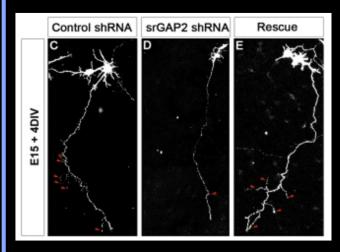
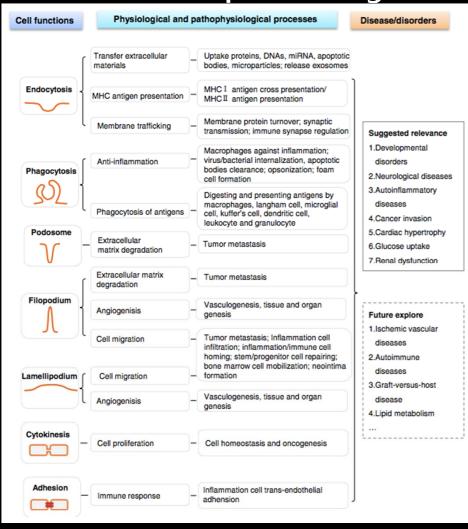
srGAP2 – neuronal migration (epilepsy/schizophrenia)

Supplementary Movie 5
Control EGFP-expressing neurons
Imaged at E15 + 3DIV
1 frame every 16 minutes
Total duration: 8 hours 48 minutes

Supplementary Movie 6 srGAP2-EGFP- expressing neurons Imaged at E15 + 3DIV 1 frame every 16 minutes Total duration: 10 hours 24 minutes



Tightly regulated spatially and temporally



Limitations of *in vivo* techniques

- Unable observe nanoscale membrane dynamics
- Cannot directly observe assembly/activity of membrane-bound remodeling proteins
- No current tools to measure lipid dynamics or the physical characteristics of membrane at sites of deformation

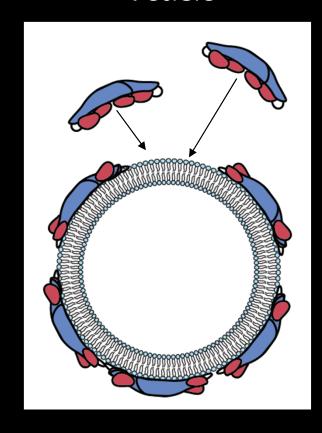
How to directly observe membrane remodeling?

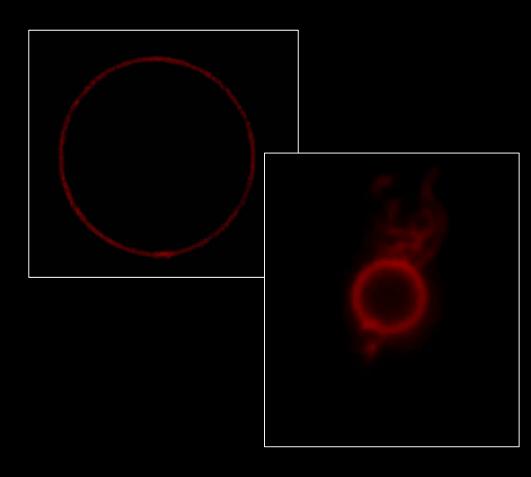
- In vitro simplified system, purified components
 - Manipulate lipid composition
 - Protein density
 - Directly visualize membrane remodeling
 - Evaluate/measure the important physical parameters affecting membrane curvature

Directly observing membrane binding and deformation

Giant Unilamellar Vesicle

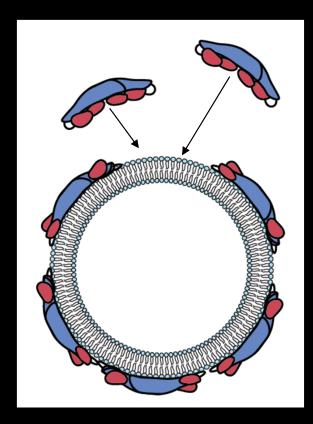
F-BAR - SNAP549





Directly observing membrane binding and deformation

Giant Unilamellar Vesicle F-BAR - SNAP549

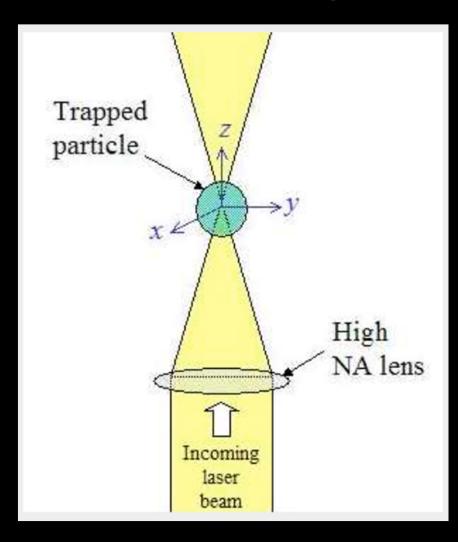


Protein mobility – detecting stable scaffolds

Nwk^{F-BAR}-coated vesicles **Jndeformed** Individually bound -Fast recovery **Deformed** Assembled -

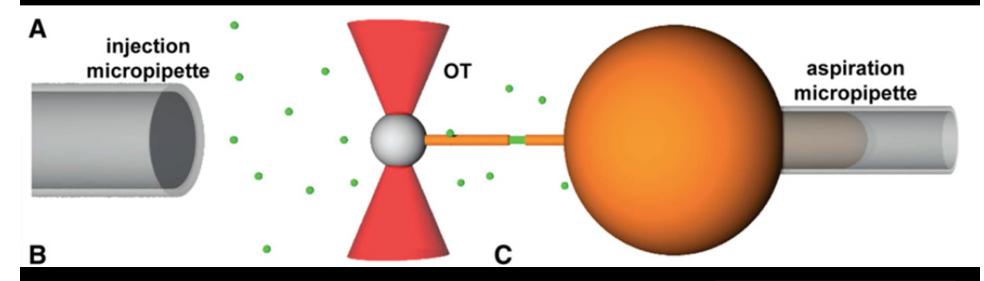
Slow recovery

Optical Trap



- Highly focused laser beam
- Exerts small (pN) forces
- "Beam Waist" has a strong electric gradient
- Objects get stuck in center, where electric force is greatest

Optical Trap + Giant Vesicles



Injector delivers labeled protein locally

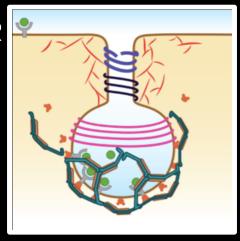
Trapped bead sticks to membrane, forms membrane tubule

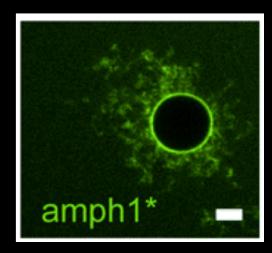
Aspirator sets membrane tension and holds vesicle in place

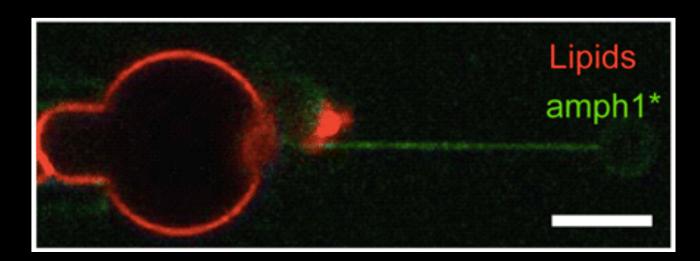
Morlot 2012

Protein activity is dependent on protein density/ membrane shape

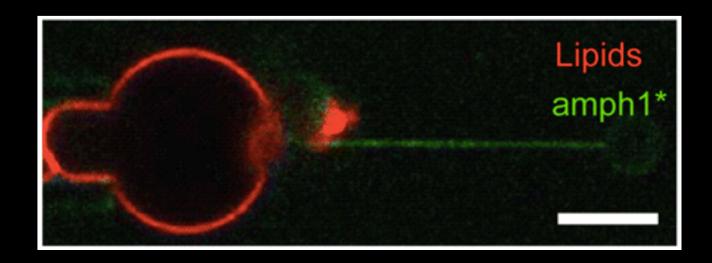
- Mechanical action of BAR domains as function of:
- Amphyphysin
 - Tubulates membranes
 - Important for endocytosis



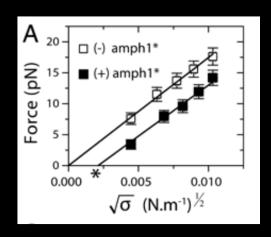


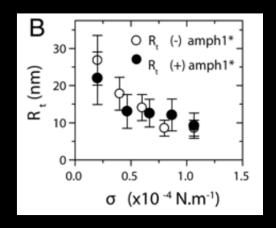


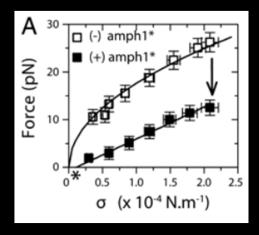
Curvature preference independent of protein concentration

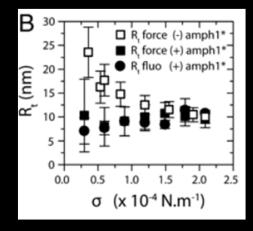


Mechanical impact on membrane depends on protein density

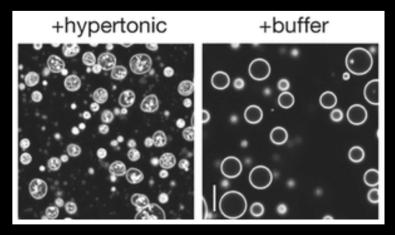


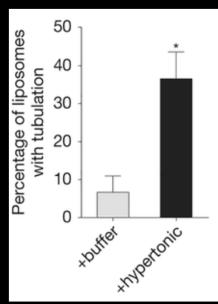


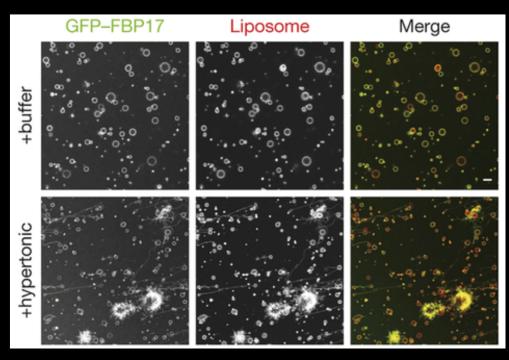




Membrane tension



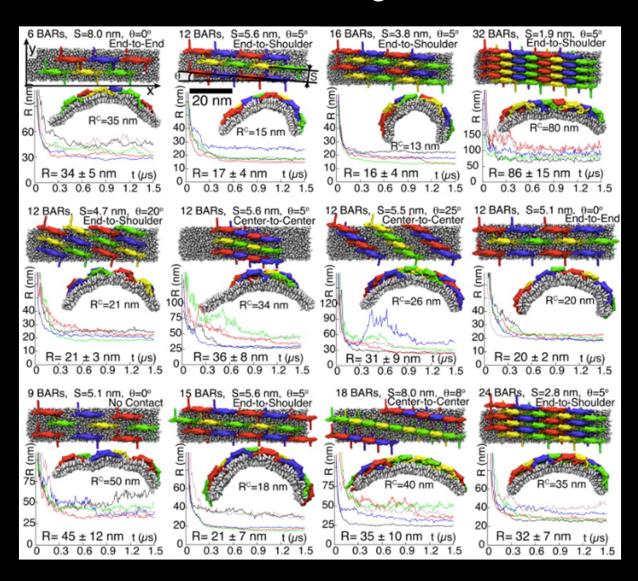




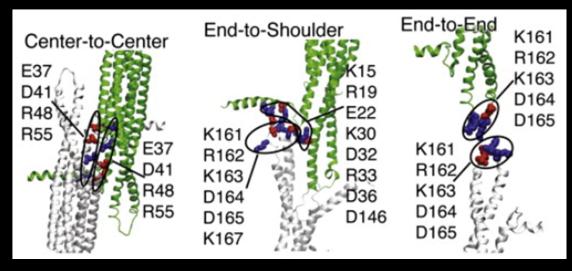
Overall: physical parameters regulate membrane remodeling

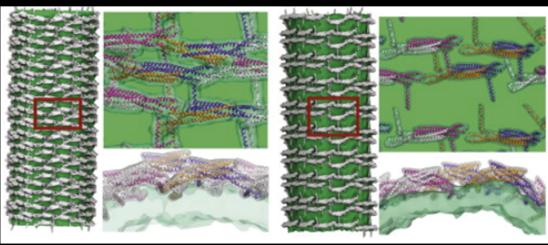
- Protein density
- Membrane tension
- Membrane shape
 - composition

Molecular dynamic simulations



BAR domains can assemble differently dependent on intermolecular interactions





Questions that cannot be answered experimentally (yet)

- What roles do lipid dynamics and composition play?
- How do these physical parameters affect stability and/or assembly of individual BAR domains on the membrane?
- How do all these factors come together to induce curvature in vivo?