

EXCITED-STATE REACTION DYNAMICS OF BACTERIORHODOPSIN STUDIED BY FEMTOSECOND SPECTROSCOPY

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The photodynamics of bacteriorhodopsin were studied by transient absorption and gain measurements after excitation with femtosecond pulses at 620 nm. With probing pulses at longer wavelengths ($\lambda > 770$ nm) the previously reported formation of the J intermediate (with a time constant of 500 ± 100 fs) was confirmed. With probing pulses around 700 nm, a faster process with a relaxation time of 200 ± 70 fs was observed. The data analysis strongly suggests that this kinetic constant describes the reactive motion of the polyatomic molecule on its excited-state potential energy surface, i.e. one observes directly the incipient isomerization of the retinal molecule. The minimum of the S_1 potential energy surface reached in 200 fs lies approximately 13300 cm^{-1} above the ground state of bacteriorhodopsin and from this minimum the intermediate J is formed with a time constant of 500 fs.

1. Introduction

Photosynthetic activity in halobacteria is based on the membrane protein bacteriorhodopsin (BR), which acts as a light-driven proton pump. BR contains only one chromophore, a retinal molecule, which is linked via a protonated Schiff's base to lysine 216 in the primary structure of the protein. In the functionally active light-adapted form of BR the retinal molecule adopts the all-trans configuration. The photochemical reaction of BR has been studied intensively during the past decade [1], but only recently, the primary reaction leading to the first ground-state photoproduct could be resolved [2-5]. Light quanta promote the BR molecule to the first excited electronic state (S_1). From the bottom of the S_1 potential surface the transition to the photoproduct J takes place within 430 fs and a subsequent reaction leads to the intermediate photoproduct K with a time constant of 3 ps [4-6]^{#1}. The primary re-

action is an all-trans to 13-cis [7], presumably 13-cis, 14s-cis photoisomerization [8]. In the current interpretation the reactive motion during the initial steps proceeds on the S_1 potential surface as well as on the ground-state (S_0) surface. Until now, no time-resolved experimental observation of the excited-state reactive motion has been reported.

We apply improved femtosecond time-resolved methods to the study of the excited-state dynamics of bacteriorhodopsin. Transient absorption and gain experiments are described which reveal the motion of the retinal molecule on its excited-state potential energy surface.

2. Material and methods

Bacteriorhodopsin (as purple membranes) was prepared from *Halobacterium halobium* according to the published procedure [9]. Samples had optical densities of 20-30 at $\lambda = 570$ nm and were placed in a rotating cell of 300 μm optical path. The BR sample was kept in the light-adapted form by appropri-

^{#1} In ref. [4] we published a time constant of 5 ps for the J to K transition. Our more recent femtosecond experiments indicate a somewhat shorter time constant of 3 ± 1 ps.

ate background illumination. Time-resolved excite and probe experiments were performed using 80 fs pulses from a colliding pulse mode-locked (CPM) laser/amplifier system operated at a repetition rate of 7 kHz [10,11]. 10% of the output of the laser/amplifier system served as the exciting pulse. The excitation wavelength was $\lambda=620$ nm and the excitation energy 200 nJ. Approximately 10% of the BR molecules within the illuminated area were excited by each pulse. The main part of the energy of the laser output generated a broad continuum in a 1 mm thick ethylene glycol jet [12]. A small fraction of the femtosecond continuum of 10 nm width selected via interference filters gave the probing light pulse. The time resolution of the experiment was determined by the cross-correlation function between exciting and probing pulses. The cross-correlation widths τ_c (fwhm) slightly depended on the selected wavelength of the probing pulses. A typical value was $\tau_c=120$ fs at $\lambda=850$ nm.

Transmission changes of the sample induced by the excitation pulses were recorded as a function of time. In order to guarantee a continuously fresh sample in the experiment the cuvette (8 cm diameter) rotated at 12.5 Hz. In this way each pulse illuminated a new portion of the sample and after one revolution the intermediate photoproducts were

reconverted to the starting condition, i.e. the light-adapted BR.

3. Results

Fig. 1 shows the absorption and fluorescence of light-adapted BR. The excitation and probing wavelengths are indicated by arrows. The absorption band is broad and the excitation wavelength of $\lambda=620$ nm lies on its red wing. At wavelengths $\lambda > 650$ nm residual absorption is smaller than required to explain the absorbance decrease seen in fig. 2 by simply bleaching the ground-state absorption. The fluorescence band is also broad. It peaks around 730 nm and decays slowly towards the near infrared; at 850 nm the fluorescence intensity exhibits 50% of its peak value.

Results of the time-resolved experiments are presented in fig. 2 for four probing wavelengths. The experimental results are shown as circles; the solid curves are calculated using a simplified level scheme, where excitation light populates a molecular state with reduced absorption. This level decays with the time constant $\tau=200$ fs to a further species of increased absorption (figs. 2a and 2b). The curves in figs. 2c and 2d are calculated using a decay time τ of

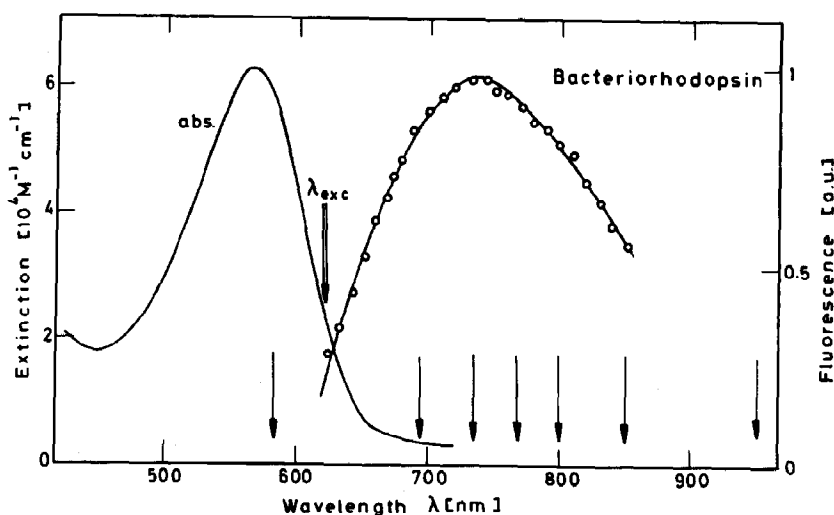


Fig. 1. Absorption and fluorescence emission spectra of light-adapted bacteriorhodopsin. The excitation and probing wavelengths of the femtosecond experiments are indicated by the arrows.

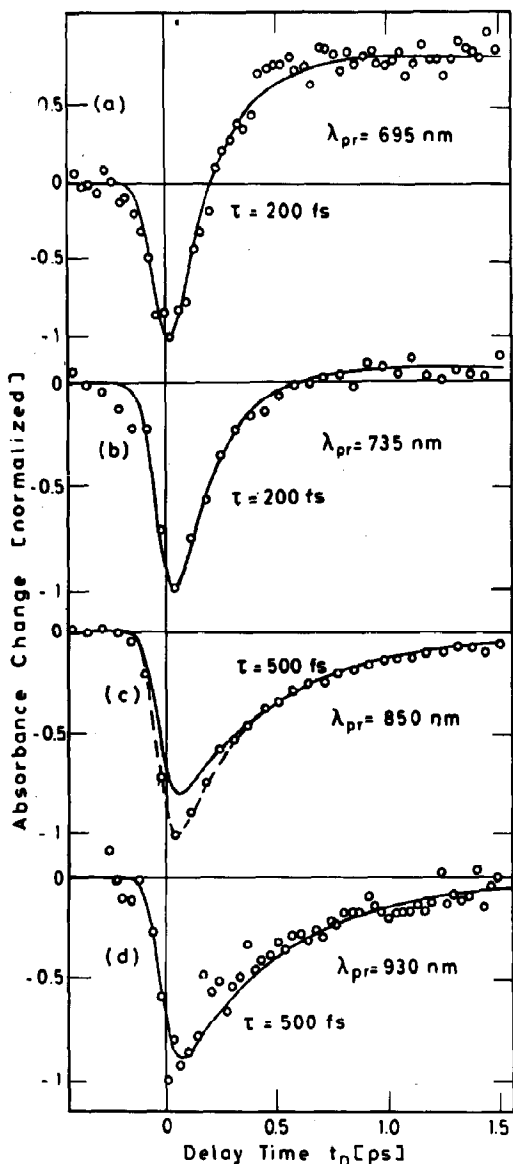


Fig. 2. Time-resolved absorption induced by exciting femtosecond pulses at $\lambda = 620$ nm detected at probing wavelengths $\lambda = 695$ nm (a), 735 nm (b), 850 nm (c) and 930 nm (d). The experimental points are normalized to the peak absorption changes. The solid curves are calculated using the model given in the text with decay times of 200 fs (a,b) or 500 fs (c,d). The broken curve takes into account a double-exponential decay with both time constants.

500 fs and a negligibly increased absorption at later times. Each curve is normalized to the maximal changes, which correspond to transmission changes

of the order of several percent. At the shortest wavelength, $\lambda = 695$ nm (fig. 2a), a rapid reduction of the absorption to negative values around time zero turns with a time constant of 200 fs to a level of enhanced absorption. Because BR has no appreciable absorption at $\lambda = 695$ nm (see fig. 1), the observed negative absorption at short delay times must be due to light gain by the excited BR molecule. At a probing wavelength of $\lambda = 735$ nm (as well as at 695 nm) (fig. 2b) the early gain disappears with a time constant of 200 fs and a small increased absorption due to the product states appears concomitantly. At 850 nm (fig. 2c) a gain decay kinetics with $\tau = 500$ fs is measured (solid curve). A small contribution of the 200 fs process explains the differences between the experimental data points and the calculated (solid) curve. A further red-shift of the probing light pulse to 930 nm makes the 200 fs process undetectable and the gain curve can be fitted by a single 500 fs time constant (fig. 2d). Measurements were taken at $\lambda = 770$ and 800 nm but are not shown in fig. 2. At 770 nm a double-exponential decay is composed of contributions from the 200 and 500 fs kinetics with equal amplitudes. The data at 800 nm can be fitted by the same time constants with an amplitude contribution of the 200 fs process of approximately 25%.

When the same experiments were performed in the absorption region of BR, e.g. at 585 nm, an absorbance decrease recovered in 500 ± 100 fs. No distinct contribution from a faster component was detected. This result is in agreement with our previous observation at $\lambda = 620$ nm of a 430 ± 50 fs decay time [2].

The observed absorption changes are related to population changes of the ground state S_0 and the excited state S_1 of BR. Considering the momentary population densities N_0 of S_0 and N_1 of S_1 , three processes are of interest here: (i) absorption from S_0 proportional to N_0 ; (ii) excited-state absorption from S_1 proportional to N_1 ; (iii) stimulated emission (amplification or gain) from S_1 proportional to $N_1 - N_0$.

4. Discussion

4.1. General consideration

Before presenting a model for the primary reac-

tion in BR we summarize our experimental observations. We find: (i) negative absorption (amplification or gain) with a relaxation time of 200 fs for probing wavelengths between 695 and 850 nm (energy between 14400 and 11800 cm^{-1}); (ii) negative absorption with a lifetime of 500 fs for probing wavelengths $\lambda > 735$ nm (the onset of the 500 fs component is between 735 and 770 nm, i.e. the high-frequency limit is between 13600 and 13000 cm^{-1}); (iii) two time constants of 200 and 500 fs in the overlapping wavelength range of (i) and (ii).

The following interpretation takes advantage of current knowledge on fast processes on excited electronic surfaces. The potential energy surfaces of the retinal chromophore of bacteriorhodopsin for the ground state S_0 and for the first excited state S_1 are shown as a function of reaction coordinate in fig. 3. The latter is related to the isomerization of the retinal molecule.

During excitation with the pump pulse the retinal chromophore is excited to Franck–Condon states on the S_1 potential surface. These states are rapidly depopulated with the time constant τ_1 by reactive motion along the S_1 potential surface. As a result of this

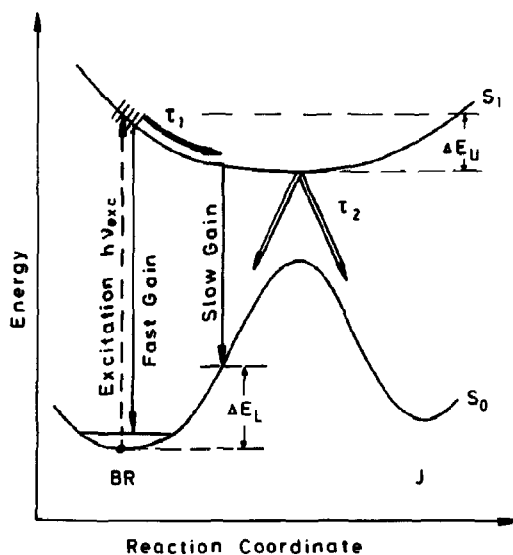


Fig. 3. Model of the ground-state (S_0) and excited (S_1) energy surfaces as a function of the reaction coordinate. The first motion leads the system with the time constant of 200 fs to the bottom of the S_1 potential surface. The reaction proceeds to the ground-state photoproduct J or the initial state of BR with the time constant of 500 fs.

fast motion the gain from the Franck–Condon states (fig. 3) is very short-lived, $\tau_1 = 200$ fs. The relaxation from the minimum of the S_1 potential surface has a longer time constant, $\tau_2 = 500$ fs, and is seen as a slow gain at smaller energies ($\lambda > 770$ nm).

4.2. The excitation process

The excitation follows the Franck–Condon principle, i.e. the optical transition proceeds vertically without changing the nuclear coordinates of the molecule. The broad absorption spectrum (see fig. 1) and the steep slope of the S_1 potential surface (see below) suggests an inhomogeneously broadened transition (a wide S_0 potential minimum). Light at a certain wavelength is absorbed by a number of ground-state configurations and a hole is burned in the corresponding population. This hole evolves in time with the thermal motion of the retinal molecules. The hole vanishes within a period of a low-lying vibrational level within 50–100 fs. On the other hand, the molecules on the S_1 potential surface are – immediately after excitation – in the Franck–Condon states, depicted schematically by the shaded area in fig. 3. At that point various normal modes are displaced from their equilibrium position on the S_1 potential surface. One can understand this state as a coherent superposition of different normal modes (a wave packet) [13]. The energy of this Franck–Condon state is higher than the bottom (equilibrium position) of the S_1 surface by ΔE_u . Experimentally, at wavelengths close to the excitation wavelength ($\lambda_{pr} \approx \lambda_{exc}$) one observes both the depletion (hole) in the ground-state population N_0 and the change in the excited-state population N_1 . Using shorter probing wavelengths, $\lambda_{pr} \ll \lambda_{exc}$, the filling of the hole in the ground-state population N_0 and possible excited-state absorption (N_1) from the Franck–Condon state are seen. At longer wavelengths, $\lambda_{pr} \gg \lambda_{exc}$, gain and excited-state absorption from the Franck–Condon state are observed (both proportional to N_1).

4.3. Motions on the S_1 potential surface

The evolution of the system after excitation may be described as follows. The initial wave packet consists of different low- to medium-frequency vibra-

tional modes. The excited modes move the molecule along the reaction coordinate. No energy exchange of the retinylidene moiety, e.g. by collision with solutes, is possible within the protein matrix and thus the process proceeds undisturbed. The wave packet spreads out due to the rapid dephasing processes in the excited electronic state [14] and the energy stored in the Franck-Condon states of the molecules distributes over many degrees of freedom of the large retinal molecule. At the minimum of the potential energy surface most of the energy ΔE_u is quasi-thermalized. The system stays within an energy kT^* at the bottom of the S_1 state. A transient temperature $T^* = 400$ K is estimated from the relaxed energy ΔE_u [15].

A simple classical treatment allows us to estimate the reaction time from the Franck-Condon state to the relaxed state. We assume that an energy of $\Delta E_u \approx 2500 \text{ cm}^{-1}$ (see below) is available for the reactive motion. The initial reaction is believed to be a 13-cis double-bond rotation or a 13-cis, 14s-cis isomerization [7,8,16,17]. The initial rotation in both cases requires (for rotation angles up to 60°) a mass transfer which is equivalent to the movement of a group of atoms (consisting of one carbon, one nitrogen and several hydrogens) by less than 1.5 \AA . The time t necessary to move the atoms over a distance x is estimated to be $t = (2mx^2/\Delta E_u)^{1/2}$ ^{#2}. In the present case one calculates $t = 200$ fs. The exact agreement with the experimental short relaxation time must be considered fortuitous on account of the uncertainty of the different factors entering the estimate of t .

4.4. The relaxed S_1 state

At the minimum of the S_1 surface the system has access to a wide range of the reaction coordinate via thermal motions. Three major processes start from this relaxed state: Light emission or gain in a broad frequency range and internal conversion leading to the ground-state surfaces of either the initial state or the product state J [18].

We believe that the longer time constant of 500 fs is related to the decay of this relaxed state. This interpretation is supported by the fact that the slow

gain is at longer wavelengths than the fast gain (fig. 3). In addition, we note that the 500 fs process is the only slow gain process observed. Information on the S_1 potential surface is obtained from the following energy considerations:

$$h\nu_{\text{exc}} = E_g + \Delta E_L + (\Delta E_u - kT^*) .$$

Here E_g is the high-frequency limit of the slow gain, $E_g = 13000 \text{ cm}^{-1}$ (770 nm) and ΔE_L marks the energy of the terminal state on the S_0 potential surface of the gain at 13000 cm^{-1} . An upper limit of the energy ΔE_u (dissipated during the S_1 motion) may be calculated from the equation given above with the excitation energy $h\nu_{\text{exc}} = 16100 \text{ cm}^{-1}$ and a lower limit for ΔE_L of several $kT \approx 500 \text{ cm}^{-1}$. With these numbers we estimate an upper limit of $\Delta E_u \leq 2800 \text{ cm}^{-1}$.

A final remark should be added to the time-resolved experiments in the absorption region of BR. In these experiments [2] BR was excited and probed at the same wavelength $\lambda = 620$ nm and no fast component was seen. The reduced time resolution ($t_p = 150$ fs) and the coherent artifact around time zero presumably prevented the detection of the faster component.

Interesting femtosecond experiments with improved time resolution have been performed by Mathies and Shank in the absorption region of BR around 620 nm [19]. Their time-resolved spectra are caused by transient population changes in the excited electronic state as well as in the ground state.

5. Conclusion

We followed the fast photochemical reaction of bacteriorhodopsin on the excited-state potential surface and observed the motion out of the initially populated Franck-Condon region proceeding in 200 fs. Subsequently, the system reaches a relaxed state at the minimum of the S_1 potential surface which decays in 500 fs. The experimental results indicate that only a small amount of energy $\Delta E_u < 2800 \text{ cm}^{-1}$ is available for the excited-state reactive process. This finding places the bottom of the S_1 potential state at least 13300 cm^{-1} above the ground state of the light-adapted bacteriorhodopsin.

^{#2} A linear dependence of the potential energy $E(x)$ was assumed.

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