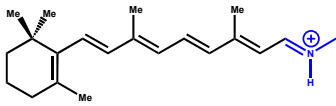


Same chromophore - different absorption maximum



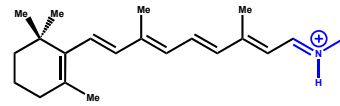
Length of the conjugation

Electrostatic interaction with the binding pocket

Steric interaction with the protein

Protonation / deprotonation

Same chromophore - different absorption maximum



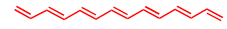
Length of the conjugation

Shorter chain (blue shift)

Longer chain (red shift)



Short wavelength



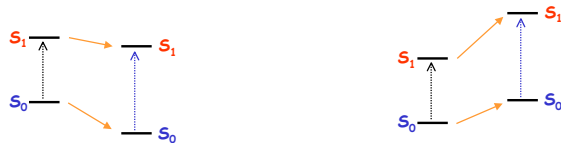
Longer wavelength

Same chromophore - different absorption maximum

Electrostatic interaction with the binding pocket

BLUE SHIFT (shorter wavelengths):

- Ground state stabilization
- Excited state destabilization



Same chromophore - different absorption maximum

Electrostatic interaction with the binding pocket

RED SHIFT (longer wavelengths):

- Ground state destabilization
- Excited state stabilization



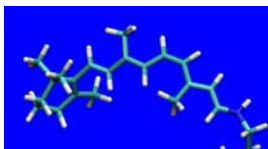
Very important

Same chromophore - different absorption maximum

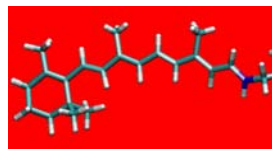
Steric interaction with the protein

Twisting the bonds in the backbone (blue shift)

Straightening the backbone (red shift)



Rhodopsin (500 nm)



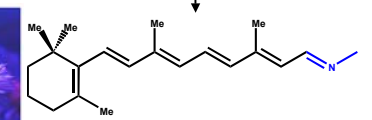
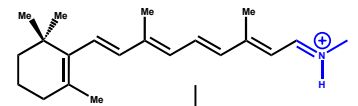
Bacteriorhodopsin (568 nm)

Same chromophore - different absorption maximum

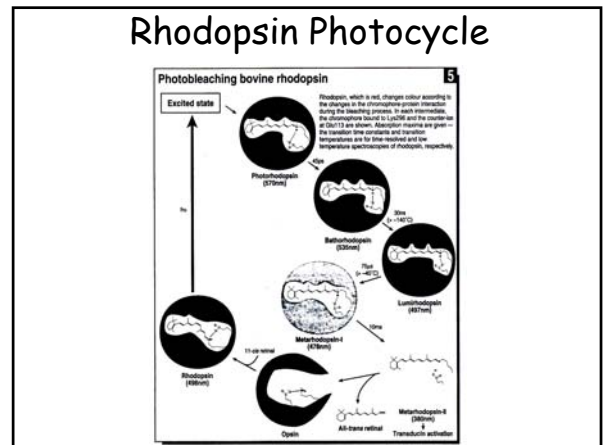
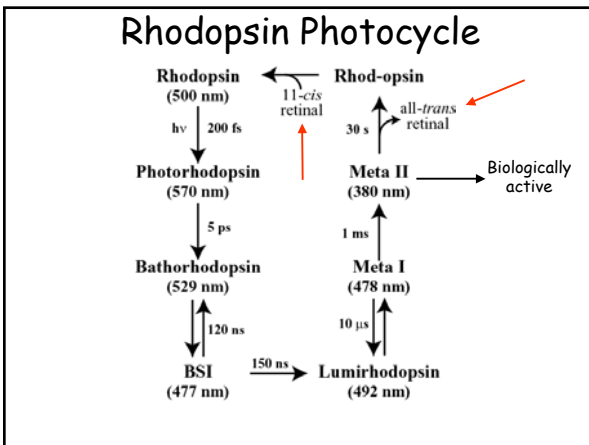
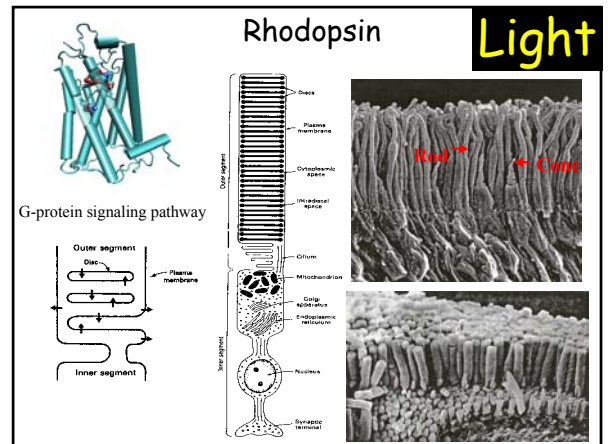
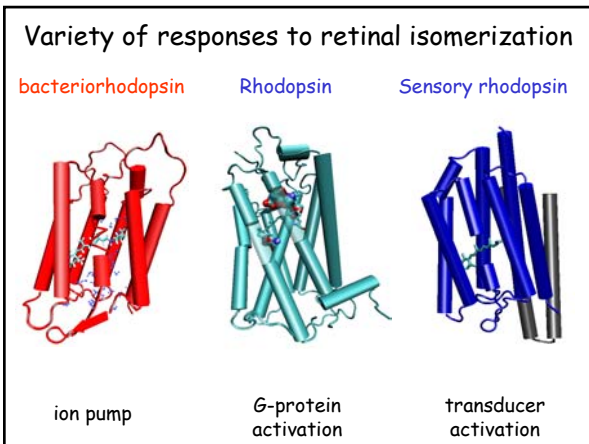
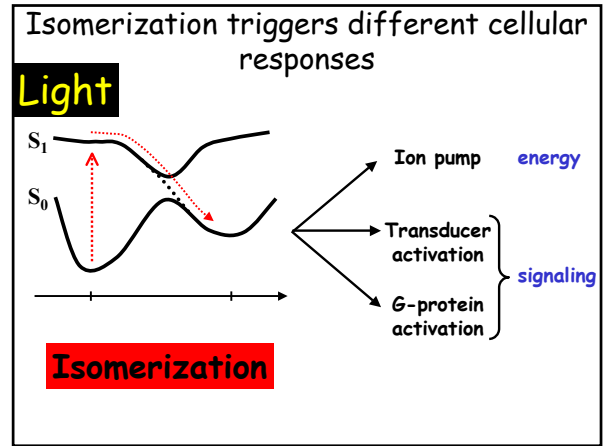
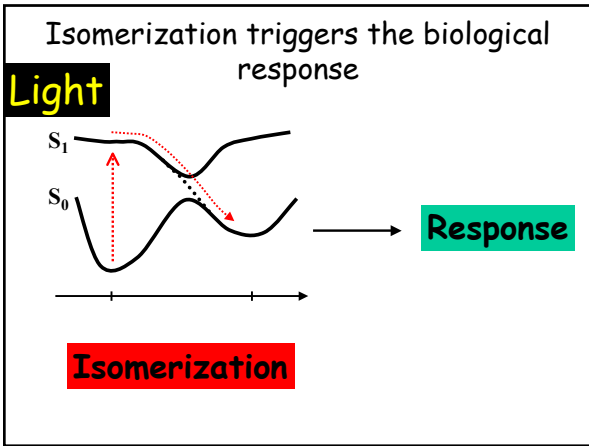
Protonation / deprotonation



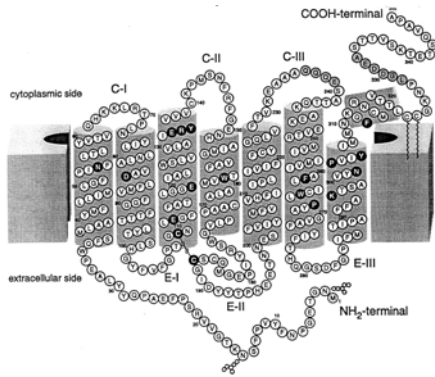
UV vision birds, honeybee



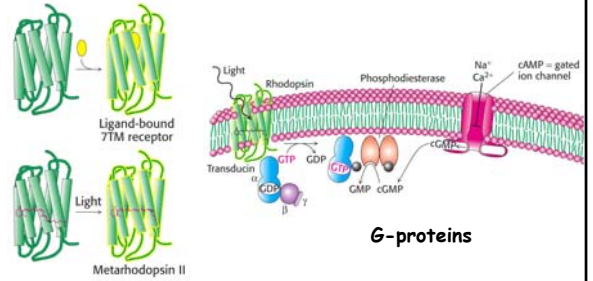
Strong blue shift



Rhodopsin Architecture



Signaling pathway of rhodopsin



Rhodopsin: $D = 0.4 \mu\text{m}^2 \cdot \text{s}^{-1}$ (effective signaling)

Lipid molecules: $D = 1 \mu\text{m}^2 \cdot \text{s}^{-1}$

G-protein coupled receptors

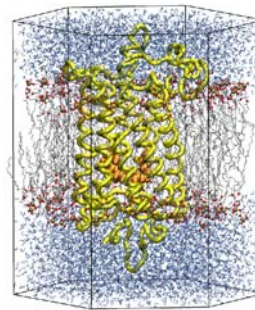
More than 1000 types of receptor specific for:

- odorants
- Photon
- Neurotransmitter
- Hormones
- Calcium
- Peptides
- ...

50% of drugs in the market act on GPCRs, which are very difficult to crystallize ☹

Importance of rhodopsin structure

Simulation of rhodopsin in membrane



Protein: pdb file, 1HZX

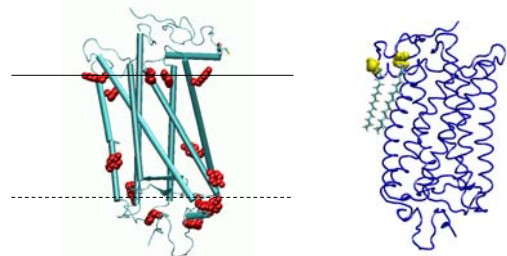
Lipids: POPC bilayer (artificial)

Total: ~40,000 atoms

NAMD2, CHARMM27, PME

1 ns equilibration, 10 ns relaxation after isomerization

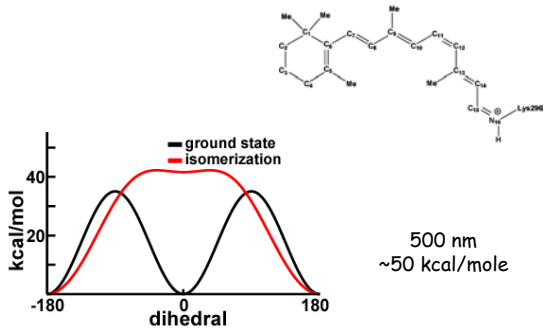
Embedding the protein in membrane



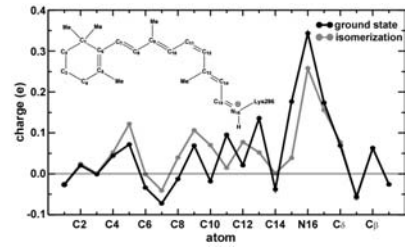
1. Tyrosine residues

2. Palmitoyls connected to Cys322 and Cys323

Inducing isomerization

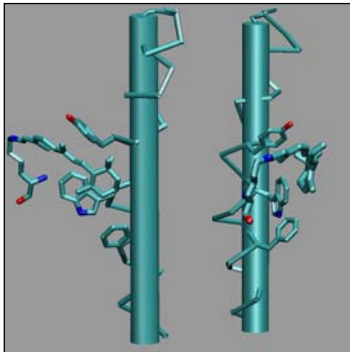


Retinal Charge Distribution

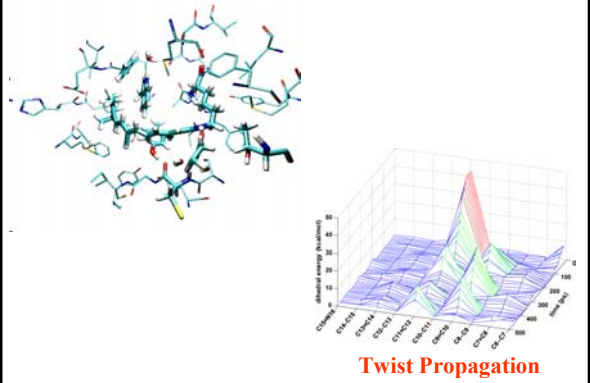


QM/MM derived partial atomic charges

In the ground state, the ring of the chromophore blocks the rotation of helix VI

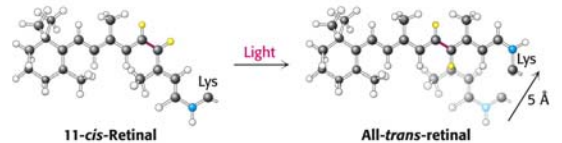
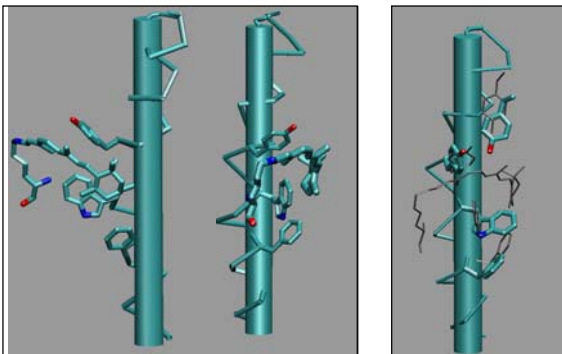


Retinal Isomerization



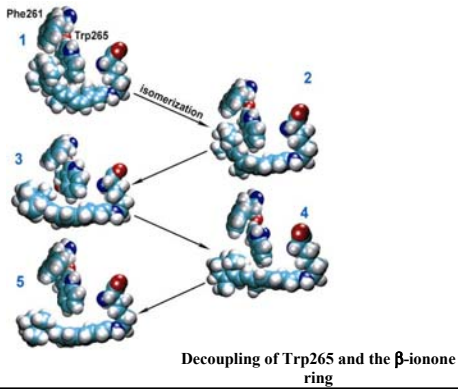
Twist Propagation

In the ground state, the ring of the chromophore blocks the rotation of helix VI



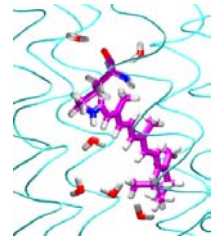
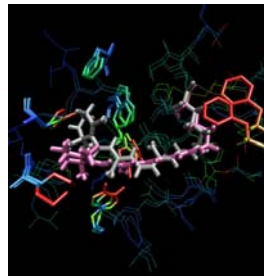
Text book figures not quite right!

Retinal Isomerization



Rhodopsin

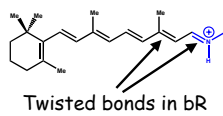
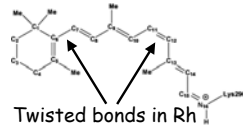
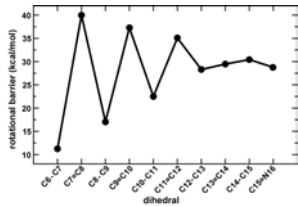
Bacteriorhodopsin



Isomerization 11-cis \rightarrow all-trans

Isomerization: all-trans \rightarrow 13-cis

Isomerization Barriers and bond selectivity



Counterion and bond selectivity

