

A BLUE SHIFT OF PROTONATED RETINAL SCHIFF BASE. A MODEL STUDY FOR BACTERIORHODOPSIN

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ABSTRACT: A blue shift was observed in the absorption maxima of protonated retinal Schiff base. The shift is due to electrostatic interaction of a non-conjugated positive charge, located in the vicinity of the ionone moiety, with the polyene chromophore.

Visual pigments consist of a chromophore covalently bound to an apoprotein, opsin. Biochemical extraction studies have shown that in all pigments the chromophore is 11-cis retinal,^{1,2} the parent aldehyde (retinal) of vit A, which is bound to the ε-amino terminal of a lysine residue of the apoprotein opsin via a protonated Schiff base (SBH⁺) linkage.^{3,4} The 11-cis retinal SBH⁺ formed from n-butylamine has its absorption maximum at 440 nm in methanol, whereas the maxima of visual pigments from various sources have maxima as far to the red as 580 nm. These red shifts (in cm⁻¹) from 440 nm, which are due to the effects of the protein environment were called "opsin shift"⁵

The purple membrane is a novel light energy transducing membrane which constitutes part of the plasma of *Halobacterium halobium*. The color of the purple membrane is due to a pigment which has its absorption maxima at 560 nm, and has all-trans retinal as its chromophoric group linked to a lysine⁶ through a protonated Schiff base.

Recently, Nakanishi^{5,7} proposed an external point charge model, which places, in addition to a counter-anion near the Schiff base iminium nitrogen, a second negative charge. In bovine rhodopsin, where the opsin shift is 2730 cm⁻¹, the second negative charge is located in the vicinity of carbons 12-14 of the chromophore. In bacteriorhodopsin (opsin-shift-4870 cm⁻¹), the second negative charge is located in the vicinity of the ionone moiety.

In order to find out if a non-conjugated charge in the vicinity of the ionone ring can affect the absorption maxima of retinyl-iminium polyene, we synthesized compounds 1, 2 and 3.

β-Ionone was brominated with NBS in refluxing CCl₄ for 1 hour and then was reacted at room temperature in THF with dimethylamine for 12 hours, to give amino-keto 4.⁸ Condensation with sodium salt of triethyl phosphonoacetate at 25°C gave amino-ester 5. Reduction with diisobutyl-aluminium hydride followed by oxidation with MnO₂ afforded amino-aldehyde 6.⁹ A second condensation of 6 with sodium salt of diethyl 4-phosphono-3-methyl-crotononitrile gave nitrile

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Table: λ_{\max} /nm of Protonated Schiff Bases

Chromophore	pH	λ_{\max}	Solvent	$\Delta\nu/\text{cm}^{-1}$ ^a
<u>1</u>	8	443	EtOH	
<u>1</u>	2 ^b	430	EtOH	683
<u>1</u>	8	467	CHCl ₃	
<u>1</u>	2 ^b	444	CHCl ₃	1109
<u>2</u>	8	378	EtOH	
<u>2</u>	2 ^b	337	EtOH	3218
<u>2</u>	8	393	CHCl ₃	
<u>2</u>	2 ^b	345	CHCl ₃	3540
<u>3</u>	8	343	CHCl ₃	
<u>3</u>	2 ^b	304	CHCl ₃	3740

^a $\Delta\nu$ between pH 8 and pH 2 in the same solvent.

^b Acidified with HCl.

observed for compound 1 suggests that in the excited state the density of positive charge at carbons 8-10 is larger than at carbon 5. The increasing blue shifts observed in compounds 1, 2 and 3 (by protonation of the amino group), suggest that the density of positive charge at carbon 5 in the excited state increases as the polyene becomes shorter. In a shorter chromophore, the charge is distributed between a smaller number of carbons, resulting in larger density of positive charge at carbon 5.

The spectroscopic information obtained encourages the synthesis of a compound carrying a negative charge at carbon 5, which will allow the evaluation of the external point charge model for bacteriorhodopsin.

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References and Notes

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8. ^1H n.m.r. (CDCl_3) δ 1.05 (6H, s), 1.79 (3H, s), 2.22 (6H, s, $(\text{CH}_3)_2\text{N}$), 2.32 (3H, s), 3.08 (1H, t, J 8 Hz, 4-H), 6.11 (1H, d, J 16.5 Hz, 8-H), 7.25 (1H, d, J 16.5 Hz, 7-H).
9. ^1H n.m.r. (CDCl_3) δ 1.03 (6H, s), 1.75 (3H, s), 2.23 (6H, s, $(\text{CH}_3)_2\text{N}$), 2.32 (3H, s), 3.09 (1H, t, J 8 Hz, 4-H), 5.95 (1H, d, J 8 Hz, 10-H), 6.17 (1H, d, J 16.2 Hz, 8-H), 6.75 (1H, d, J 16.2 Hz, 7-H), 10.14 (1H, d, J 8 Hz, 11-H).
10. ^1H n.m.r. (CDCl_3) δ 1.04 (6H, s), 1.76 (3H, s), 2.02 (3H, s), 2.33 (3H, s), 3.09 (1H, t, J 8 Hz, 4-H), 5.96 (1H, d, J 8 Hz, 14-H), 6.2-6.9 (4H, m, 7-H, 8-H, 10-H, 12-H), 10.11 (1H, d, J 8 Hz, 15-H).
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