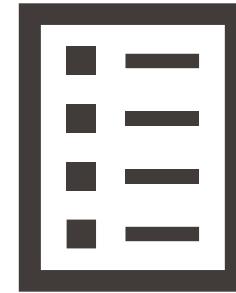


# Nanoscale organization of multiple GPI-anchored proteins in living cell membranes

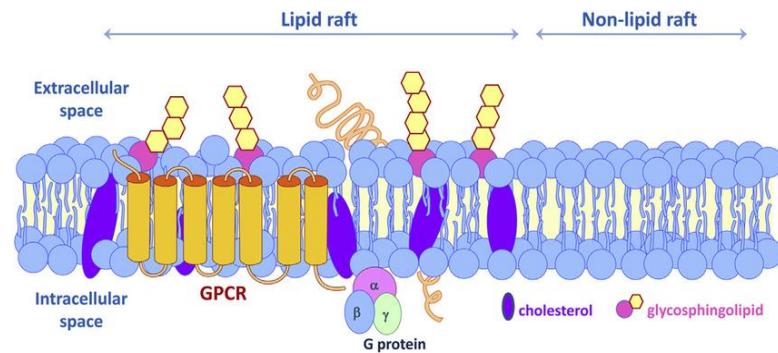
Charpié Coralie  
Dias Gomes Vera  
Richard Jonas  
Reuse Coralie

# Plan of the presentation:

- Introduction and definitions
- Aim of the study
- Methods used
- Explanation of the results
- Conclusion
- Key takeaways from the study

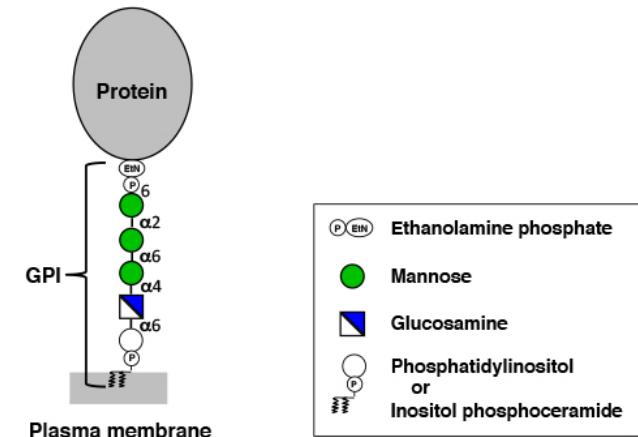


- **Definition** : lipid microdomains within the cell membrane
- **Role** : organizing membranes components
- **Structure** : enriched in lipids (glycosphingolipids and cholesterol) and proteins (as GPI-APs)
- **Functions** : transmit signal, facilitate transport, membrane stability and resistance, cell communication



# GPI-APs (glycosylphosphatidylinositol-anchored proteins)

- Protein attached to the outer surface of the cell membrane
- **Structure of GPI group:** lipid and sugar chain
- **Application :** can be used as markers for membrane rafts
- **Interest :** form small clusters that could represent a form of raft



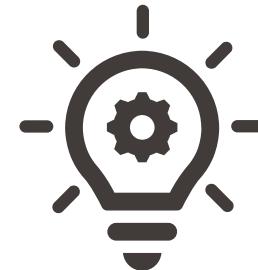
# Evolution of the concept of "rafts"

- **Initially** : large and stable structures acting as platform for gathering proteins and lipids
- **New hypothesis** : small and dynamic structure able to change shape and size

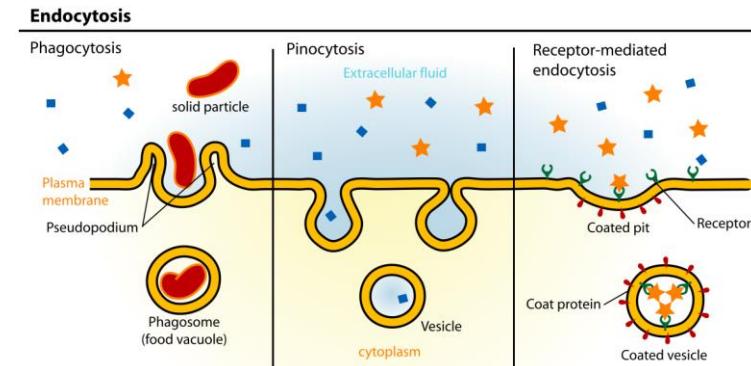


# Aim of the study

- Investigate the nanoscale organization of GPI-APs.
- Provide direct evidence of lipid raft-like structures in living membranes.
- Clarify raft's role in cellular functions.

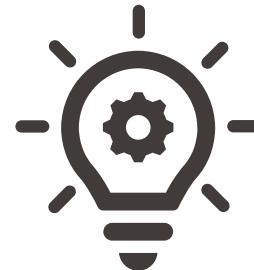


- Process by which a cell absorbs substances by surrounding them with its membrane to form a vesicle that enters the interior of the cell.



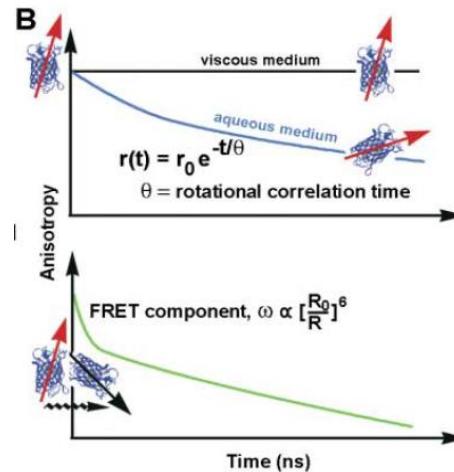
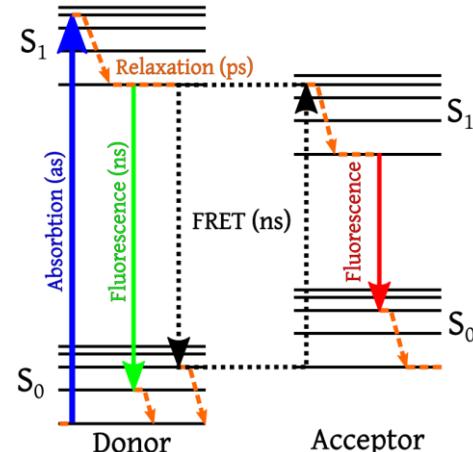
# Aim of the study

- Investigate the nanoscale organization of GPI-APs and their connection to the concept of lipid rafts.
- Determine informations about clusters.
- Seeks to provide direct evidence of lipid raft-like structures in living membranes.
- Clarify raft's role in cellular functions.



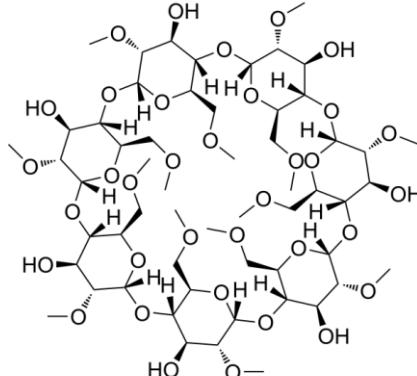
# Methods used

- Fluorescence and electron **microscopy**
- homo-Fluorescence Resonance Energy Transfer (**FRET**) microscopy: to deduce protein clustering.
- **time-resolved anisotropy decay experiments**: to measure intermolecular distances.
- **photobleaching** experiments combined with **theoretical modeling**: to analyze cluster size and density.

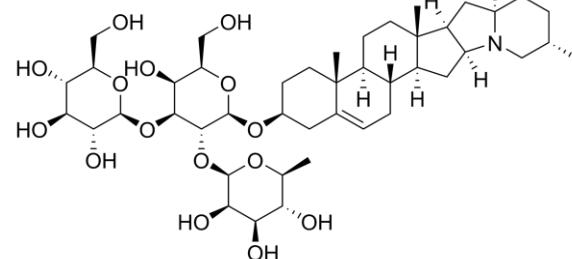


# Methods used

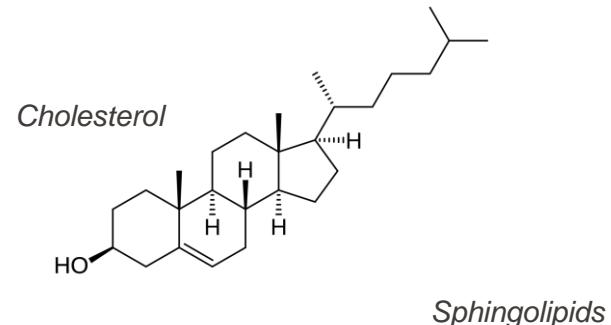
- **Cholesterol and Sphingolipids depletion:** role of lipids in maintaining GPI-AP organization.



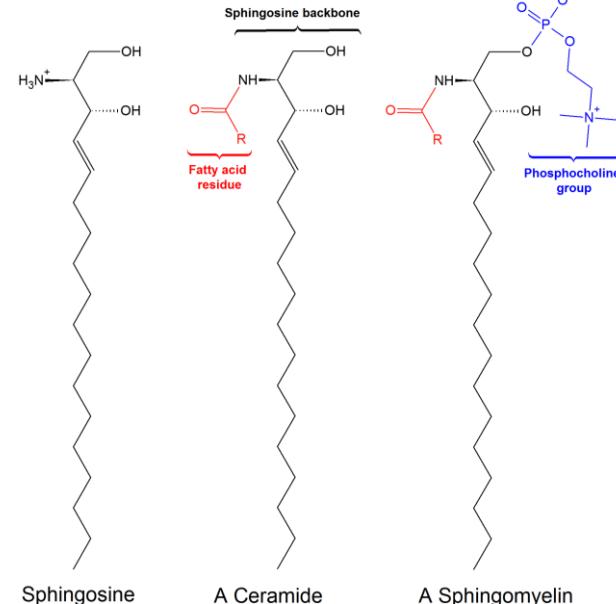
*methyl- $\beta$ -cyclodextrin*



*saponin*



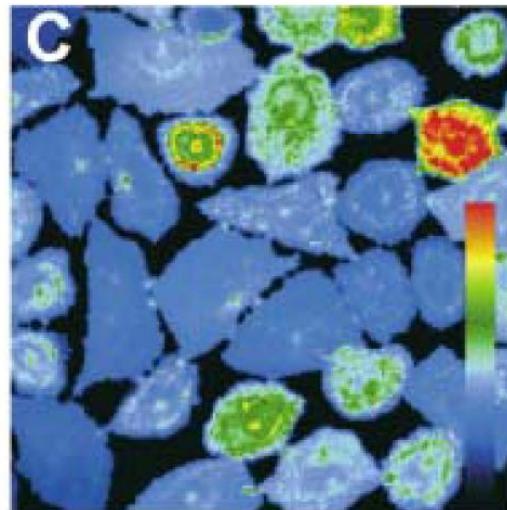
*Sphingolipids*



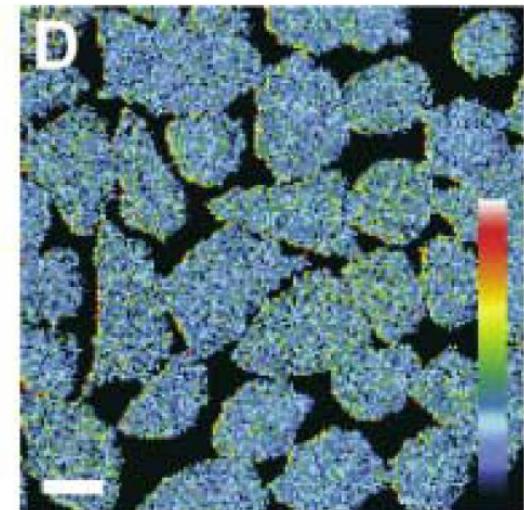
# Methods used

- **Hetero-FRET**: to detect clustering between distinct GPI-AP species.
- **Antibody-mediated crosslinking**: to reorganize clusters.

Total intensity image

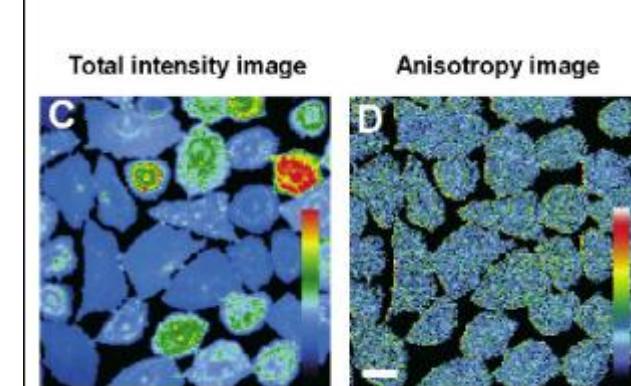


Anisotropy image



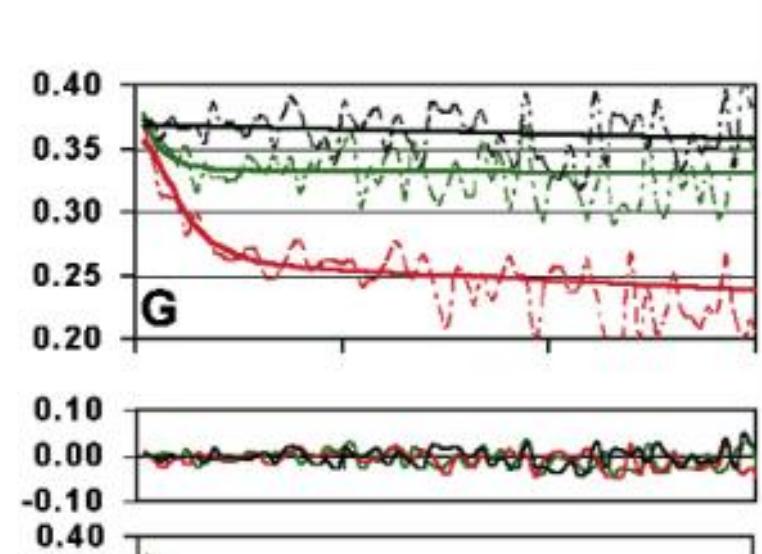
# The presence of GPI-APs as clusters at the cell surface

- GFP-GPI shows fluorescence
- Reduced anisotropy values
- Homo-FRET vs rotational mobility



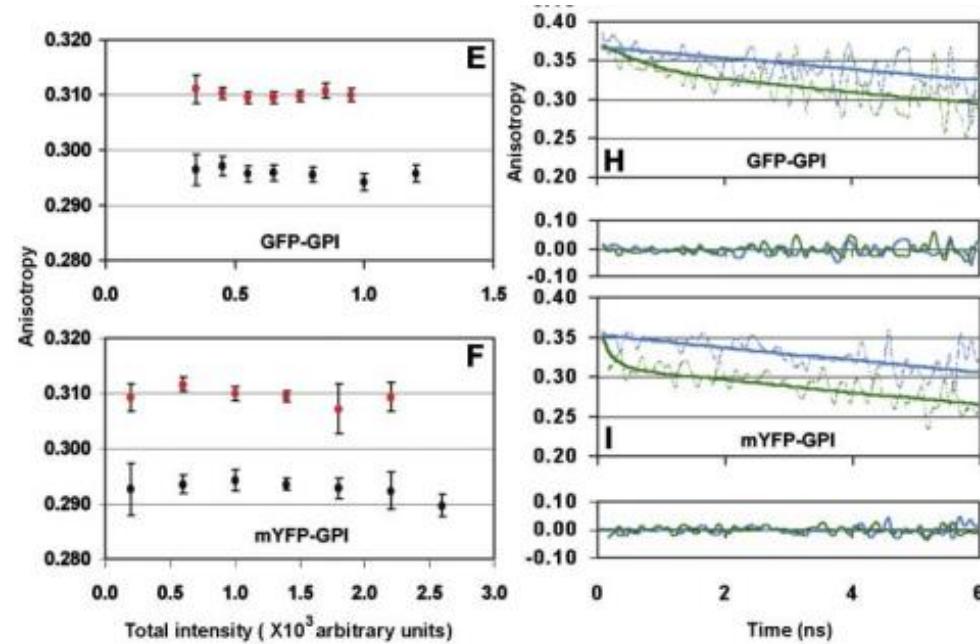
# Decrease comes from FRET which proves clustering of GFP-GPI

- The decay is absent for GFP molecules in free solution (Black line)
- Fast decay is present upon cross linking with glutaraldehyde (Red line)
- Not sensitive to viscosity increase (Green line)
- Occurs at rate much faster than the lifetimes of fluorophores
- Reducing the fluorophores density increase the steady-state of anisotropy



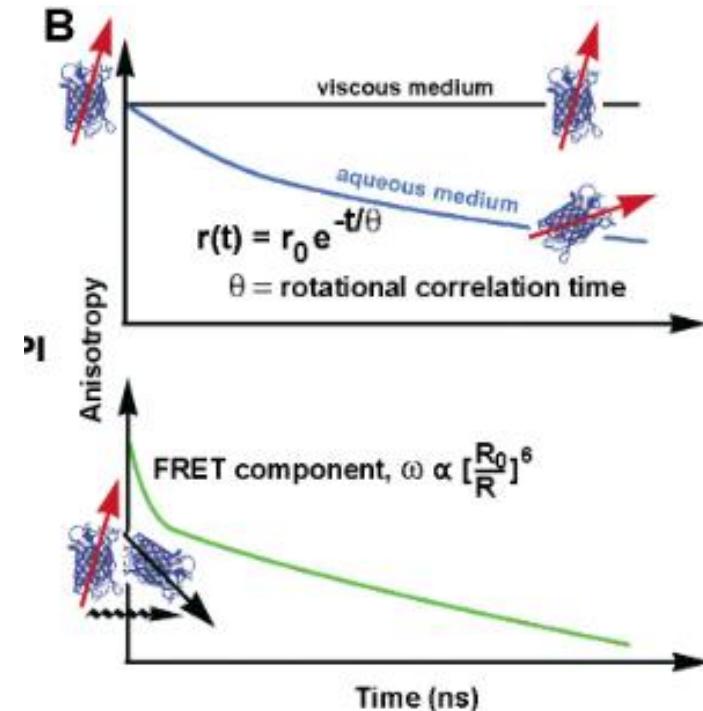
# Origin of FRET between GPI-AP species

- Not from protein-protein interaction
- Loss of homo-FRET with cholesterol depletion (Figure H & I)
- Constant anisotropy (Figure E & F)
- Disparition of FRET dependance when changing the anchor



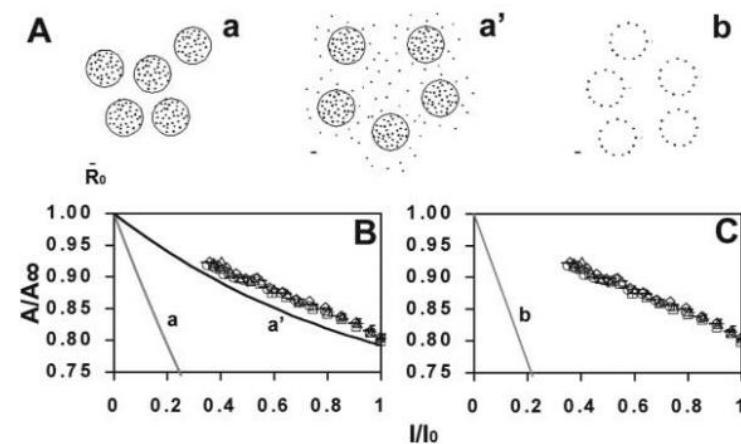
# GPI-APs are present in extremely high-density structures

- Utility of the previous result
- More than 10% in high-density structure,  $3.53 \pm 0.455$  nm
- Comparison with 0.3 nm crosslinker
- mYFP-GPI faster decay
- PFL-FR-GPI same
- So GPI-anchor are responsible of high density structure < 4nm

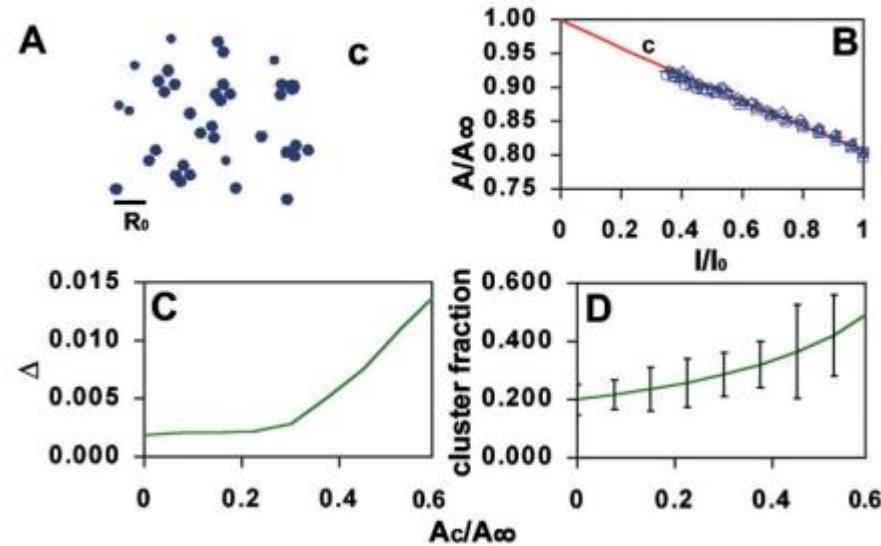


# Three different models for describing spatial distribution of proteins

- Model A : GPI-APs are uniformly distributed in large domains (10 times larger than their molecular size,  $\sim 50$  nm).
- Model A' : a fraction of GPI-APs are in large domains, the rest are dispersed as monomers on the surface.
- Model B : GPI-APs are concentrated on the periphery of the domains.



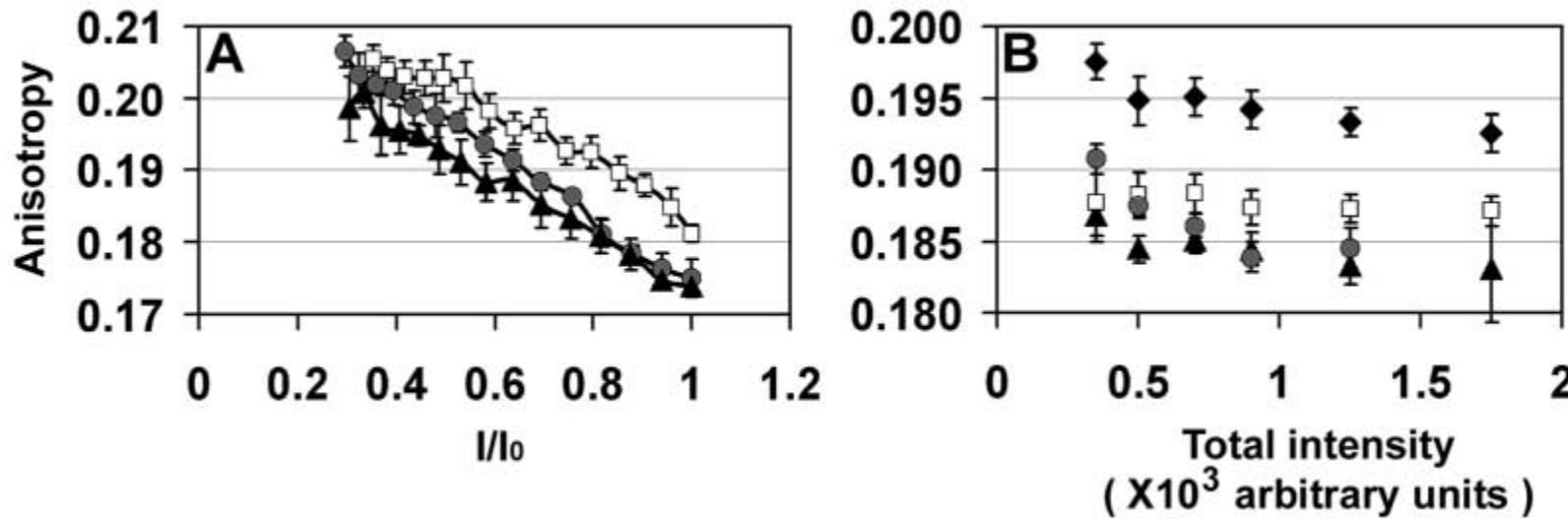
- ~20-40% of proteins are in cluster smaller than  $R_0$
- Protein close to each other which favors FRET  $\sim 5\text{nm}$
- Remaining proteins are monomers distributed on the cell surface
- Membrane have clusters and monomers



# Organisation of GPI-APs into clusters

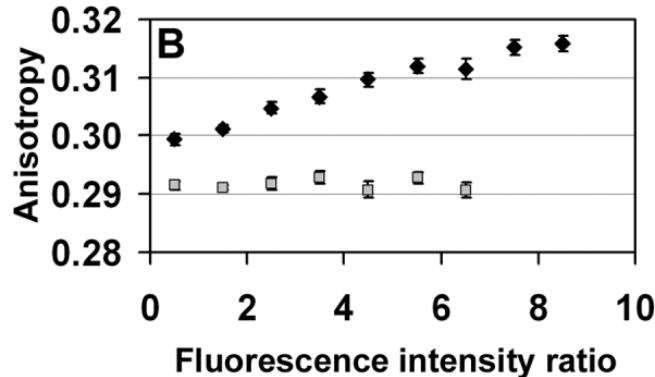
- **Homo-FRET detected :**
  - ~30% of GPI-APs are likely to be organized in very compact clusters
  - Small clusters and/or little proteins inside it
- **Hetero-FRET undetected :**
  - No signal between different fluorophore
  - No large cluster
- **Confirmation of the observation:**
  - Big cluster forced with Aerolysine Y221G
  - Measurable Hetero-FRET
  - So absence of hetero-FRET was due to the size and fraction of the cluster
- **Conclusion**
  - ~20-40% of clusters
  - $\leq 4$  molecules per cluster

# Cholesterol and sphingolipid depletion differentially affect GPI-AP clustering

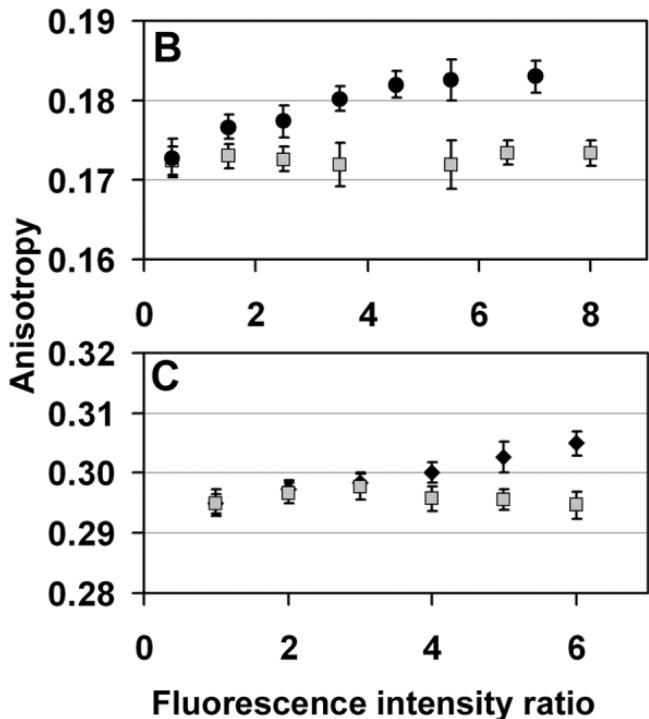
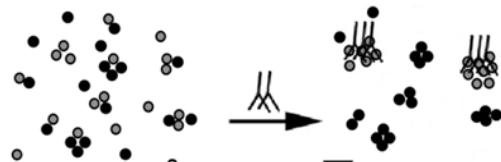


- Black triangle : untreated
- Gray circle : sphingolipid-depleted
- Open square : cholesterol-depleted
- Black diamond : depleted of both cholesterol and sphingolipid

# Multiple GPI-APs inhibit the same nanometer-sized cluster

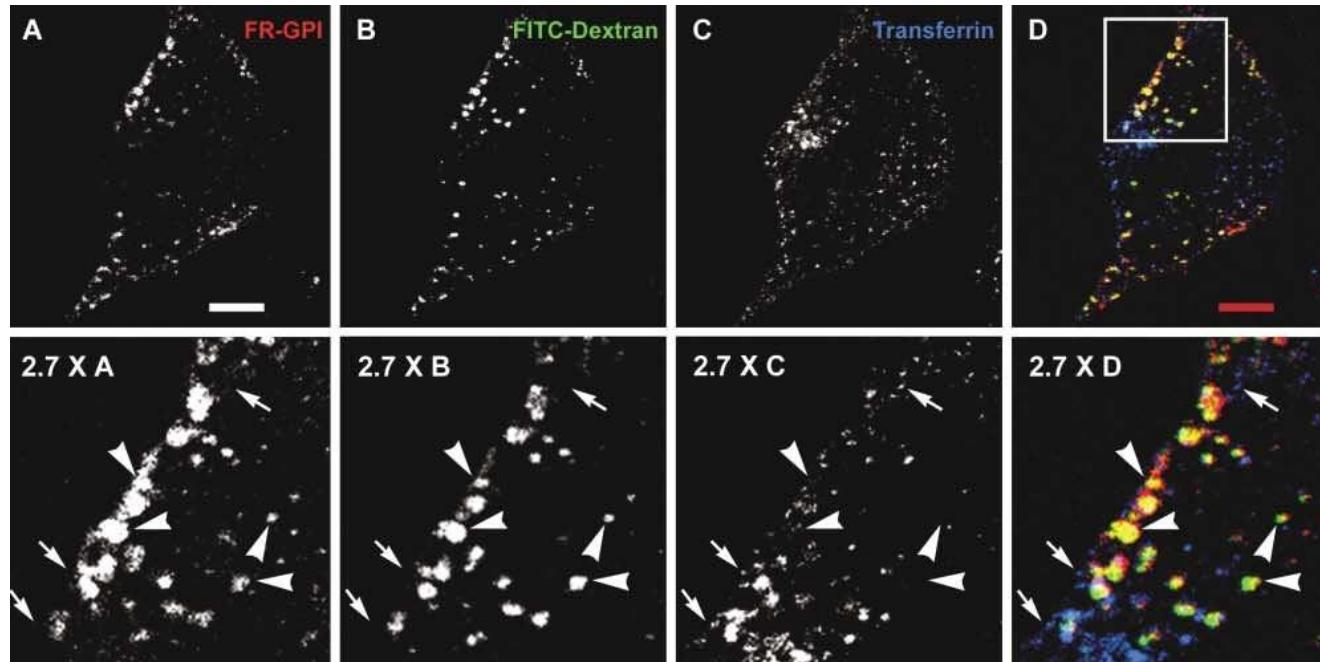


- Black diamond : FR-GPI
- Open square : FR-TM



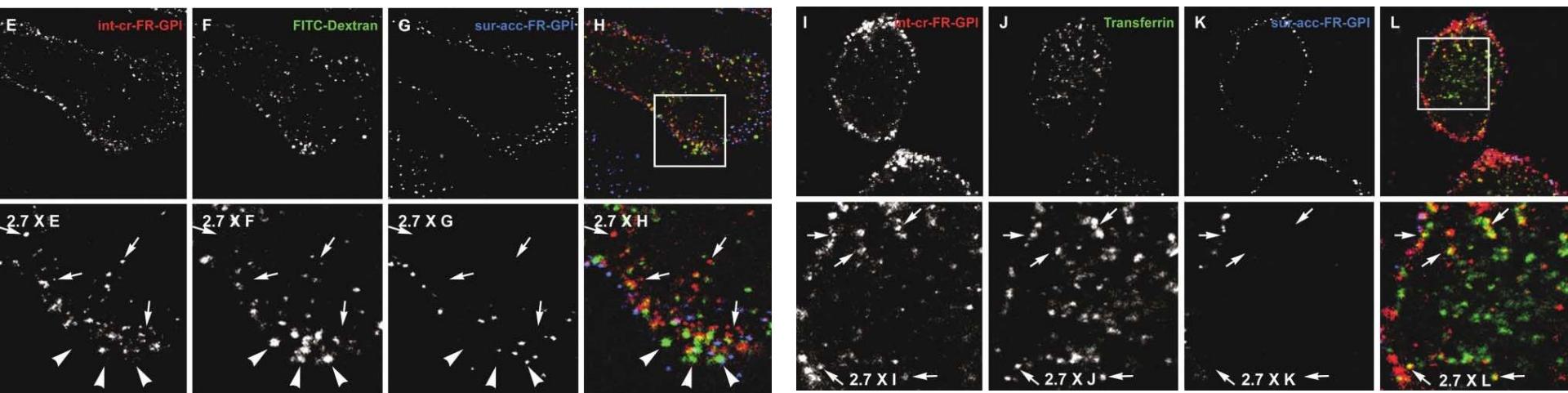
- Black circle : FR-GPI
- Black diamond : GFP-GPI
- Open square : crosslinking with DAF

# Antibody-crosslinking alters endocytic routing of crosslinked proteins



- FITC-dextran : GPI-AP-specific endocytic compartments, or GEECs
- Transferrin : clathrin-mediated pathway

# Antibody-crosslinking alters endocytic routing of crosslinked proteins



- FITC-dextran : GPI-AP-specific endocytic compartments, or GEECs
- Transferrin : clathrin-mediated pathway
- Tertiary fluorescent antibody : to distinguish between internal and surface-bound FR-GPI

# Conclusion:

- Possibility to measure the structures
- 20-40% of GPI-APs form small clusters of 3-4 molecules
- Remaining of GPI-APs are monomers
- Cholesterol is essential
- Sphingolipids play an indirect role
- Multiple GPI-APs can be found in the same cluster
- Antibody crosslinking alters endocytic fate



# Key Takeaways from the Study



# Thank you for listening !