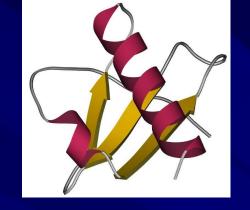
### PROTEIN PHYSICS

### LECTURES 11-12

- Fibrous proteins and their functions
- Membrane proteins and their functions

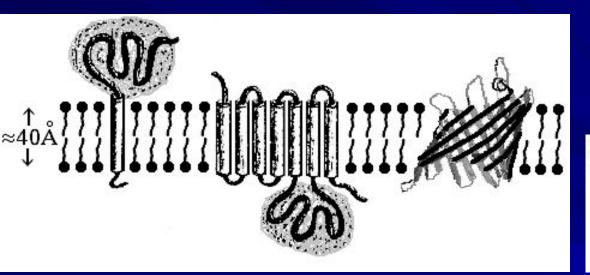
- Fibrous proteins: building blocks
- Membrane proteins: transmitters



### Globular proteins

00•00••0000••0•0•00

quasi-random



# Membrane proteins

Hydro- | Hydrophobic philic block block

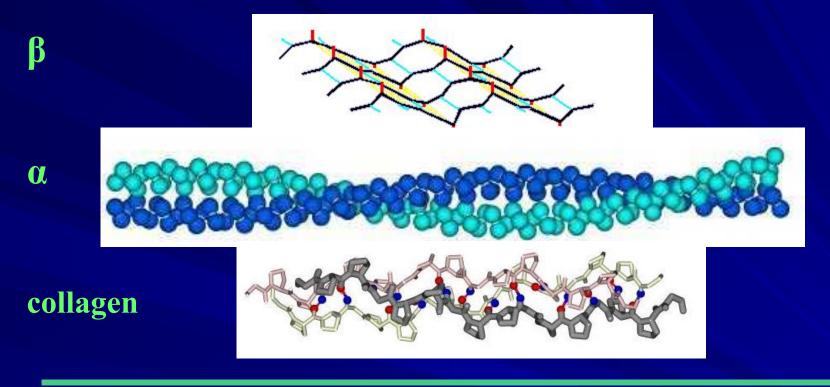


#### Fibrous proteins

•00•000•00•000•000 | <u>repeat</u> | <u>repeat</u> |

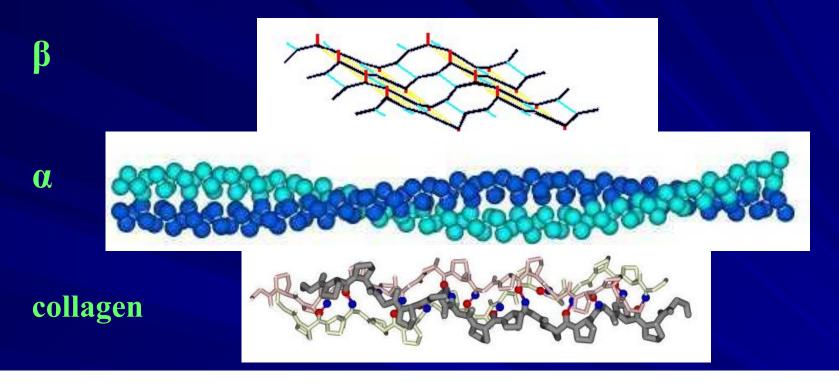
H-bonds (NH:::OC) & hydrophobic forces

### Fibrous proteins: regular building blocks

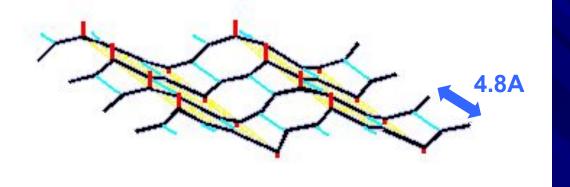


Here, we will not consider fibrous proteins made of globules (actin, etc.)

# Fibrous proteins: regular building blocks



PROTEINS	SEQUENCES	
Globular	00•00•00•0•0•0•00•000•0•0•0•0•0•0	quasi-random
Membrane	Hydro-   Hydro-    phobic philic	blocks
<u>Fibrous</u>	•00•000•00•000•00•00•00•00•00•000•00	repeats



β

#### Silk fibroin

$$\sim (\sim^{\text{Ser}}_{\text{Gly}}\sim^{\text{Ala}}_{\text{Gly}}\sim^{\text{Ala}}_{\text{Gly}})_{8}\sim \times \sim 50$$

Ser\_Gly\_Ala\_Gly\_Ala\_Gly

3.5 Å

Ser\_Gly\_Ala\_Gly\_Ala\_Gly

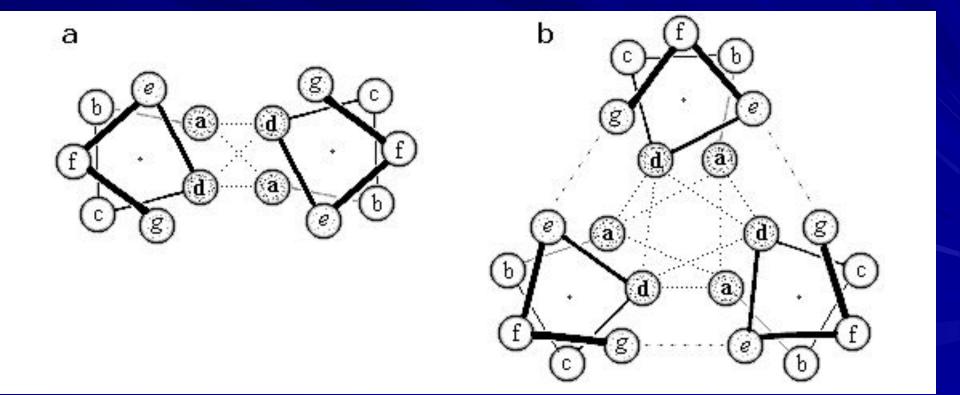
5.7 Å

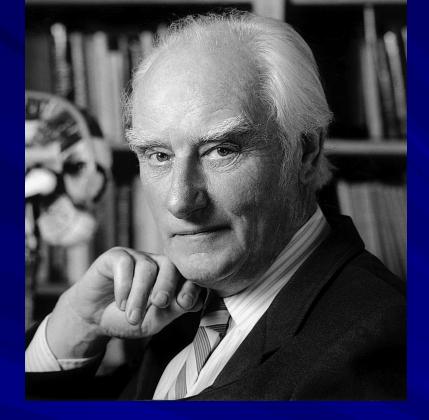
Ser\_Gly\_Ala\_Gly\_Ala\_Gly

Ser\_Gly\_Ala\_Gly\_Ala\_Gly



# α-helical coiled-coil



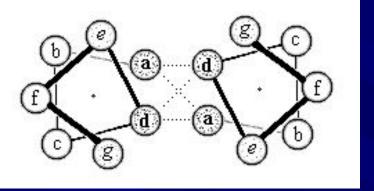


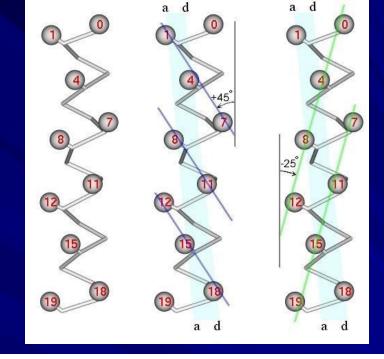
Francis Harry Compton **Crick** (1916 – 2004) Nobel Prize 1962 for DNA structure, 1953

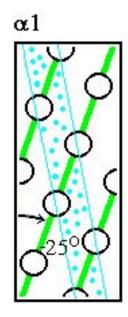
Coiled coil structure: F. Crick, 1952

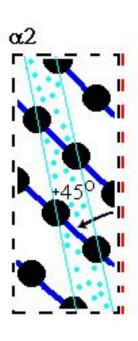
C. Chothia, M. Levitt, D. Richardson, 1977

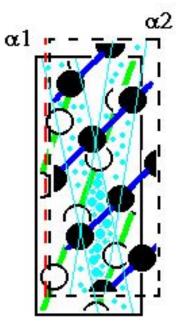
### α-helix packing

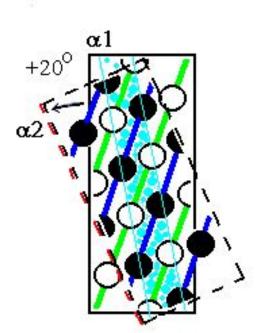




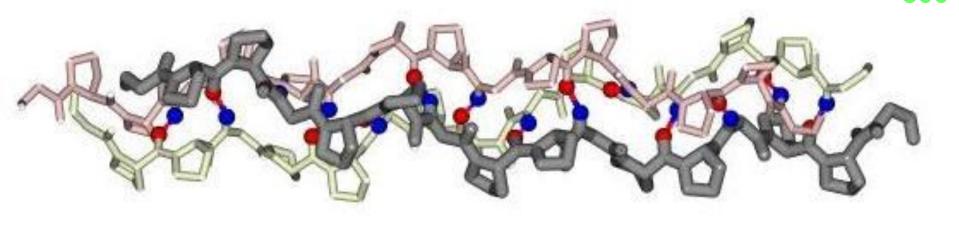


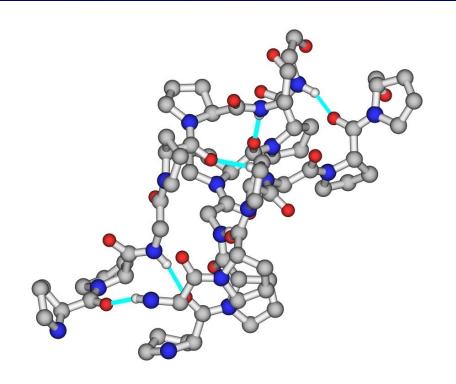


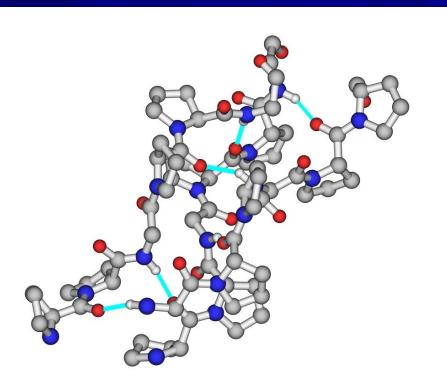


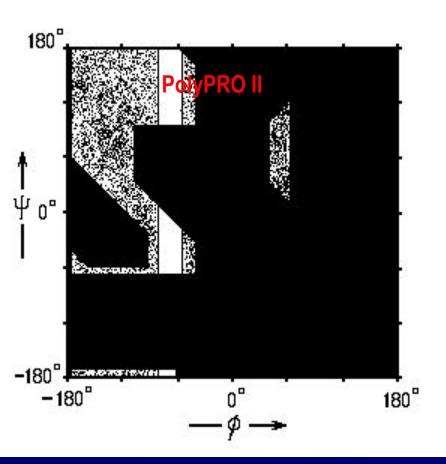


# collagen triple helix: 3 chains ≈ [Gly-X-Pro]<sub>≈500</sub>

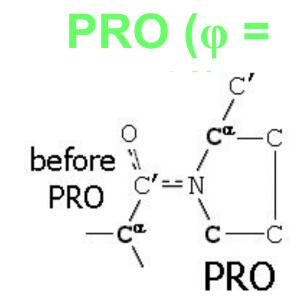


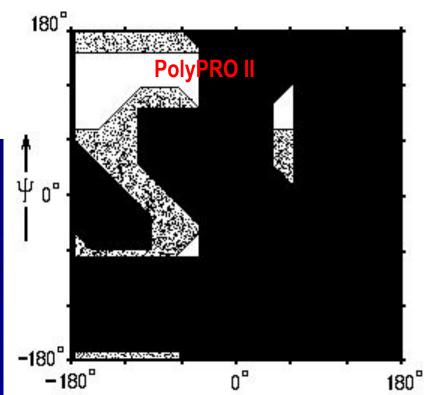




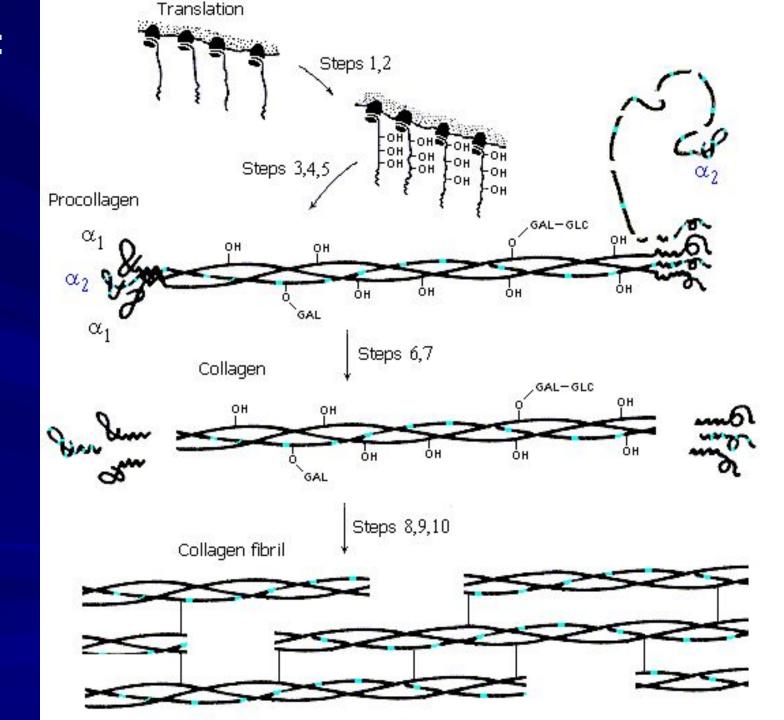


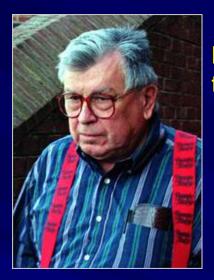
### **Before PRO**





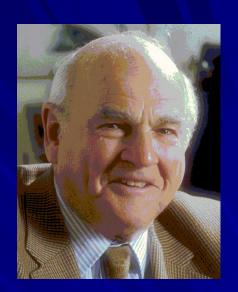
### Collagen: assisted folding

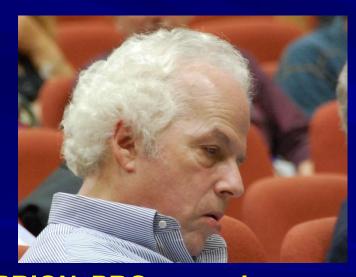




Kuru: a mysterious disease, later demonstrated to be infectious prion disease.

Daniel Carleton **Gajdusek** (1923 –2008) Baruch Samuel **Blumberg** (1925 – 2011) Nobel Prize 1976





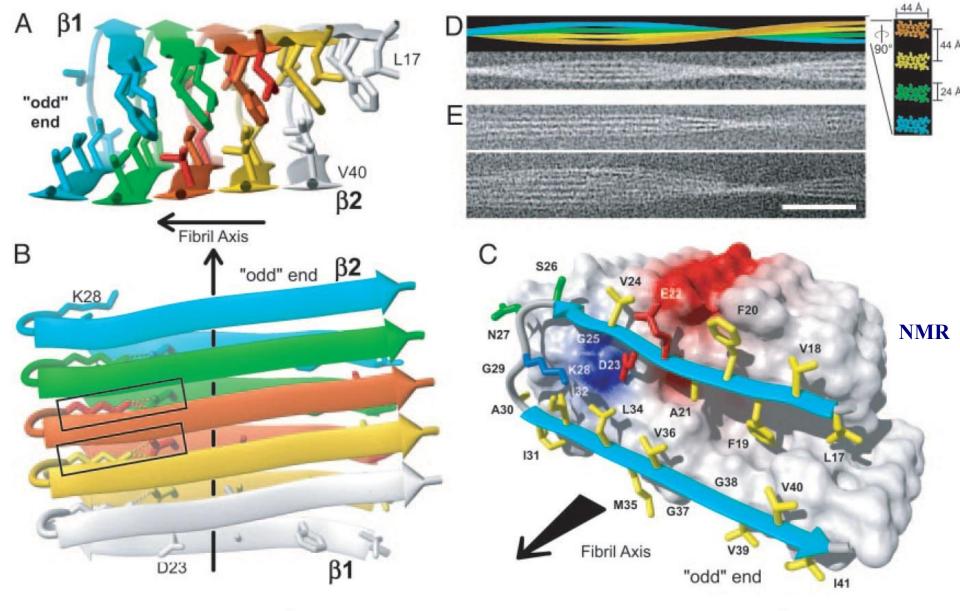
PRION: PROtein and Infection

Stanley Benjamin **Prusiner**, 1942 Nobel Prize 1997



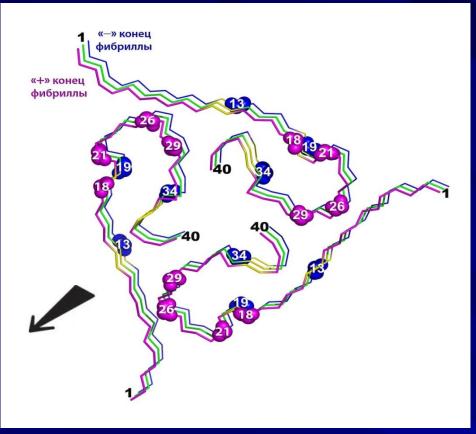
Studies of amyloid formation

Christopher Martin **Dobson**, 1949 Royal Medal 2009



3D structure of Alzheimer's amyloid- $\beta$ (1–42) fibrils

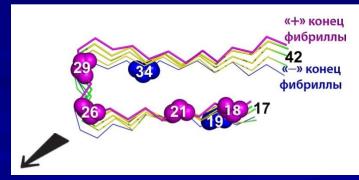
T.Lührs, C.Ritter, M.Adrian, D.Riek-Loher, B.Bohrmann, H.Döbeli, D.Schubert, R.Riek. *PNAS* 102:17342-17347 (2005)

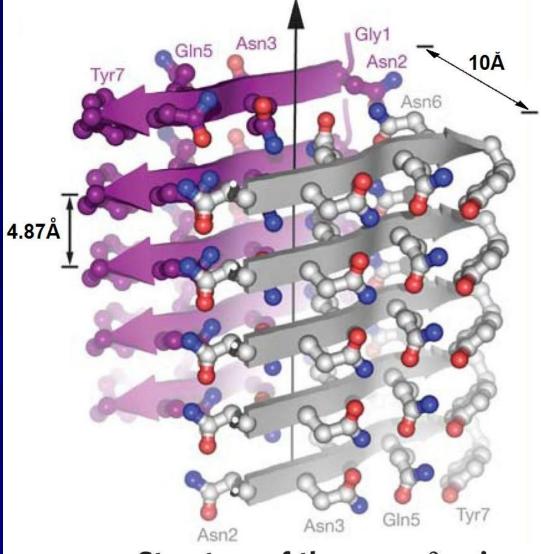


Lu J.X., Qiang W., Yau W.M., Schwieters C.D., Meredith S.C., Tycko R. Molecular structure of β-amyloid fibrils in Alzheimer's disease **brain tissue**. Cell 154:1257-1268 (2013) .

# VARIABILITY OF STRUCTURES

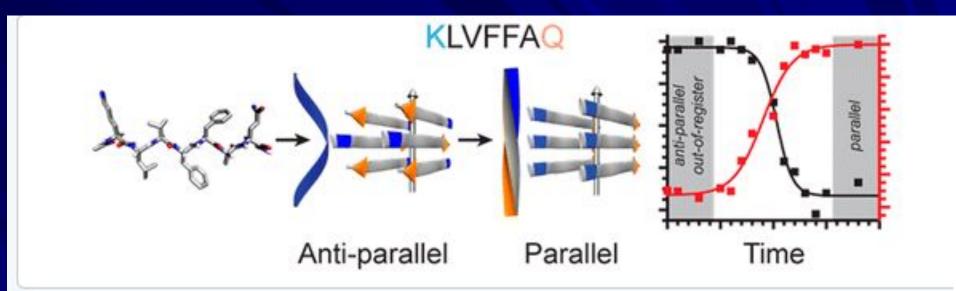
Lührs T., Ritter C., Adrian M., Riek-Loher D., Bohrmann B., Döbeli H., Schubert D., Riek R. 3D structure of Alzheimer's amyloid-beta(1-42) fibrils. PNAS 102:17342-17347 (2005).





# Structure of the cross-β spine of amyloid-like fibrils

R.Nelson, M.R.Sawaya, M.Balbirnie, A.Ø.Madsen, C.Riekel, R.Grothe, D.Eisenberg *Nature* **435**:773-778 (2005) X-RAY



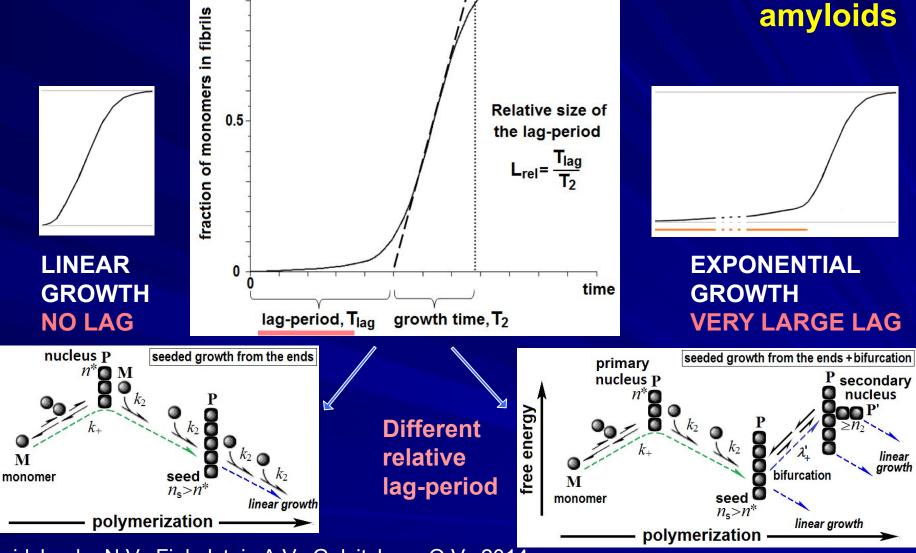
In contrast to an expected Ostwald-like ripening of amyloid assemblies, the nucleating core of the Dutch mutant of the Aβ peptide of Alzheimer's disease assembles through a series of conformational transitions. Structural characterization of the intermediate assemblies by isotope-edited IR and solid-state NMR reveals unexpected strand orientation intermediates and suggests new nucleation mechanisms in a progressive assembly pathway.

JACS→ Volume·136,·Issue·43 → October·29,·2014

Kinetic·Intermediates·in·Amyloid·Assembly

¶

Chen·Liang, Rong·Ni, Jillian·E.·Smith, W.·Seth·Childers, Anil·K.·Mehta, and David·G.·Lynn

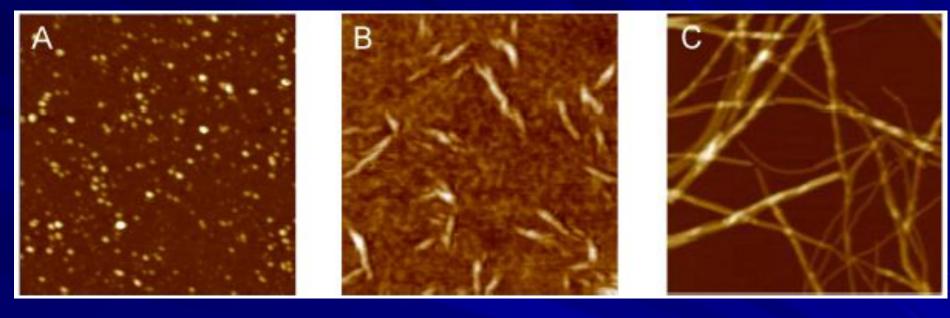


**Growth of** 

Dovidchenko N.V., Finkelstein A.V., Galzitskaya O.V. 2014.

free energy

How to determine the size of folding nuclei of protofibrils from the concentration dependence of the rate and lag-time of aggregation. I. Modeling the amyloid photofibril formation. *J. Phys. Chem. B*, 118:1189-1197.



Oligomers Protofibrils Mature amyloid fibrils

Atomic force microscopy

Relini A., Marano N., Gliozzi A. 2014.

Misfolding of amyloidogenic proteins and their interactions with membranes

Biomolecules, 4, 20-55.

# Natively non-structured fibrous proteins:

# **Elastin:**

Matrix protein.

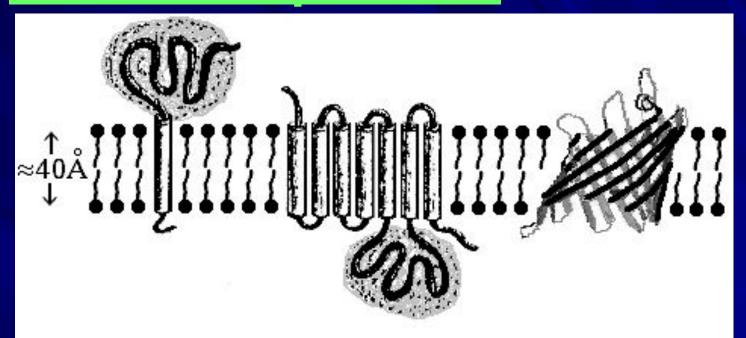
**Short repeats.** 

Poor secondary structure.

Chains are linked by chemically modified Lys residues.

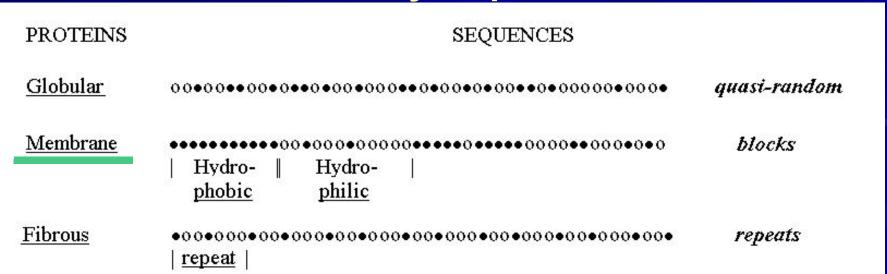
Like in rubber.

### Membrane proteins: transmitters



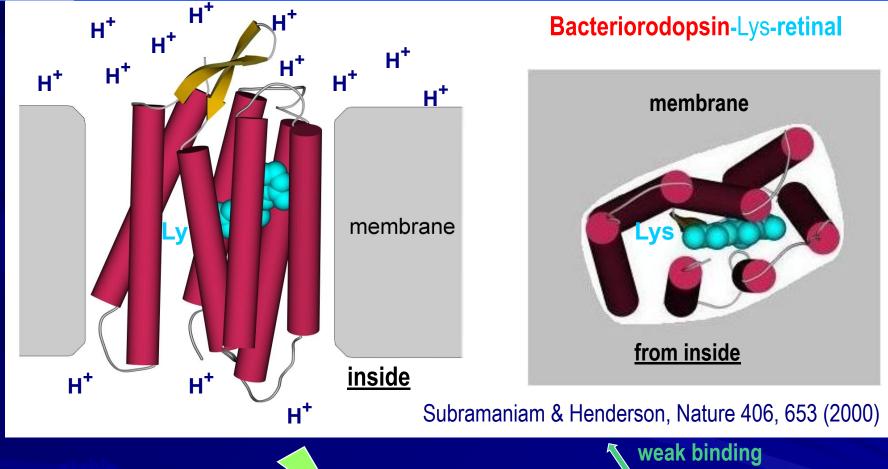
heads (polar) tails tails heads (polar)

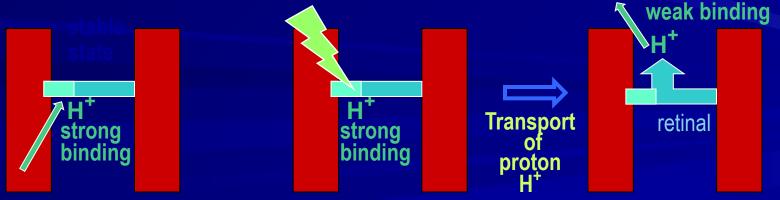
#### H-bonds & hydrophobics

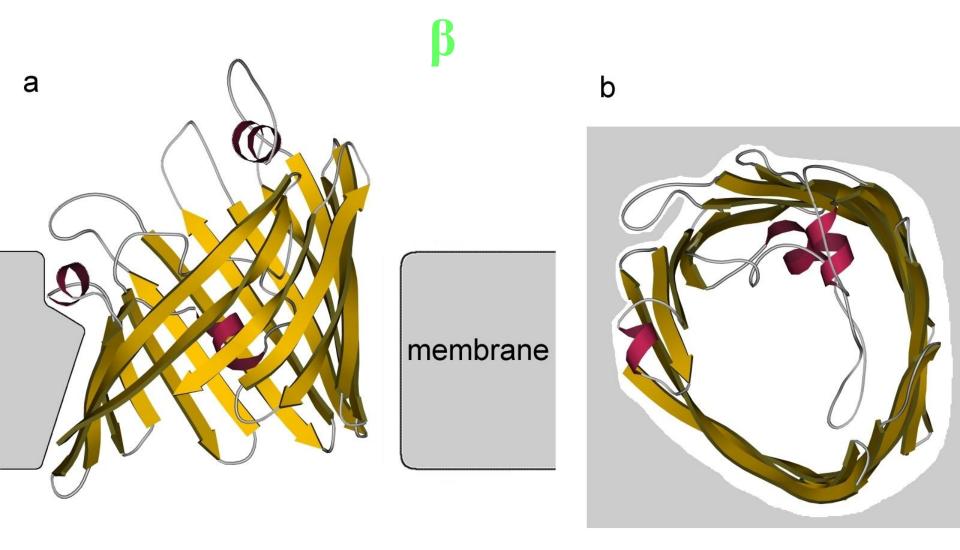


#### **Bacteriorodopsin** (O) with retinal:

the simplest transporter machine with a light-induced conformational change

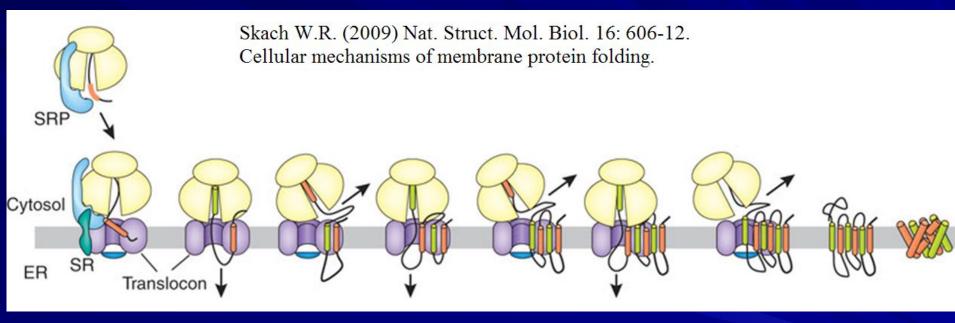




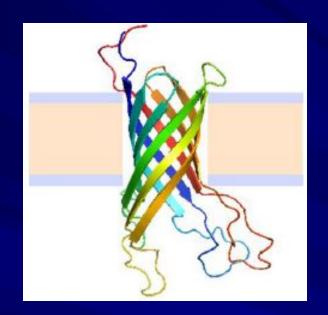


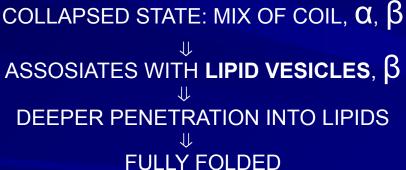
# Porin Transport of polar molecules

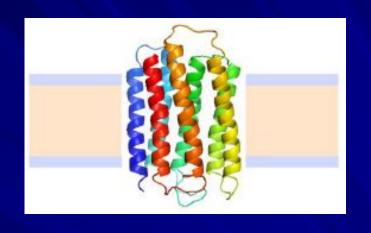
# Membrane protein *in vivo*: Folding is assisted by "directing factors" - chaperones



# MANY OF **SIMPLE** MEMBRANE PROTEINS REFOLD *IN VITRO*IN THE PRESENCE OF PHOSPHOLIPID VESICLES OR SURFACTANT MICELLES

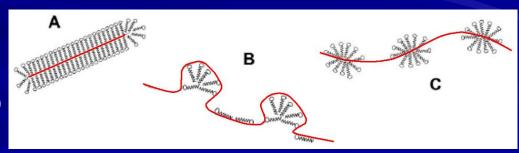


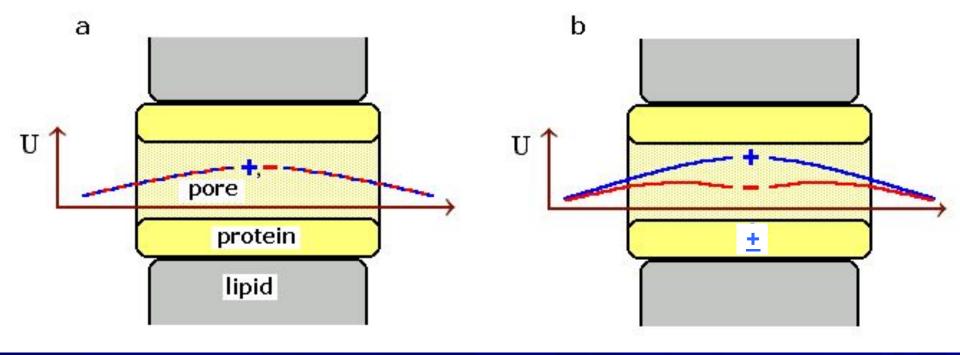




INDEPENDENT  $\alpha$ -HELICES  $\downarrow \downarrow$  ASSEMBLE IN LIPID TO FULLY FOLDED

**DIFFICULT TO STUDY:**DENATURED STATES OF MEMBRANE
PROTEINS ARE **DIVERSE & COMPLICATED** 



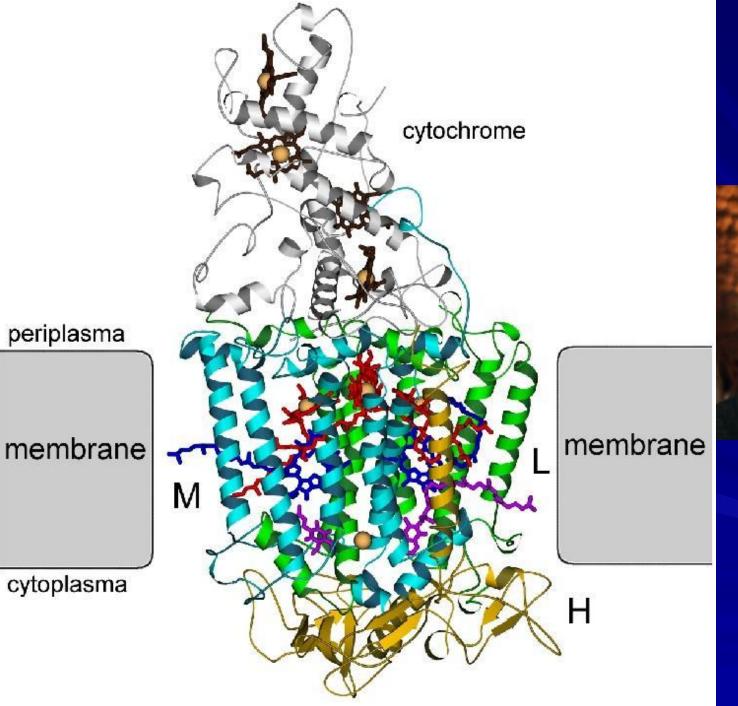


#### **Pore in membrane: SELECTIVITY**

Free energy of a charge in the non-charged non-polar pore:

~ 
$$q^2$$
 / [( $\epsilon_{MEMBR}$   $\epsilon_{WATER}$ )<sup>1/2</sup>  $\Gamma_{PORE}$ ] ~

~ 20 kcal/mol / r<sub>PORE</sub>(Å)



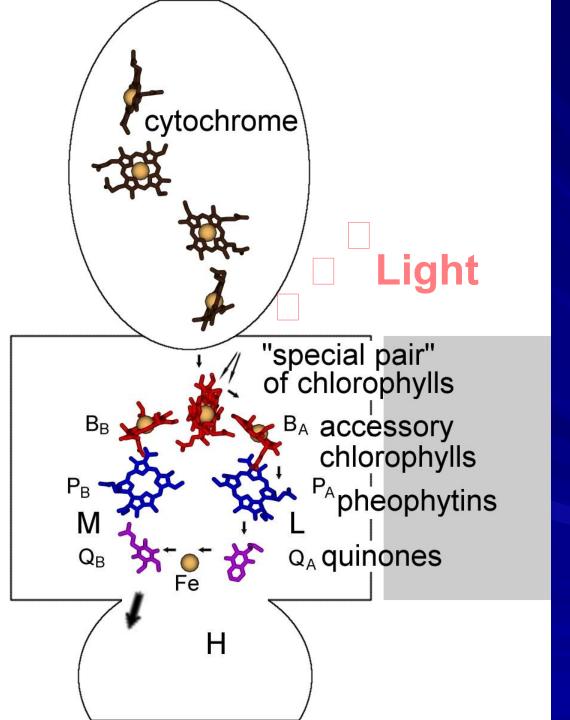
periplasma

cytoplasma

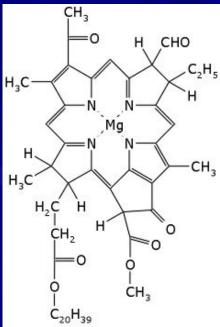
### Photosynthetic center



Robert Huber, 1937. Nobel prize 1988



# Pigments in photosynthetic center: Electron transfer



#### chlorophyll

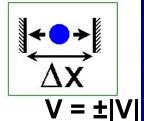
### **Tunneling**



### **Atom** ≈ 1Å ⇒ Attenuation of electron density: P(X) ~ 10<sup>-X(Å)</sup>

Heisenberg's uncertainty:

$$\Delta(mv)\cdot\Delta x \triangleq \hbar \frac{Planck's}{const}$$



Energy of localization in  $\Delta x$ :

$$E = mv^2/2 \sim (\hbar^2/m)/(\Delta x^2)$$

**DELOCALIZATION LEADS TO** MORE STABLE STATE OF C

**T-independent** Frequency of vibrations (attacks):  $V = \pm |V| f \sim 10^{15}/sec$ Successful attacks:  $f_{SUCCS.}(x) \sim P(x) \cdot f$ , e.g.:  $f_{SUCCS.}(5A) \sim 10^{-5+15} \sim$ 

 $\sim 10^{10}/\text{sec}$