

FTIR spectroscopy of bacteriorhodopsin microcrystals at Beamline 1.4

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Bacteriorhodopsin (bR) is the sole protein component of the purple membrane of *Halobacterium salinarium*¹. The function of bR *in vivo* is to convert solar energy into a pH gradient across the cell membrane which the organism uses to drive ATP synthesis². Bacteriorhodopsin undergoes a light-induced cycle of physicochemical changes for every proton it pumps out of the cell. The photocycle of bR has been well-characterized by both visible and IR spectroscopy. The major intermediates are identified as the K, L, M, N and O intermediates, and each has a distinct visible color and a distinct IR spectrum³. The Schiff base that connects the side chain of Lys 216 to the retinal molecule buried within the core of the apoprotein⁴ is deprotonated upon formation of the M intermediate, and reprotonated when the M intermediate decays. Since access to the Schiff base switches from the extracellular side of the membrane to the cytoplasmic side between these two proton transfer events, the M intermediate is of particular interest.

High resolution x-ray diffraction experiments on microcrystals of bR have recently become possible, through the discovery by Landau and Rosenbusch that the solubilized protein can be crystallized from the bicontinuous lipid-water gel that is formed by mono-olein⁵. Structural studies on intermediate states of the photocycle thus become a high priority, allowing the visualization of the structural changes that are responsible for converting light energy into a proton-motive force.

Previously we collected high resolution x-ray diffraction data from crystals of wild type bR trapped in both the L and M states by illumination at low temperature, using Fourier Transform IR (FTIR) spectroscopy to confirm the identity of the photointermediate⁶. This year we turned our attention this year to the F219L mutant of bR, which forms N, the intermediate following M, more readily than does the wild type protein⁷. We have succeeded in characterizing two photointermediates of the bacteriorhodopsin mutant F219L by means of FTIR spectroscopy. At 213K an early phase of the M photostate is trapped, whereas at 173K the IR signature strongly suggests that the L photostate is trapped. Further studies of this mutant are planned, with an eye towards trapping the N intermediate.

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