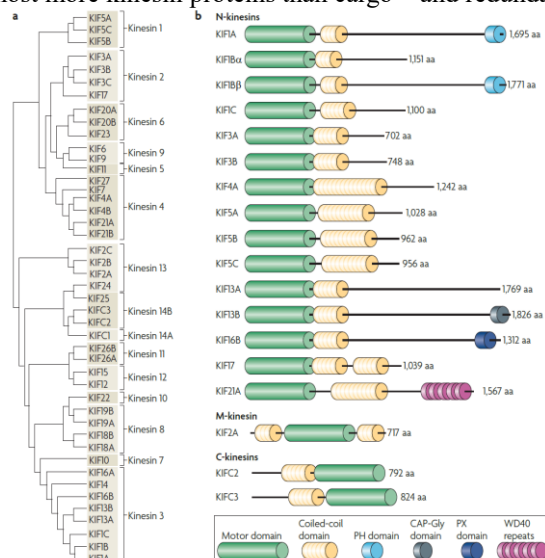
Pair of globular heads
Bind microtubule
ATP-hydrolysing
Neck / stem and tail
Tail binds cargo

Move toward plus end of microtubule

Plus end directed

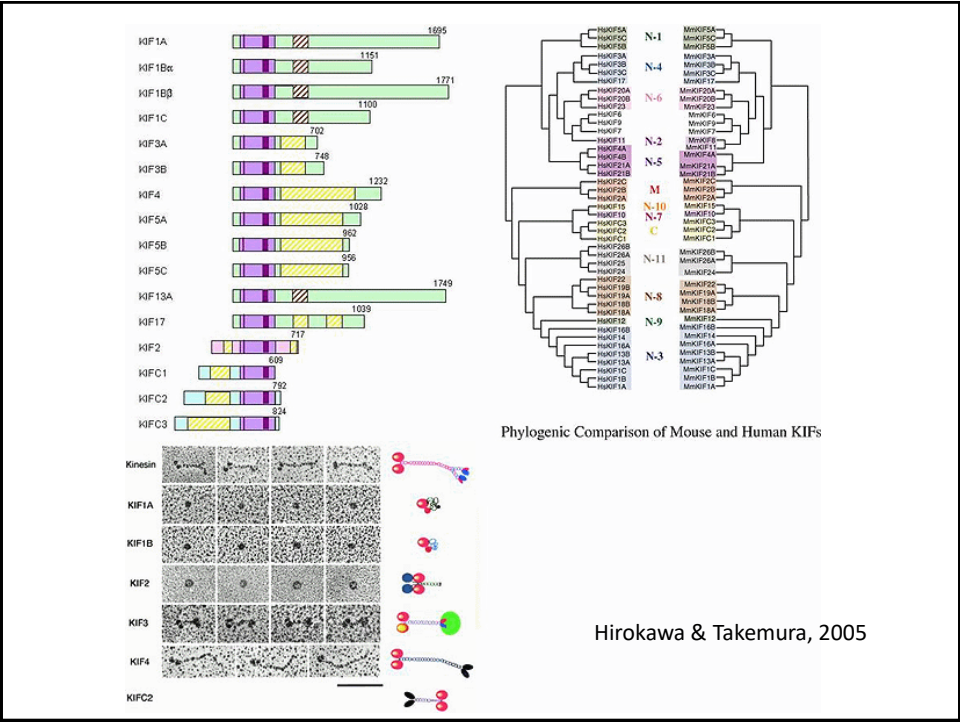
Kinesins: a family of ~45 members in mammals

Conserved motor domain fused to variable domains
almost more kinesin proteins than cargo – and redundancy



Hirokawa, 2009

Figure 1 | The structure and phylogeny of major mouse kinesins. **a** | A phylogenetic tree of all 45 kinesin superfamily (also known as KIF) genes in the mouse genome, which are classified into 15 families^{10,11,13,15-18}. **b** | The domain structure of



Hirokawa & Takemura, 2005

Different kinesins to transport different cargoes in axonal and dendritic transport

Hirokawa • Motors and mRNA Transport in Dendrites
J. Neurosci., July 5, 2006 • 26(27):7139–7142

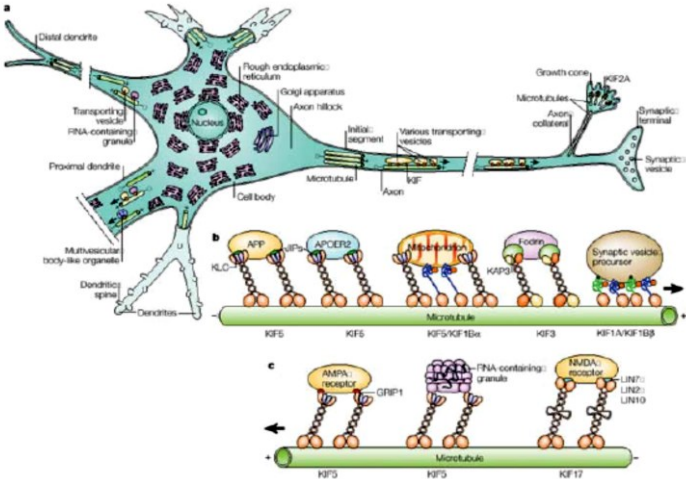
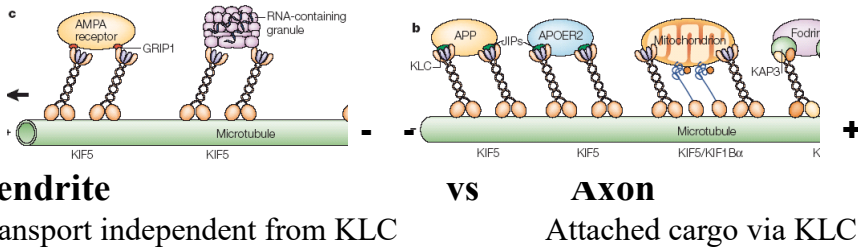
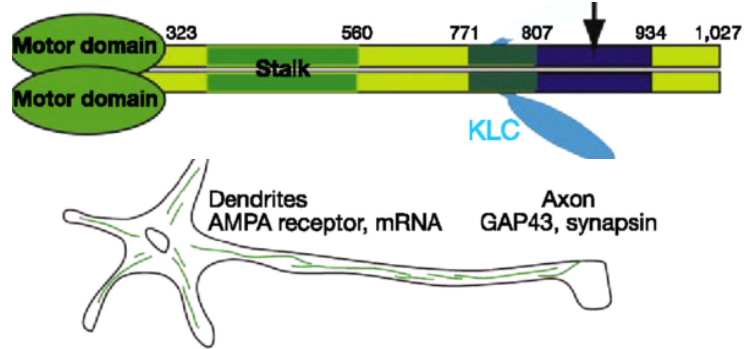


Figure 1. KIFs and cargoes for axonal and dendritic transport. **a**, Microtubule polarity in the axon and dendrites. **b**, KIFs and their cargoes in the axon. **c**, KIFs and their cargoes in dendrites. mRNA and granules composed of a large number of proteins are transported by KIFs. Reproduced with permission from Hirokawa and Takemura (2005).

Model proposed by Hirokawa (Nat Rev Neurosci. 2005 6:201-214)



Adaptor-proteins for cargoes transport by kinesins

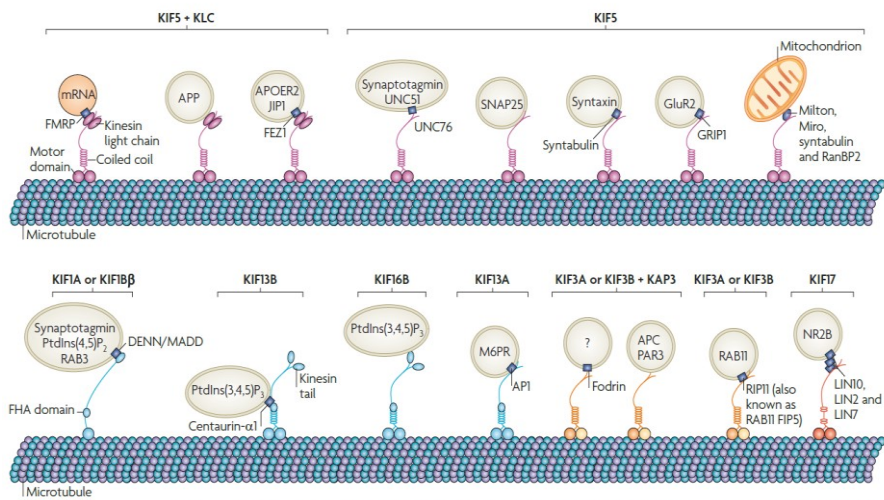


Figure 3 | Kinesins, cargoes and molecules involved in cargo recognition. Major transport mechanisms that are based on

Horokawa, 2009

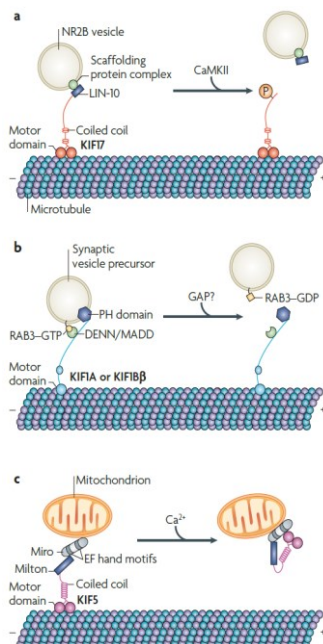


Figure 4 | **Regulation of kinesin-cargo binding by three different mechanisms.** **a** | Phosphorylation. The kinesin 2 family motor KIF17 (also known as OSM3) transports vesicles that carry NMDA (N-methyl-D-aspartate) receptor subunit 2B (NR2B) towards the plus end of microtubules through a scaffolding protein complex⁶⁵. Phosphorylation of KIF17 by the Ca^{2+} /calmodulin-dependent protein kinase CaMKII disrupts the direct interaction between KIF17 and its interacting partner LIN10 (the mouse homologue of *Caenorhabditis elegans* LIN-10), leading to the unloading of NR2B-carrying vesicles¹⁰². **b** | The Rab GTPase cycle. The kinesin 3 family motors KIF1A and KIF1B transport RAB3-carrying synaptic vesicle precursors through the adaptor protein mitogen-activated protein kinase-activating death domain (MADD; also known as DENN and shown as DENN/MADD)¹¹⁵. DENN/MADD recognizes the GTP-bound form of RAB3 on vesicles. Because the membrane-binding capacity of the pleckstrin homology (PH) domain of KIF1A and KIF1B is insufficient to transport cargo¹¹⁵ when RAB3 is inactivated by GTPase-activating proteins at the axon terminus, RAB3-GDP-carrying vesicles are released. **c** | Calcium. The kinesin 1 family motor KIF5 transports mitochondria through interaction with the Milton-Miro complex¹⁹. Miro has two EF hand motifs, which sense intracellular Ca^{2+} levels. In response to high Ca^{2+} influx, these motifs bind to the motor domain of KIF5 to inhibit its activity. As a result, mitochondria motility is inhibited.

KIF1A and KIF1B transport synaptic vesicle precursors containing synaptic vesicle proteins such as synaptotagmin, synaptophysin, and Rab 3A.

KIF1B and KIF5 (KIF5A, KIF5B, and KIF5C) transport mitochondria in the anterograde direction

KIF5 also transports other cargoes, including vesicles that contain APPs (amyloid precursor proteins) and vesicles containing APOER2 (apolipoprotein E receptor 2)

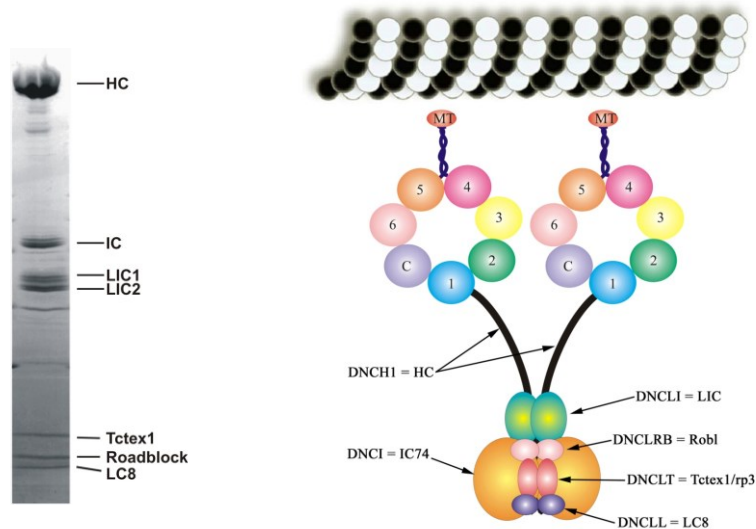
KIF5 participates in slow axonal transport of cytoskeletal proteins

KIF3, mainly composed of KIF 3A/KIF3B heterodimers and an associated protein KAP3, transports vesicles associated with fodrin and is important for neurite extension.

In dendrites, transport of NMDA receptors is mediated by the molecular motor **KIF17**;

Motor Proteins:

Cytoplasmic Dynein: Polypeptides and Structure



One Type of Cytoplasmic Dynein Motor Domain (HC) But Multiple Mammalian Cargo Binding Subunits

Intermediate Chains (dimer) – 2 genes

5 - 6 alternative splice variants

Phosphorylated isoforms

Light Intermediate Chains (dimer) - 2 genes

Phosphorylated isoforms

May be alternative splice variants

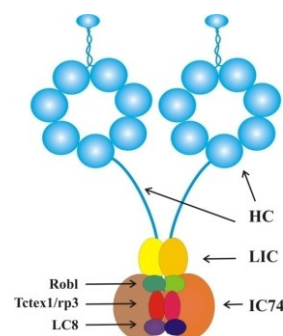
Heavy Chains

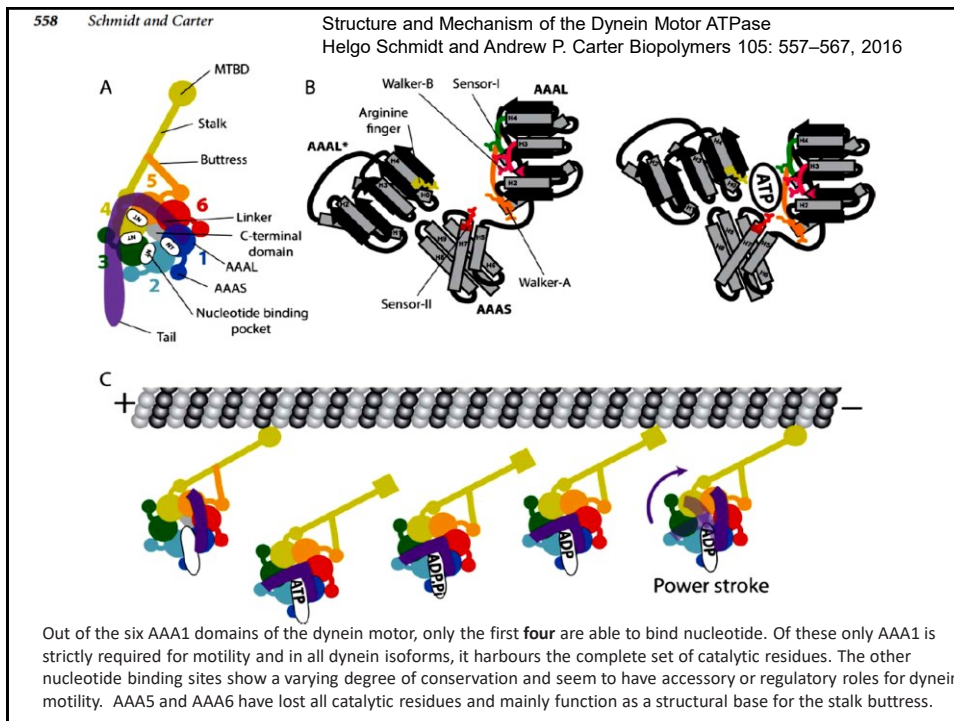
Tctex1 Family (dimer) - 2 genes

Roadblock Family (dimer) - 2 or more genes

LC8 Family (dimer) - 2 or more genes

No evidence for cytoplasmic dynein LC phosphorylation

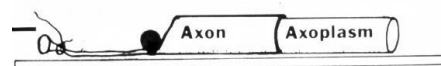




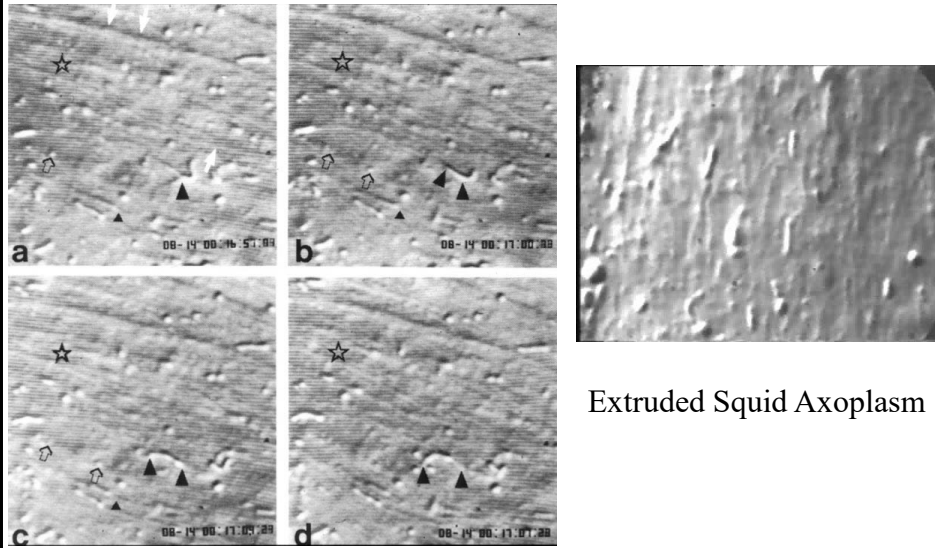
MICROTUBULE-BASED MOTOR PROTEINS & THE MECHANISM FOR FAST AXONAL TRANSPORT

Fast Axonal Transport

Extrusion of Axoplasm from squid giant axon

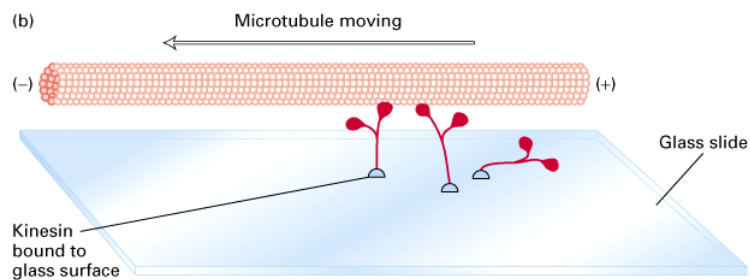


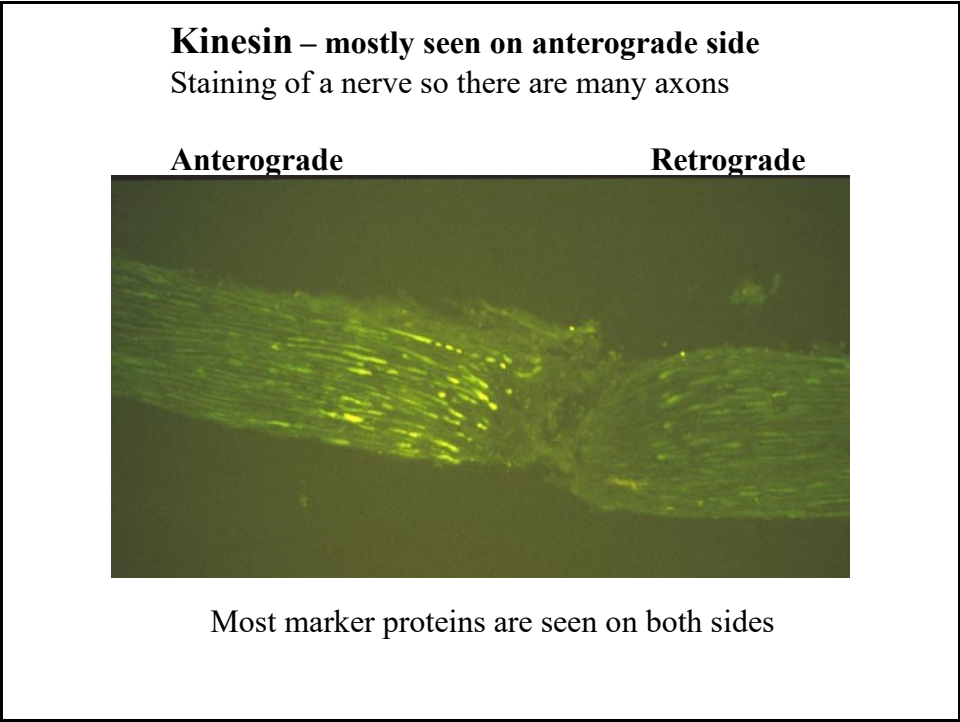
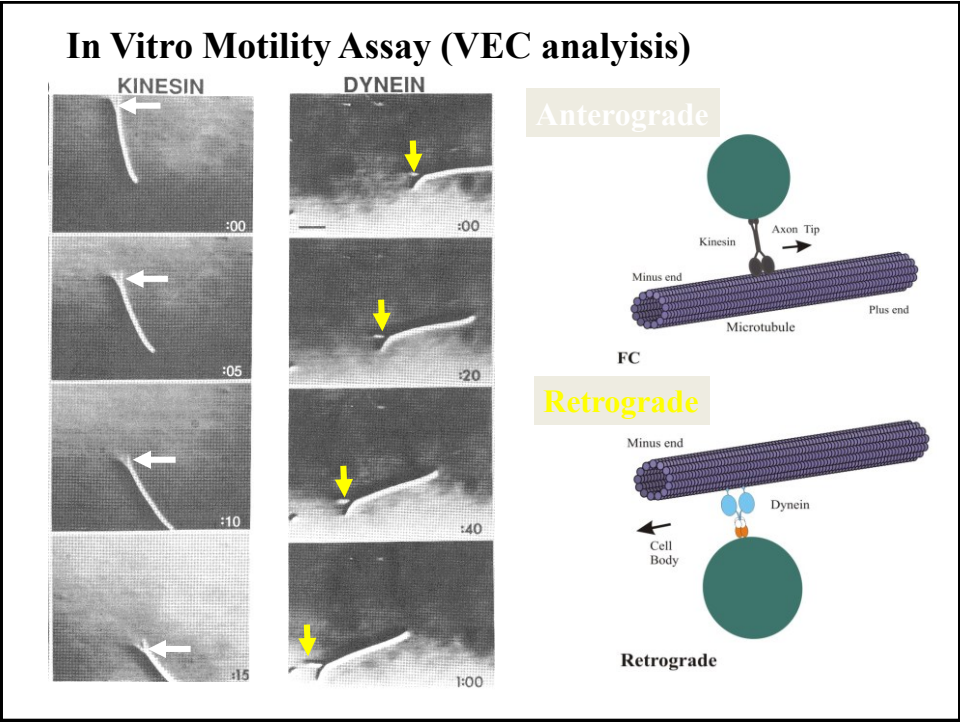
1. Time Lapse analysis of extruded Axoplasm (VEC=Video-enhanced contrast Images)



In Vitro Motility Assay

MTs move (almost glide) over surface of a motor protein coated coverslip (requires ATP).

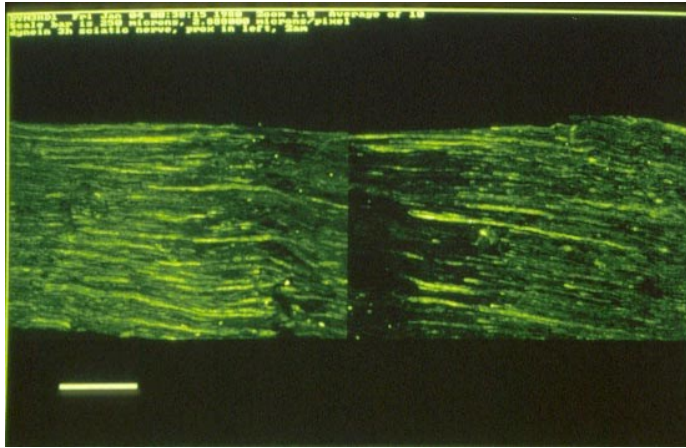




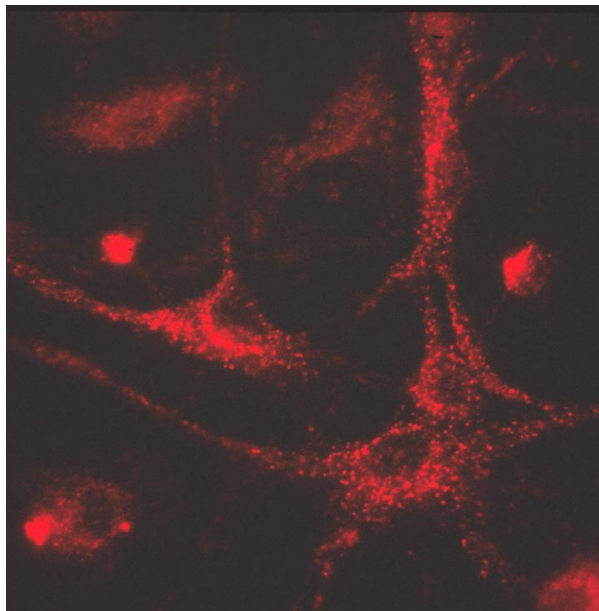
Cytoplasmic Dynein seen on both sides

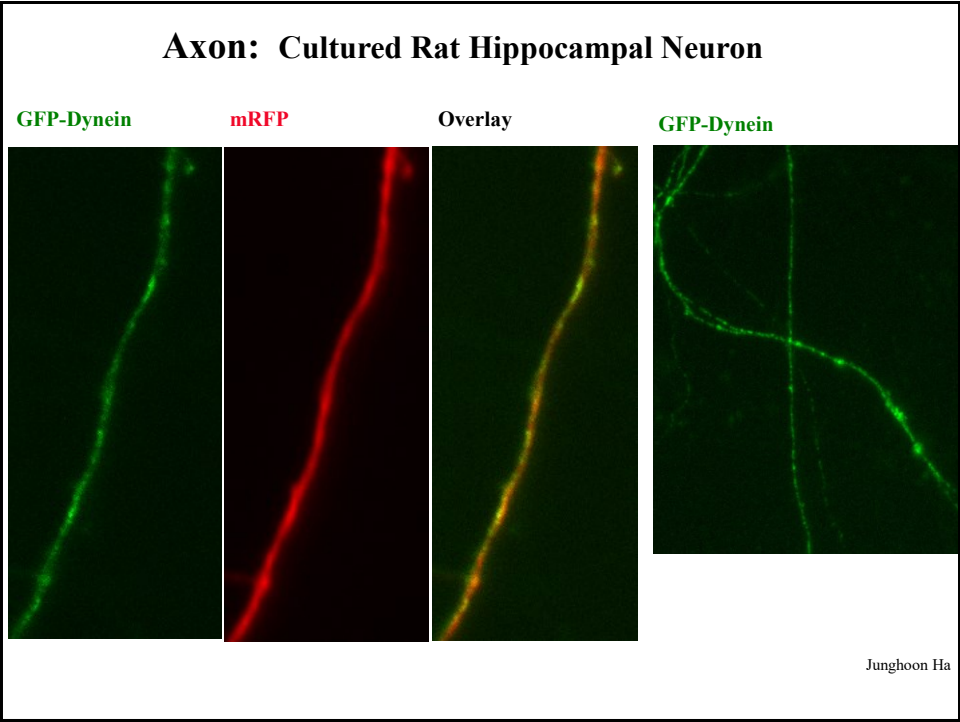
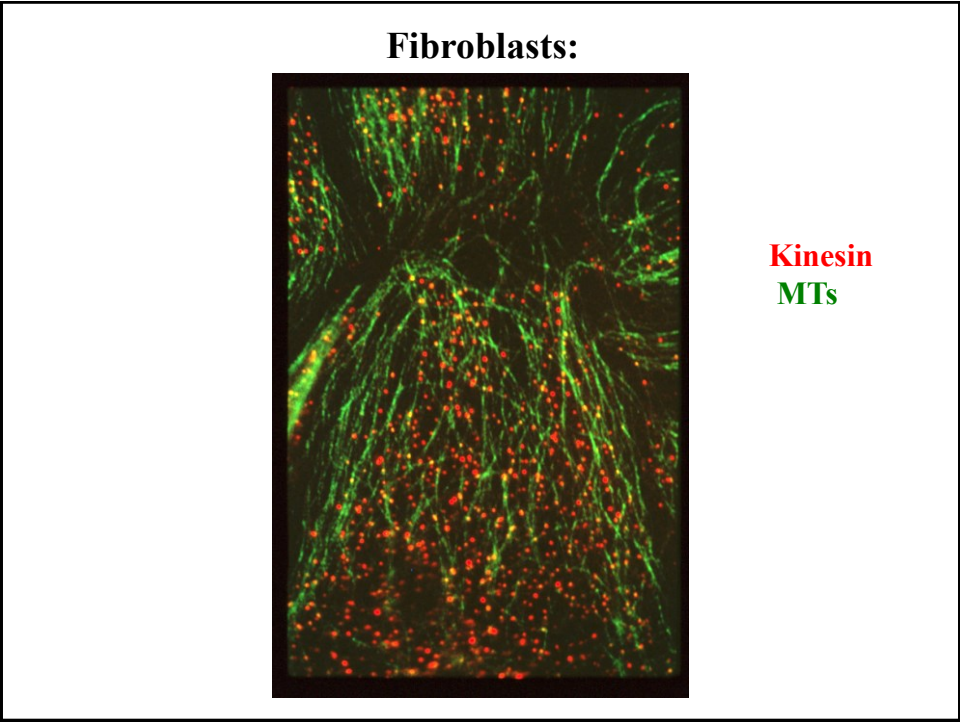
Anterograde

Retrograde

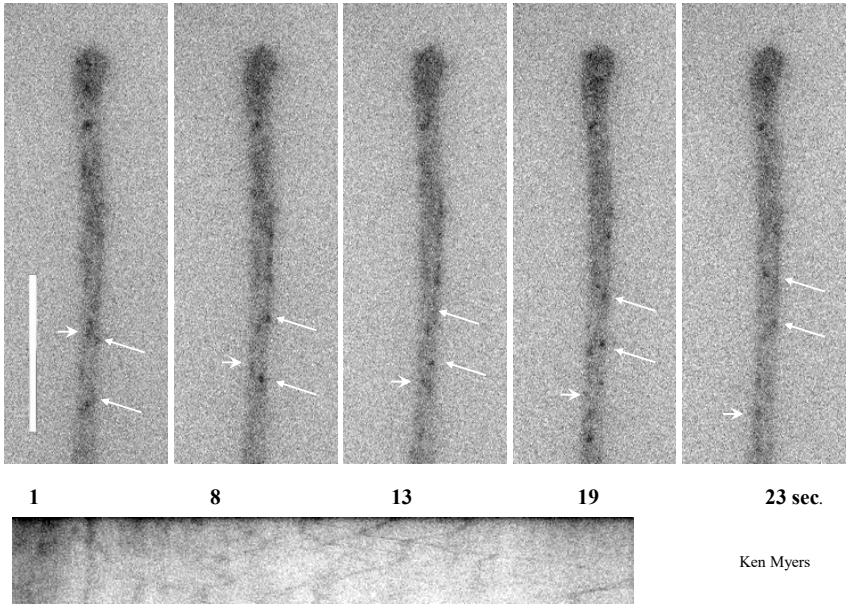


Kinesin in Cultured Glia

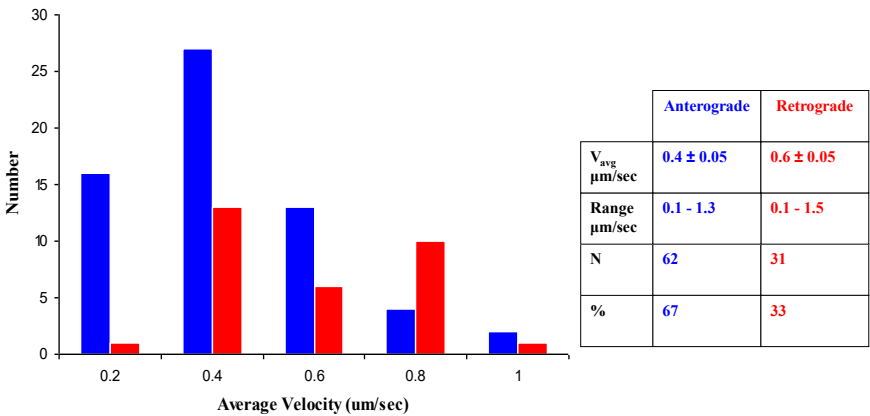




Movement of GFP-Dynein Puncta in Both Directions

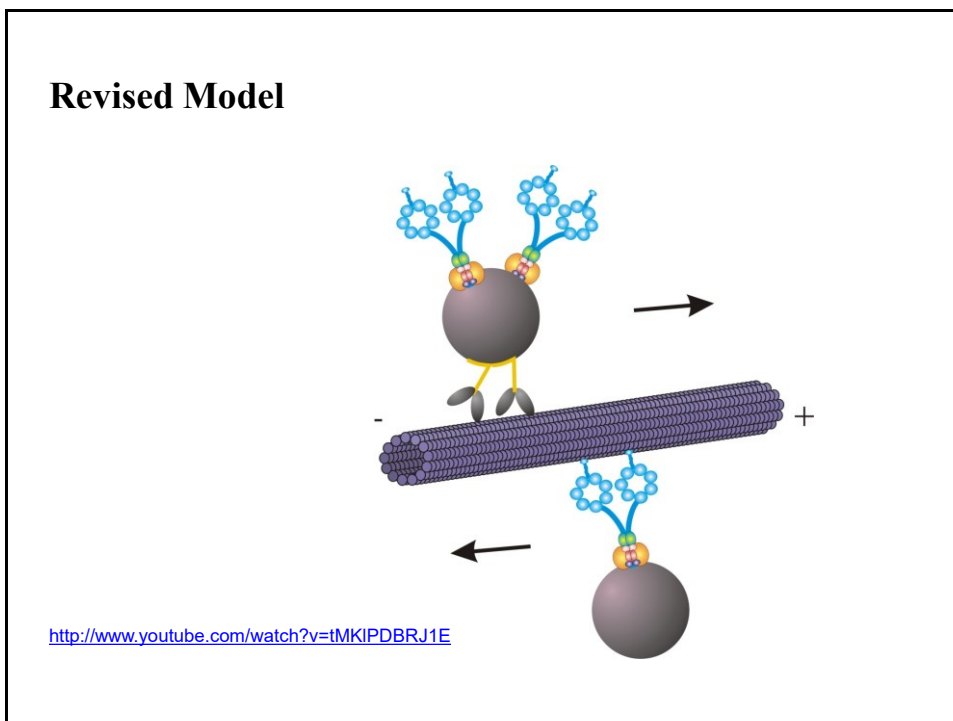
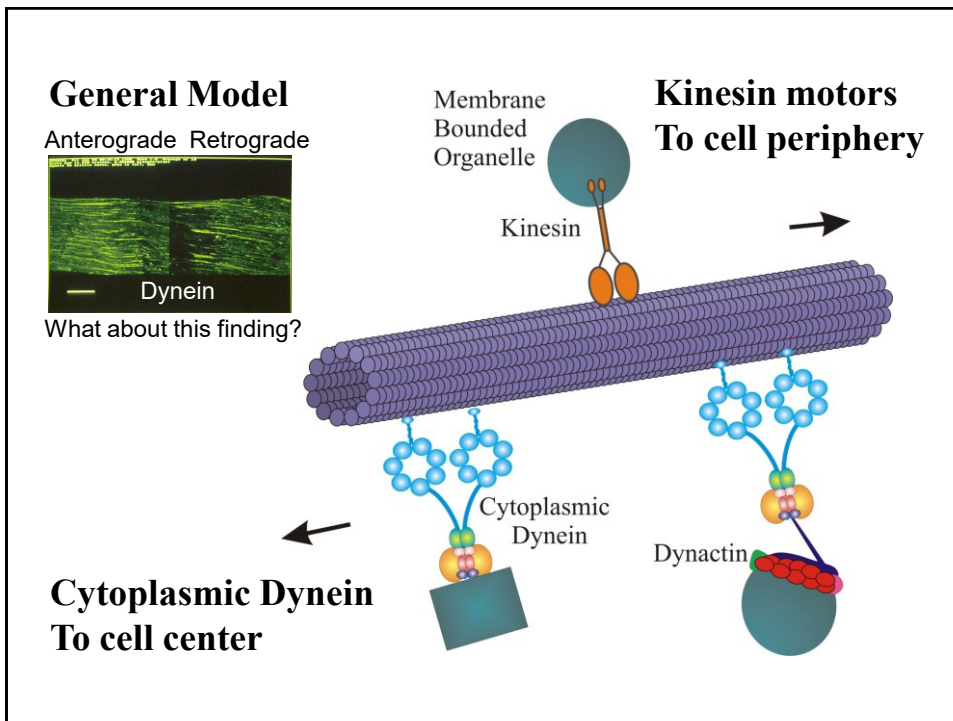


Characterization of Dynein Puncta Motility in PC12 Cell Neurites



Anterograde
Retrograde

Junghoon Ha



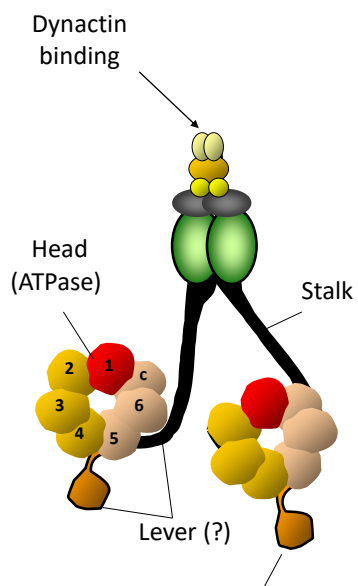
Dynein Can Shift Gears

Dynein

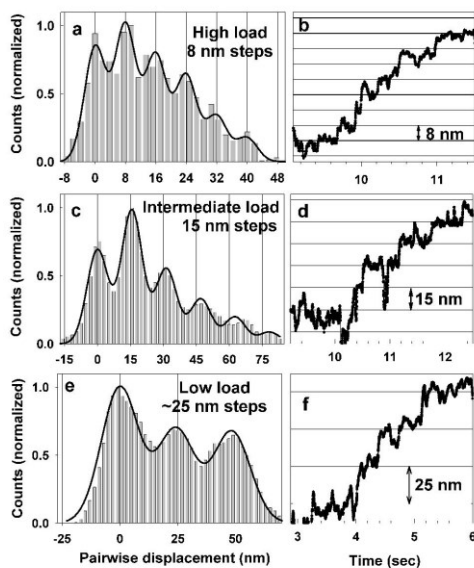
Roop Mallik¹, Brian Carter¹, Stephanie Lax², Stephen King², Steven Gross¹

¹UC Irvine

²Univ. of Missouri-Kansas City



Step Size as a Function of Load at High [ATP]



High load
8nm steps

Intermediate load
15 nm steps

Low load
~25 nm steps

Complication !

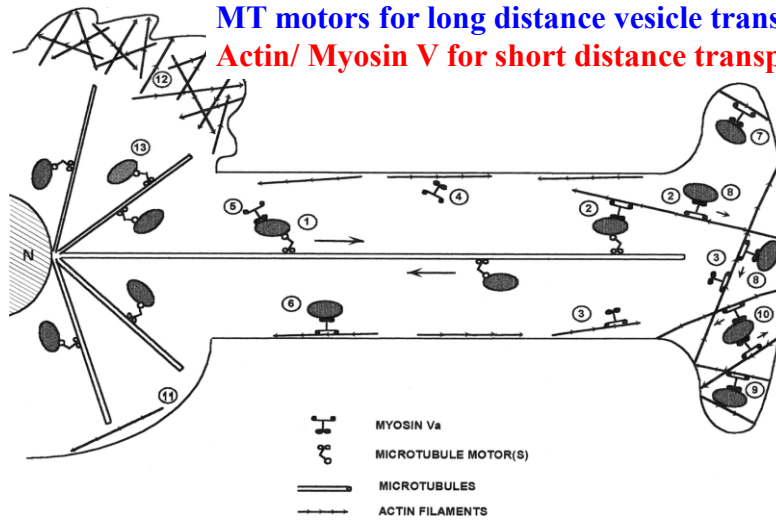
Mouse “dilute” Mutation is Myosin V

It has neurological defects

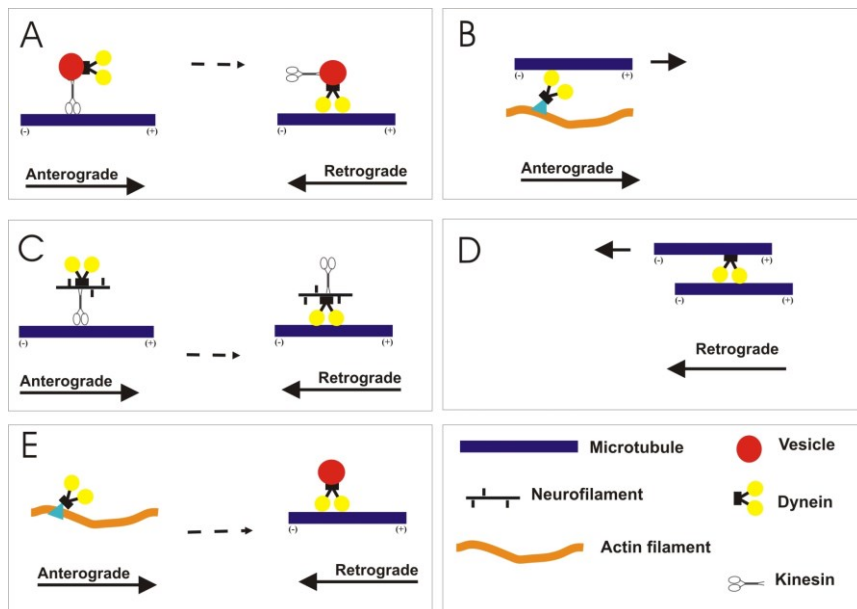
Revised Model:

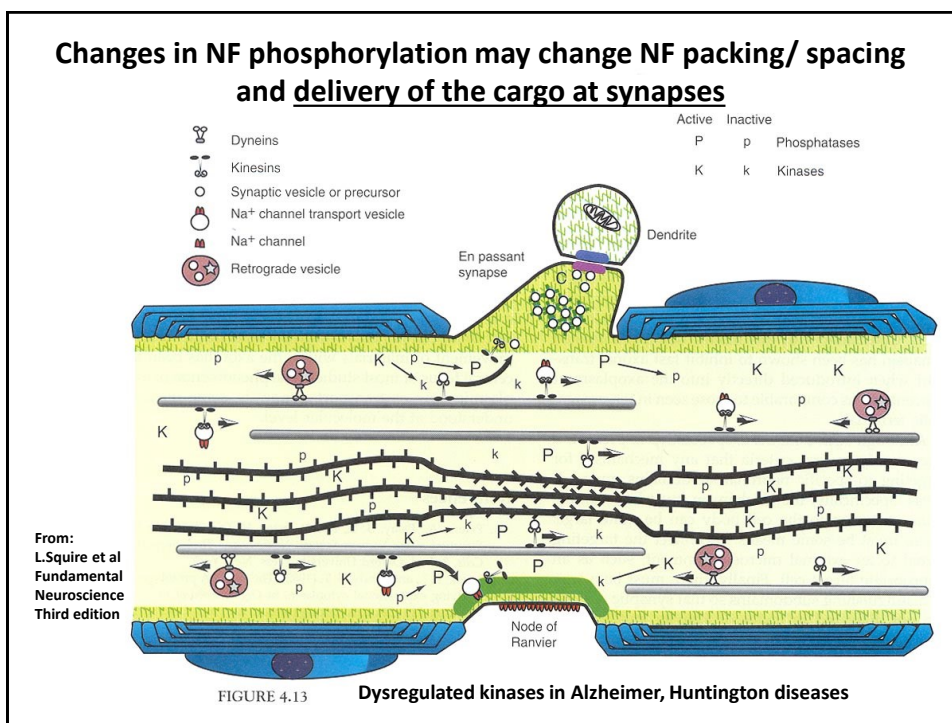
MT motors for long distance vesicle transport

Actin/ Myosin V for short distance transport



Exercise: what is Fast and what is Slow Transport ?





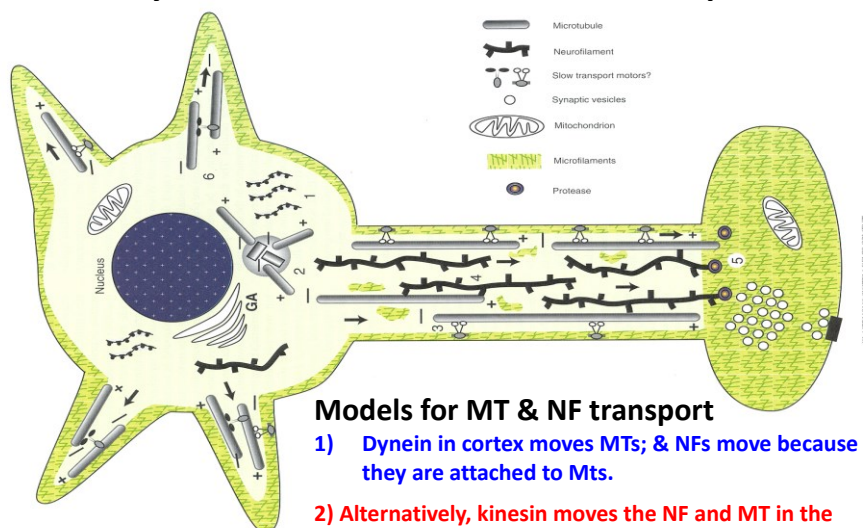
Summary: Axonal Transport 1

- Many proteins made in soma must be transported to axon and axon terminal
 - repair axolemma, for gated ion channel proteins, as enzymes or neurotransmitters
- Fast anterograde axonal transport
 - either direction up to 400 mm/day for organelles, enzymes, vesicles and small molecules

Summary Axonal Transport 2

- Fast retrograde for recycled materials and pathogens
- Slow axonal transport or axoplasmic flow
 - moves cytoskeletal and new axoplasm at 1-6 mm/day for normal housekeeping, and also for repair and regeneration in damaged axons

Roles of Dyneins and Kinesins in MT&NF transport



Models for MT & NF transport

- 1) Dynein in cortex moves MTs; & NFs move because they are attached to Mts.
- 2) Alternatively, kinesin moves the NF and MT in the anterograde direction and dynein moves them in the retrograde direction.
- 3) Or a combination of both models.