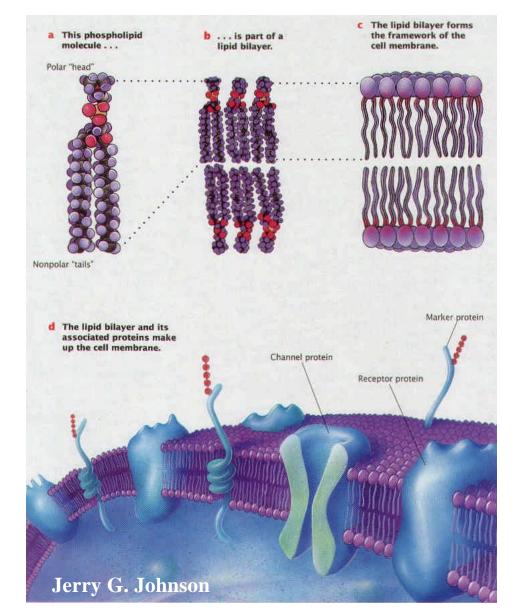
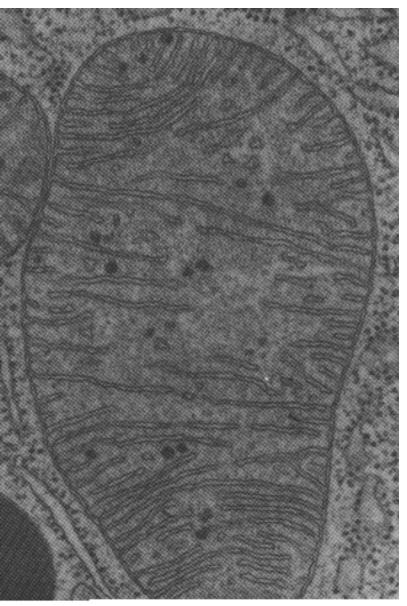


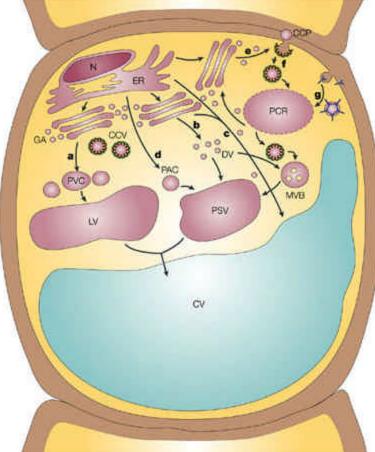
Replication cycle of an influenza virus

Lipid membrane



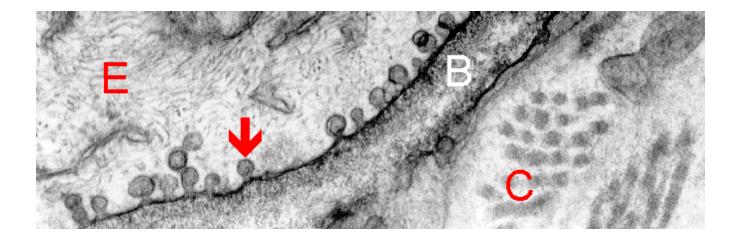


Lipid membrane fusion Basic biological process: cell trafficking



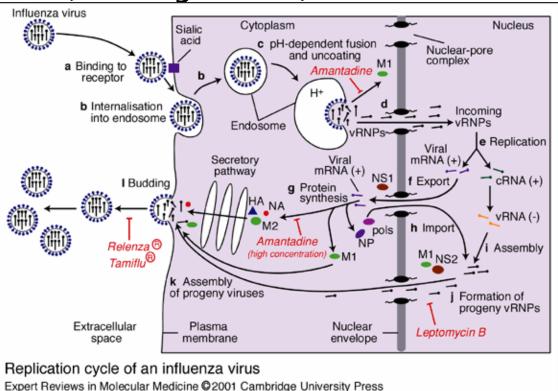
www.cepceb.ucr.edu

Lipid membrane fusion Basic biological process: cell trafficking exocytosis (synaptic release)

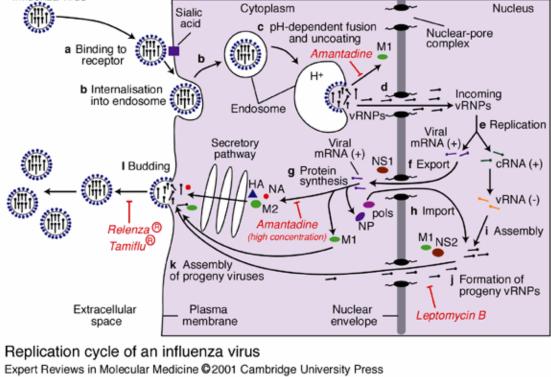


Lipid membrane fusion Basic biological process: cell trafficking synaptic release (exocytosis)

viral entry



Lipid membrane fusion Basic biological process: cell trafficking synaptic release (exocytosis) Influenza virus Cytoplasm Sialic viral entry acid c pH-dependent fusion Nuclear-pore and uncoating complex a Binding to M1 íЩ Amantadine receptor drug delivery b Internalisation

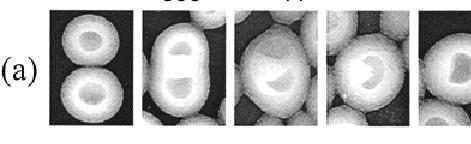


Fusion: experimental facts

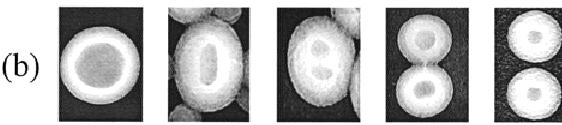
- Short timescales *micro-seconds*
- Small length scales *nano-meters*
- No direct observation of intermediates
- Indirect evidence via dependence on:
 - lipid composition
 - mechanical/electrical stress
 - temperature
- Microscopic mechanism is not well understood
- Universality(?): fusion peptides or other "machinery" are not necessary for fusion of lipid membranes

Not only in lipids! Simple amphiphiles: ^A ^{8-21nm} ³⁻⁵ nm Polymersomes

200nm



PS₃₀₀-PAA₄₄



Luo and Eisenberg, Langmuir (2001)

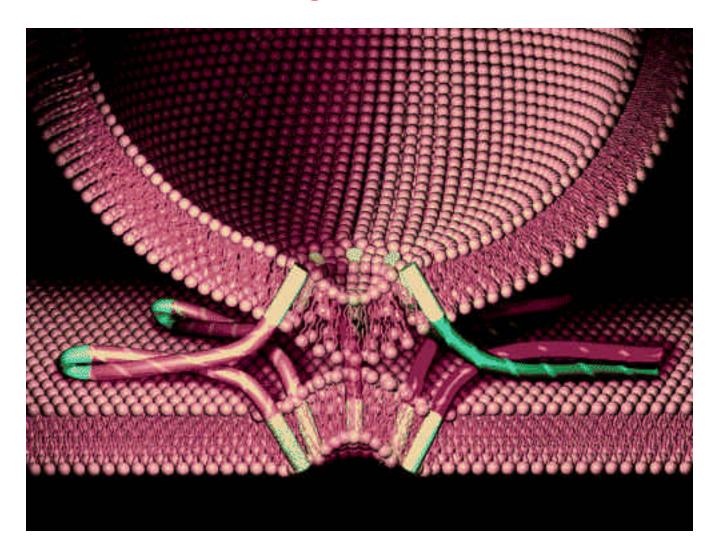
Discher et al., Science (1999)

PBD₅₀-PEO₃₀

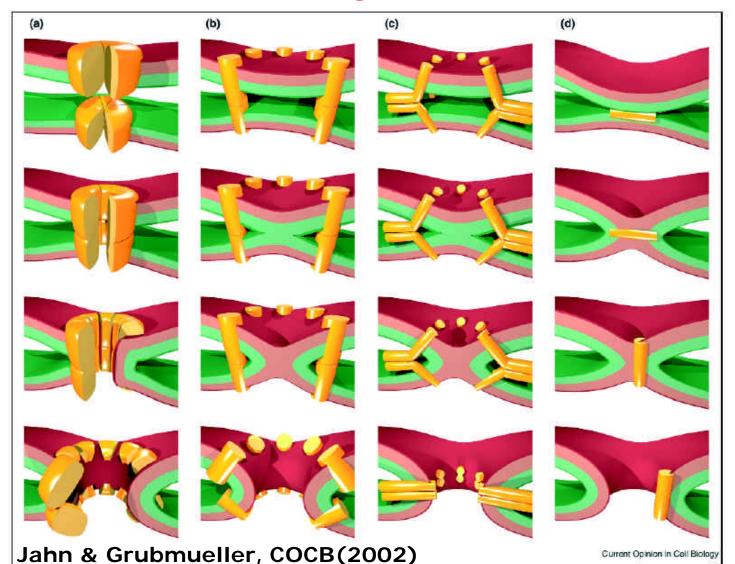
В

20nm

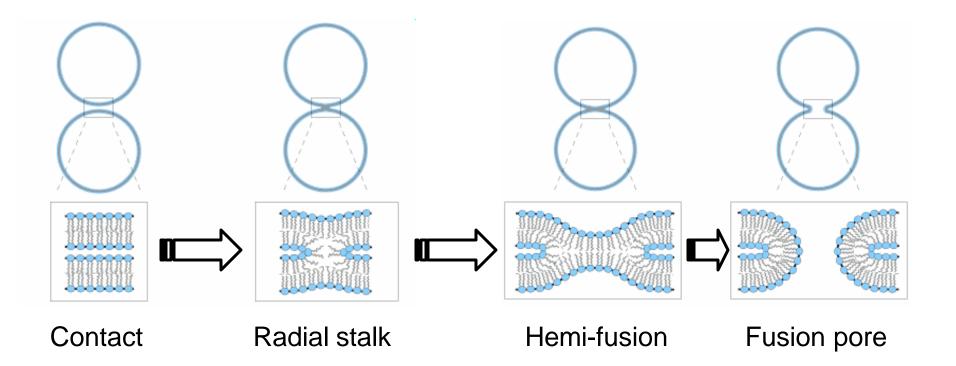
How do things fuse? A biologist's view



How do things fuse? A biologist's view



How do things fuse? A physicist's view



http://www.bnl.gov/bnlweb/pubaf/pr/2002/bnlpr091202.htm

Conceptual difficulty

Membrane stability: long lived holes must be difficult to form.

Membrane fusion: long-lived holes must be easy to form.

Modelling membranes

-ab initio: MD [1nm, 10ns]

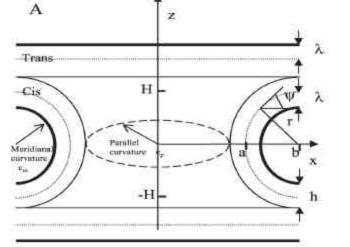
-coarse-grained: MD, MC, (D)DFT, (D)SCFT, DPD

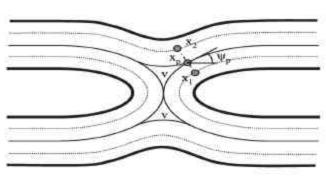
-phenomenological: 2D elasticity theory

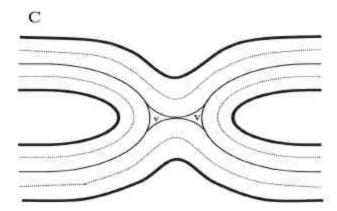
First attempt: Kozlov & Markin (1983) Phenomenological approach

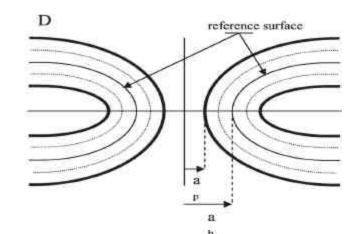
Bending elastic (free)energy + Void energy

B

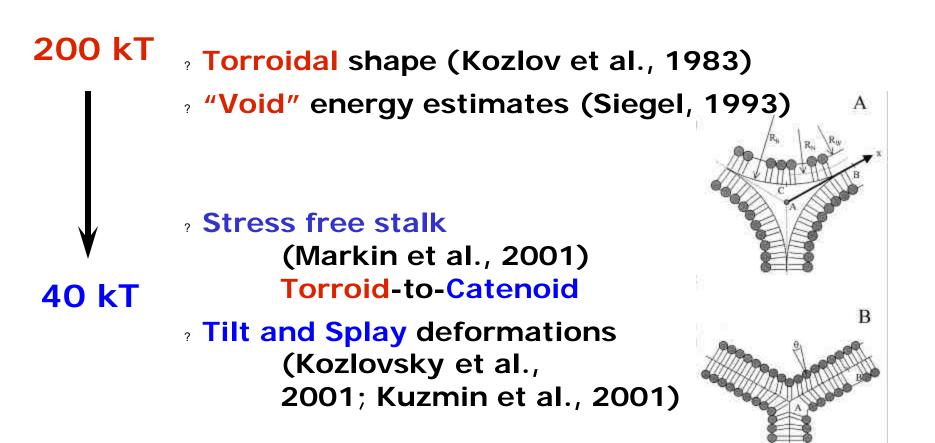








"Energy crisis"



Simulating fusion

System: diblock co-polymer + solvent

Algorithm: Bond Fluctuation Model

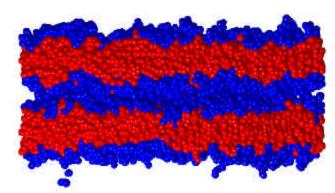
Lattice Monte Carlo

Advantages:

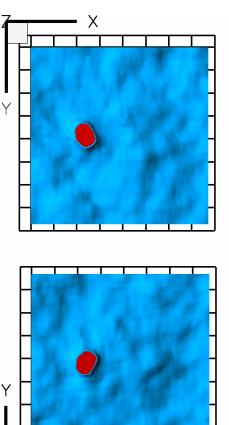
fast (fusion does happen!)

large system size (~250 nm²,2500 "lipids")

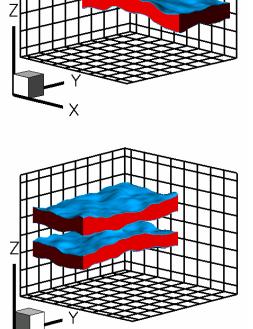
diffusive dynamics (local moves)



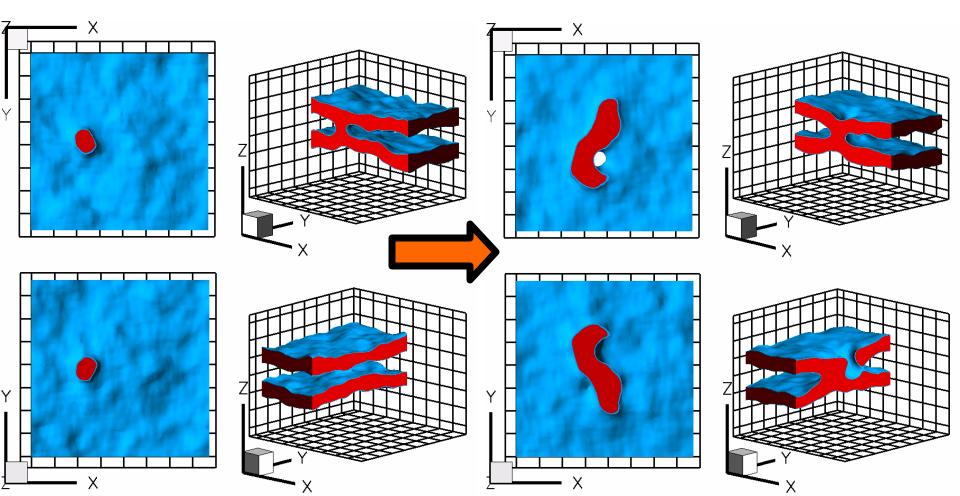
Stalk formation



Х

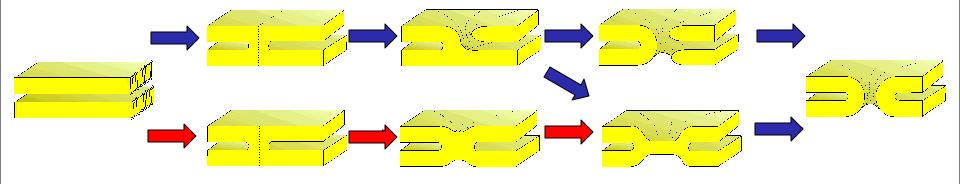


Stalk expansion + hole formation



Fusion Mechanisms

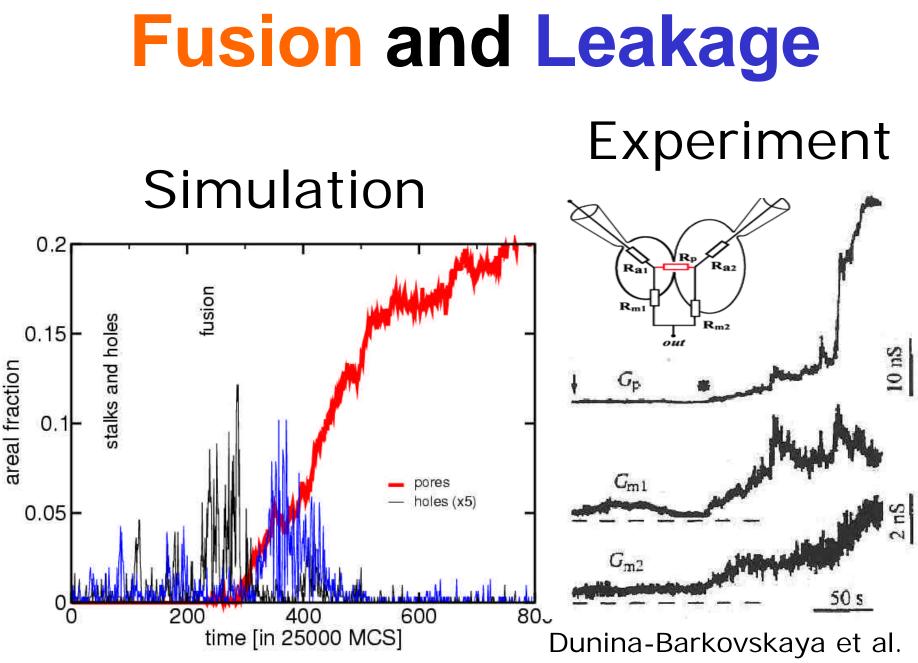
Mechanisms observed in simulation



Hypothesized "stalk" mechanism

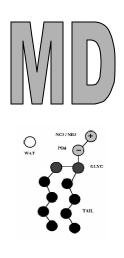
Can we distinguish these mechanisms through the indirect experiments?

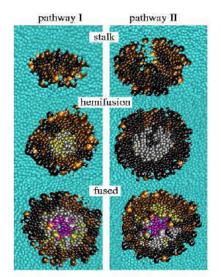
Consequences???



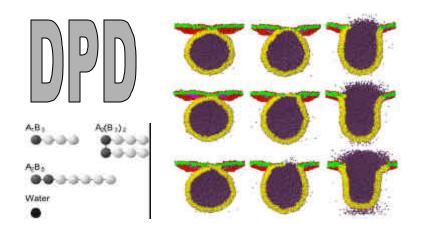
Memb.Cell.Biol. (2000)

Many models – same result...

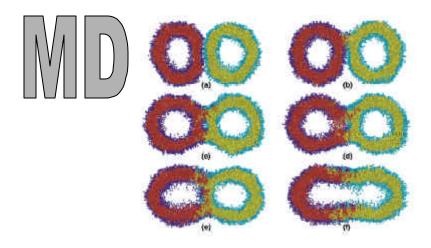




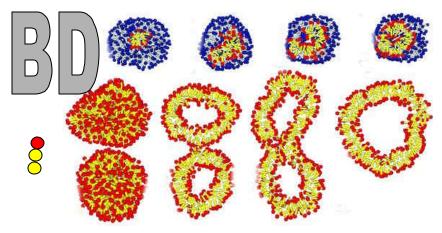
Marrink and Mark, JACS (2003)



Shillcock, Nat.Mat. (2005)



Stevens, Hoh and Wolf, PRL(2003)



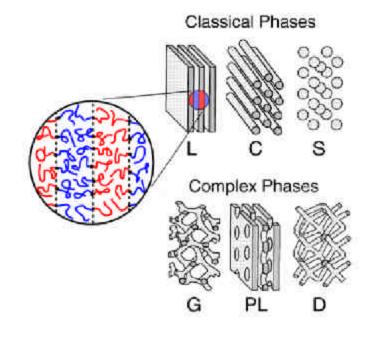
Noguchi and Takasu, JCP(2001)

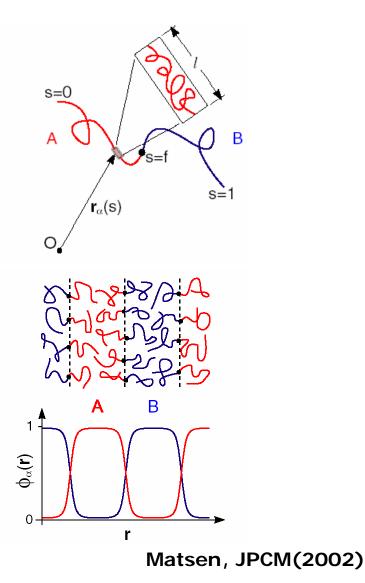
- Why no stalk expansion?
- Why transient holes form? (Isolated bilayer is very stable)

- Need to compare Free Energy of possible intermediates and transition barriers
- Hard to get Free Energy from MC simulations
- The model is simple enough to be treated with microscopic theory (not phenomenological)

Self-Consistent Field Theory

- Homopolymer-diblock blend
- Flory-Huggins interaction
- Incompressible
- Gaussian chain statistics







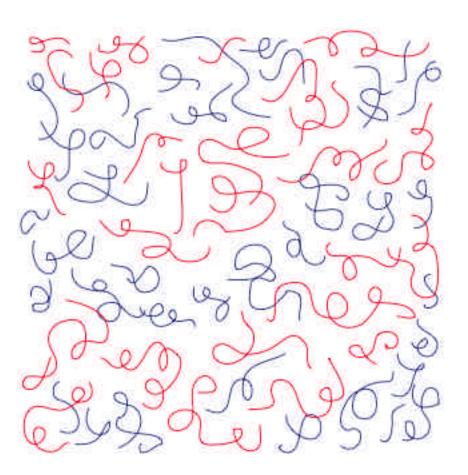
- □ Same microscopic polymer model as in MC
- Proven to be reliable (agrees with MC)
- Provides structural and energetic information
- Intermediates ([meta]stable configurations)
- Transition states (saddle points) can be found
- Reaction coordinate constraint

$$\frac{\mathcal{H}[w_A, w_B, \phi_A, \phi_B, \xi]}{k_B T \Phi} = -Q_s[w_A] - zQ_a[w_A, w_B] \\ + \chi N \int dV \ \phi_A(\mathbf{r})\phi_B(\mathbf{r}) \\ + \int dV \ (\phi_A(\mathbf{r})w_A(\mathbf{r}) + \phi_B(\mathbf{r})w_B(\mathbf{r})) \\ + \int dV \\ \psi(a) = \chi N \overline{\phi}_B(\mathbf{r}) - \mathbf{s} \mathbf{r}) \\ \psi(a) = \chi N \overline{\phi}_A(\mathbf{r}) + \overline{\xi}(\mathbf{r}) \\ 1 = \overline{\phi}_A(\mathbf{r}) + \overline{\phi}_B(\mathbf{r}) \\ \mathbf{r}) = \int_0^1 ds \ q_s(\mathbf{r}, s)q_s(\mathbf{r}, 1 - \mathbf{r}) \\ + z \int_0^z ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)q_c^{\dagger}(\mathbf{r}, s) \\ \psi(\mathbf{r}) = z \int_f^1 ds \ q_c(\mathbf{r}, s)$$

 $\Omega = \mathcal{H}[\bar{w}_A, \bar{w}_B, \bar{\phi}_A, \bar{\phi}_B, \bar{\xi}]$

From Particles to Fields

• Partition function Particle-based $Z = \int D[\mathbf{R}] e^{-U_0 - U_1}$ atonovich Hubbard-Trans nation Field-based $Z = \int D[W] e^{-H[W]}$



If you hold a cat by the tail you learn things you cannot learn any other way. --Mark Twain Reaction coordinate constraint

Quantum chemistry

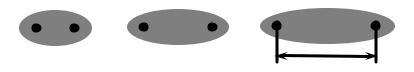
• Born-Oppenheimer approx.

(fast electrons/slow nuclei)

Electronic wave function



- Electronic ground state
- Inter-nuclear distance

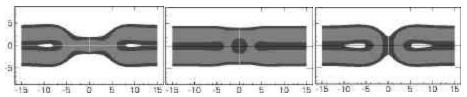


Membrane fusion

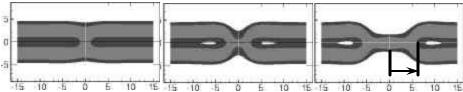
• Quasi-static approx.

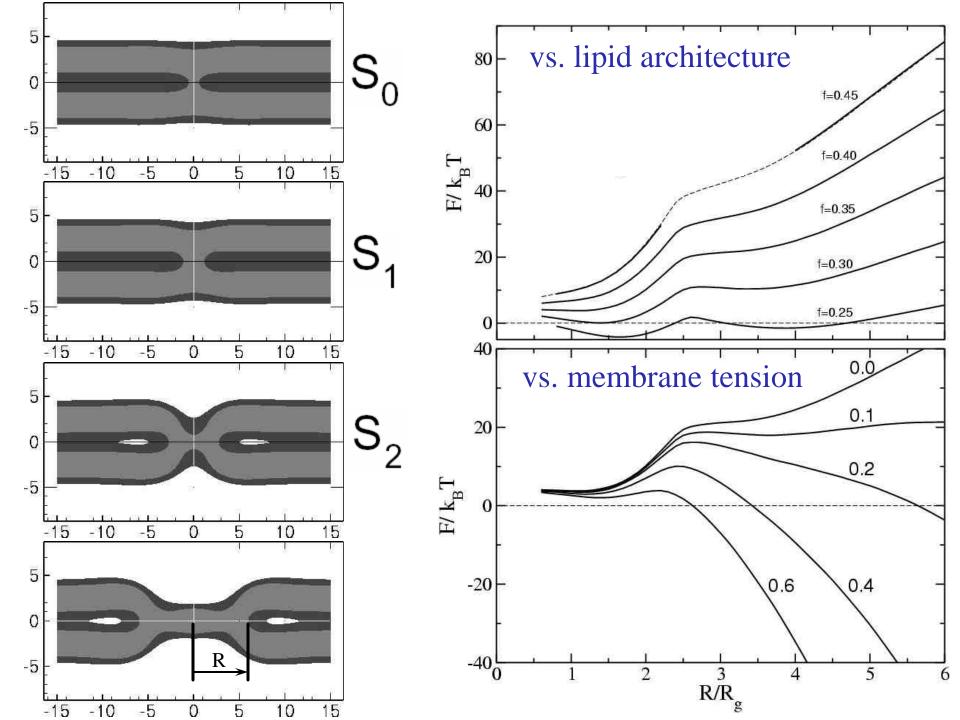
(fast individual/slow collective)

• Interface topology

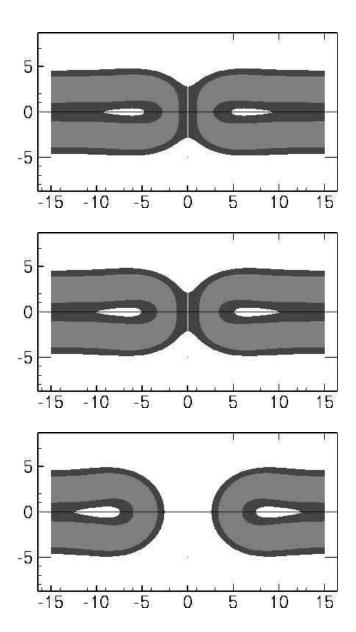


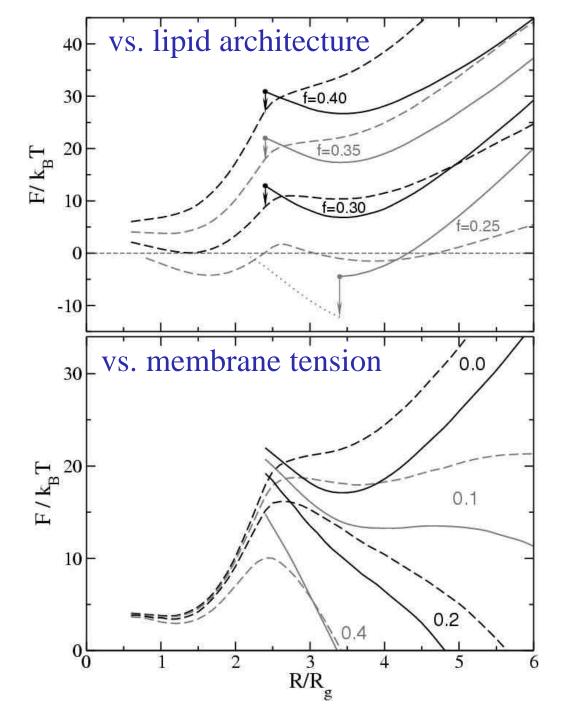
- Lowest free-energy state
- Structure size



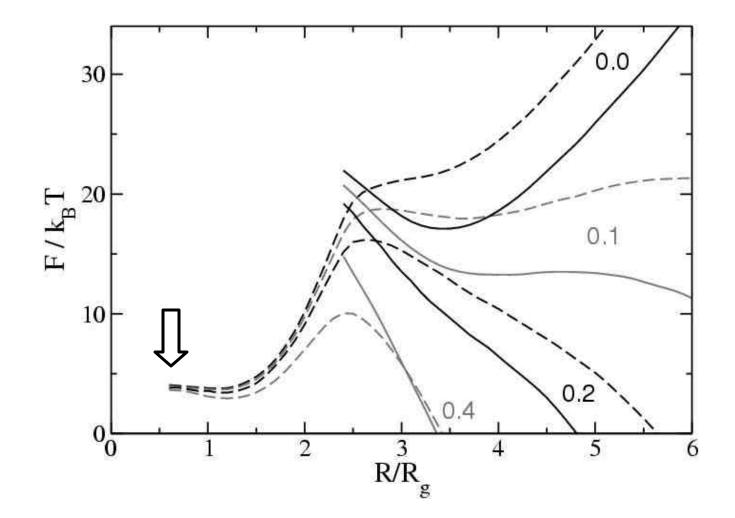


Fusion pore



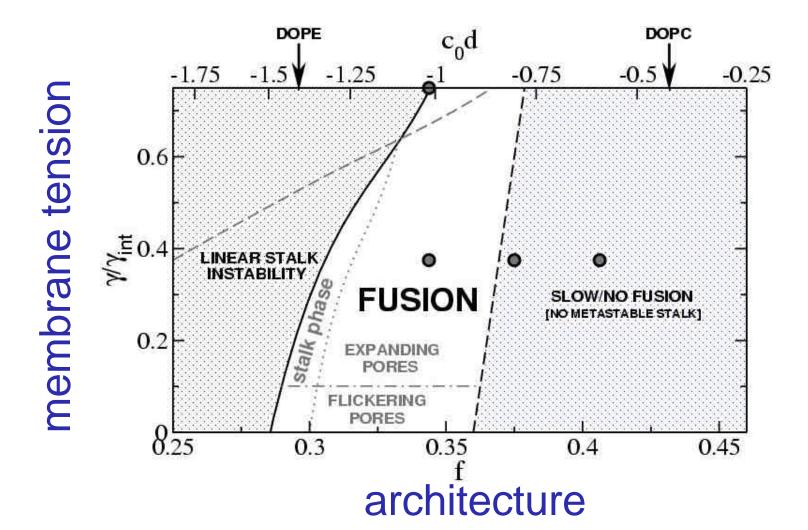


Consequences Major barrier: radial stalk expansion

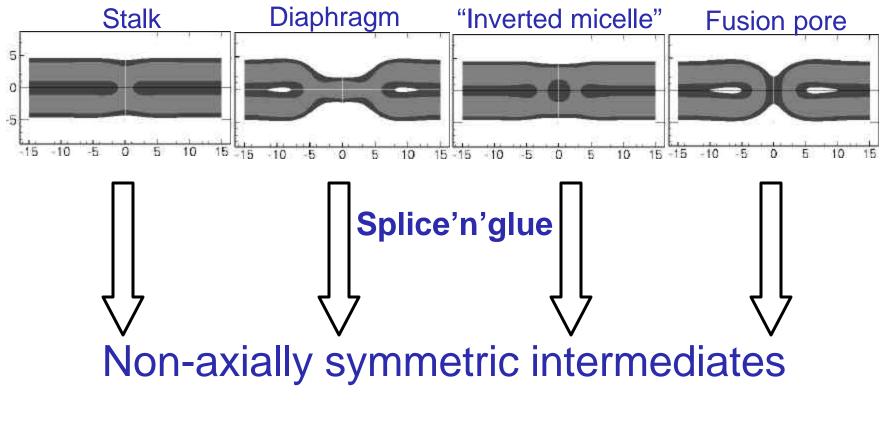


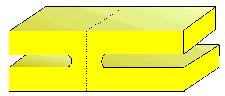
Consequences

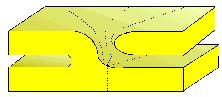
Major barrier is stalk expansion Fusion is limited (regulated)

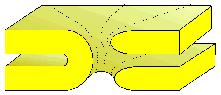


The alternative mechanism

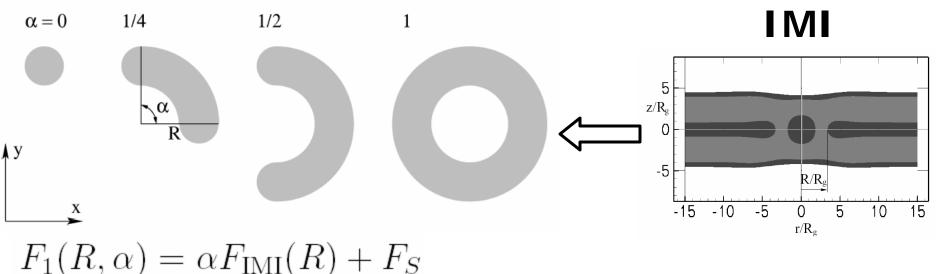




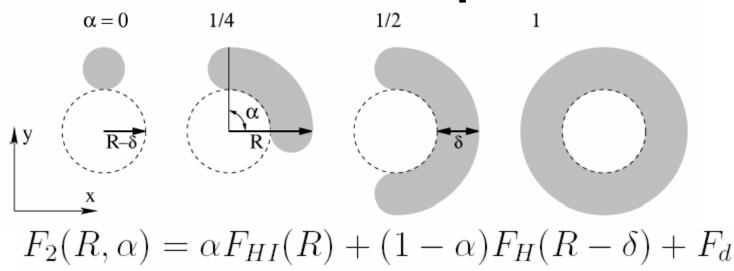




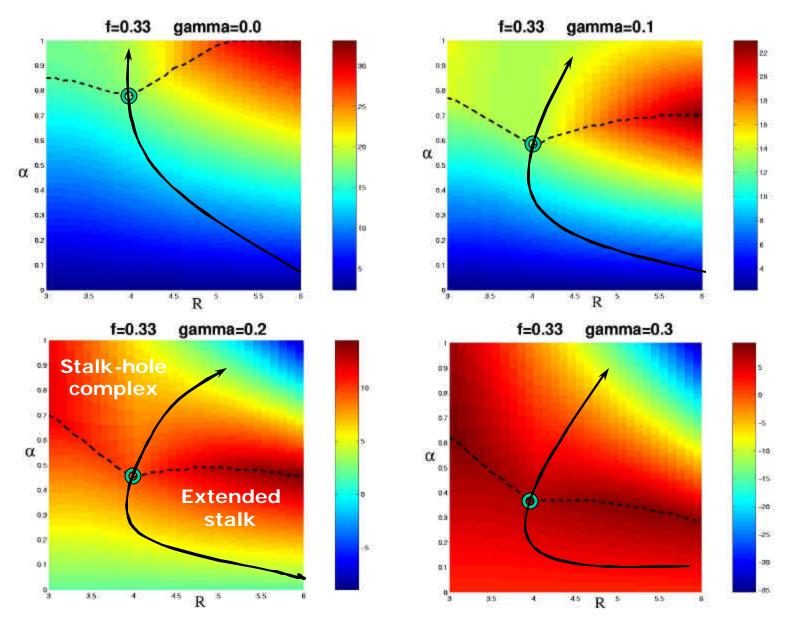
1: Extended stalk



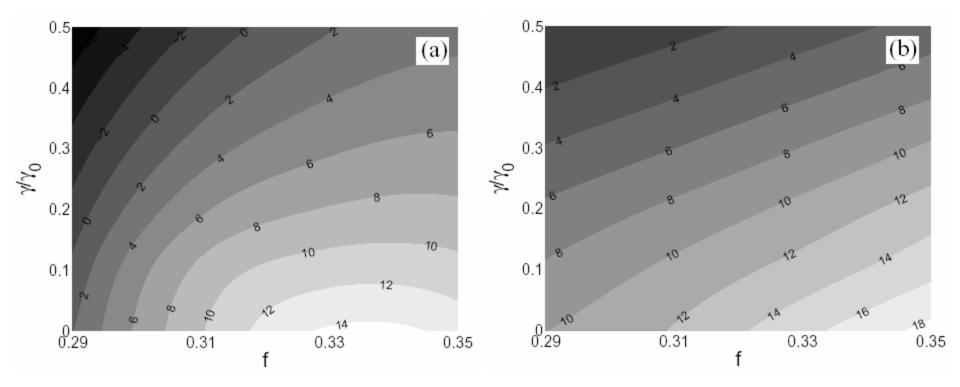
2: Stalk+hole complex



Free energy "landscape"



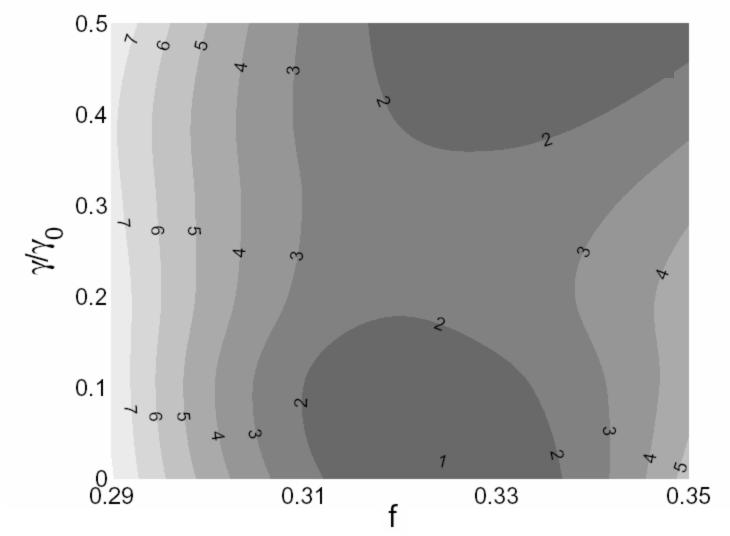
Free energy barrier



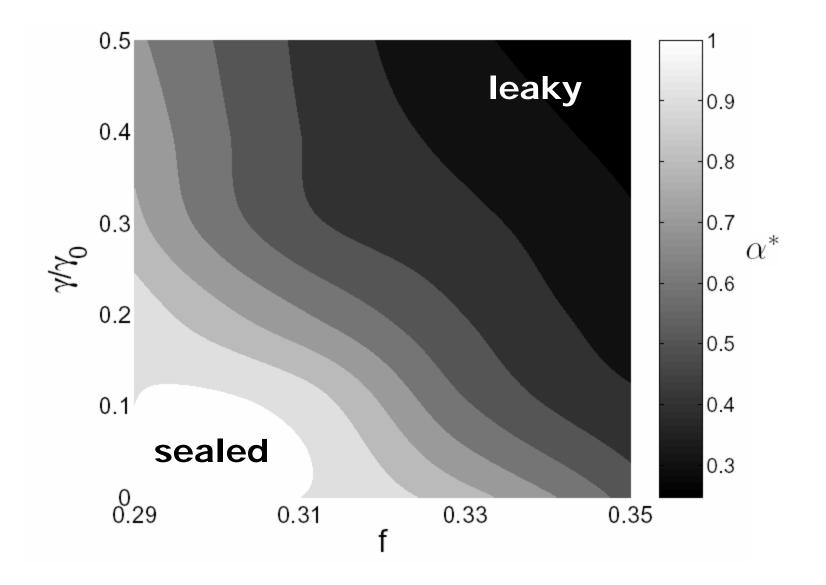
New mechanism

Old mechanism

Free energy barrier difference between old and new mechanisms



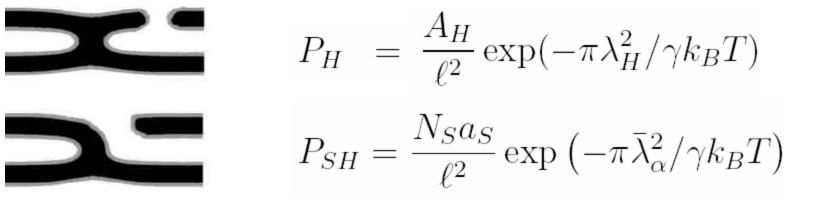
Leakiness of Transition State

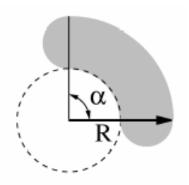


Resolving stability vs. fusion paradox or bare vs. screened holes

 $F_H = 2\pi\lambda_H R - \pi\gamma R^2$

Probability to nucleate supercritical hole





Effective line tension

 $\lambda_H \to \bar{\lambda}_\alpha \equiv \alpha \lambda_{SH} + (1 - \alpha) \lambda_H$

Resolving stability vs. fusion paradox or bare vs. dressed holes

Under simulation conditions Under experimental conditions



 $P_{SH}/P_H \sim 14$

Resolving stability vs. fusion paradox or bare vs. dressed holes

Under simulation conditions Under experimental conditions



 $P_{SH}/P_H \sim 14$



Membrane can both be stable and undergo fusion

What is really going on in biological fusion?

- -Multi component membranes [lipid mixtures, cholesterol, asymmetry]
- -External constraints [vesicle size, volume]
- -Role of fusion peptides -More experiments!

Thanks to:

L.Chernomordik (NIH) F.Cohen (Rush MC) M.Kozlov (Tel Aviv U) B.Lentz (UNC) D.Siegel J.Zimmerberg (NIH)

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> NSF: DMR 987864 and 0140500 DFG: Bi 314/17 Computer time: NIC Jülich HLR Stuttgart and Mainz Office/Home desktop