

Visual transduction

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References



- Webvision: The Organization of the Retina and Visual System (https://www.ncbi.nlm.nih.gov/books/NBK52768/)
- The Molecular Design of Visual Transduction (https://www.biophysics.org/Portals/0/BPSAssets/ Articles/Phototransduction.pdf)
- Adaptation of Rod Photoreceptors to Light and Dark (http://photobiology.info/Rozanowska2.html)

Lecture outline



- Visual transduction (dim vs. bright light)
 - Components (cells and molecules)
 - Mechanisms of activation, amplification, and termination
- Color blindness

Basics of human vision







More on rod cells

- Rod cells consist of four regions: The inner segment, the cell body (contains the cellular organelles), <u>and the synaptic region</u>.
- 2. The outer segment contains the biochemical machinery needed for visual transduction. The components of the phototransduction enzyme cascade are packed into stacks of membranous vesicles ("disks").



hodopsi

outer seament

The dark current



- Most neurons maintain a resting membrane potential (-60 to -70 mV). When excited, they open cation channels causing depolarization and opening of voltage-gated Ca²⁺ channels at the synapse. Ca²⁺ ions flow in and promote fusion of synaptic vesicles, which release neurotransmitter.
- Rods and cones work "backwards". At rest (in darkness), rods and cones are depolarized to -35 to -45 mV.





Generation of vision signals

The players



- Rhodopsin (opsin + pigment molecule)
- Transducin
- Phosphodiesterase
- Na⁺-gated channels
- Regulatory proteins

Rhodopsin

Opsin is a single polypeptide chain with seven helical segments that span the membrane.



Opsin

11-cis retinal

The chromophore (11-cis-retinal)





Light absorption by rhodopsin





Rhodopsin intermediates



- By itself, 11-cis retinal absorbs near UV light. But opsin perturbs the distribution of the electrons exciting its electrons with less energy (i.e., longer wavelength light).
- •The chromophore converts the energy of a photon into a conformational change in protein structure.
- •Rearrangements in the surrounding opsin protein convert it into the active R* state, an intermediate known as metarhodopsin II.



Transducin → Phosphodiesterase (PDE)





G proteins are heterotrimeric, consisting of α , β , and γ subunits. In its inactive state, transducin's α subunit has a GDP bound to it.

Activation of phosphodiesterase





- PDE is a heterotetramer that consists of a dimer of two catalytic subunits, α and β subunits, each with an active site inhibited by a PDE γ subunit.
- The activated transducin α subunit-GTP binds to PDE γ and relieves the inhibition on a catalytic subunit.

Guanylate Cystase

PDE 5



GMP



cGMP

cGMP-gated channels





Levels of calcium ions are reduced, too.





When the channels close, Ca²⁺ ceases to enter, but extrusion through the exchanger continues, so intracellular [Ca²⁺] falls.



Creating an image

- The large potential difference travels as an electrical impulse down the rod cell to the synaptic terminal and is then transferred to an adjoining nerve cell.
- The nerve cell carries this impulse all the way to the brain.
- The brain then determines where the nerve impulse originated and interprets the image.









Signal amplification



Facilitation of transduction





- 1. 2-dimensional surface
- 2. Low in cholesterol and high content of unsaturated fatty acids
- <u>Cooperativity of binding</u>: The binding of one cGMP enhances additional cGMP binding and channel opening (n = ~3)
- 4. Since multiple cGMP molecules are required to open the channel, it will close when only one or two cGMP molecules leave the channel, making it easily to shut down by absorption of light.

Overall, a single photon closes about 200 channels and thereby prevents the entry of about million Na+ ions into the rod cells.





Mechanism I Unstable all-trans rhodopsin complex





Mechanism II Arrestin binding



Arrestin



Mechanism II (cont.) GRK1 and recoverin



- GRK1 is more active at low [Ca²⁺]. Why?
- In the dark, Ca²⁺ ions bind to a protein called recoverin allowing it to anchor to the membrane, bind to GRK1, and inhibit it.
- In contrast to Ca²⁺-free recoverin that does not bind to GRK1, the recoverin-Ca²⁺ complex binds to the N terminus helix of GRK1,



Mechanism III Intrinsic GTPase activity of G protein





- •Transducin has an intrinsic GTPase activity that hydrolyzes GTP to GDP.
- •Upon hydrolysis of GTP to GDP, transducin α subunit releases the PDE γ subunit that re-inhibits the catalytic subunit.
- •Transducin α -GDP eventually combines with transducin $\beta\gamma$

Mechanism IV Facilitation of GTPase activity of G protein



- GTP hydrolysis is slow intrinsically, but it is accelerated by the GAP (GTPase Activating Protein) complex.
 - To ensure that transducin does not shut off before activating PDE, transducin and the GAP complex have a low affinity for each other, until transducin α-GTP binds PDEγ.



The inhibition of the Gα subunit by GTP hydrolysis and, hence, dissociation from PDE is the rate limiting step in the recovery of rod response to light.

Mechanism V Guanylate cyclase

- In the dark, guanylate cyclase-activating proteins (GCAPs) bind Ca²⁺ blocking their activation of guanylate cyclase.
- A decrease in intracellular [Ca²⁺] causes Ca²⁺ to dissociate from GCAPs leading to full activation of guanylate cyclase subunits, and an increase in the rate of cGMP synthesis.



Mechanism VI Ca-calmodulin and cGMP-gated channels





- In the dark, Ca²⁺-Calmodulin (CaM) binds the channel and reduces its affinity to cGMP and shuts it down.
- During visual transduction, the decrease in intracellular [Ca²⁺] causes CaM to be released, the affinity towards cGMP increases, and the channel reopens in response to the slightest increase to cGMP.

 Note: Ca²⁺-Calmodulin (CaM) also binds to GRK1 and inhibits it.



Adaptation to light/dark conditions

Arrestin/recoverin/transducin distribution





- In dark, the outer segment contains high levels of transducin and recoverin and low levels of arrestin (low inhibition; ready to be activated).
- In light, it is the opposite (high inhibition; ready to be inactivated).





Color vision



Cone photoreceptor proteins

short-wave cone

Relative absorbance



How different are they?





•Cone opsins have similar structures as rhodopsin, but with different amino acid residues surrounding the bound 11-cis retinal; thus, they cause the chromophore's absorption to different wavelengths.

- •Each of the cone photoreceptors vs. rhodopsin ≈ 40% identical.
- •The blue photoreceptor vs. green and red photoreceptors = $\approx 40\%$ identical.
- •The green vs. red photoreceptors > 95% identical.

Three important aa residues





A hydroxyl group has been added to each amino acid in the red pigment causing a λ_{max} shift of about 10 nm to longer wavelengths (lower energy).

Rods vs. cones



 Light absorption, number, structure, photoreceptors, chromophores, image sharpness, sensitivity (amplification)



Sharpness and sensitivity of viewing images depends on the brain determining the number and location of the photoreceptor cell(s) that passes an impulse to any given nerve fiber.



Color blindness



Chromosomal locations



- The "blue" opsin gene: chromosome 7
- The "red" and "green" opsin genes: X chromosome
- The X chromosome normally carries a cluster of from 2 to 9 opsin genes.
- Multiple copies of these genes are fine.

Red-green homologous recombination



Between transcribed regions of the gene (inter-genic)



Within transcribed regions of the gene (intra-genic)



Genetic probabilities





Spectral tuning





- The substitutions at positions 277 and 285 account for about 20 nm of the difference in peak sensitivity.
- Serine (S) vs. alanine (A) at position 180 produces a measurable shift in the spectrum.

Pedigree









Red blindness

Green blindness





https://www.buzzfeed.com/crystalro/red-color-vision-test Only People Who Can See RED Really Well Can Read These Words



