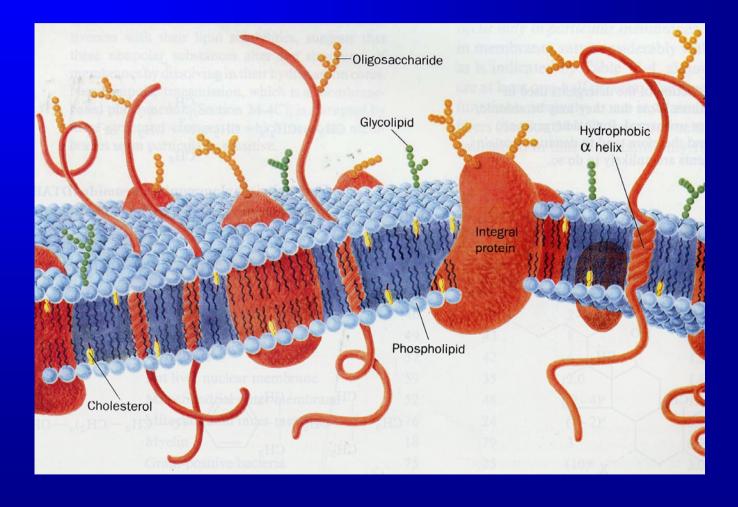
Lipidne mikrodomene

struktura

1

Singer-Nicholson fluid mosaic model of a biological membrane organization (1972)

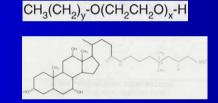


Only part of biological membranes is <u>solubilized</u> after treatment at low T ($\leq 4^{\circ}C$) with:

Triton X-100 (NP-40)

Brij-58

CHAPS

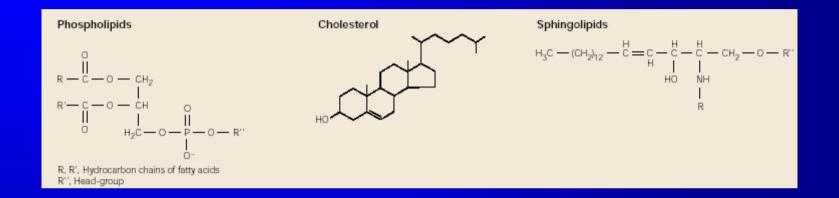


The remaining membranes are <u>soluble</u> in:

octyl glucoside

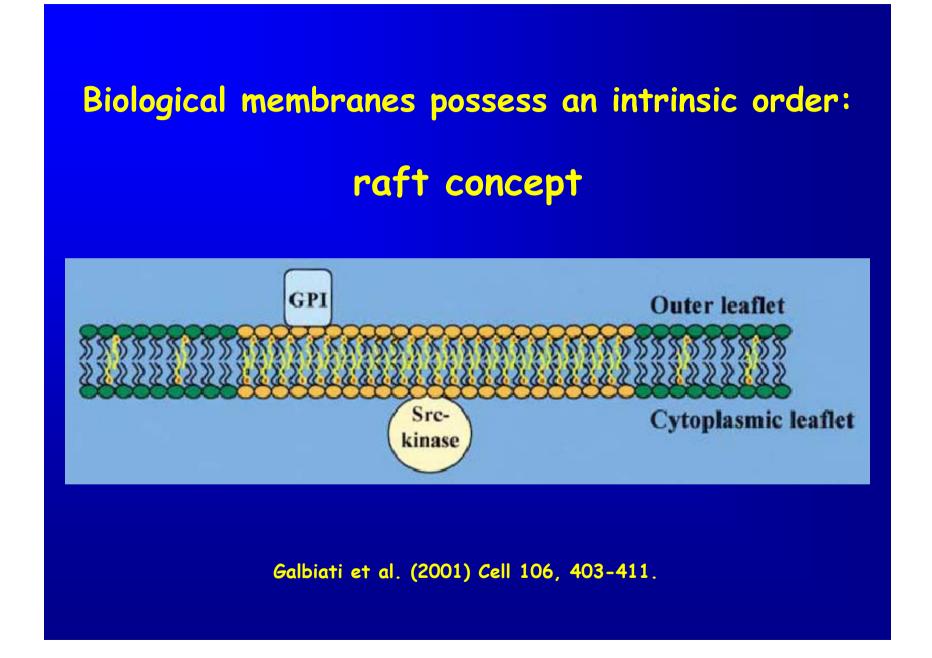
above mentioned detergents at higher T

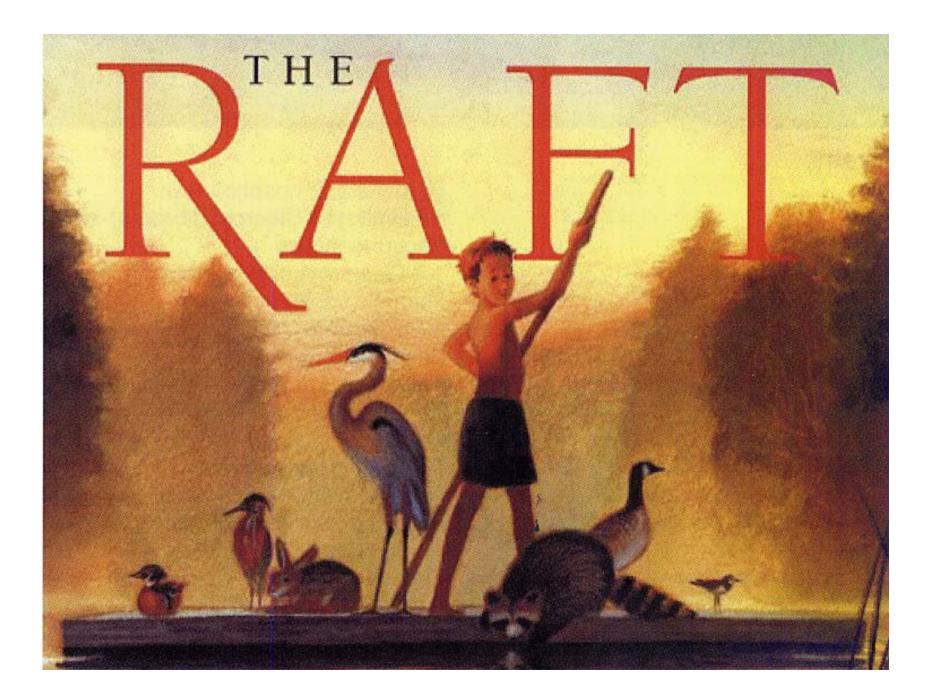
Basic lipid structures



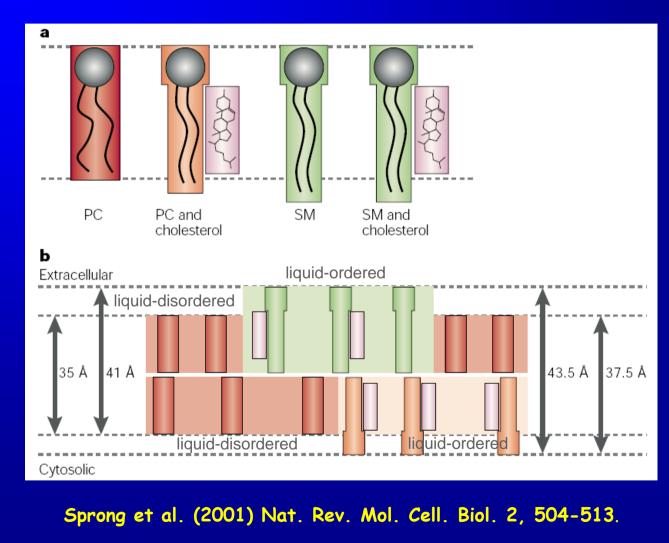
Lipids exist in:

- gel state (semi-frozen)
 - liquid-ordered state
- liquid-disordered state (fluid mosaic)

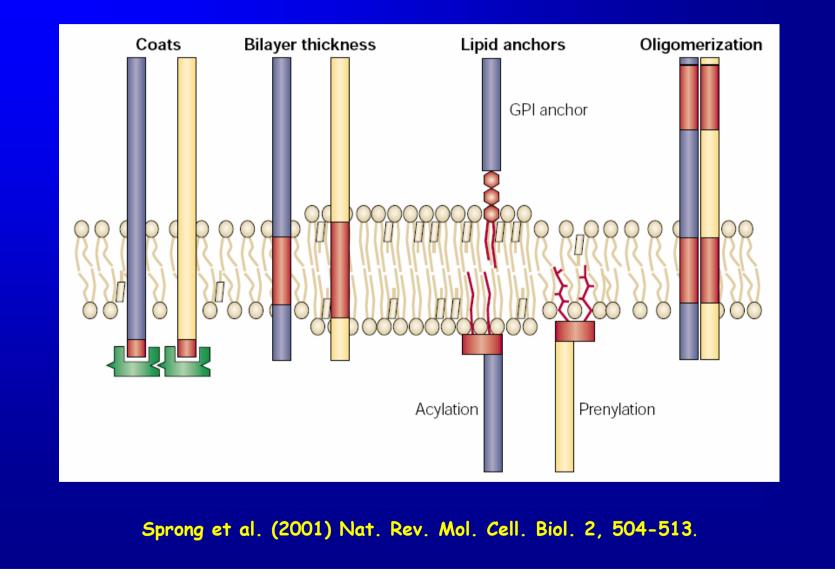




Cholesterol can induce fluid-fluid immiscibility



Lateral sorting of membrane proteins



Cholesterol is concentrated in lipid rafts



Common tools to disrupt lipid rafts

Cholesterol sequestration

- Antibiotics:
 - Filipin | Nystatin | Amphotericin
- Pore-forming agents:
 Saponin | Digitonin | Streptolysin O

Cholesterol depletion

Methyl-β-cyclodextrin

Inhibition of cholesterol biosynthesis

Lovastatin

Perturbation of raft stability

- Exogenous cholesterol
- Exogenous gangliosides
- Exogenous polyunsaturated fatty acids

Simons & Toomre (2000) Nat. Rev. Mol. Cell Biol. 1, 31-40.

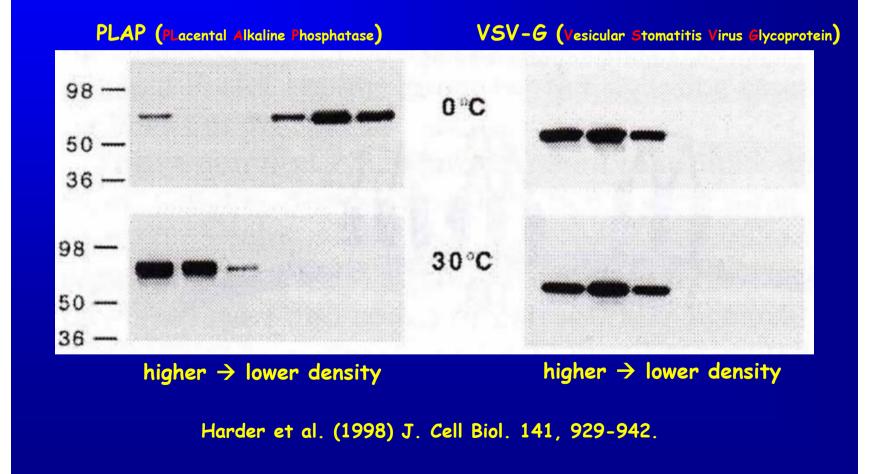
Techniques to identify lipid rafts

Approach*	Information available	Live cells	Comments	References
Flotation of detergent– resistant membranes (DRMs)	Identifies putative raft association Identifies possible raft proteins	No	 Easy to do Most common approach for identifying putative proteins involved in signalling Artefacts possible Weak associations with rafts are difficult to detect 	1, 4, 7
Antibody patching and immunofluorescence microscopy	Identifies putative raft association	No	 Easy to do Common approach Better than flotation for detecting weak raft associations Cell-cell variability makes quantification difficult 	12, 29
Immunoelectron microscopy	Determines location of raft components	No	 Promising results Requires technical expertise 	80, 81, 88
Chemical crosslinking	Identifies native raft protein complexes	Yes	 Straightforward Choice of appropriate conditions and reagents is semi-empirical 	32
Single fluorophore tracking microscopy	Monitors the diffusion and dynamics of individual raft proteins or lipids	Yes	 Requires highly specialized equipment and expertise 	89
Photonic force microscopy	Determines the diffusion constant, size and dynamics of individual rafts	Yes	 Very informative technique Requires highly specialized equipment and technical expertise Time-consuming acquisition and analysis 	30
Fluorescence resonance energy transfer (FRET)	Detects whether two raft components are spatially close (for example, <10 nm)	Yes	 Powerful approach Choice of appropriate donor and acceptor probes is important 	31, 33

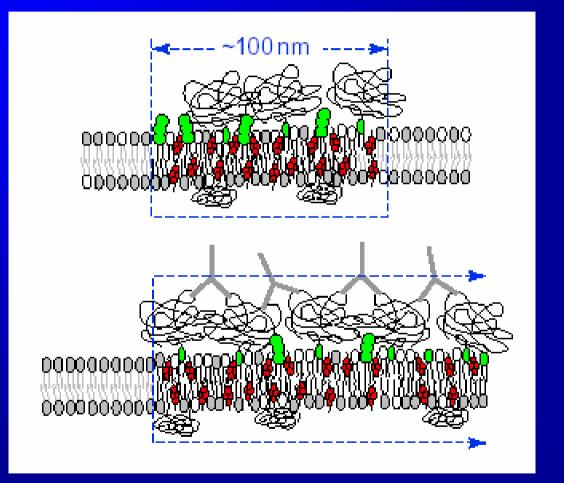
*The disruption of rafts by cholesterol depletion or sequestration is especially useful as a control for each of these approaches.

Simons & Toomre (2000) Nat. Rev. Mol. Cell Biol. 1, 31-40.

Solubilization of biological membranes in 2% (v/v) TR X-100 at 4 or 30°C followed by sucrose gradient centrifugation (flotation) analysis.

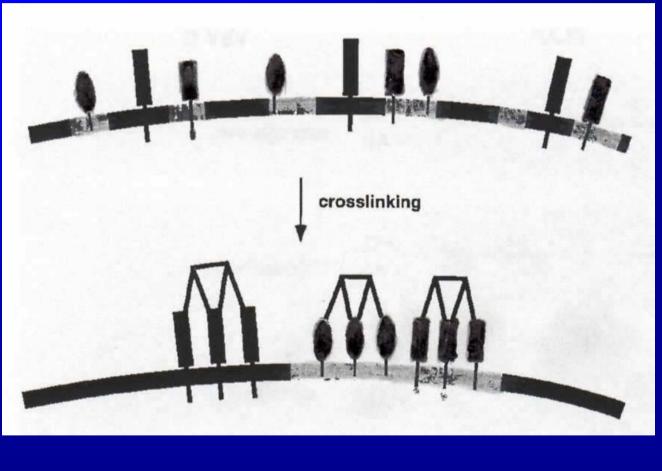


Patching (clustering) of membrane components



Jacobson & Dietrich (1999) Trends Cell Biol. 9, 87-91.

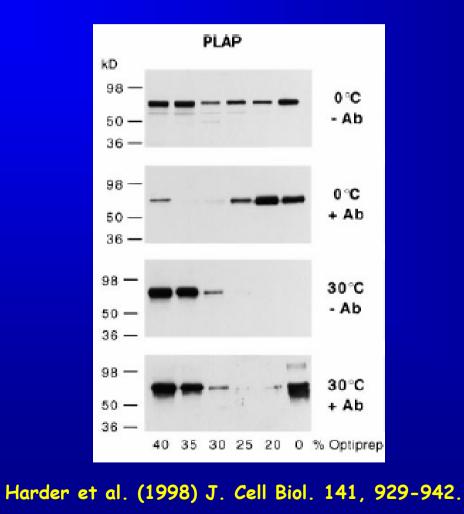
Bulk separation of membrane phases caused by clustering (patching) of membrane components



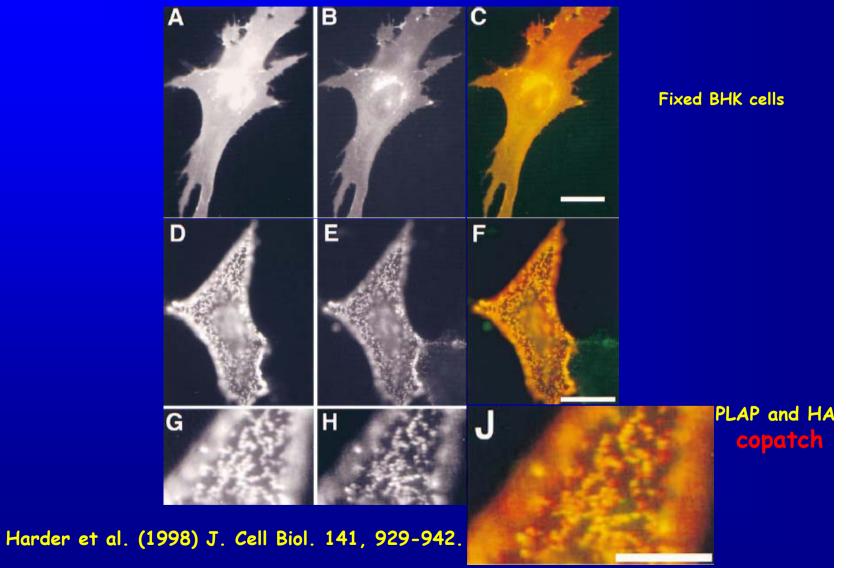
Harder et al. (1998) J. Cell Biol. 141, 929-942.

Stabilization of membrane domains by Ab crosslinking of a GPI-protein PLAP,

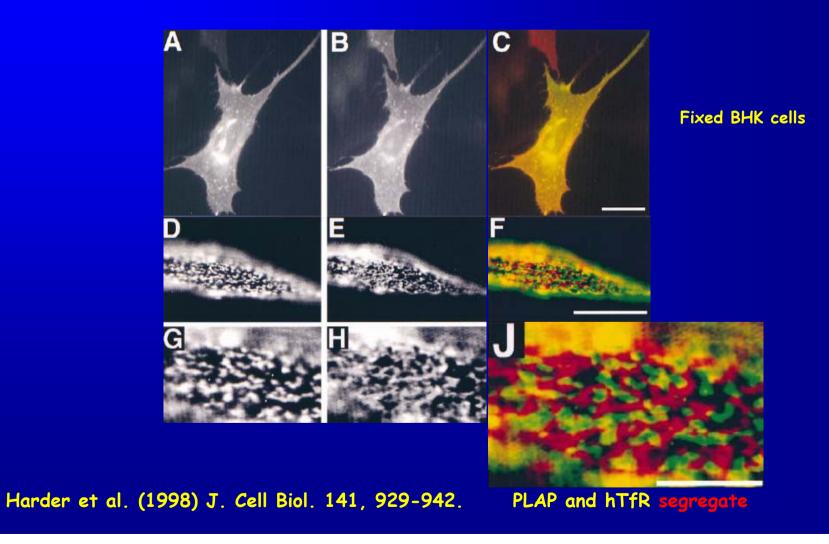
transiently expressed in nonpolarized fibroblastoid BHK 21 cells



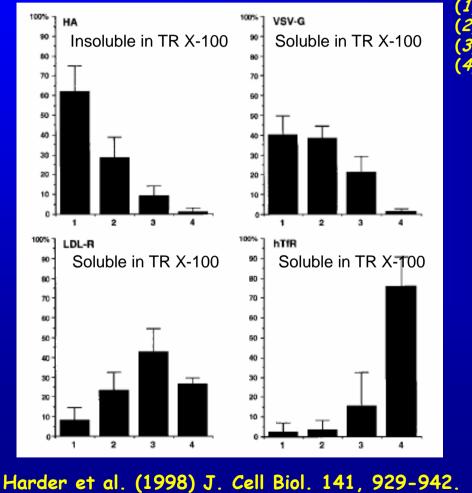
Patching of GPI-anchored PLAP (red) and influenza HA (green) transiently coexpressed in nonpolarized BHK-21 cells



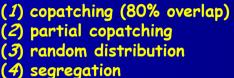
Patching of GPI-anchored PLAP (green) and hTfR (red) transiently coexpressed in nonpolarized BHK-21 cells



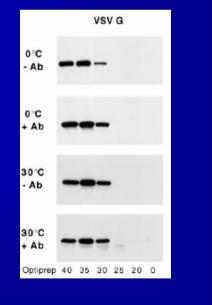
Certain proteins exhibit a weak but significant raft interaction which is not detectable by the TR X-100-solubility criterium



PLAP copatching

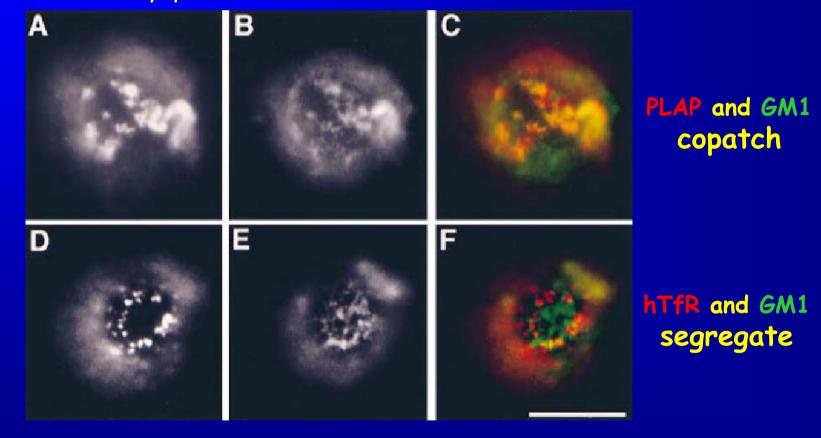


TR X-100 solubility



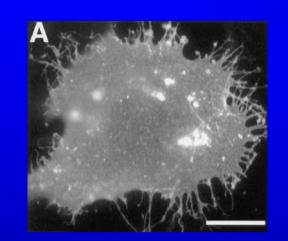
Specific involvement of lipid interactions in the copatching phenomenon

PLAP (A-C) and hTfR (D-F) were transiently coexpressed in Jurkat T-lymphoma cells.

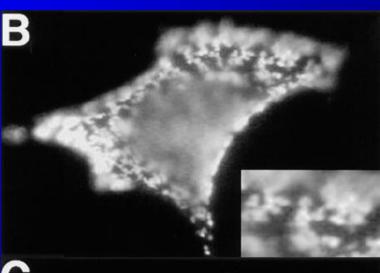


Harder et al. (1998) J. Cell Biol. 141, 929-942.

Accumulation of src-like Tyr kinase fyn in membrane domains formed by patched PLAP transiently expressed in nonpolarized BHK-21 cells.



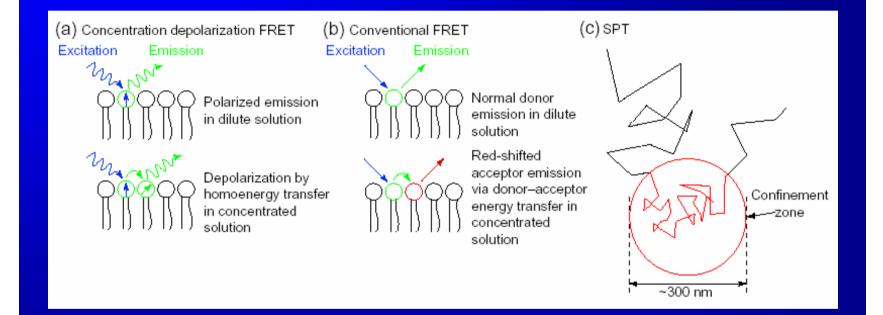
Distribution of overexpressed fyn in fixed BHK cells is even.



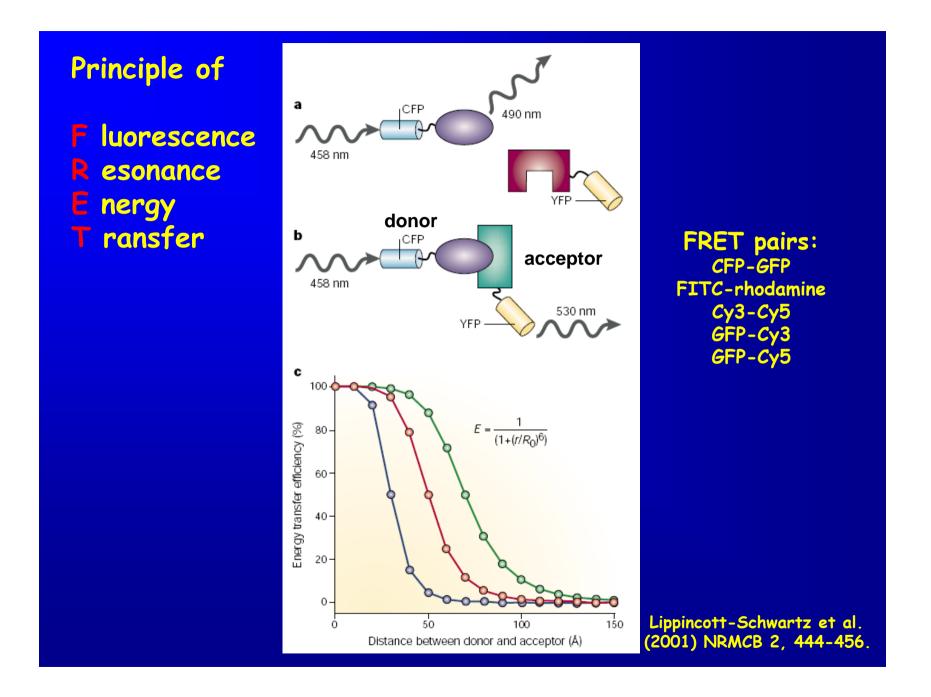
Patches of PLAP

Distribution of fyn in PLAP patched cells

Harder et al. (1998) J. Cell Biol. 141, 929-942.



Jacobson & Dietrich (1999) Trends Cell Biol. 9, 87-91.



FRET assay for detecting lipid rafts FRET as a function of donor and acceptor surface density

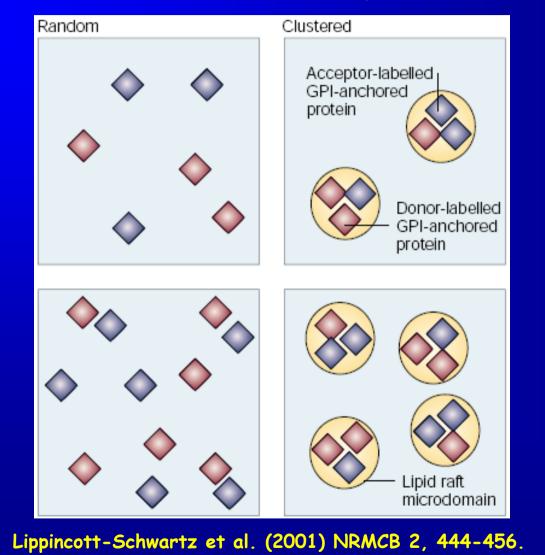


Table 1. Lipid and Protein Components of Lipid Rafts/Caveolae Lipids Cholesterol Sphingo-myelin Glyco-sphingolipids (e.g., GM₁) PIP₂ Proteins Integral/Structural Caveolins (Cav-1, -2, and -3) Flotillins (FLO-1 and -2; aka, Reggies or Cavatellins) LAT/PAG MAL/BENE Stomatins VIP36 Acylated Exoplasmic GPI-linked proteins (e.g., Thy-1, alkaline phosphatase, folate receptor) Cytoplasmic Src-family tyrosine kinases (NRTKs) G proteins eNOS H-Ras Scavenger Receptors CD 36 SRBI RAGE Other receptors Receptors Tyrosine kinases (RTKs; e.g., EGF-R, PDGF-R, Insulin-R) Hepta-helical Receptors (e.g., Endothelin receptor)

Galbiati et al. (2001) Cell 106, 403-411.

Raft distribution and trafficking is cell type-dependent

Polarized epithelial cells
accumulated in apical PM

Neurons
 accumulated in axonal PM

Lymphocytes and fibroblasts - uniform distribution

Raft nomenclature

Present raft nomenclature*

Rafts | DRMs | DIGs | DICs | GPI domains | Glycosphingolipid signalling domains | Caveolae-like domains | Microdomains | LDM | Liquid-ordered domains | DIM | GEMs | TIFF

Suggested raft nomenclature

	I. Rafts	II. Clustered rafts	III. DRMs	IV. Caveolae
Components	 Glycosphingolipids Cholesterol Lipid-modified proteins containing saturated acyl chains: GPI-anchored proteins Doubly acylated Src-type kinases Transmembrane proteins 	 Rafts clustered by: Antibody Lectin Adjacent cell proteins Physiological crosslinking proteins 	 Rafts remaining insoluble after treatment on ice with detergent‡§: Triton X-100 (most popular), Brij-58, CHAPS, NP-40 	 Raft proteins and lipids Caveolins
Properties	 50 nanometres in diameter Mobile (~10⁻⁸ cm⁻² sec⁻¹) Liquid-ordered phase 	 Large, often hundreds of nanometres to micrometres in size Often bound to cytoskeleton 	 Float to low density in sucrose or Optiprep[™] density gradients 	 Morphological 'cave-like' invaginations on the cell surface
Comments	 Native rafts are only detected in living cells 	 Clustering is used both artificially and physiologically to trigger signalling cascades 	 Non-native (aggregated) raft Variable effects depending on: Detergent type Detergent:lipid ratio Cell type 	 Raft subcategory Highly specialized

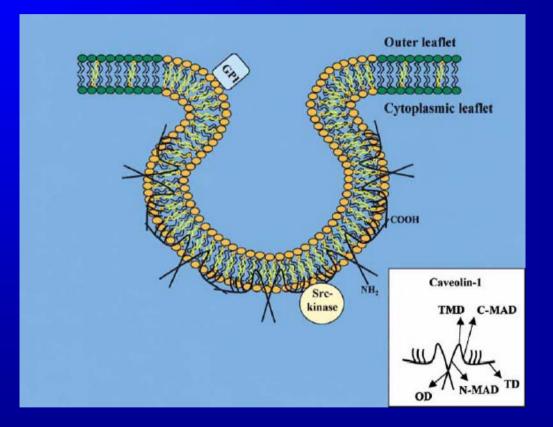
* DRM, detergent-resistant membrane; DIG, detergent-insoluble glycolipid-rich domain; DIC, detergent-insoluble complex; LDM, low-density membrane; DIM, detergent-insoluble material; GEM, glycolipid-enriched membrane; TIFF, Triton X-100 insoluble floating fraction. ‡Care should be taken when choosing solubilization conditions for co-immunoprecipitation experiments, as these popular detergents do not solubilize rafts on ice.

‡Care should be taken when choosing solubilization conditions for co-immunoprecipitation experiments, as these popular detergents do not solubilize rafts on ice. Co-localization of proteins in rafts or DRMs could be mistaken for direct protein-protein interactions if rafts are not completely solubilized.

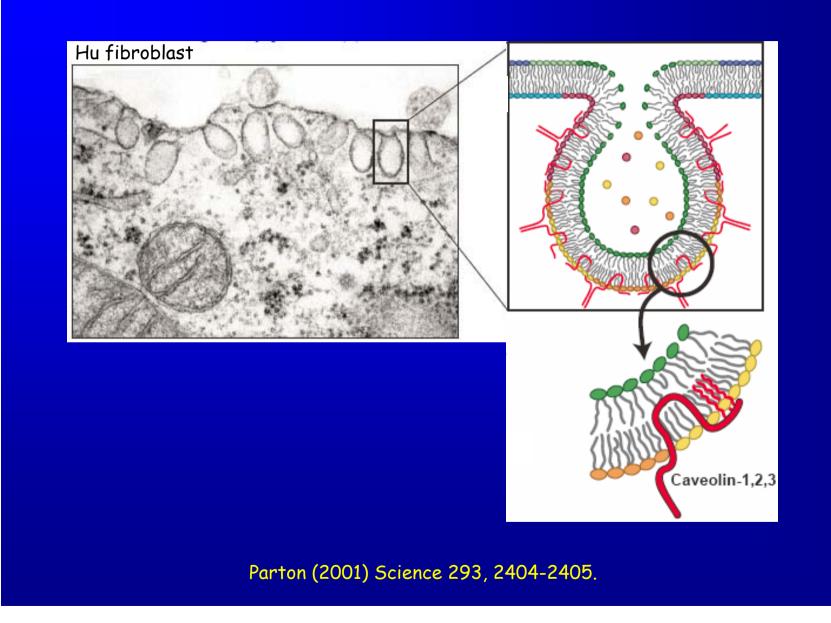
§Rafts can be solubilized in octyl glucoside or in the detergents listed above at raised temperatures.

Simons & Toomre (2000) Nat. Rev. Mol. Cell Biol. 1, 31-40.

Caveolae highly specialized raft subcategory



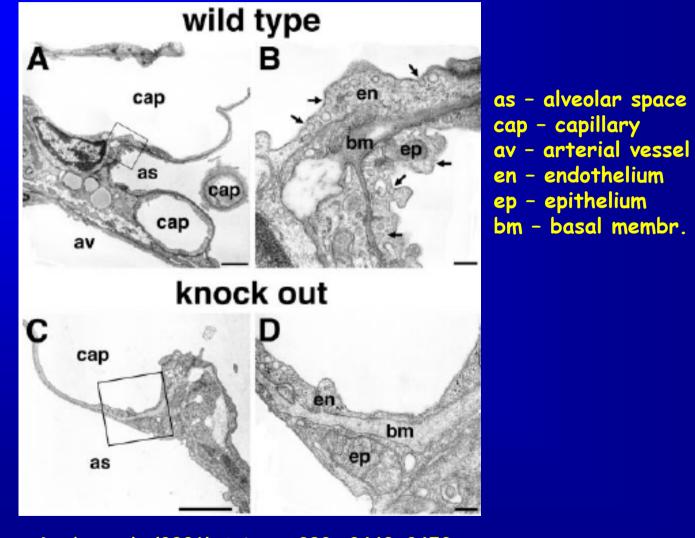
Galbiati et al. (2001) Cell 106, 403-411.



Caveolins (Cav)

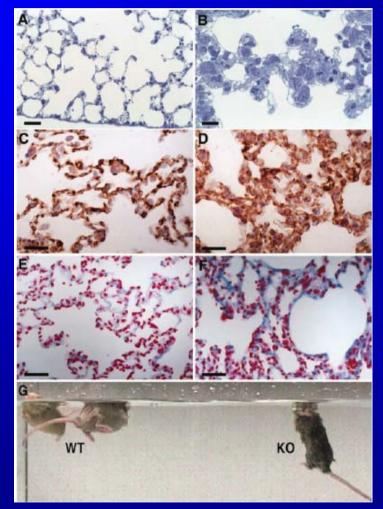
- Essential for the formation of caveolae.
- Cav gene family structurally and functionally conserved from worms (*C. elegans*) to humans.
- Cav-1(α and β), -2 and -3 in mammals (21- to 25-kDa).
- Integral membrane proteins (tri-palmitoylated).
- Cav-1 and -2 are coexpressed, Cav-3 is muscle-specific.
- Polymerize (14-16) and shape up caveolae.
- Bind cholesterol, fatty acids and interact with the broad range of signal transducing molecules (*e.g.* Tyr kinase R, eNOS, heterotrimeric G proteins).
- Not present in lymphocytes and neurons.

Disappearance of caveolae from cells in cav-1 (-/-) mice



Drab et al. (2001) Science 293, 2449-2452.

Patomorphological defects in lungs and physical disability of cav-1 (-/-) mice



Thickening of alveolar walls

caused by:

- uncontrolled endothelial cell proliferation

- increased content of extracellular fibrillar matrix (fibrosis)

results in:

physical weakness.

Drab et al. (2001) Science 293, 2449-2452.

Features of caveolin-defficient mice

Gene knockout	Features
Caveolin-1	Loss of caveolae in non-(striated) muscle cells, loss of caveolin-2 protein, pulmonary defects — exercise intolerance*, vasoconstriction or dilation abnormalities, resistance to diet-induced obesity — high serum triglyceride levels, defects in albumin uptake or transendothelial transport, defects in glycosylphosphatidylinositol-anchored protein transport
Caveolin-2	Pulmonary defects - exercise intolerance
Caveolin-3	Loss of caveolae in striated muscle, dystrophic muscle, disorganization of T-tubule network
*Proposed to be due	to the loss of caveolin-2 protein in caveolin-1-null mice.

Parton (2003) Nature Rev. MCB 4, 162-167.

Recommended reading:

- Drab, M. et al. (2001): Loss of Caveolae, Vascular Dysfunction, and Pulmonary Defects in Caveolin-1 Gene-Disrupted Mice. Science 293, 2449-2452.
- Galbiati, F. et al. (2001): Emerging themes in lipid rafts and caveolae. Cell 106, 403-411
- Jacobson, K. and Dietrich, C. (1999): Looking at Lipid Rafts? Trends Cell Biol. 9, 87–91.
- Jacobson, K., et al. (2007): Lipid rafts: at a crossroad between cell biology and physics. Nat. Cell Biol. 9, 7–14.
- Lippincott-Schwartz, J., et al. (2001): Studying protein dynamics in living cells. Nat. Rev. Mol. Cell Biol. 2, 444–456
- Riethmuller, J., et al. (2006): Membrane rafts in host-pathogen interactions. Biochim. Biophys. Acta1758, 2139–2147.
- Parton, R.G. (2003): Caveolae from ultrastructure to molecular mechanisms. Nat. Rev. Mol. Cell Biol. 4, 162–167.