

Structure and function of bovine rhodopsin

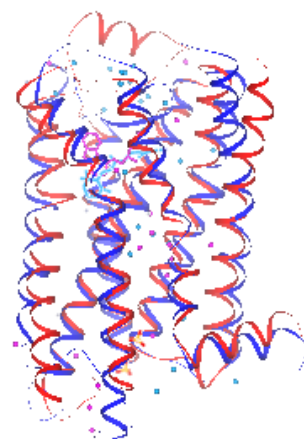
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Photoreceptor proteins of rods and cones in the retina, rhodopsin and color pigments, are the members of heptahelical G protein-coupled receptors (GPCRs). They have a covalently bound inverse agonist, 11-*cis* form of retinal (Vitamin A aldehyde), as the chromophore for capturing photons. The initial process of vision is triggered by *cis-trans* photoisomerization of the retinal protonated Schiff base attached to a lysine side chain in transmembrane helix VII. This reaction can be considered as an instantaneous conversion from an inverse-agonist to an agonist, which subsequently evokes the activation of the protein.

The first crystal structure of rhodopsin was determined at 2.8 Å resolution in 2000.¹ Since then, several new coordinates have been obtained, including the ground state at higher resolution (blue ribbon in the figure),^{2,3} two photoreaction intermediates^{4,5} and a 9-*cis*-analogue pigment.⁶ In addition, some other crystal structures have been reported in different space groups.⁷⁻⁹ These recent advances are summarized in the context of visual function, and the discussion will also include some comparison with the new structure of the human β_2 -adrenergic receptor (red ribbon).¹⁰

X-ray crystallographic and quantum chemical studies of bovine rhodopsin in P4₁ space group have been contributed by the following researchers: (alphabetical order) CA Behnke, AN Bondar, V Buss, M Elstner, P Entel, BA Fox, Y Fujiyoshi, T Hori, T Kumasaka, EM Landau, I Le Trong, M Miyano, H Motoshima, H Nakamichi, J Navarro, K Palczewski, M Schreiber, S Sekharan, Y Shichida, M Silow, RE Stenkamp, M Sugihara, DC Teller, M Yamamoto, O Weingart. Support at the beamlines of SPring-8 is gratefully acknowledged.



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