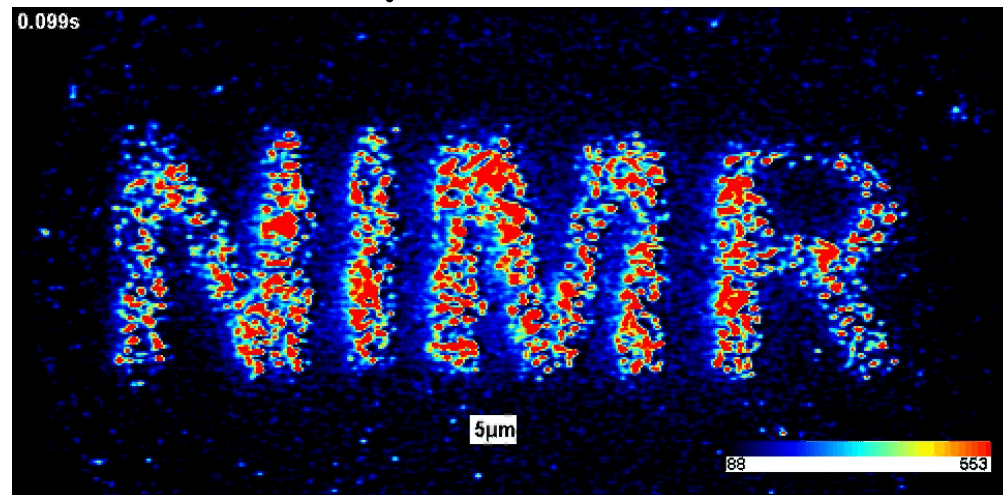


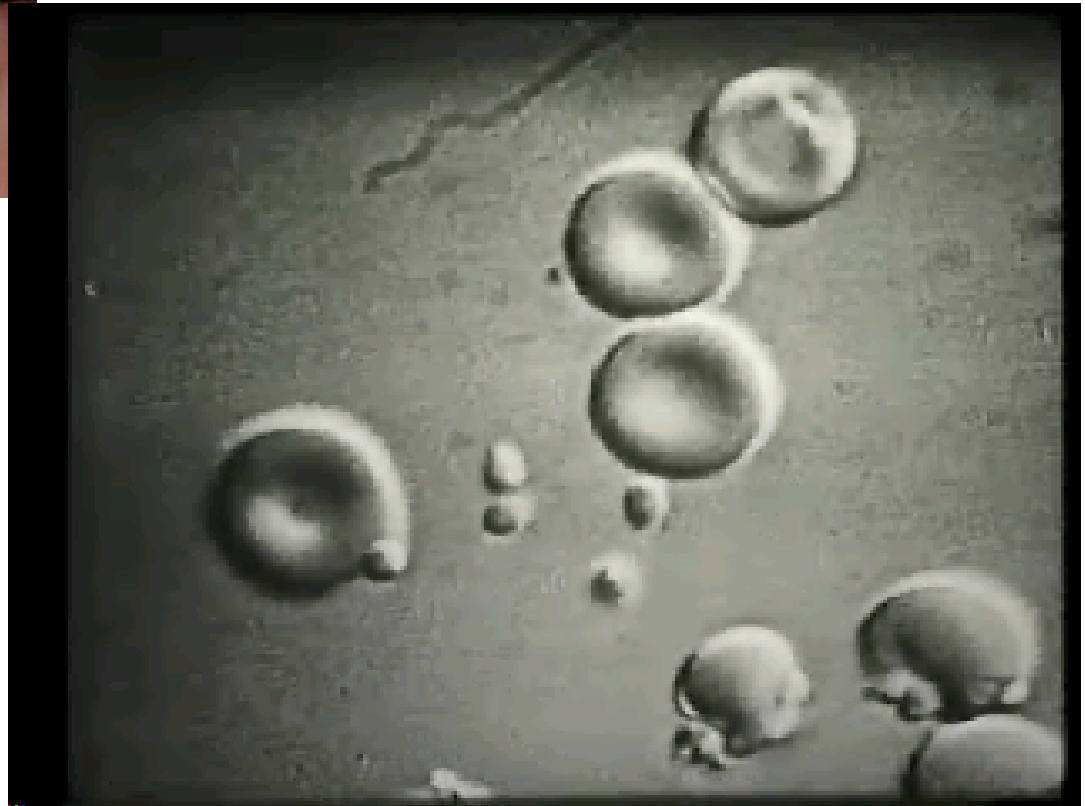
# "Some single molecule studies of myosins"

Justin Molloy  
Division of Physical Biochemistry



Motivation for the work

Molecular motors are important in disease -  
e.g. red blood cell invasion by MALARIA



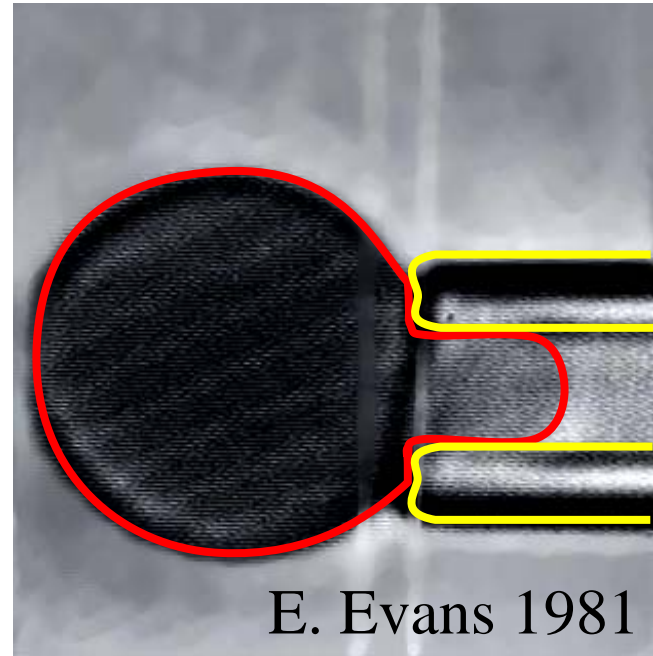
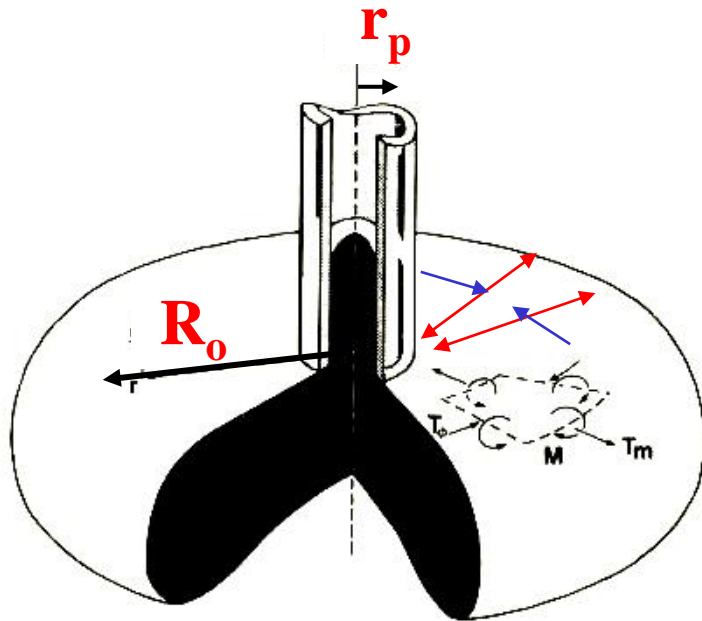
10 $\mu$ m

# Mechanical Properties of Red Blood cells

- Erythrocyte mechanical properties are well-studied
- Early work used fluid shear flow and osmotic compression
- Observation of membrane “flickering”
- More recently by direct manipulation by:
  - Pipette aspiration (1980s)
  - AFM (1990s)
  - Optical tweezers (1990s)
- Biophysics is directly applicable to malarial invasion but little mention is made of this in the literature!

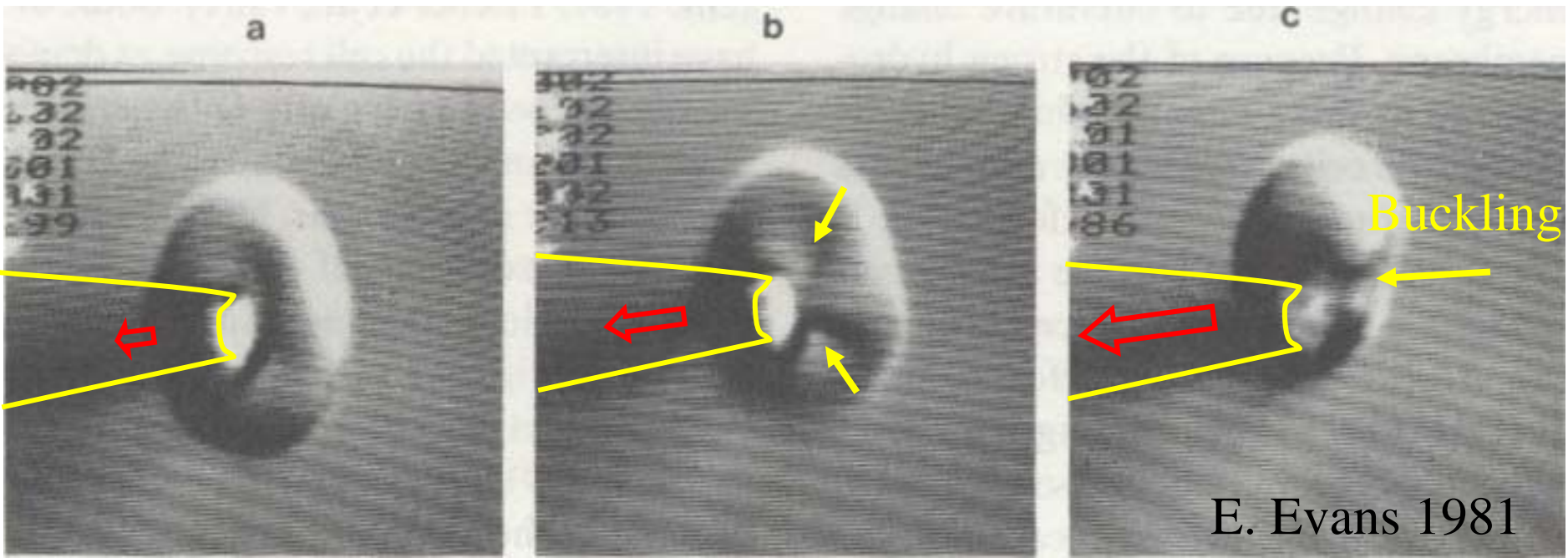
# Simple engineering treatments work well.

- Elastic sheet with boundary conditions:



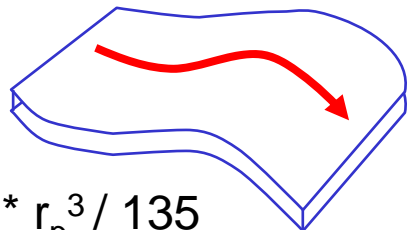
- Both **extension** and **compression** forces are generated
- Compression forces lead to buckling

# The buckling force enables the membrane elastic modulus to be estimated.



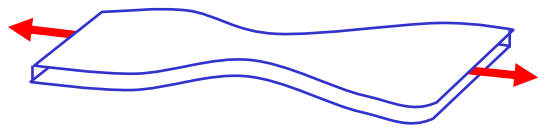
E. Evans 1981

Bending modulus (B)



When  $R_0 = 3r_p$       $B = \Delta P * r_p^3 / 135$   
 Experiment gives  $B = \underline{200 \text{ pN.nm}}$

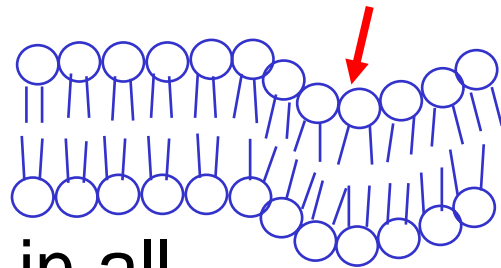
Elastic Shear modulus (H)



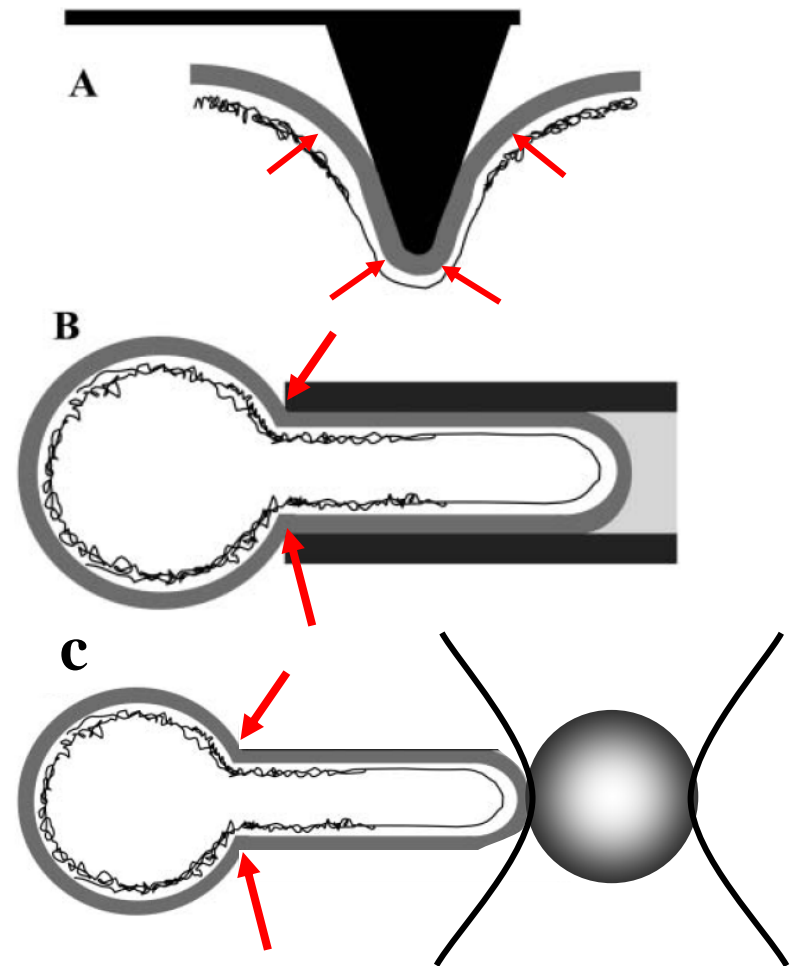
Experiment gives  $H = \underline{0.002 \text{ pN.nm}^{-1}}$

# Similar estimates can be made using AFM, and Optical tweezers

- The lipid membrane will flow freely. But it will also exert a restoring force wherever it is bent (due to its bending stiffness)



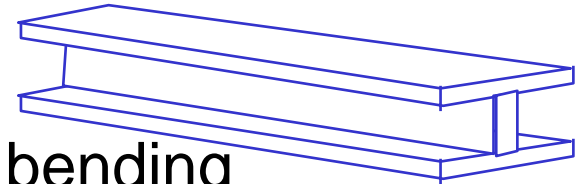
- However, in all experiments the underlying cytoskeleton is probably disrupted.



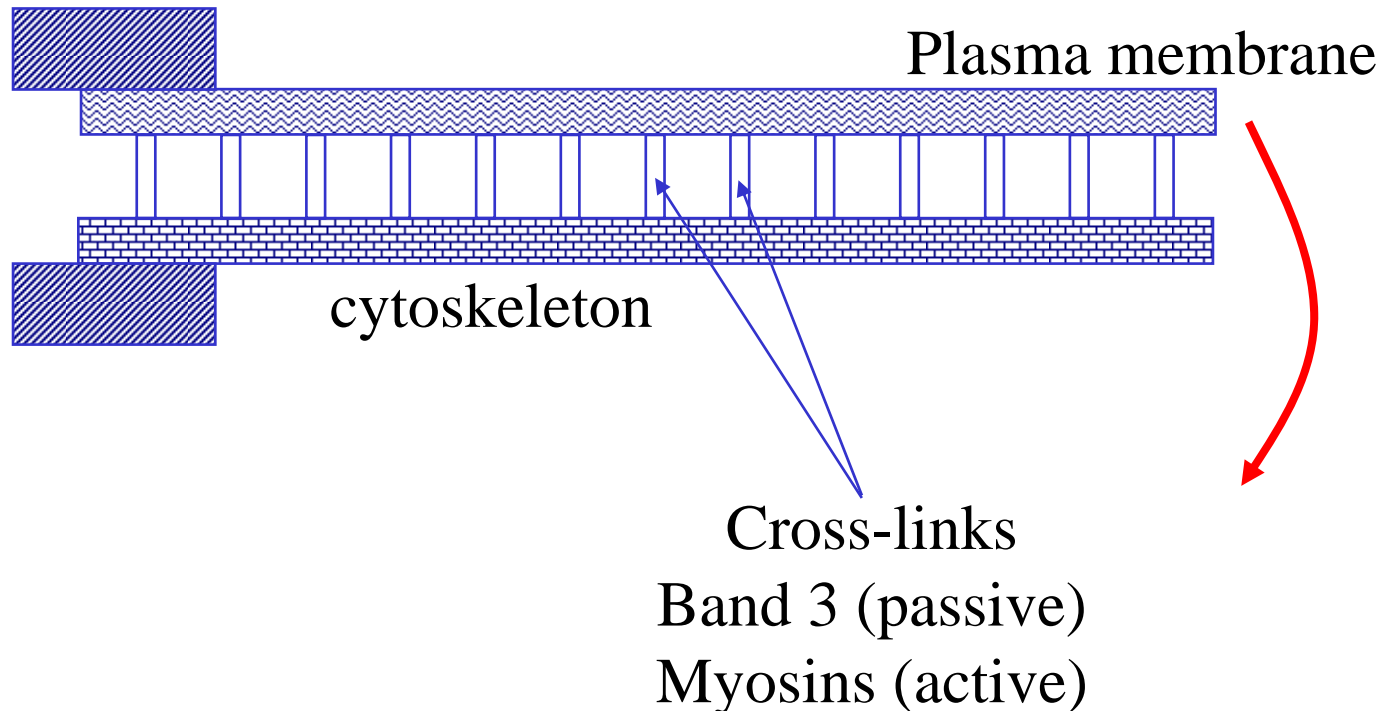
The buckling force is in the nanoNewton range

- this requires >10,000 myosin molecules

Gross mechanical properties of the plasma membrane depend upon underlying structures e.g. the cytoskeleton.

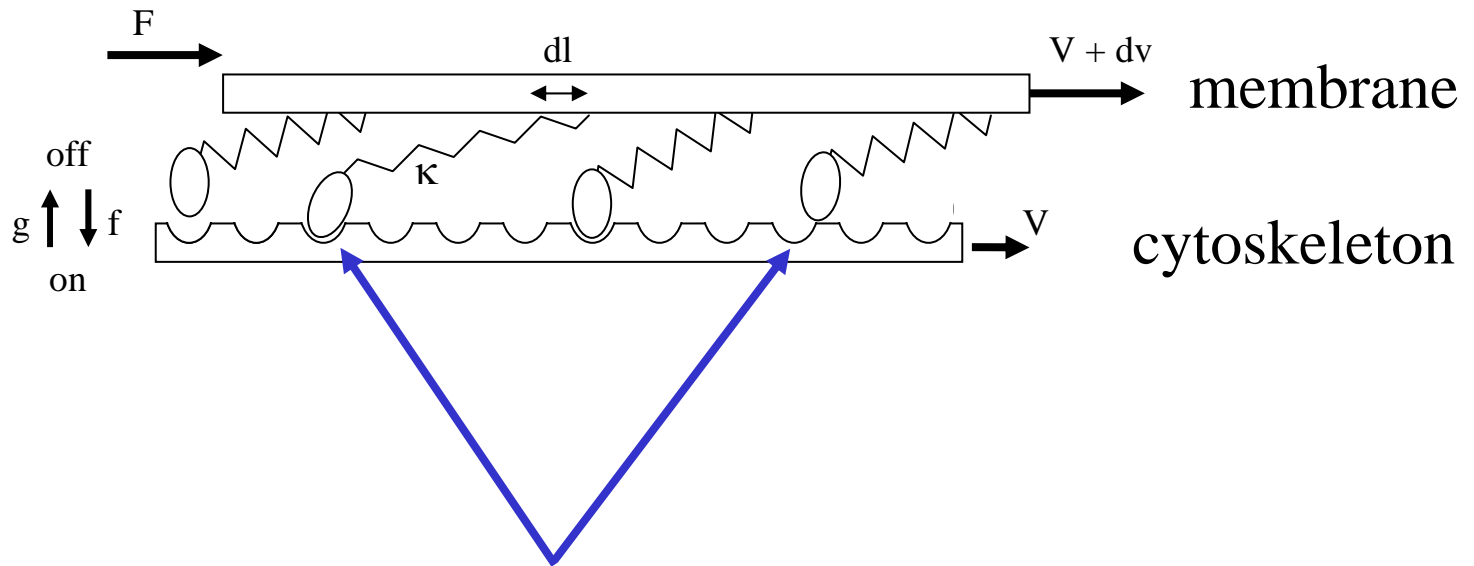


Classic “I-beam” structure – very stiff in bending





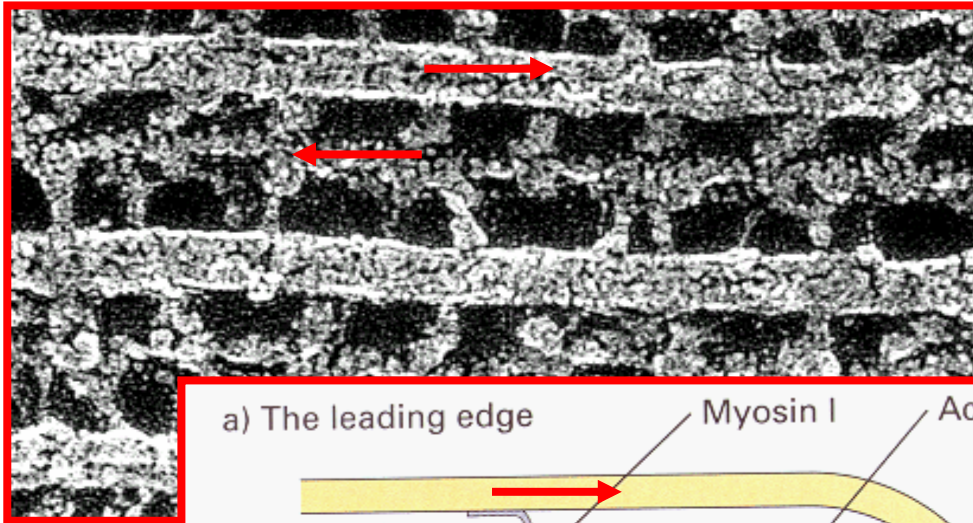
# The cross-links can be active or passive - but all are dynamic.



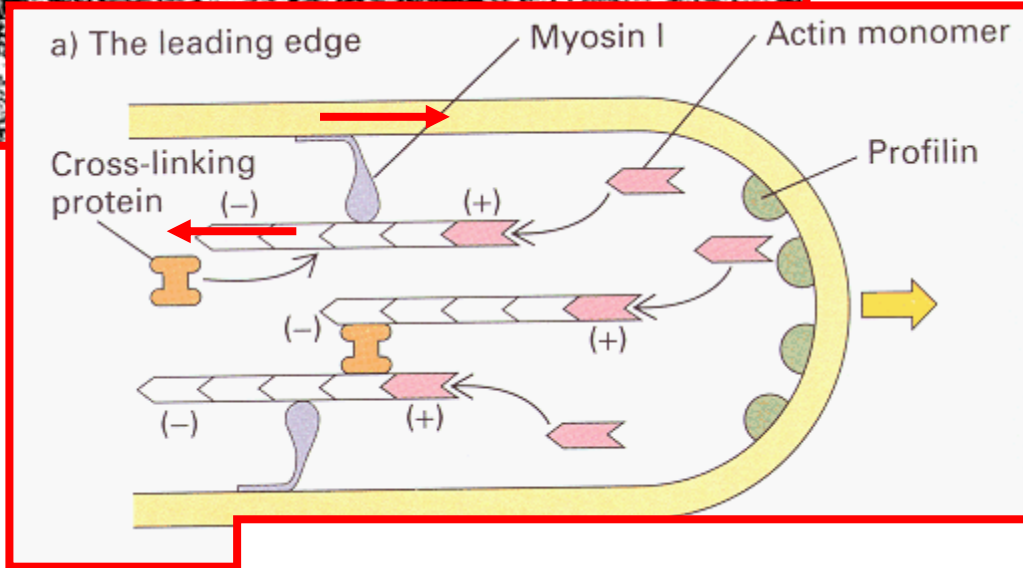
Cross-links might either.....

- remain bound and resist shearing – **elastic response**
- detach and then reattach giving rise to **viscous drag force**
- actively change shape and generate **active forces** (“cortical tension”)

# Muscle and membranes

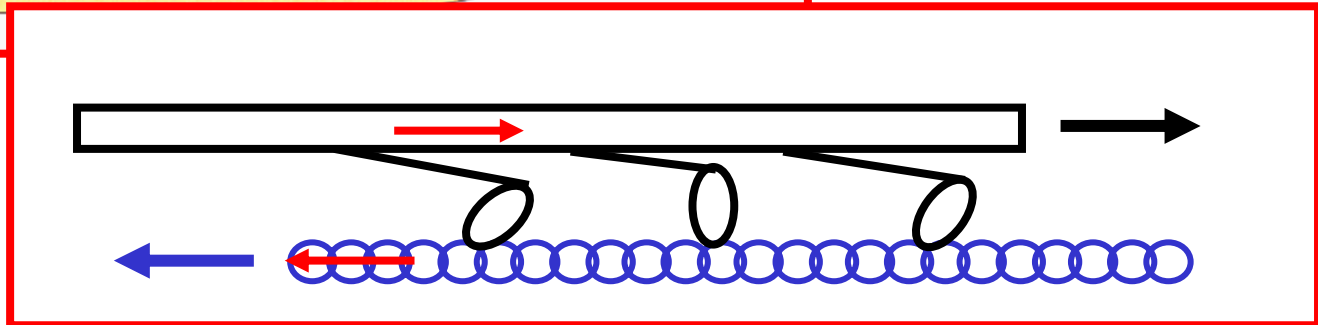


Muscle



Membrane

Model



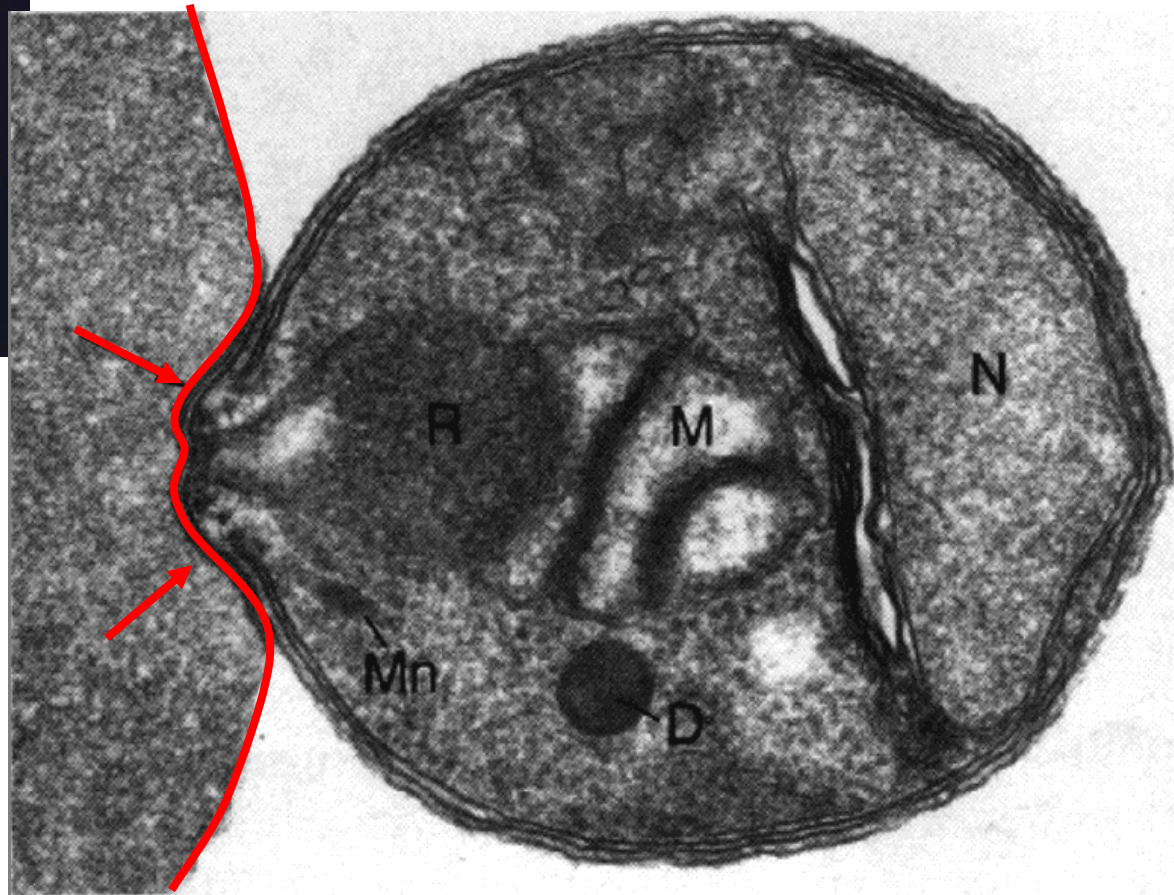
We are investigating the dynamic links made between the cytoskeleton and the plasma membrane.

- Active links power the invasion of a parasite
- Passive links resist the invasion of a parasite!
- They are responsible for rearrangement and control of cell shape (endocytosis, filopodia)
- They actively move “*integral membrane proteins*” involved in a variety of functions (e.g. receptors and channels)

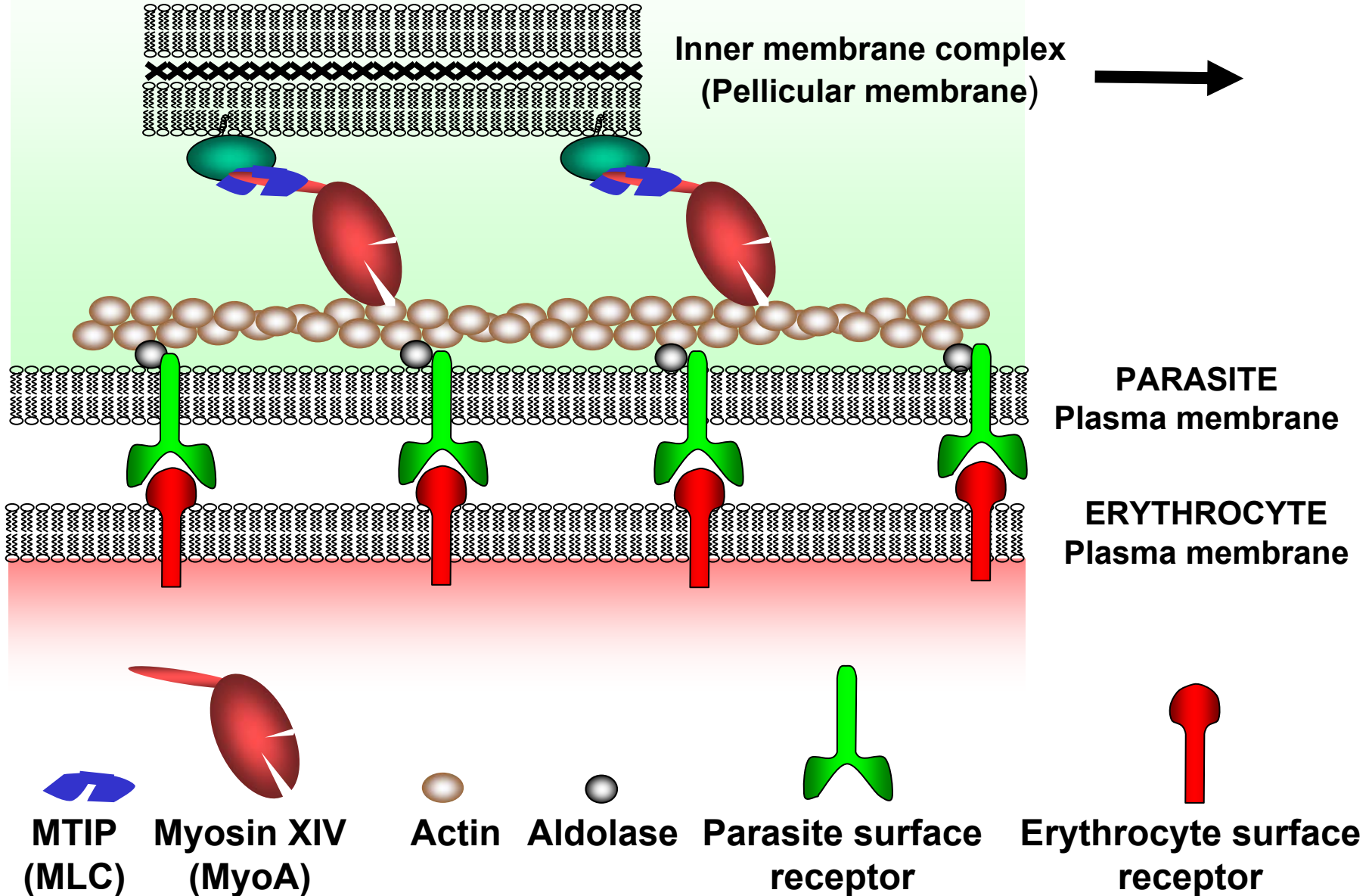
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- E.G. Cortical tension can be reduced by expression of PH domains (that interfere with cross-links)

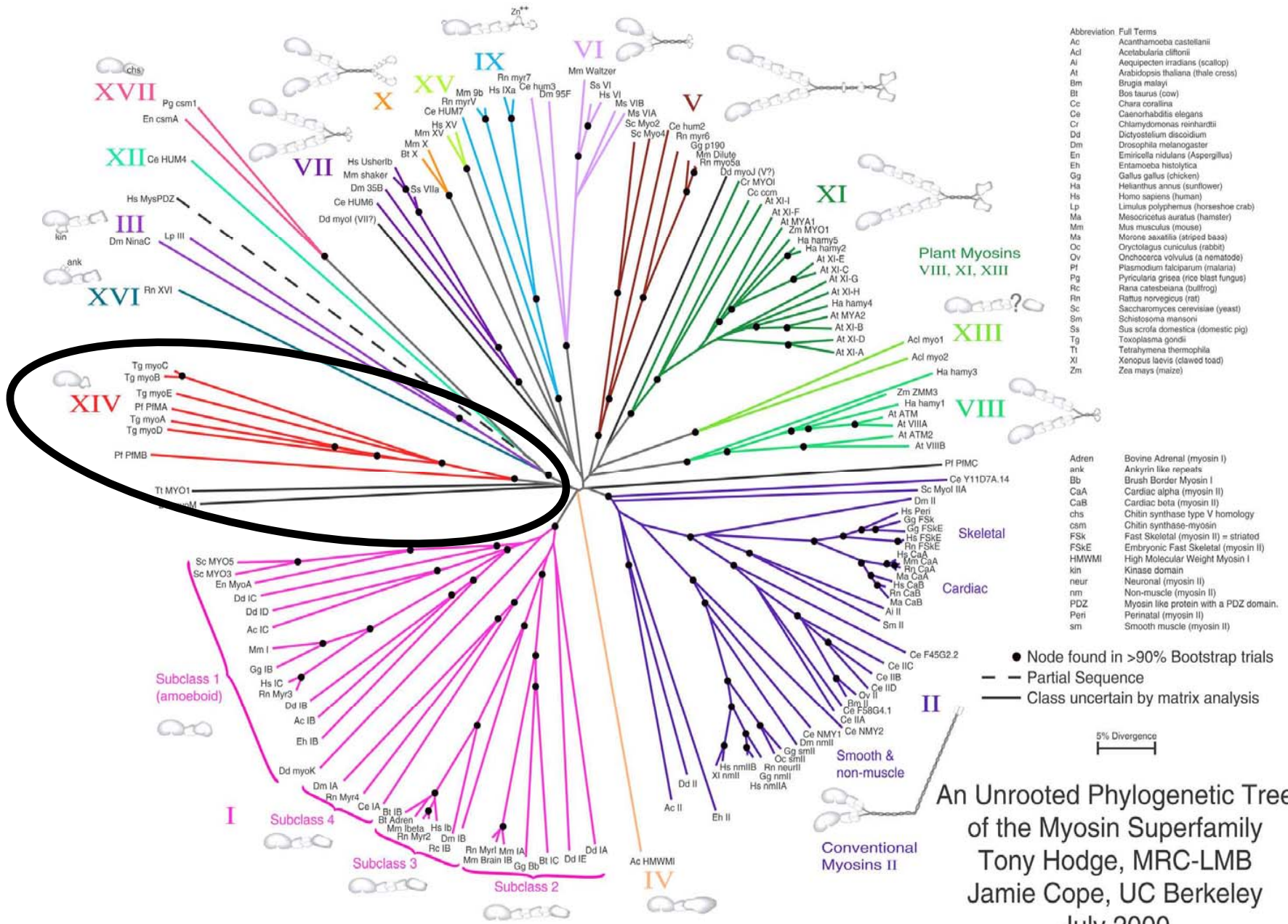
# Malarial invasion of the red blood cell



# Putative organisation of the motor complex

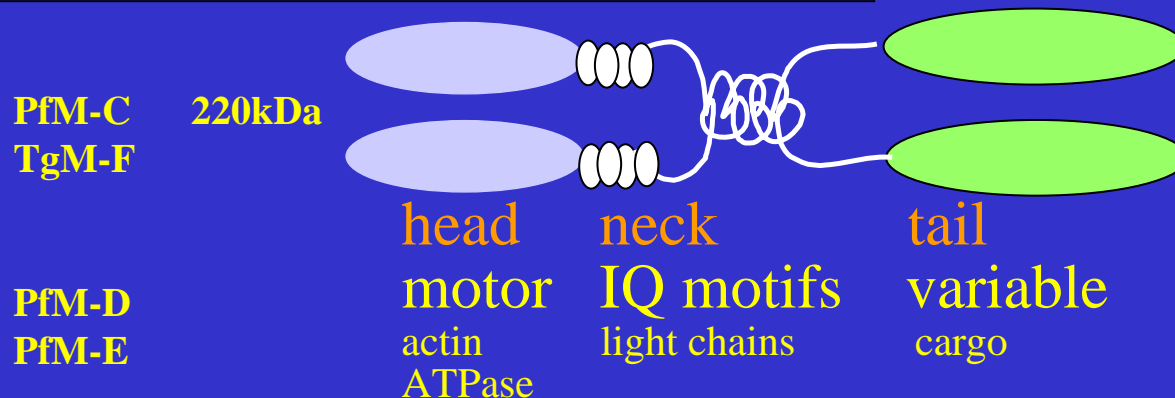
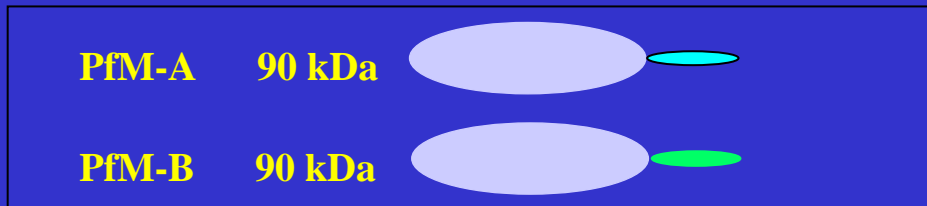
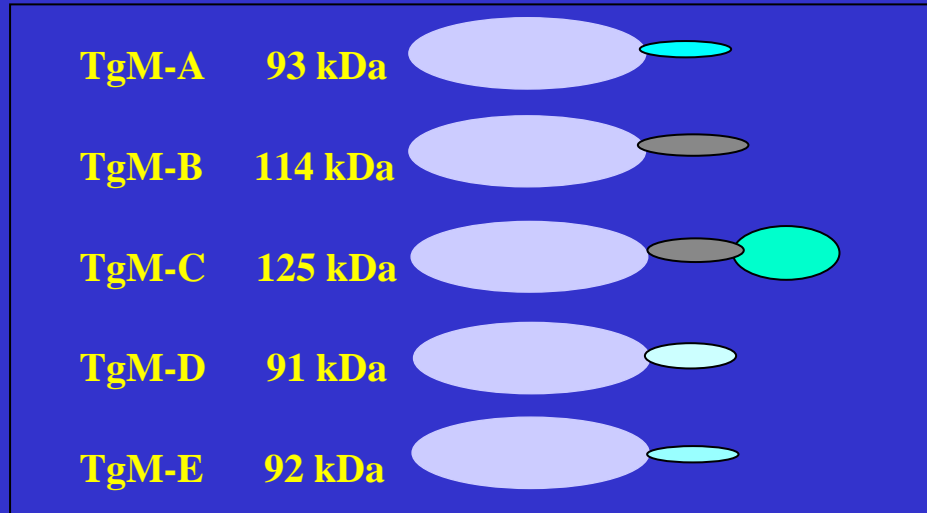


# Myosin Family Tree



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

# Class XIV Myosins



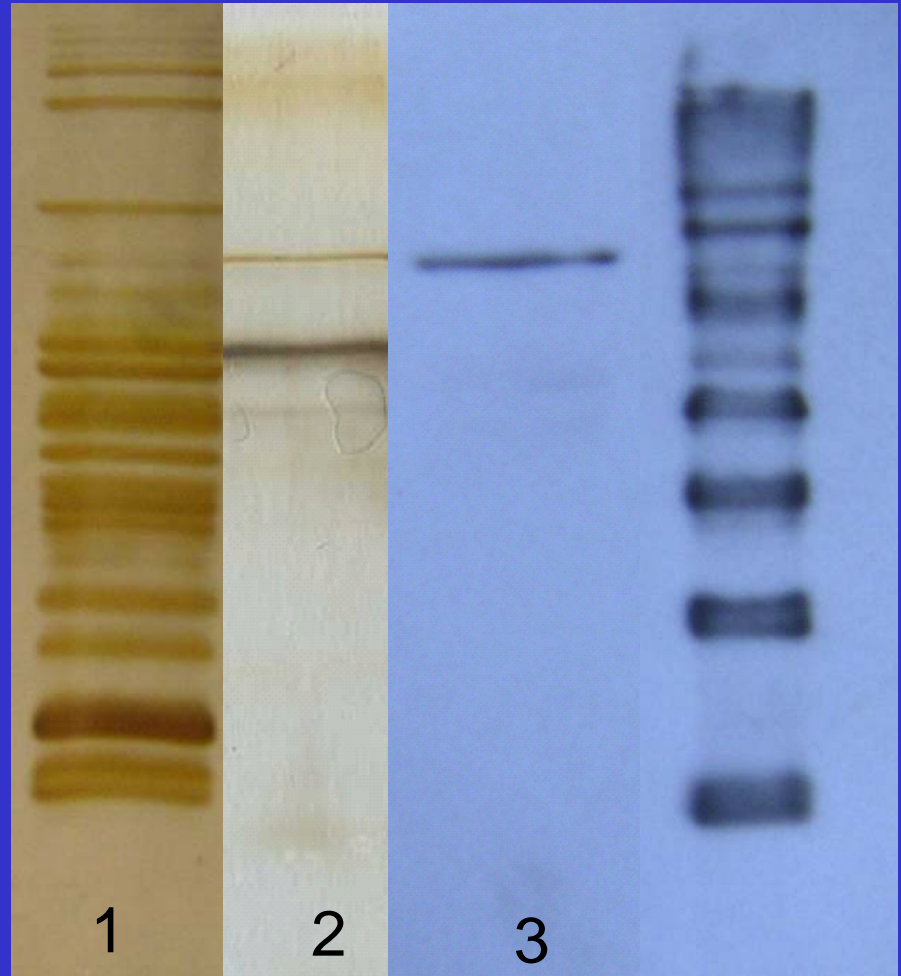
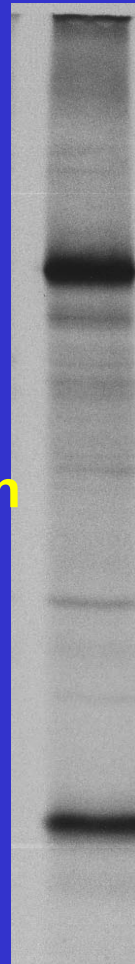
Courtesy of  
Dominique  
Soldati

# Native tissue Pfmmyo-A purification

$\text{S}^{35}$

Pfmmyo-A

MTIP



kDa

120

100

80

60

50

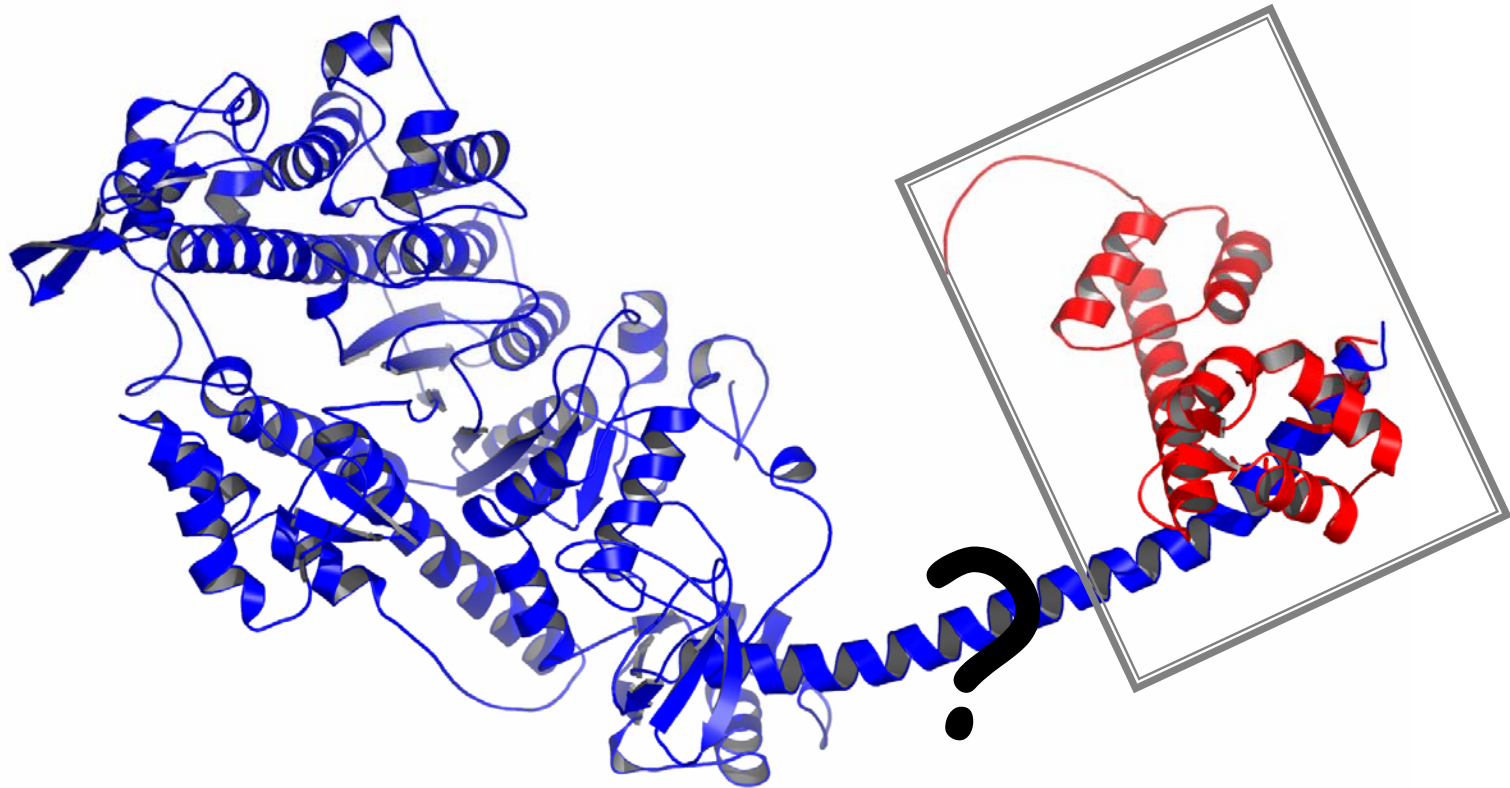
40

30

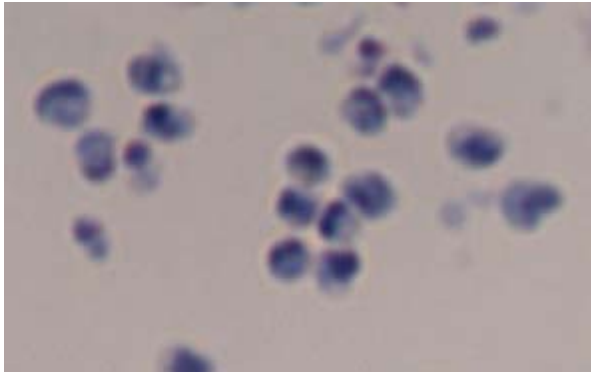
1 Crude cell lysate 2 Pfmmyo-A after peptide competition 3 Pfmmyo-A western

- MTIP immunoprecipitation
- Compete Pfmmyo-A





# Myosin XIV *in vitro* motility



Purified merozoites

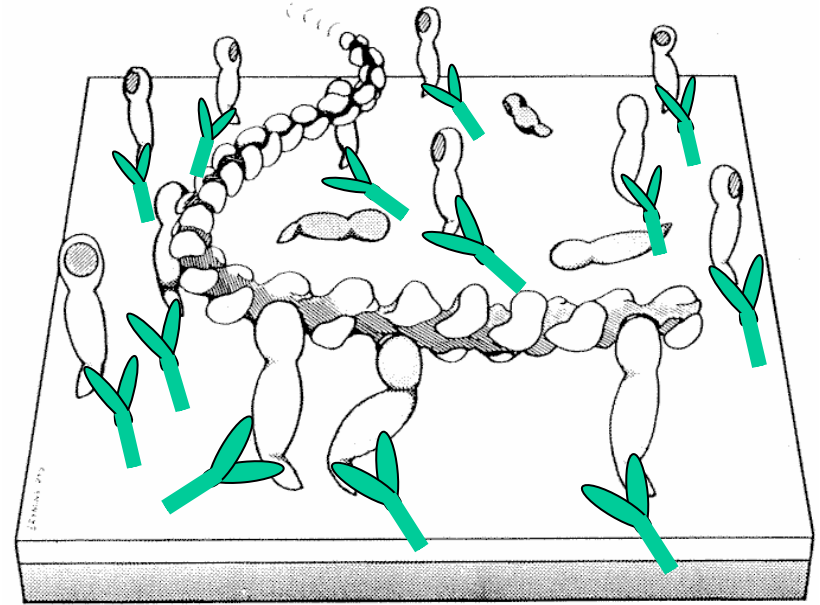
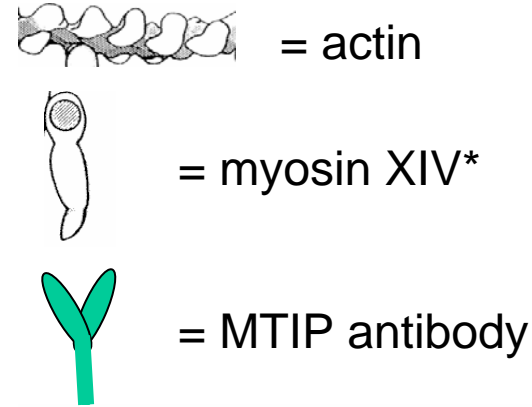


Freeze thaw cycle  
80µl hypotonic buffer  
Protease inhibitors

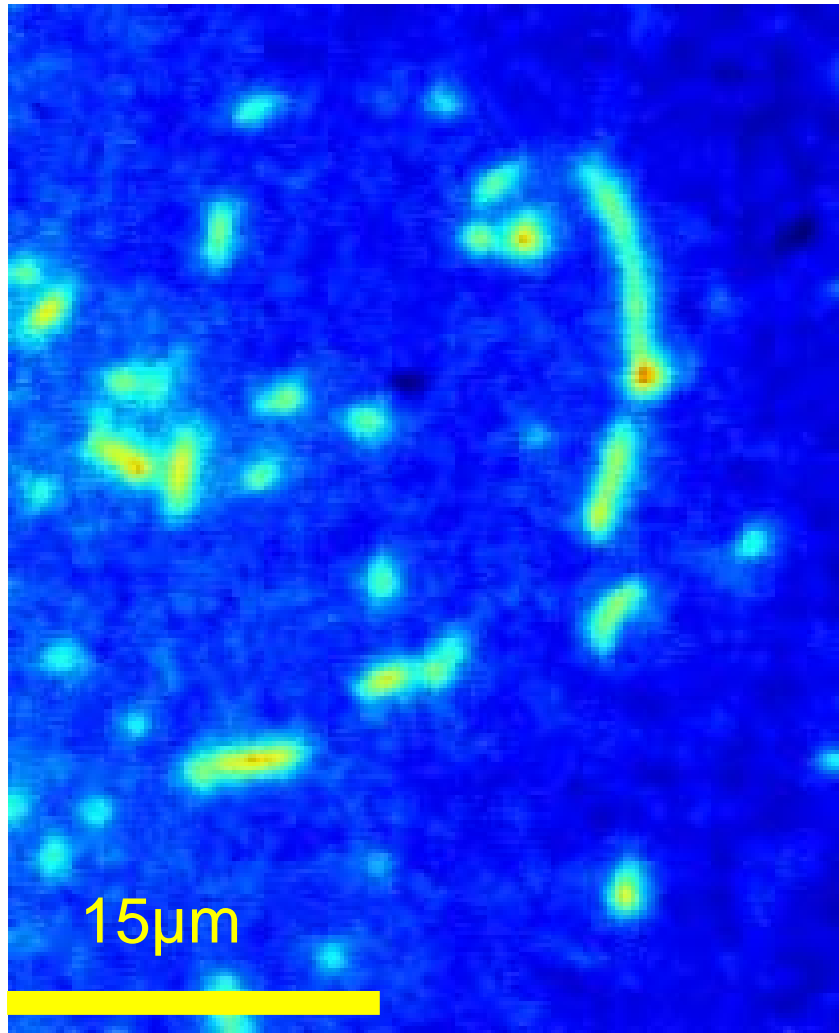
100k spin  
S/N removed



*In vitro* motility  
assay

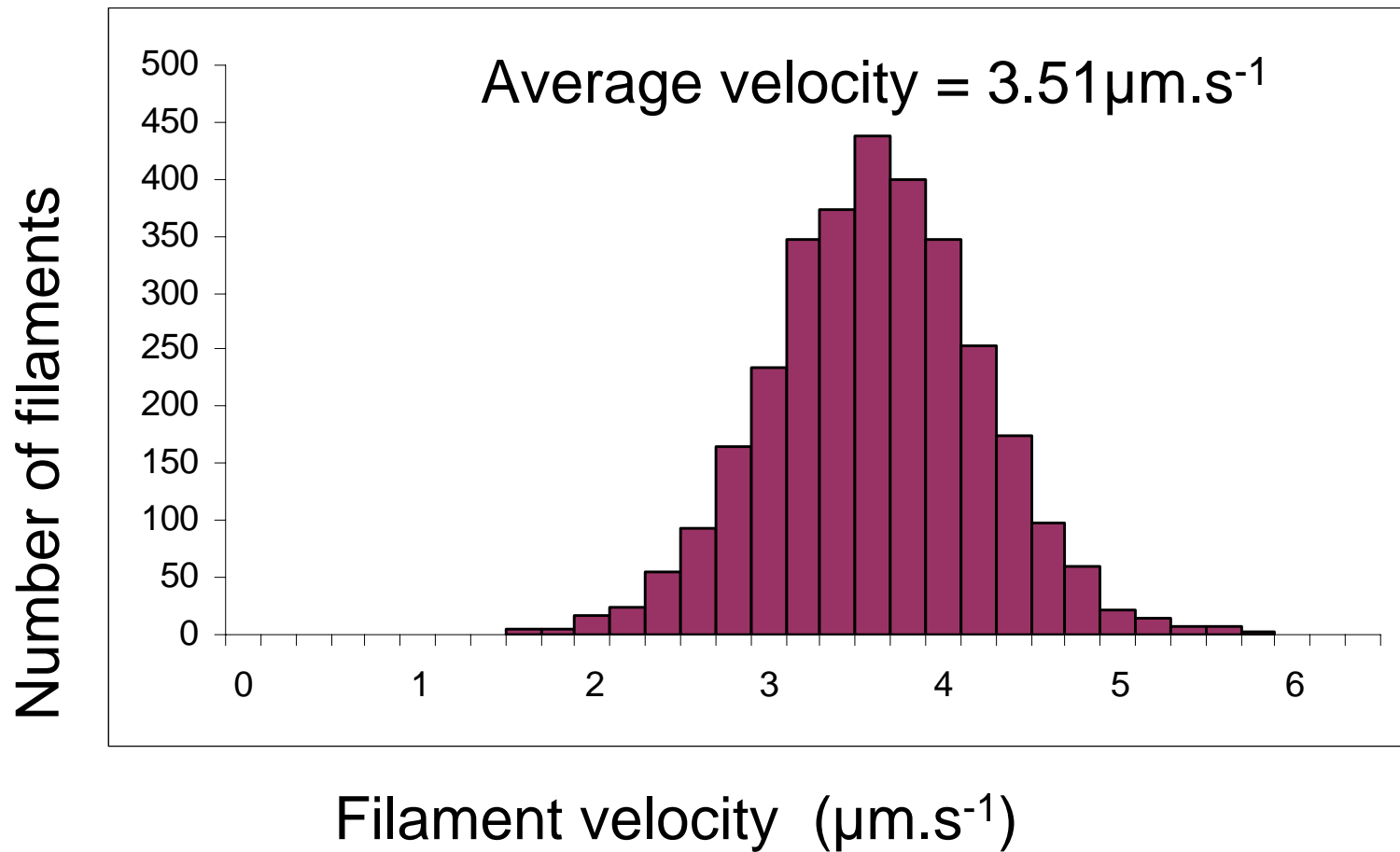


# Myosin XIV in vitro motility

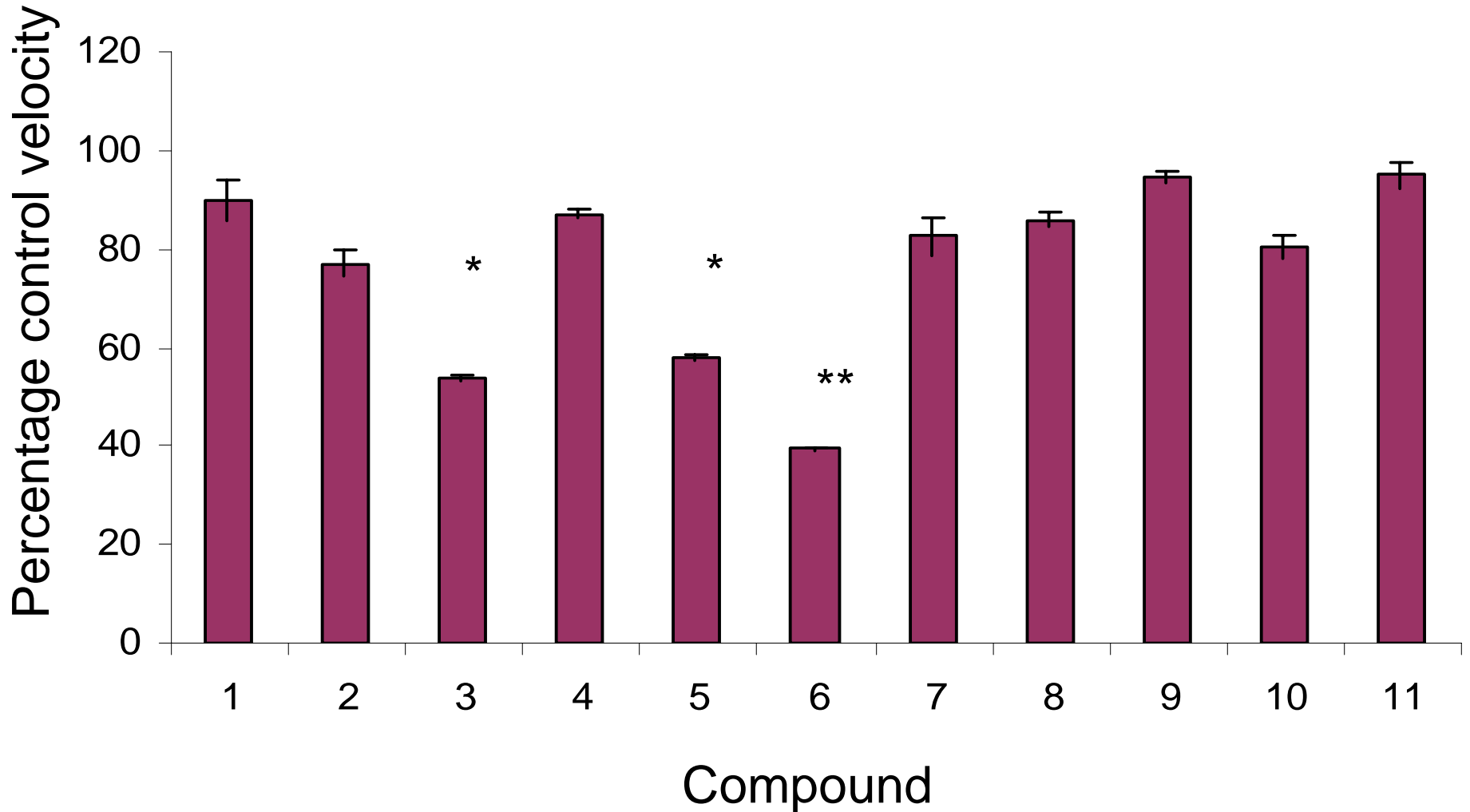


actin sliding velocity  
=  $3.51 \mu\text{m}\cdot\text{s}^{-1}$   
At  $25^\circ \text{C}$

# *Pfmyo-A in vitro* motility



# Blebbistatin analogues



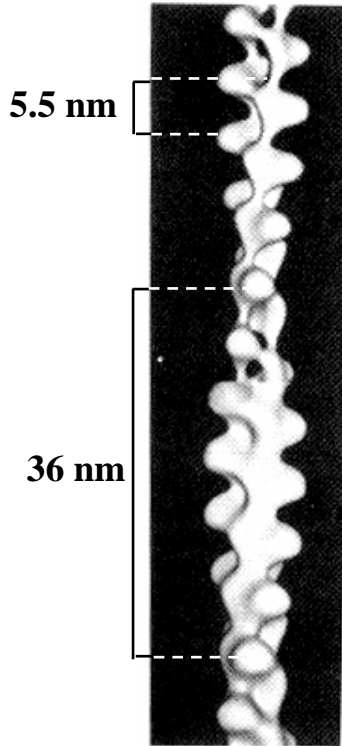
# A brief interlude on processive motors

## “Processive” and “Intermittent” motors

- Most myosins and many kinesins interact in an **“*Intermittent*”** manner with their track. They must work in teams to produce large movements and forces.
- kinesin 1, myosin V, and most DNA processing enzymes are **“*Processive*”** motors and take many steps before detaching from their track. They work as single molecules.
- Some motors (e.g. myosin VI) can modulate their properties.

ACTIN FILAMENT  
(repeats every 5nm  
and every 36nm)

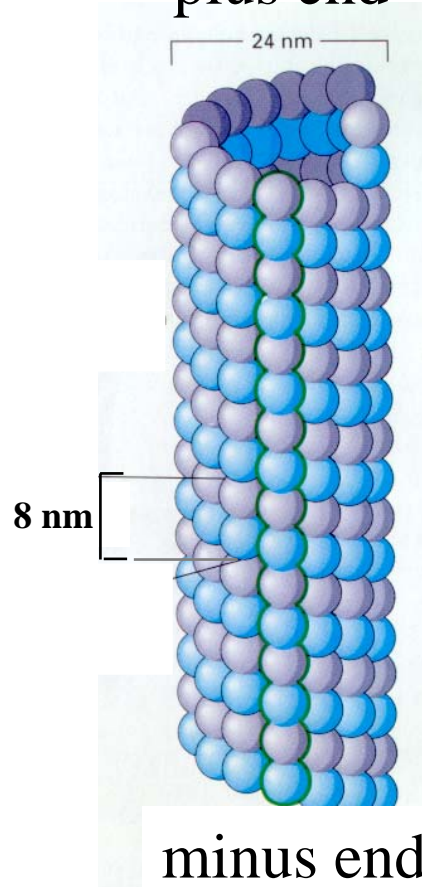
Minus end - pointed end



Plus end - barbed end

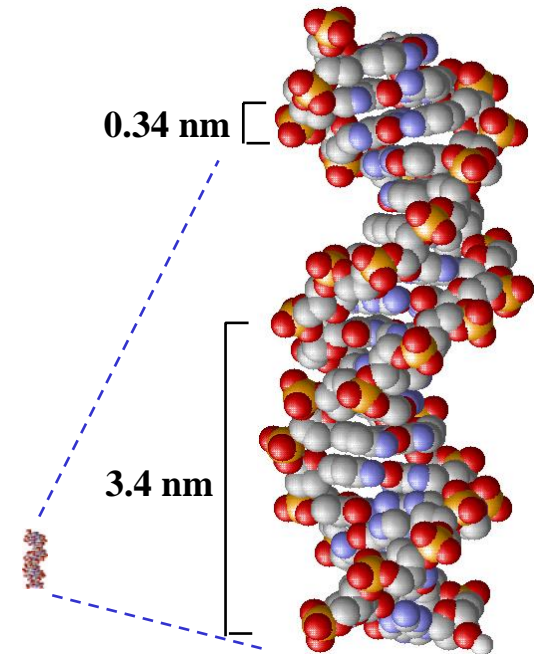
MICROTUBULE  
(repeats every 8nm)

plus end



DNA  
(repeats every 0.34nm)

3' & 5'



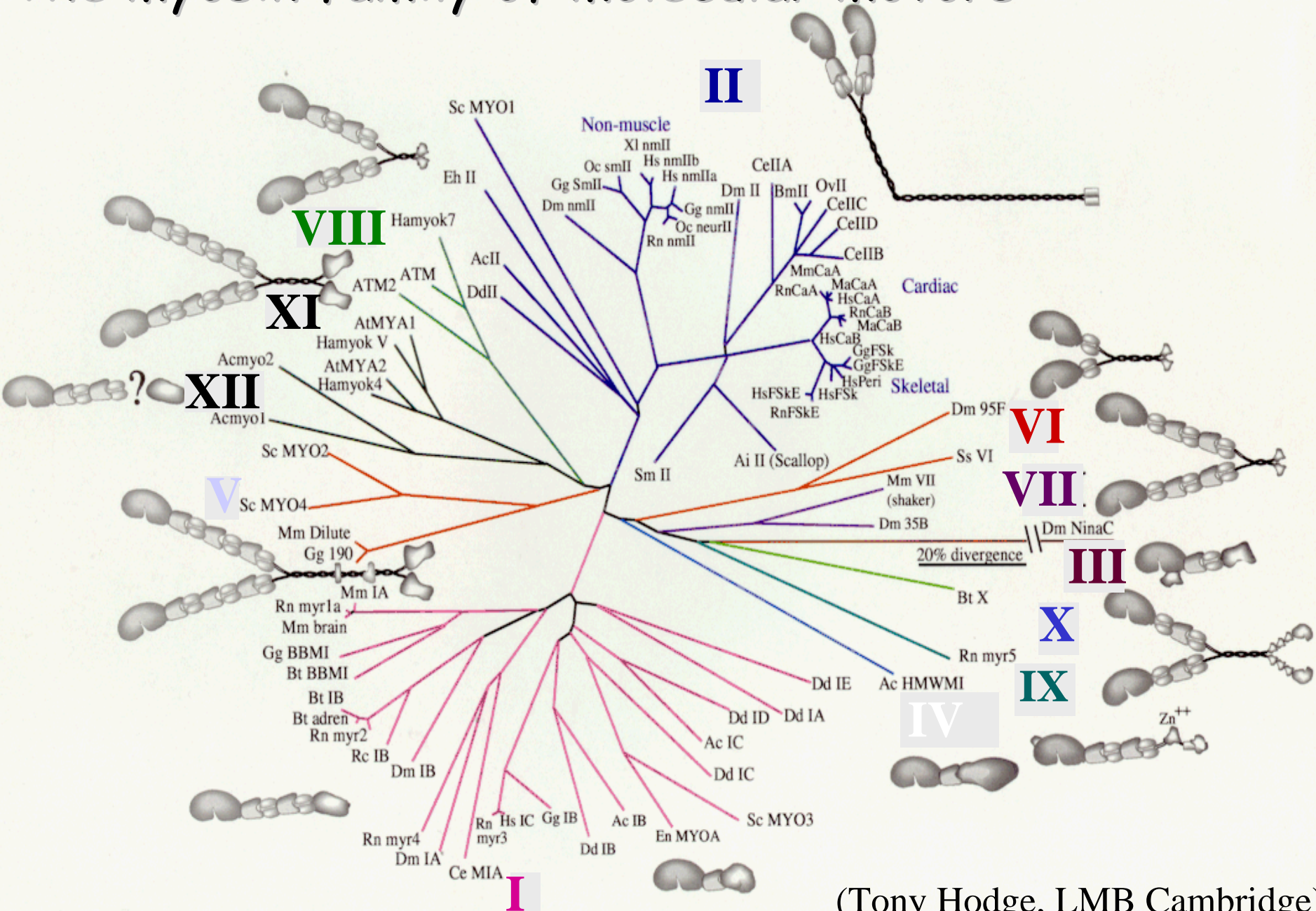
3' & 5'

(not to scale)

(Note: some cell motilities are driven by filament polymerisation)

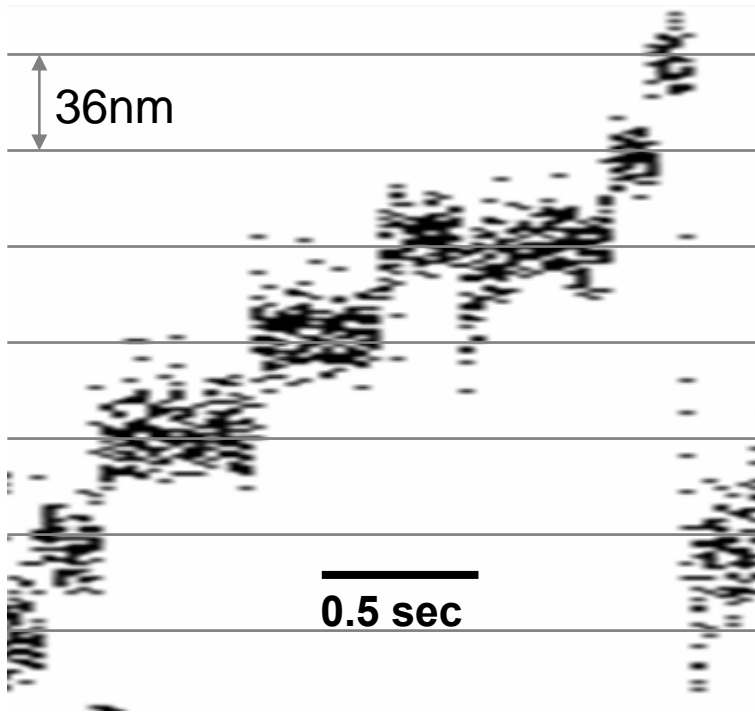
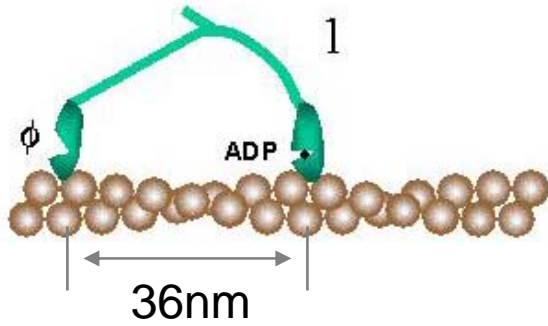


# The myosin family of molecular motors:



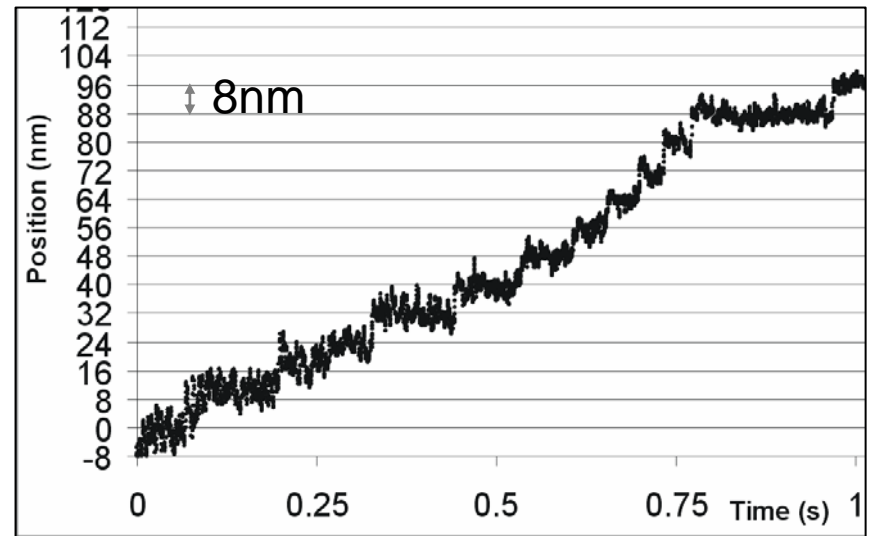
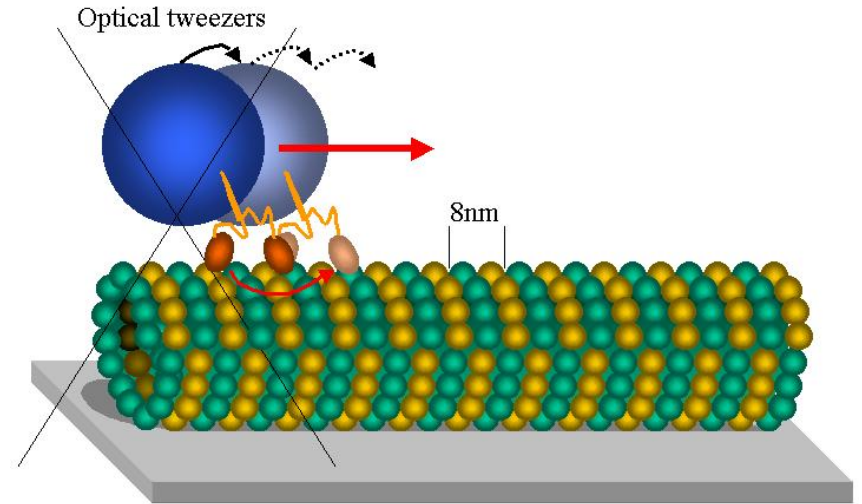
(Tony Hodge, LMB Cambridge)

# Myosin V



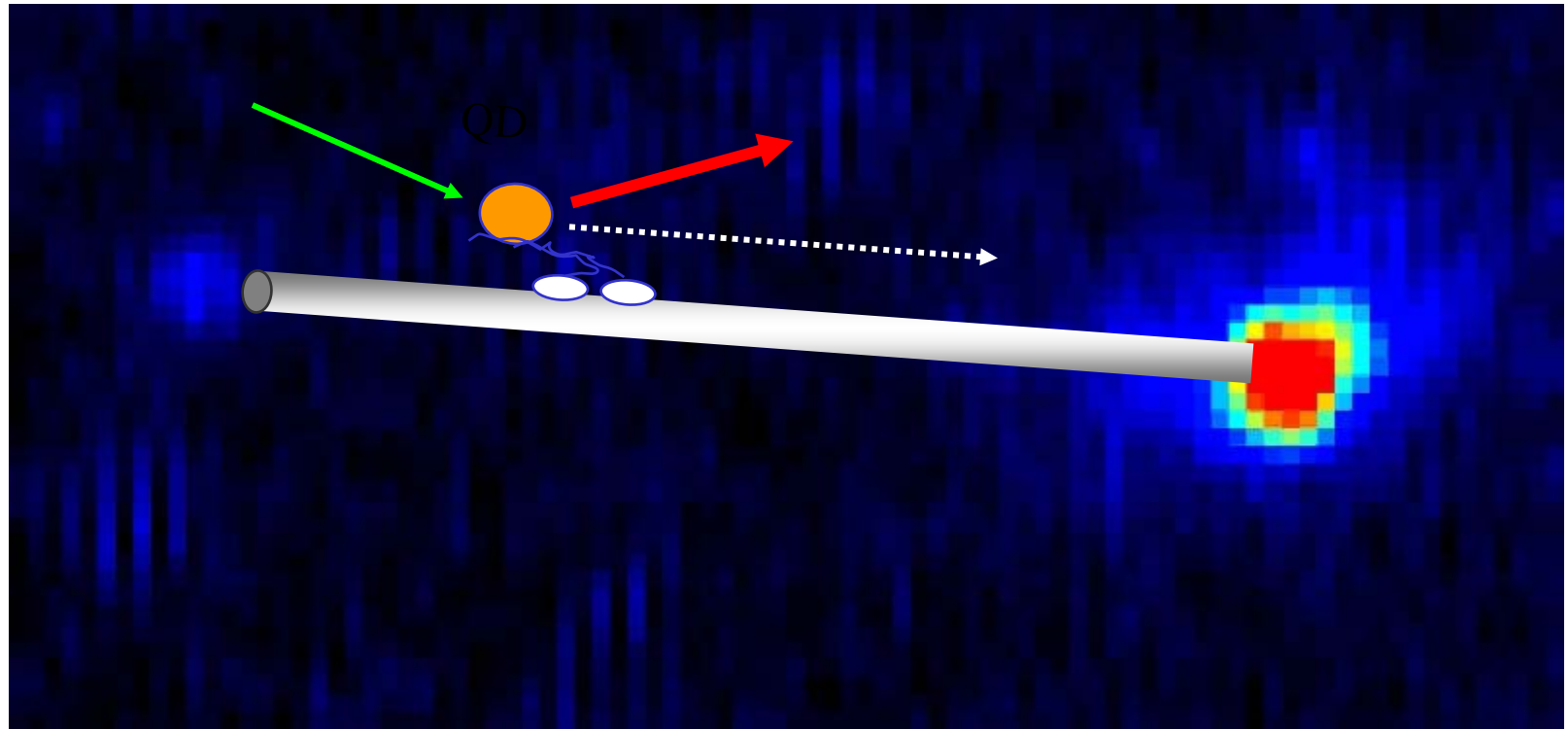
Veigel & Molloy

# Conventional kinesin



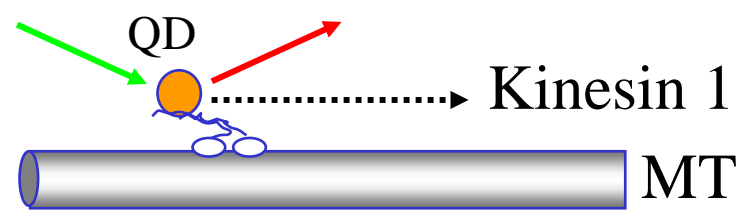
Carter & Cross

# MOVEMENT OF AN INDIVIDUAL KINESIN ON A MICROTUBULE VISUALISED USING A QUANTUM DOT



2  $\mu\text{m}$

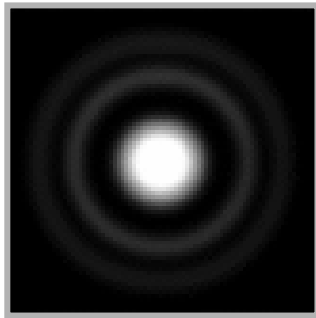
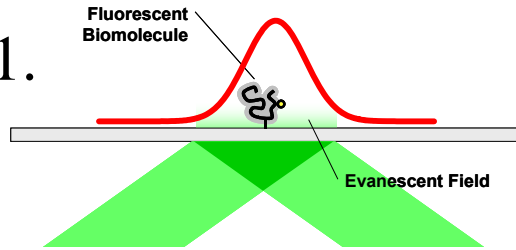
real time



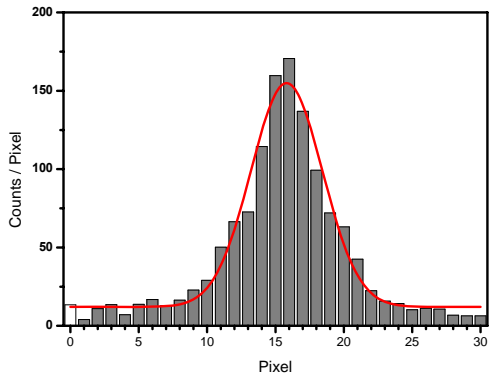
Mellor, Toba, Oiwa & Molloy  
unpublished data

# Single fluorophore localisation

1.

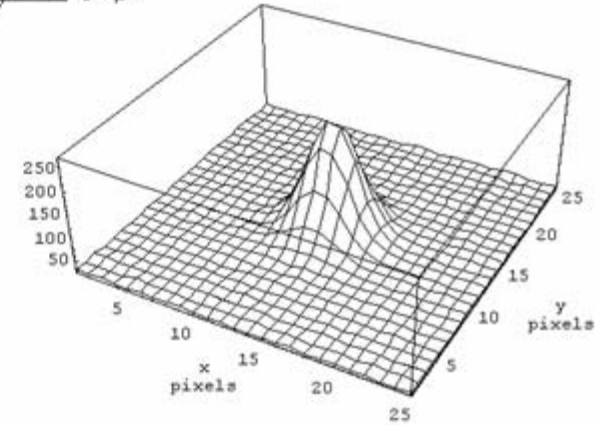
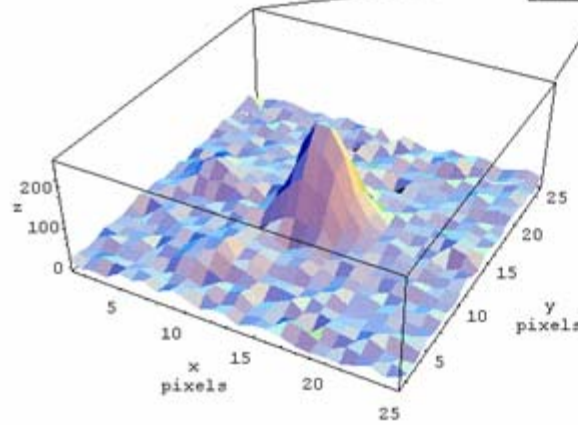
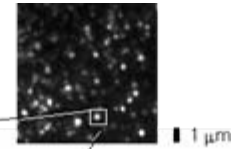


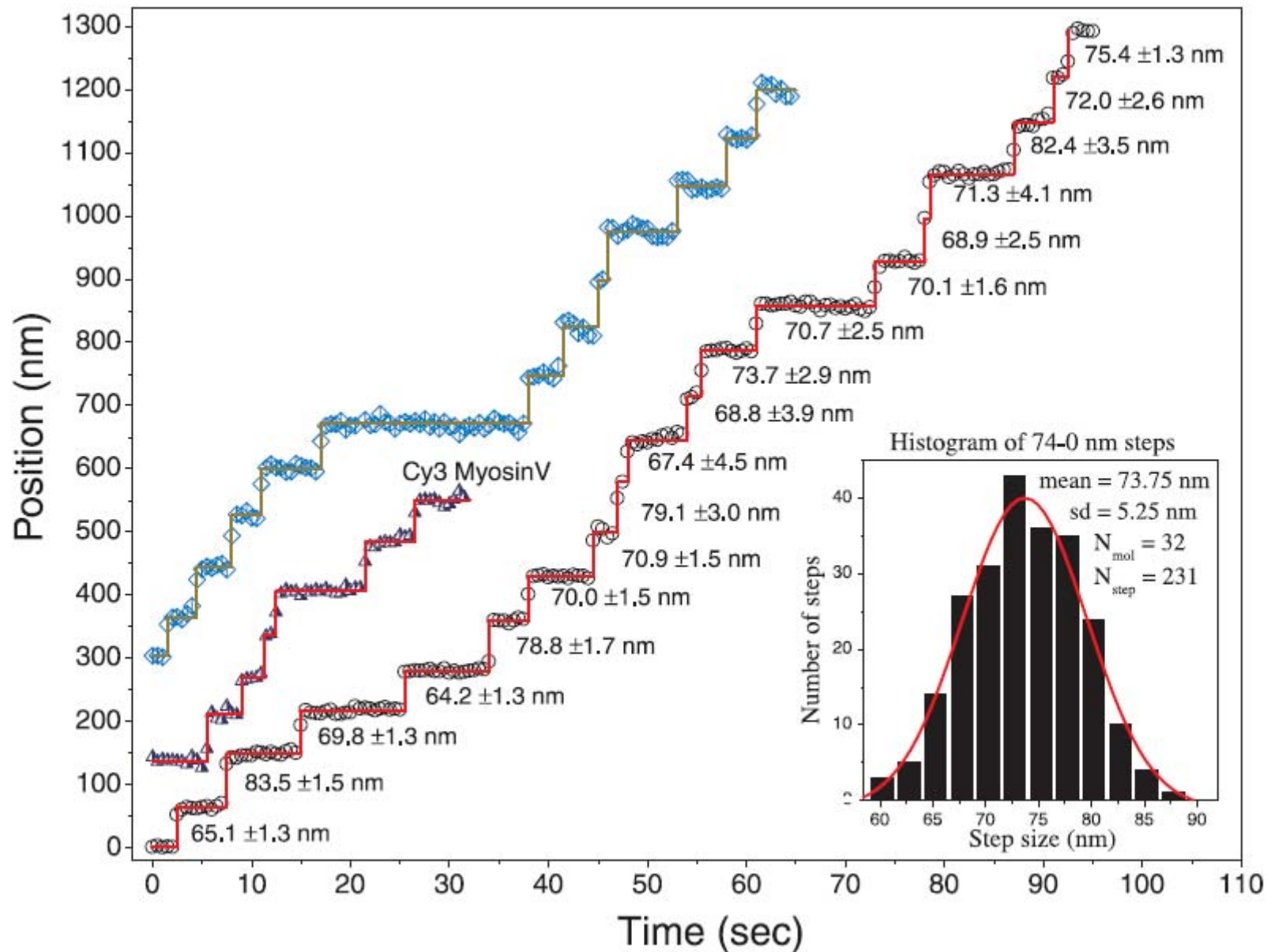
250nm



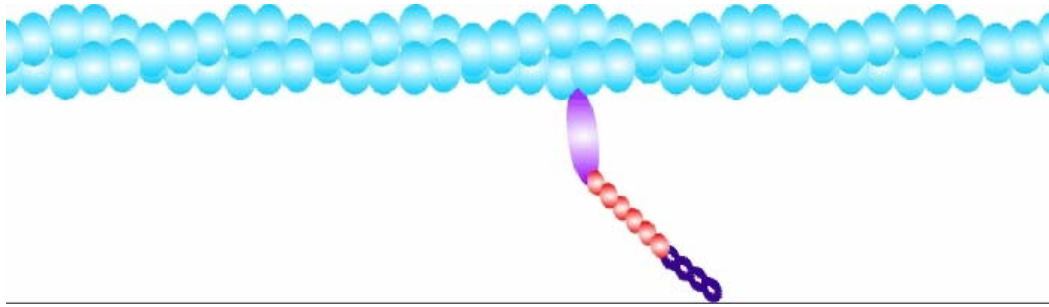
$$SEM = \sigma / \sqrt{n}$$

2.

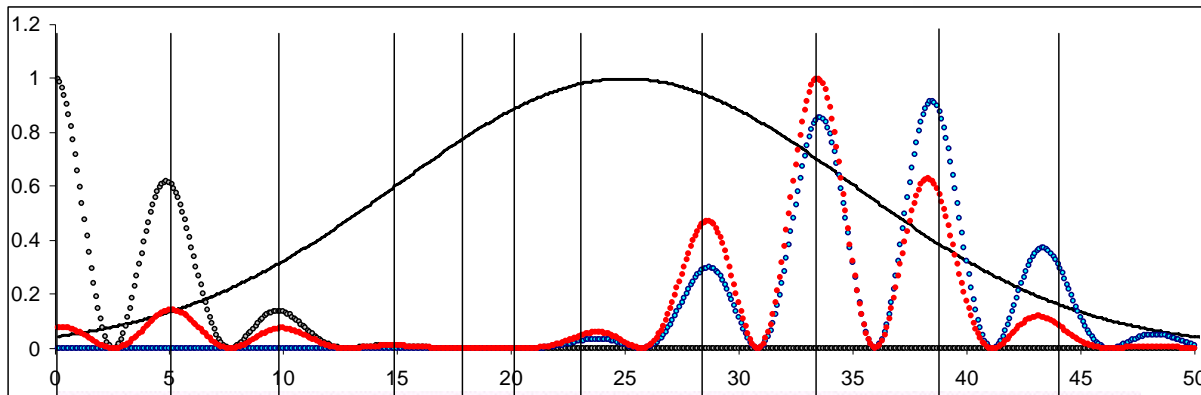
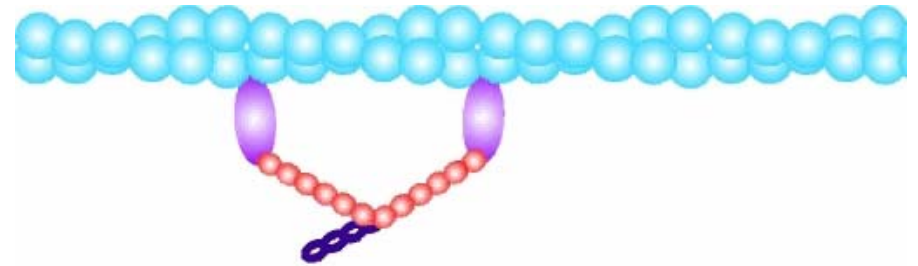




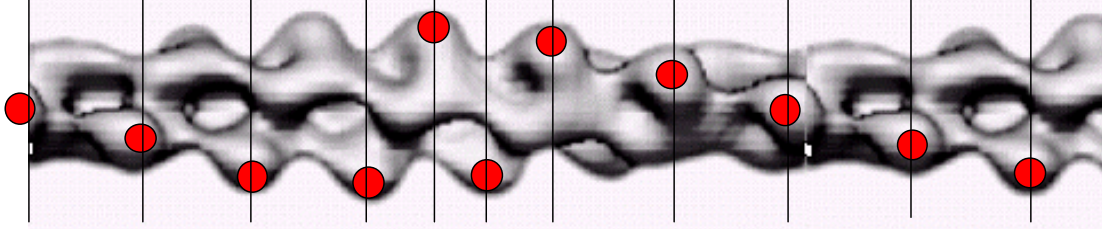
Yildiz, A., et al. (2003). "Myosin V walks hand-over-hand: Single fluorophore imaging with 1.5-nm localization." *Science* **300**(5628): 2061-2065.



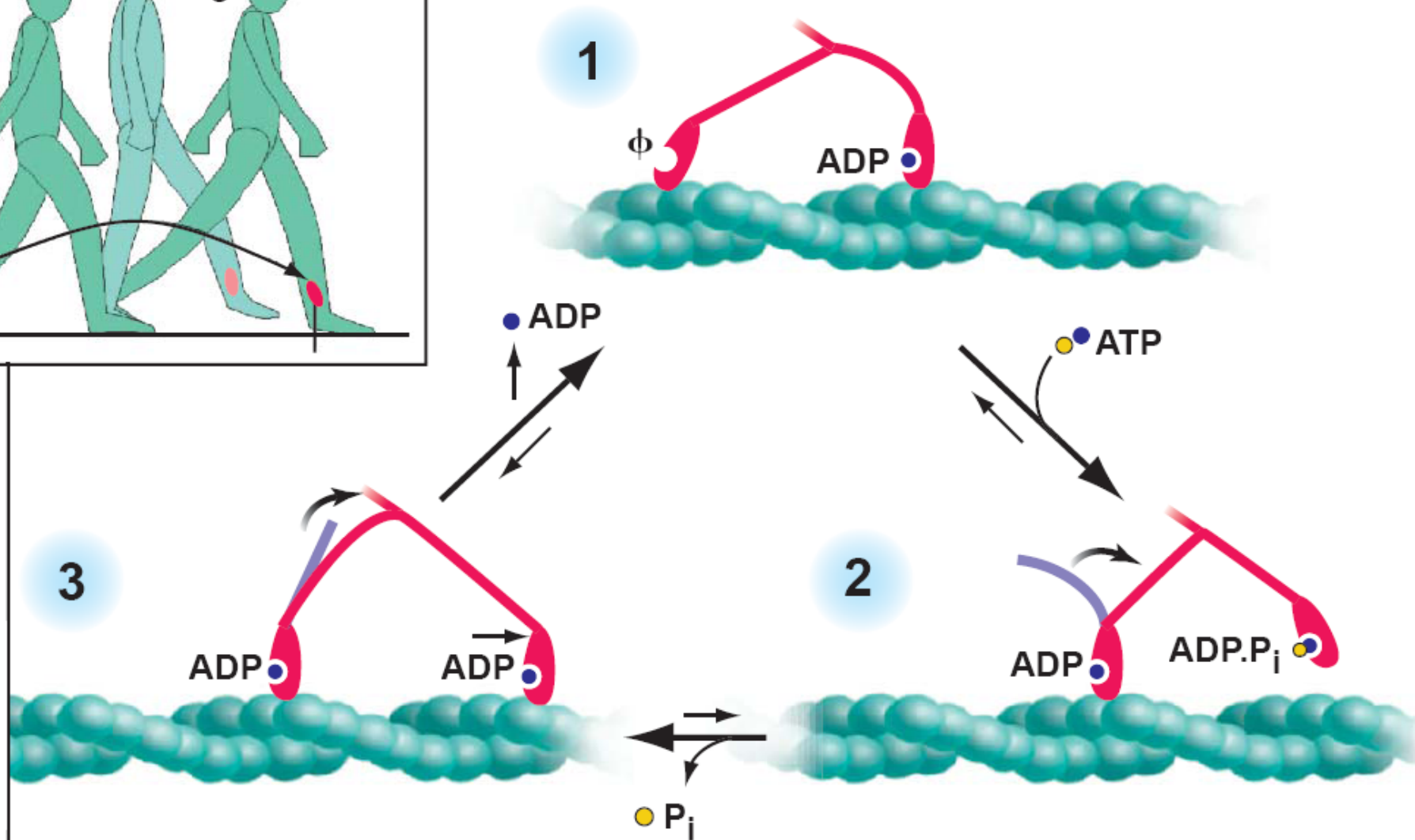
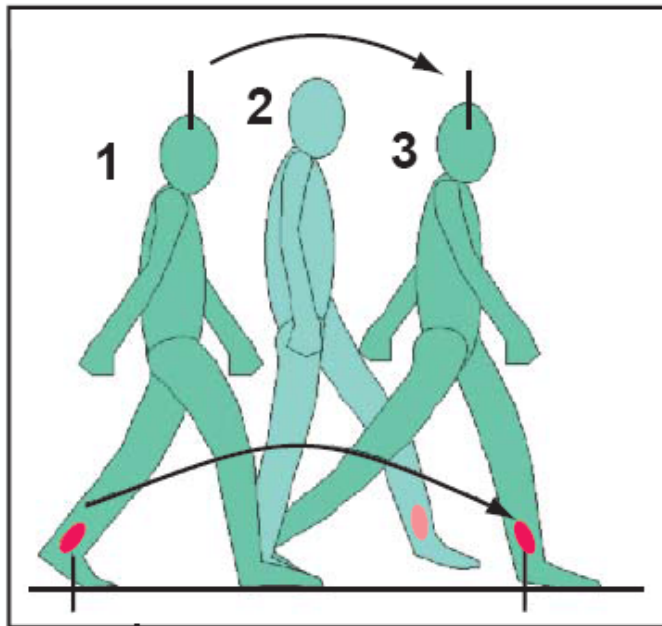
Intermittent

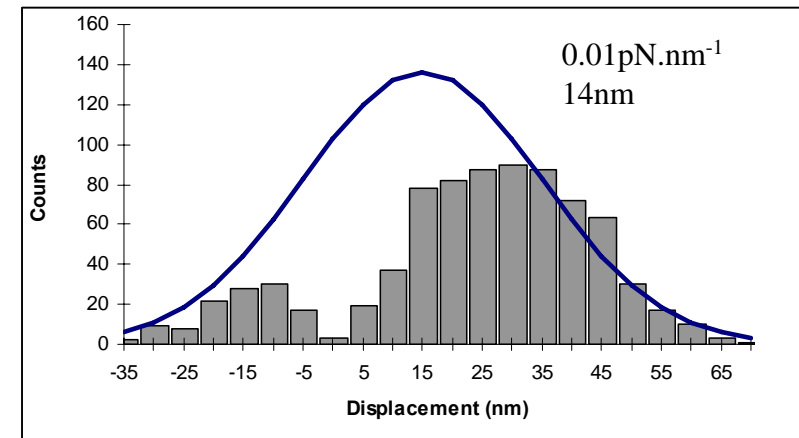
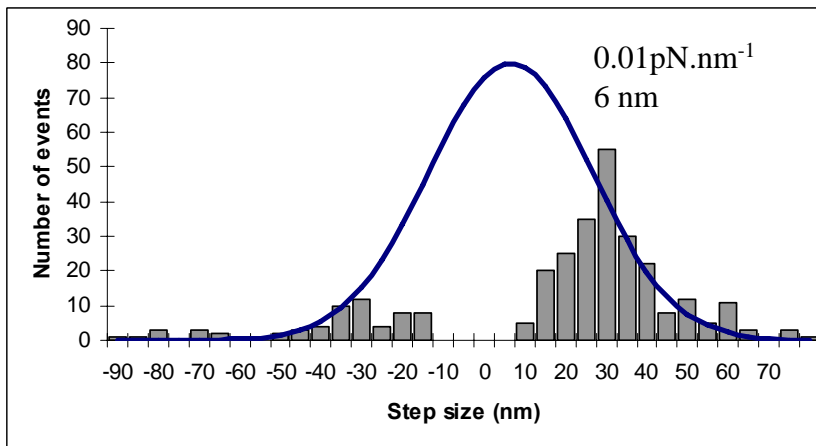
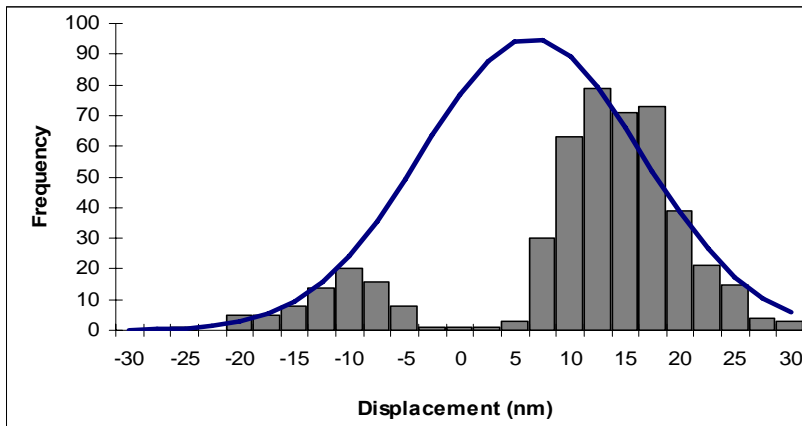


Processive



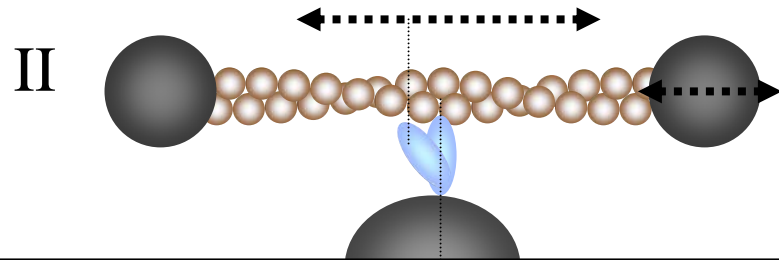
# Myosin V cycle





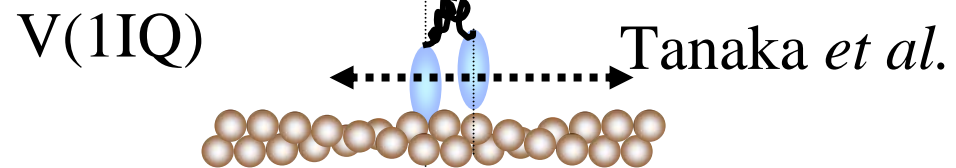
## Intermittent interactions

Myosin fixed – actin fluctuates in position



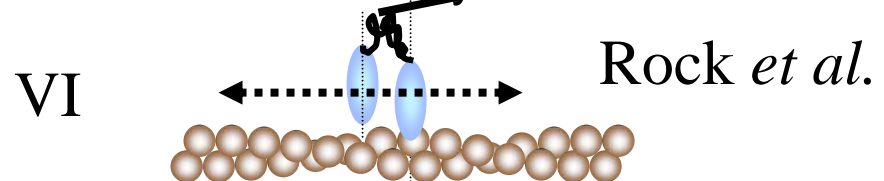
## Processive interactions

Actin fixed – free myosin head fluctuates



## Processive interactions

Actin fixed – free myosin head fluctuates





# Many myosins exert forces on membranes

## Examples:

Myosin **Ib** – membrane tensioning in the microvilli of the gut

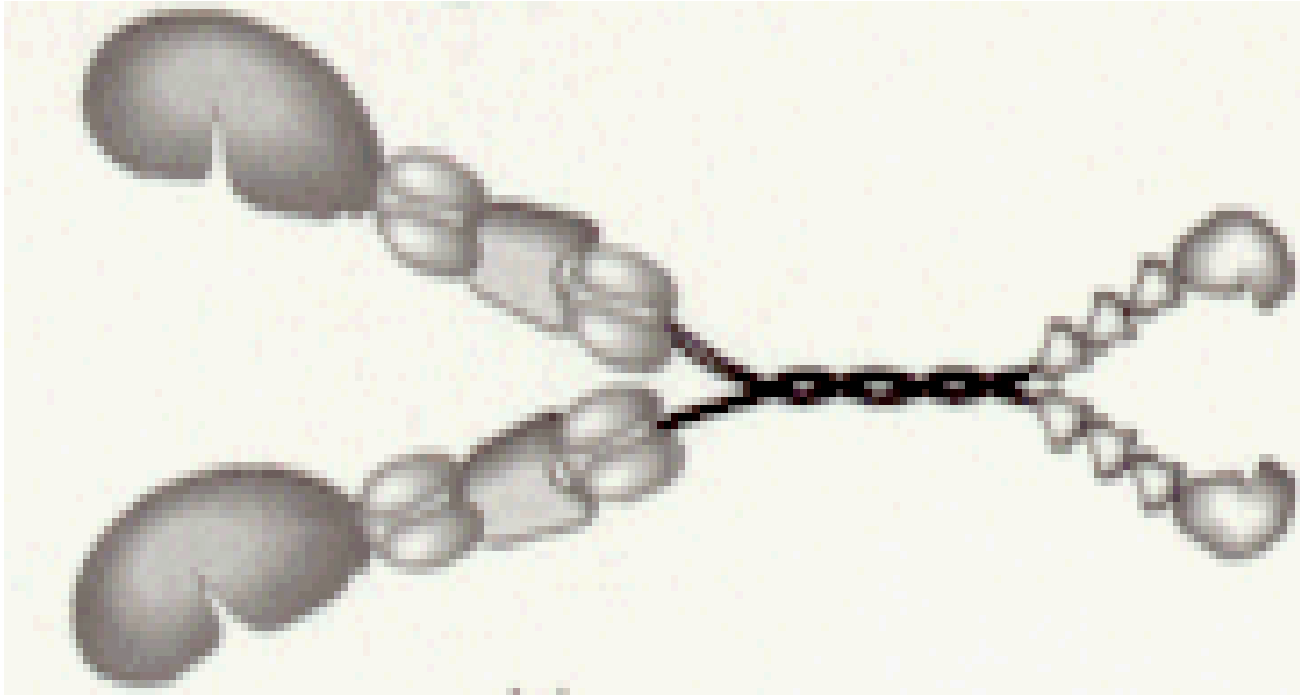
Myosin **Ic** – regulates the position of a sensory channel (in the ear)

Myosin **VI** – involved in endocytosis

Myosin **X** - inserts onto the plasma membrane via PH domains  
(function is unknown)

Myosin **XIV** – responsible for motility of Apicomplexans  
e.g. invasion motor of *Plasmodium falciparum* (malaria)

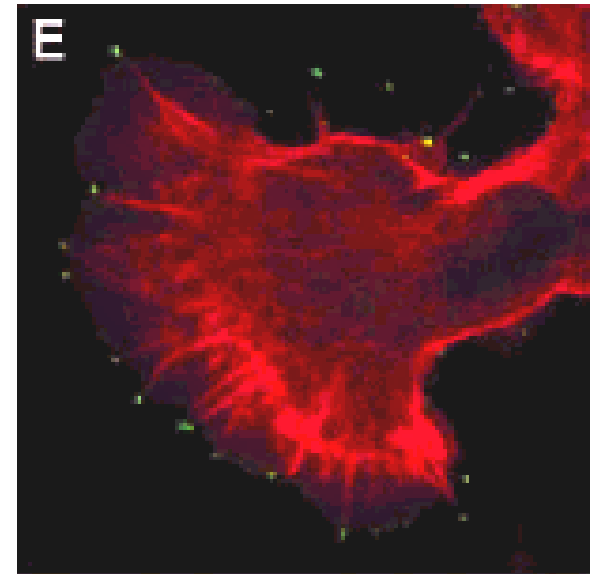
# Cellular targeting of Myosin X



Using Total Internal Reflection  
Fluorescence Microscopy

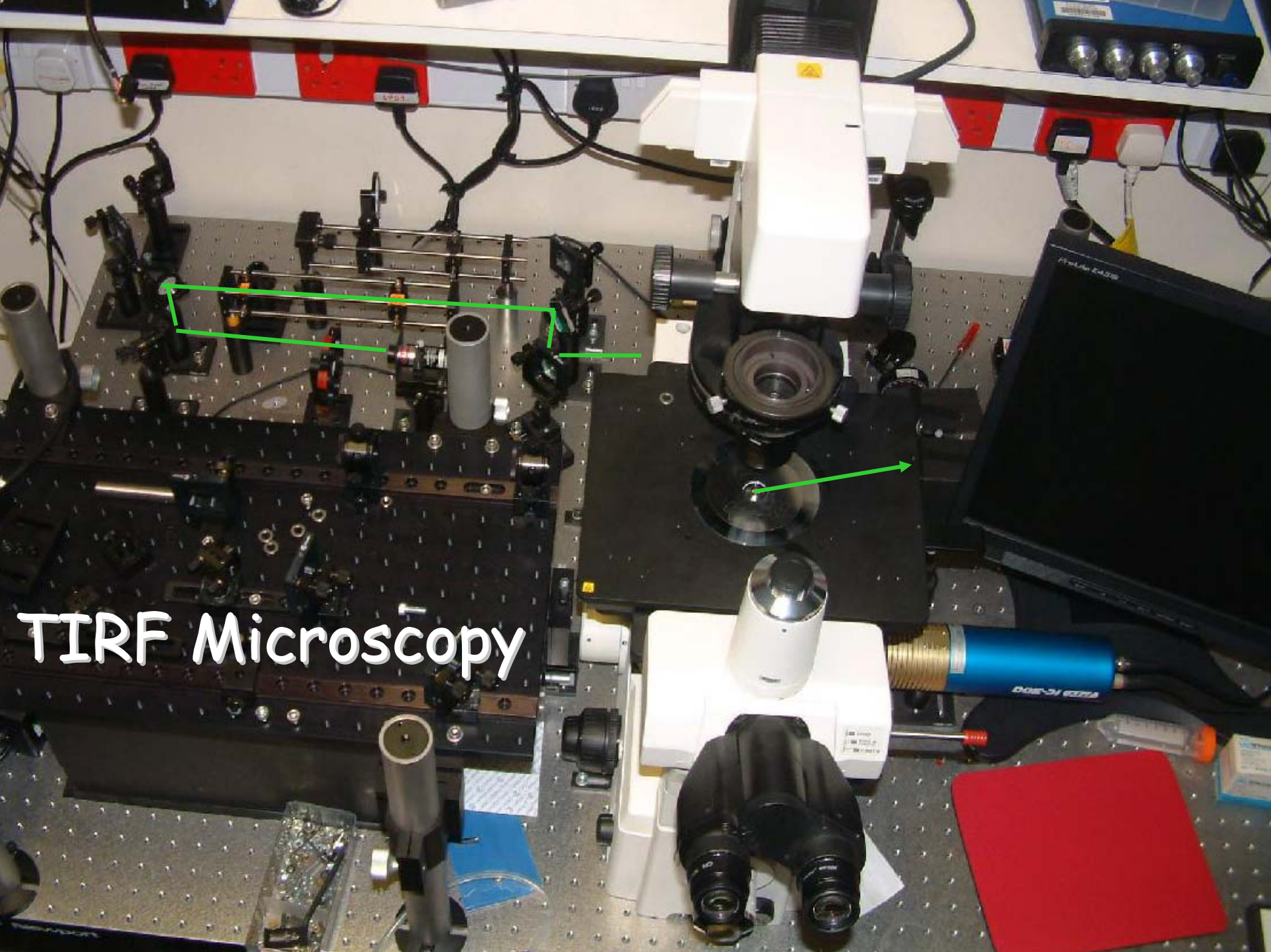
## How is myosin X targeted to its cellular location and what switches it on and off?

- The “head” of the molecule is the motor – the “tail” defines the cargo to be carried (and therefore its cellular function).
- Myosin X is targeted to the lamellipodium, to membrane ruffles and the tips of filopodial actin bundles.

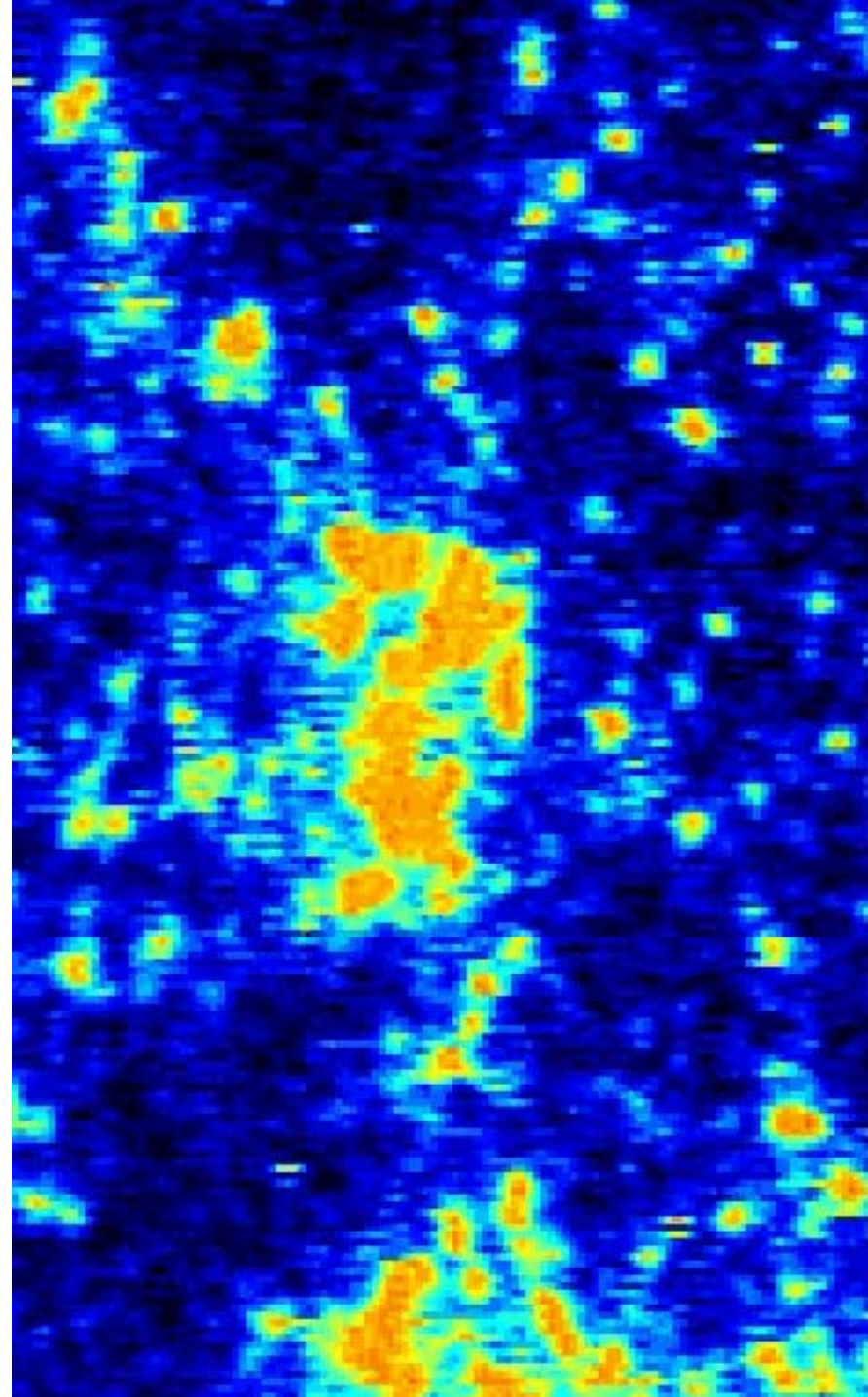
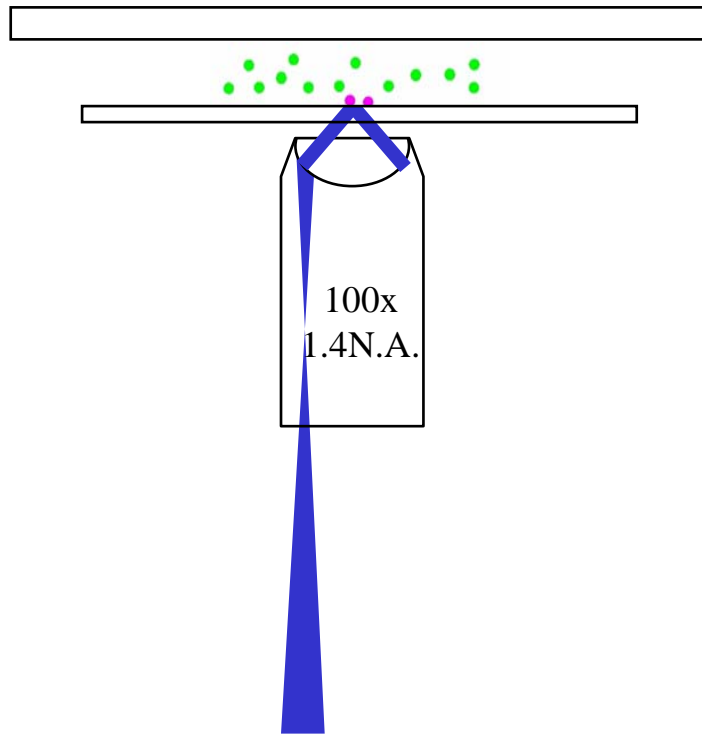


(Berg *et al.*, *J. Cell Science*, **113**:3439-3451)

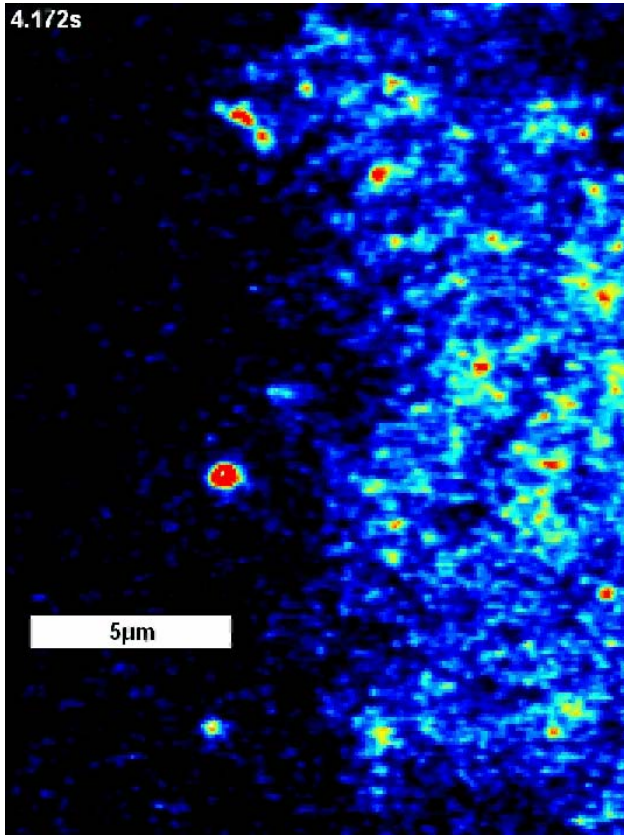
# TIRF Microscopy



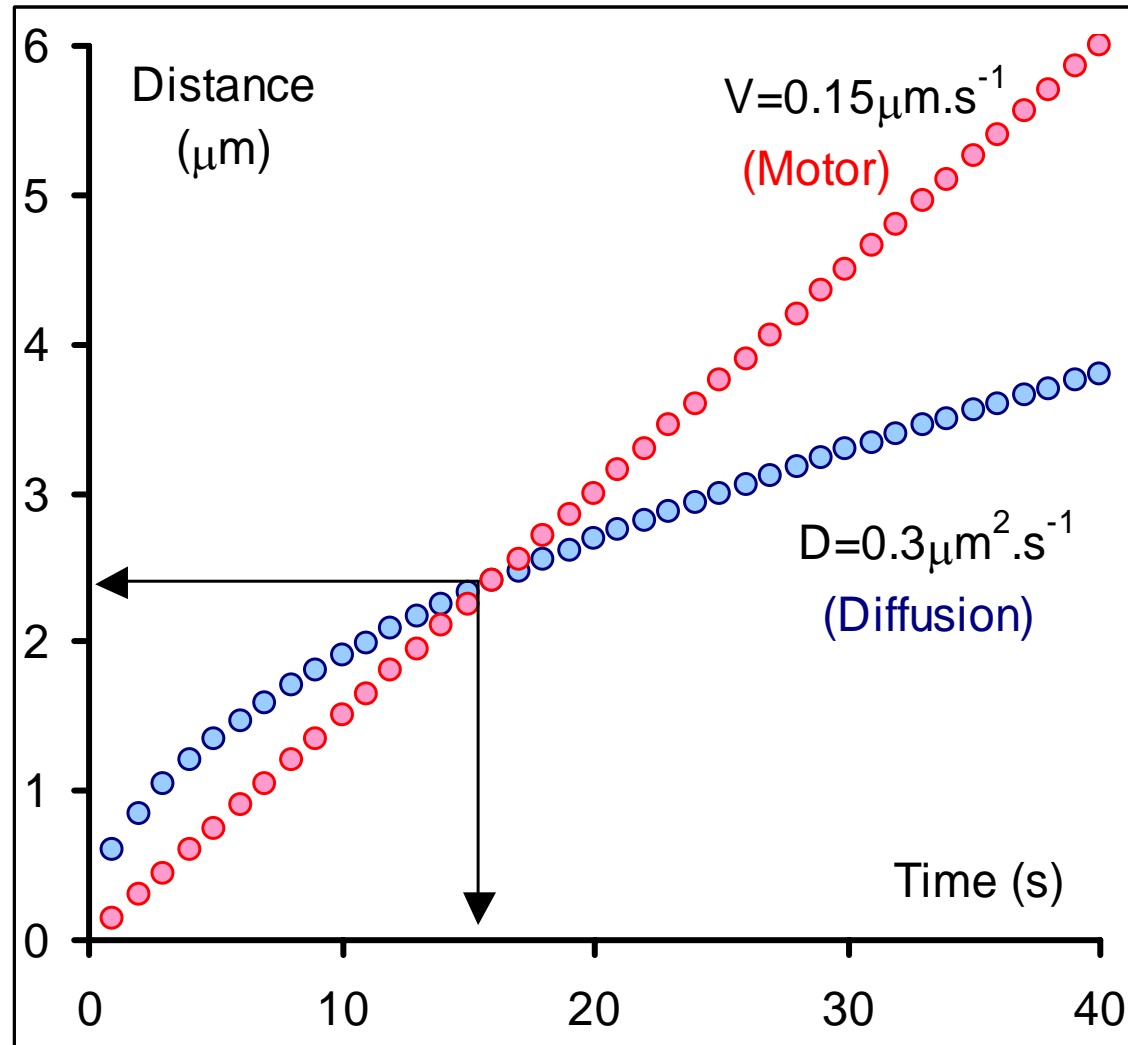
# TIRF



# Motor Properties:



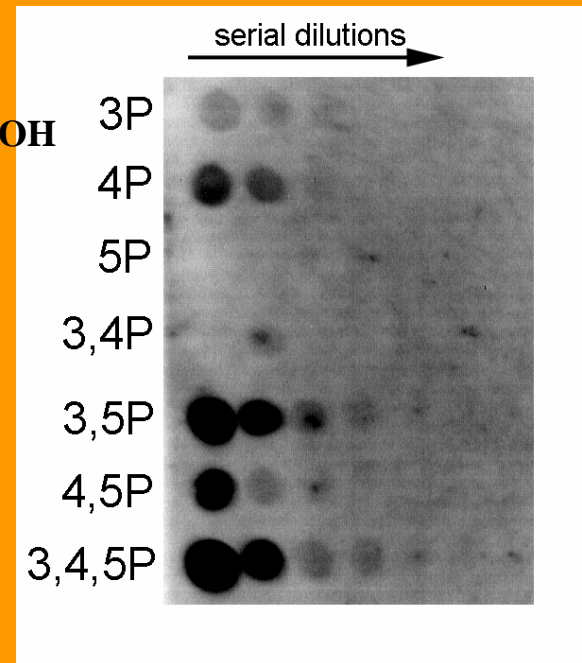
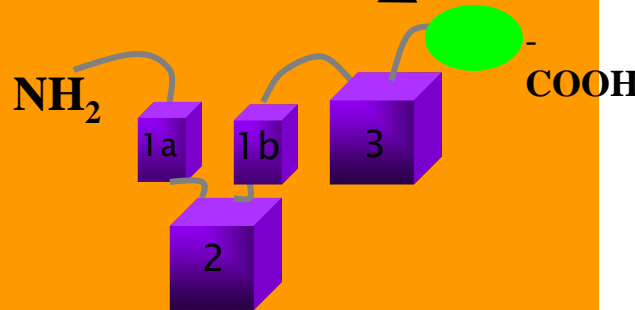
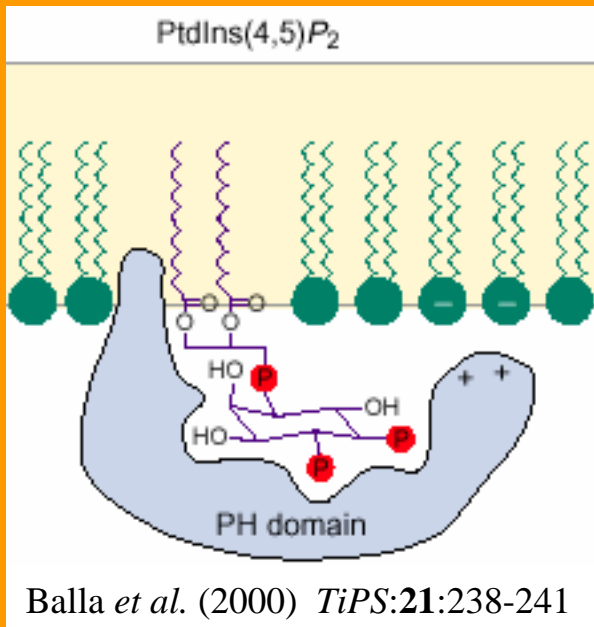
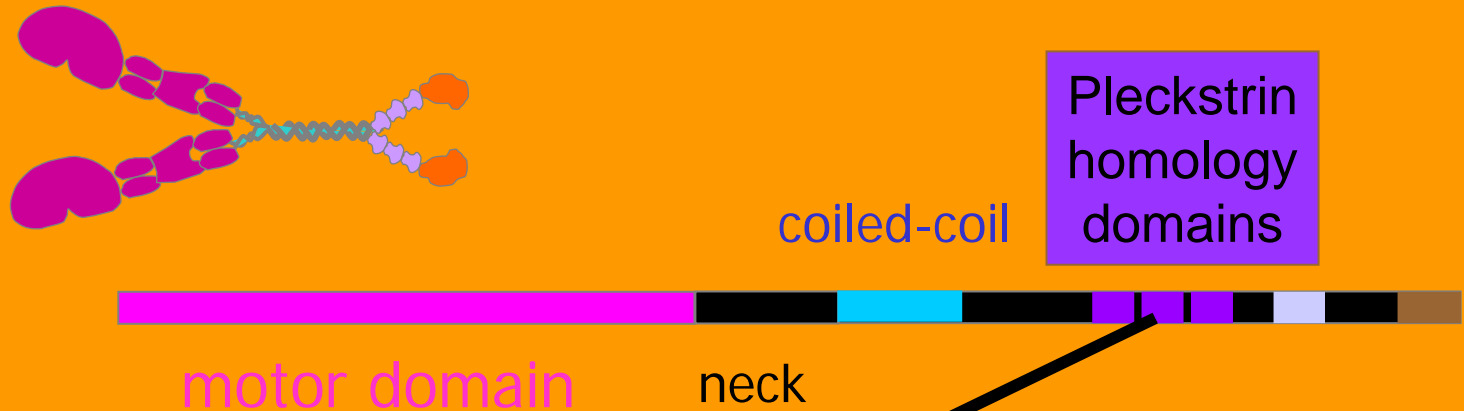
GFP-Myosin X  
expressed in human  
endothelial cells



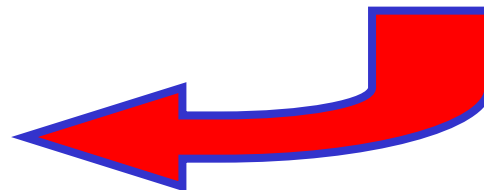
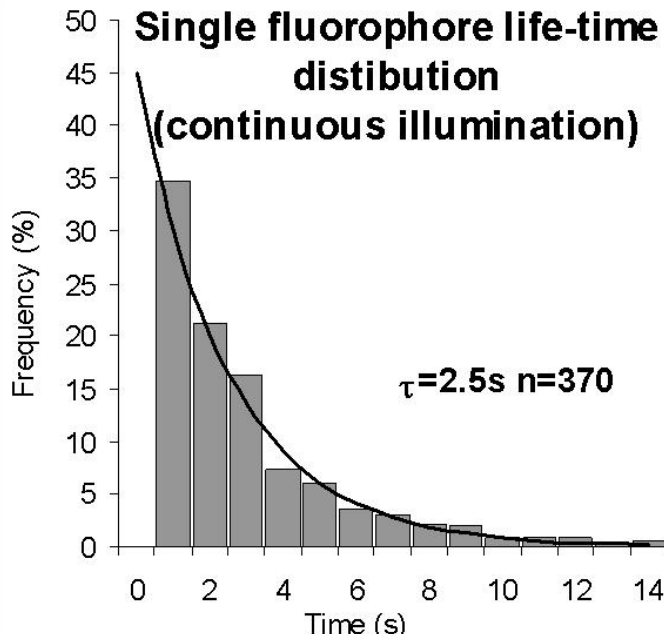
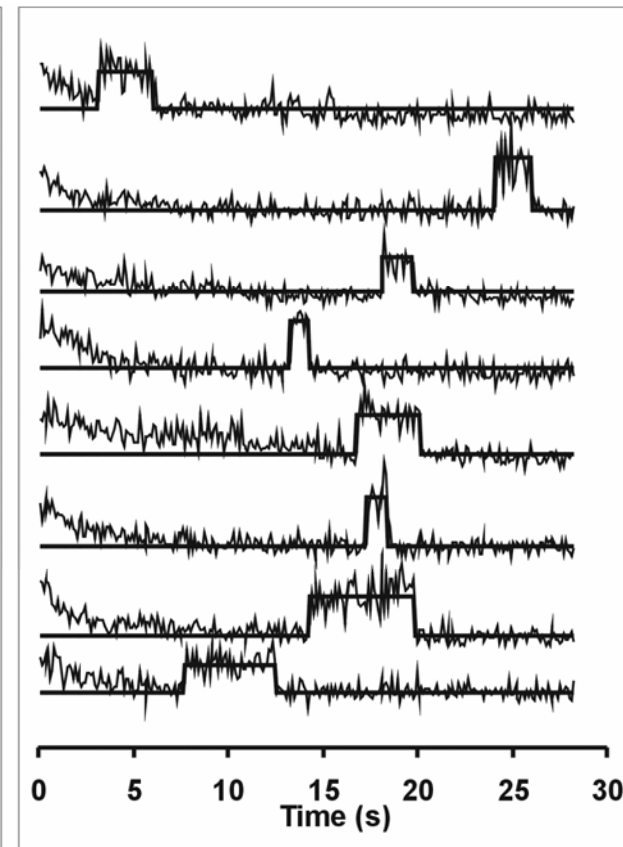
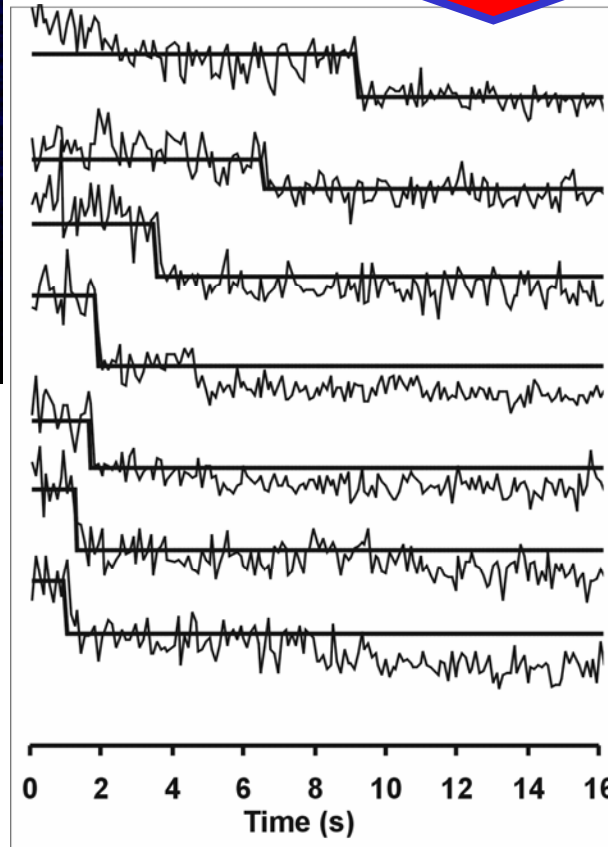
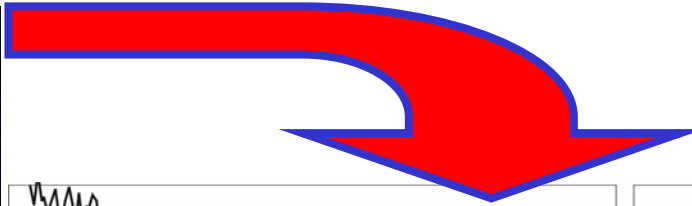
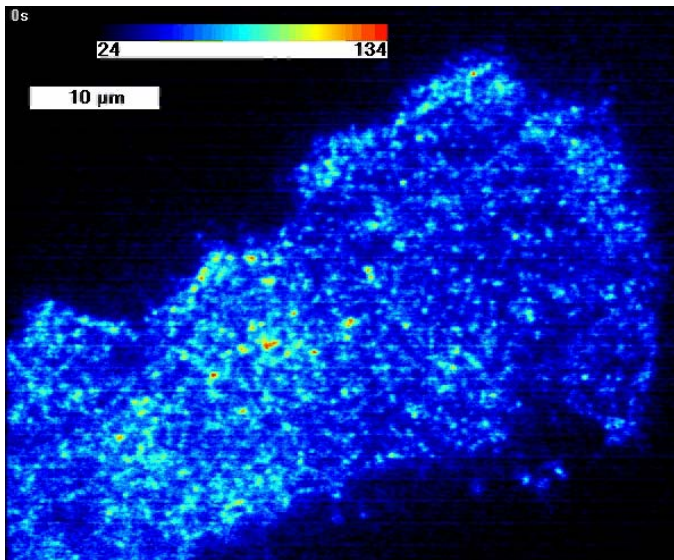
In vitro motility assay sliding velocity =  $125\text{nm s}^{-1}$   
Myosin X coated vesicles move at  $\sim 140\text{nm s}^{-1}$

# Myosin X has several membrane targeting domains in its tail.

## We have studied the PH domains.



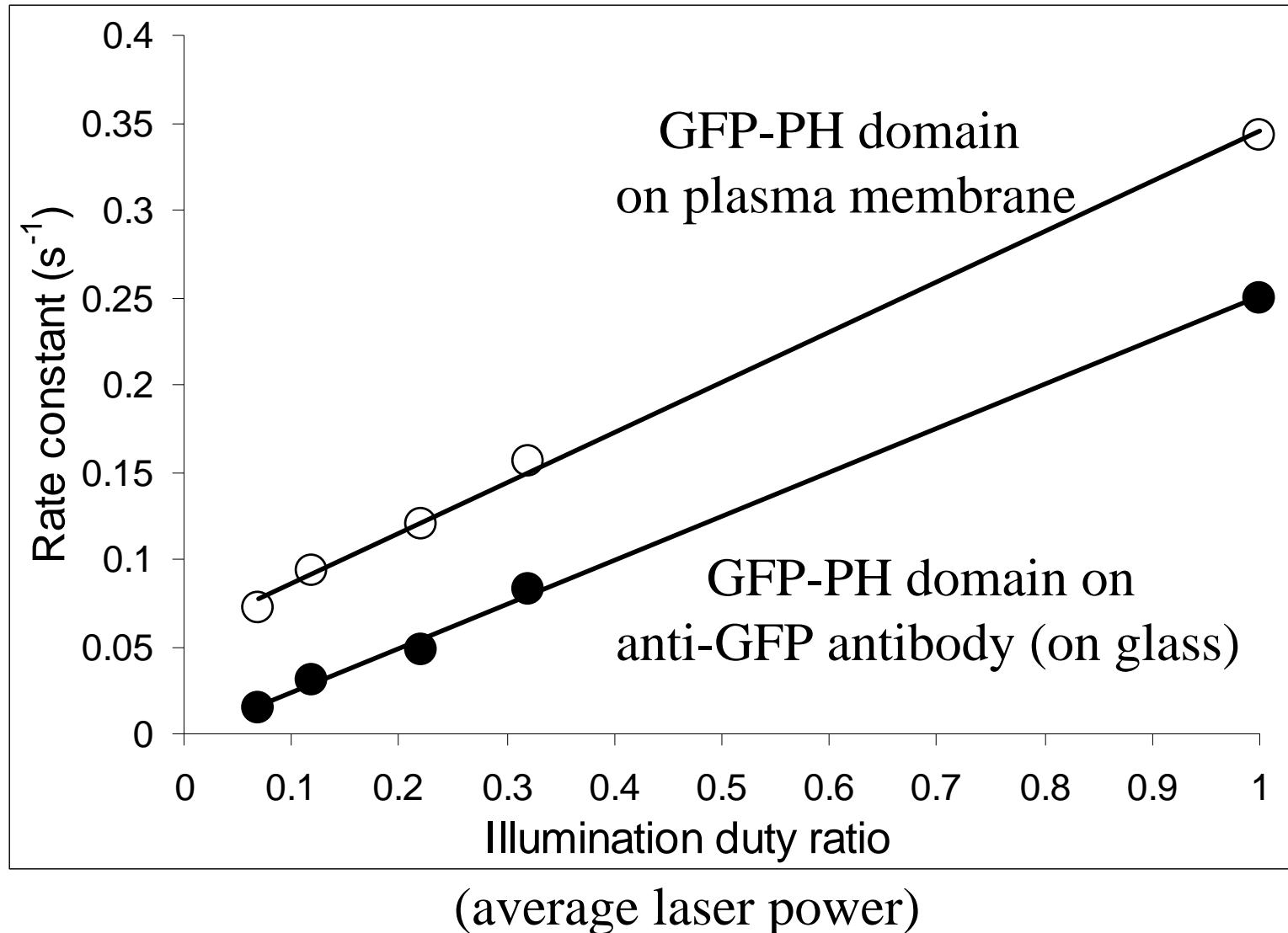
# Single fluorophore temporal trajectories

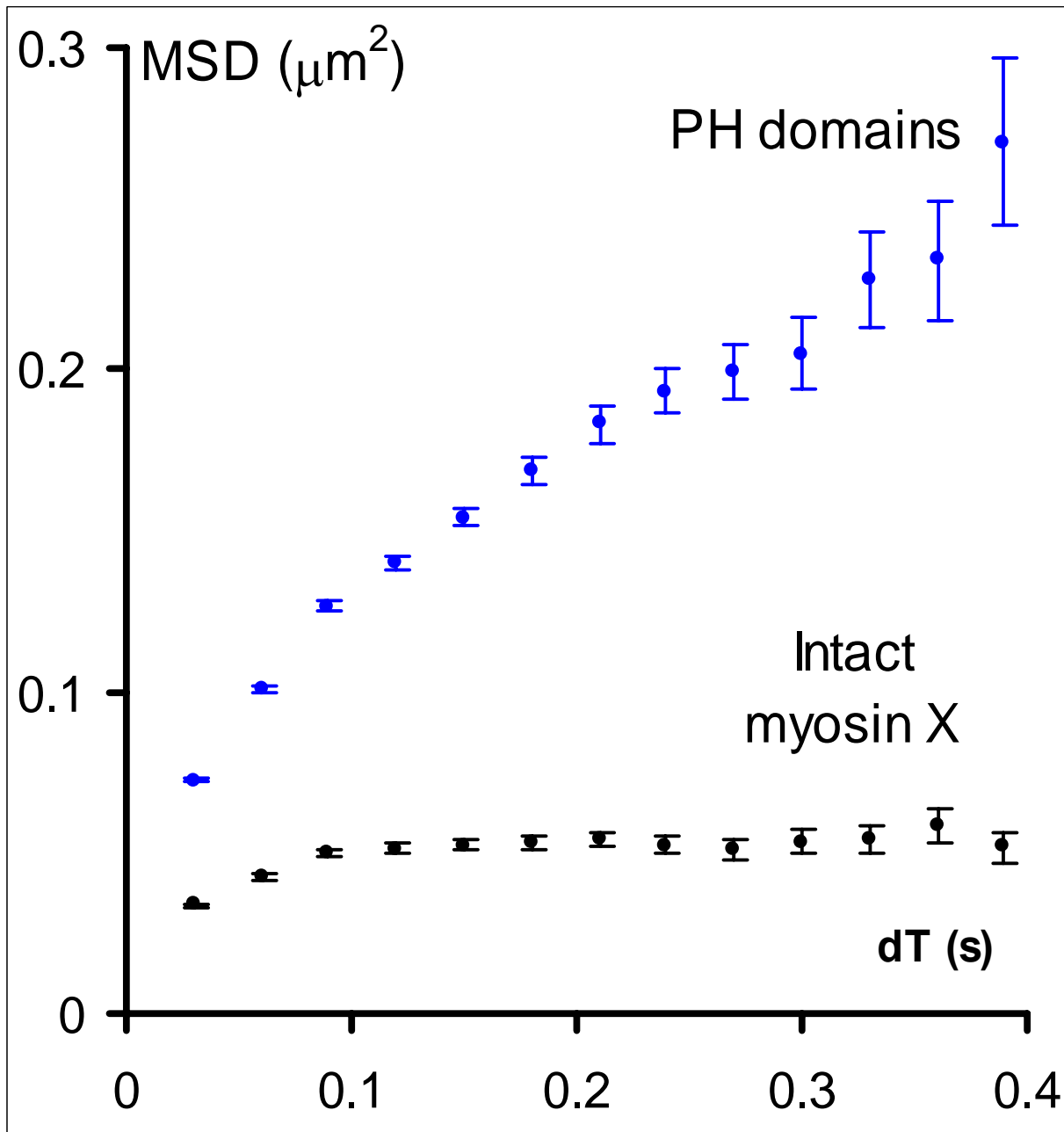


**NOTE:  
BLINKERS NOT  
SHOWN!**

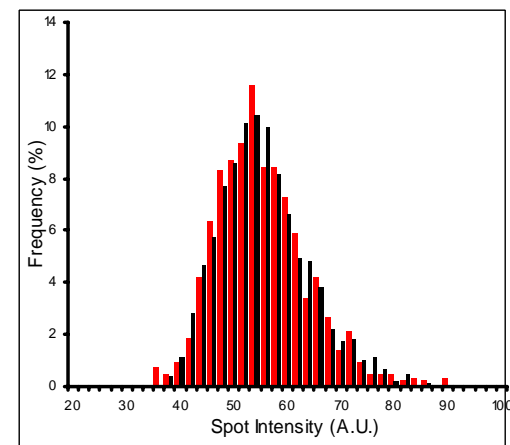
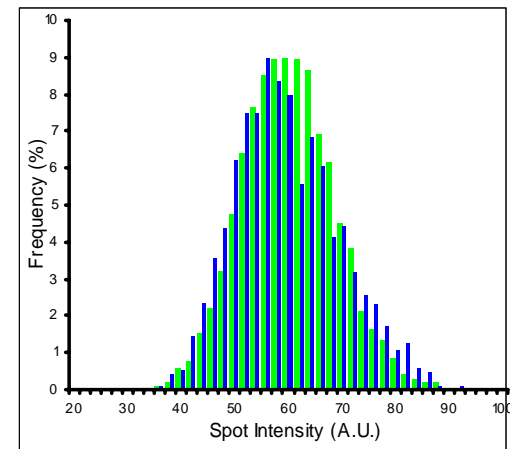


# Calculation of PH123 Dissociation Rate

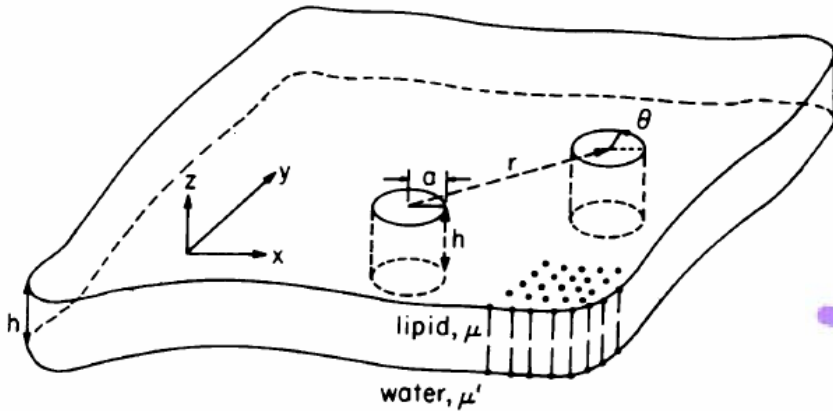




Intensity Dist.

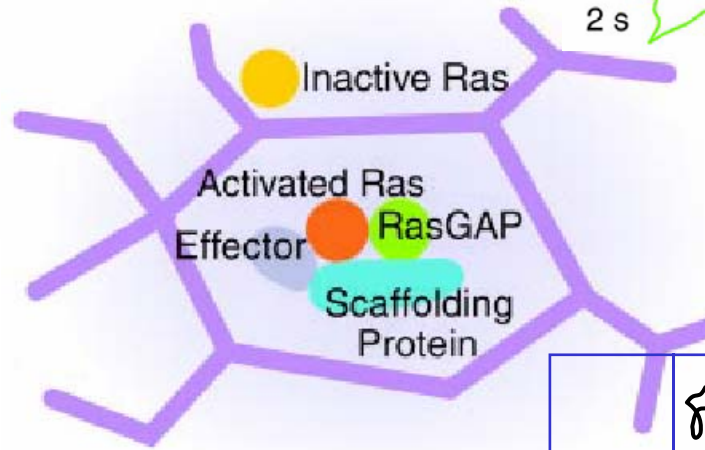
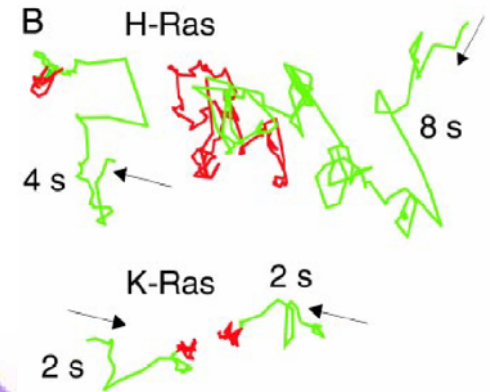
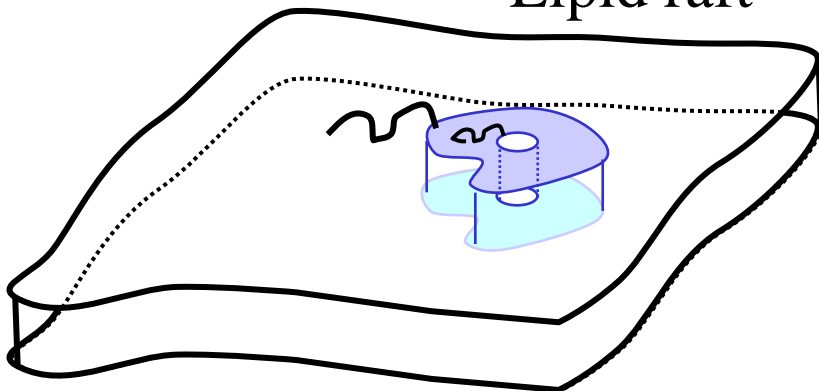


# Diffusion in membrane

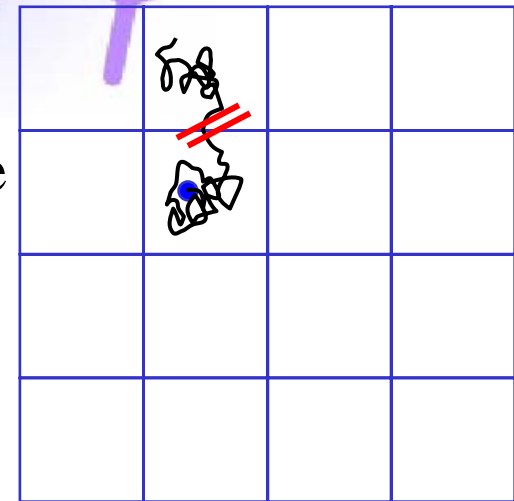


Saffman & Delbruck  
PNAS, 1975

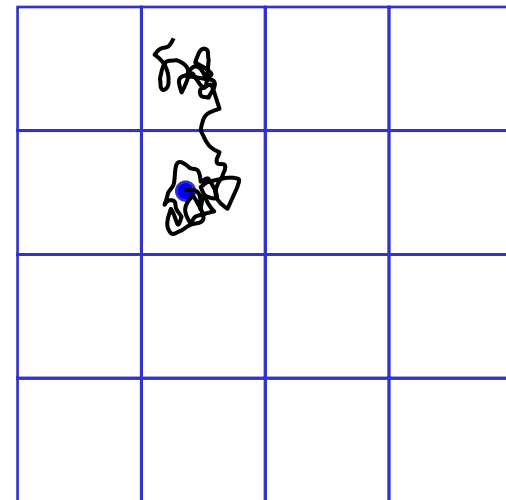
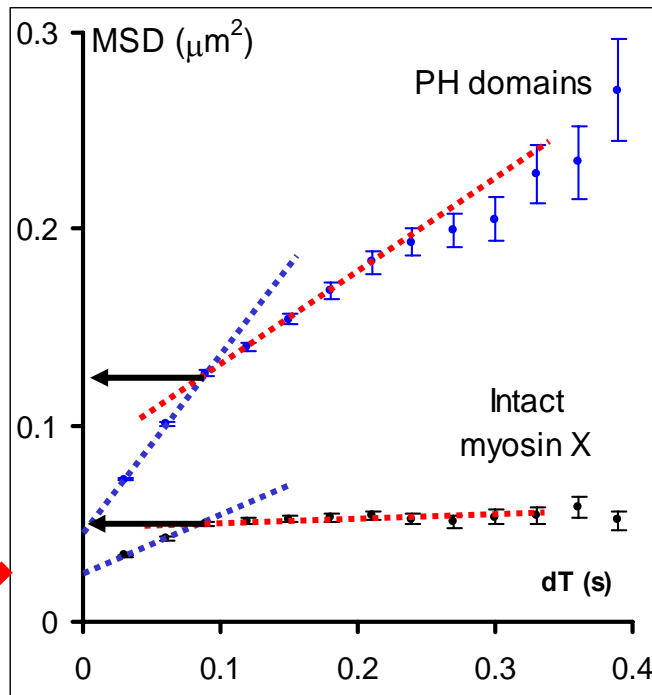
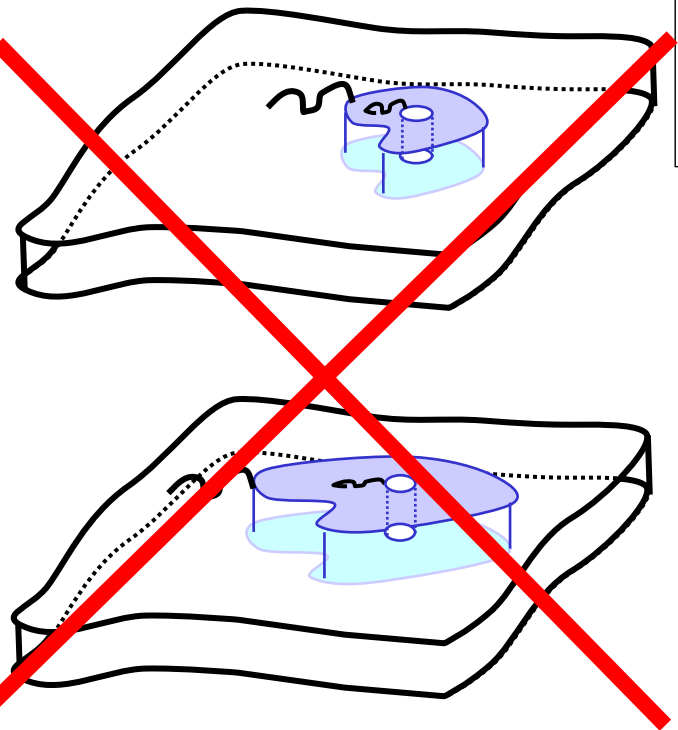
Lipid raft



Membrane Fence  
(energy barrier)

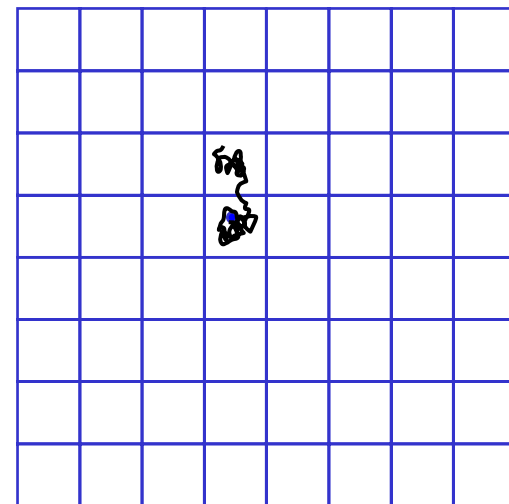


Murakoshi .. Kusumi  
PNAS, 2004



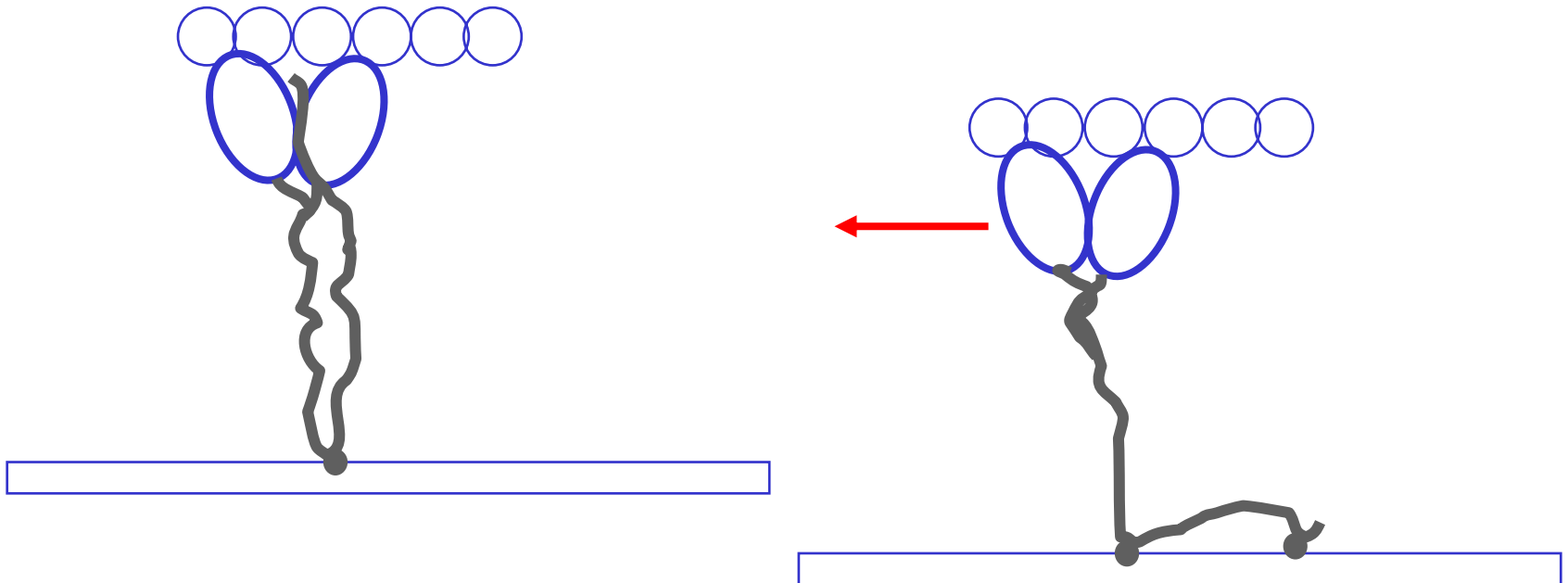
Cell size  $\sim 8\text{nm}$

Cell size  $\sim 15\text{nm}$



# Myosin X - On/Off switch

- Our model is that myosin X is switched off by its tail domain binding to the head when it is inappropriately localised.



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# Optical tweezers - single molecule mechanics

