

Electronic pathway in reaction centers from *Rhodobacter sphaeroides* and *Chloroflexus aurantiacus*

Michal Pudlak · Richard Pincak

Received: 28 June 2008 / Accepted: 9 December 2009 /
Published online: 7 January 2010
© Springer Science+Business Media B.V. 2009

Abstract The reaction centers (RC) of *Chloroflexus aurantiacus* and *Rhodobacter sphaeroides* *H(M182)L* mutant were investigated. Prediction for electron transfer (ET) at very low temperatures was also performed. To describe the kinetics of the *C. aurantiacus* RCs, the incoherent model of electron transfer was used. It was shown that the asymmetry in electronic coupling parameters must be included to explain the experiments. For the description of *R. sphaeroides* *H(M182)L* mutant RCs, the coherent and incoherent models of electron transfer were used. These two models are discussed with regard to the observed electron transfer kinetics. It seems likely that the electron transfer asymmetry in *R. sphaeroides* RCs is caused mainly by the asymmetry in the free energy levels of *L*- and *M*-side cofactors. In the case of *C. aurantiacus* RCs, the unidirectionality of the charge separation can be caused mainly by the difference in the electronic coupling parameters in two branches.

Keywords Photosynthetic bacterial reaction centers · Primary charge separation · Electron transfer · Asymmetry in electron transfer · Quantum yields · Photosynthesis · Rate constants · Charge-separating reactions · Solar energy

M. Pudlak · R. Pincak (✉)
Institute of Experimental Physics, Slovak Academy of Sciences,
Watsonova 47,043 53 Kosice, Slovak Republic
e-mail: pincak@saske.sk

M. Pudlak
e-mail: pudlak@saske.sk

R. Pincak
Joint Institute for Nuclear Research, BLTP, 141980 Dubna,
Moscow Region, Russia

1 Introduction

Photosynthesis is a reaction in which light energy is converted into chemical energy. The primary process of photosynthesis is carried out by a pigment-protein complex embedded in the membrane, that is, the photosynthetic reaction center (RC). In photosynthetic purple bacteria, the cyclic electron transfer reaction is performed by the RC and two other components: the cytochrome (Cyt) bc_1 complex and the soluble electron carrier protein. The RC is a special pigment-protein complex that functions as a photochemical trap. The precise details of the charge separation reactions and subsequent dark electron transport (ET) form the central question of the conversion of solar energy into the usable chemical energy of a photosynthetic organism. The function of the RC is to convert solar energy into biochemically usable energy. Therefore, we wish to understand which features of the RC are responsible for the rate constants of these reactions.

Insight into the molecular organization of the RC has been derived, initially, from spectroscopic studies and, subsequently, from the development and analysis of high-resolution crystal structures of several photosynthetic organisms. The first RC structurally resolved (3 Å) was of the purple bacterial RC from *Rhodospseudomonas viridis* [1]. This was soon followed by the elucidation of several other purple bacterial structures. Good progress is also being made toward achieving two- and three-dimensional structures of photosystem II (PSII) crystals. It is surprising that the structures of all the different RCs show a dimeric core with a pseudo- C_2 axis of symmetry.

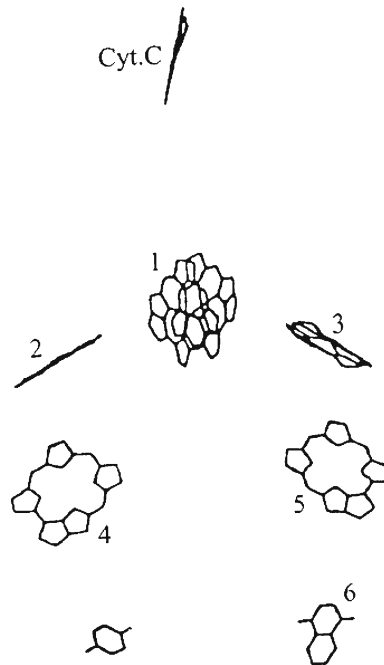
A remarkable aspect of the RC structures is the occurrence of two almost identical electron acceptor pathways arranged along the C_2 axis relative to the primary charge-separating dimer (bacterio) chlorophyll (Fig. 1). This finding posed a key question: Does electron transfer involve both branches? In the purple bacterial RC, only one branch is active, although the inactive branch can be forced into operation with modification of amino acid side chains on the active branch [2]. Moreover, the charge-separating electron transfer reactions occur with a remarkably high quantum yield of 96%, where, from two possible symmetric branches, only the branch L is active in the electron transfer. This efficiency relies on the rates of the charge-separating reactions being two to three orders of magnitude faster than the rates of the competing reactions.

The strong asymmetry imposed on primary charge separation photo-chemistry in the purple bacterial RC results from two homologous polypeptides that function as a heterodimer. A heterodimer is also involved in the core of the RCs of PSI and PSII. However, some RCs, such as heliobacteria [3] and green sulfur bacteria [4], contain two identical homodimeric polypeptides, and electron transfer is potentially bifurcated.

Genetic sequence information has greatly improved the understanding of the origin of the RC proteins. From the sequence analysis, it became clear that the purple bacteria RC is remarkably similar to that of PSII, and PSI was also discovered to have similarity with that of the green sulfur bacteria [5]. Recent structural comparisons between PSI and PSII, for example, show a distinct structural homology, which suggests that even these two RCs likely share a common ancestor [6].

After extensive studies, the rate is now established to be critically coupled to the properties of the bacteriochlorophyll monomer that lies between the donor and bacteriopheophytin acceptor (Fig. 1). The involvement of the bacteriochlorophyll monomer may give rise to multiple pathways for electron transfer [7] and can partially determine the unidirectionality of charge separation along one branch [8, 9].

Fig. 1 The RC of purple bacteria are composed of three protein subunits called *L*, *M*, and *H*. Dimer *P* is described by molecule 1. Cofactors in the *L* subunits are represented as molecules 3 (BChl_{*L*}), 5 (BPh_{*L*}) and 6 (Q_{*L*}); similarly, in the subunits *M*, BChl_{*M*} is described as molecule 2, and molecule 4 represents BPh_{*M*}. Cytochrome C serve as a source of electrons for RC



The chain located on subunit *M* is inactive in ET, and the highly asymmetric functionality, however, can be decreased by amino acid mutations or cofactor modification. We used this approach to explain the effect of individual amino acid mutation or cofactor modifications on the observed balance between the forward ET reaction on the *L* side of the RC, the charge recombination processes, and ET to the *M*-side chromophores [10–14].

The theoretical models describe the charge transfer in RCs using parameters with clear physical interpretations. Some of these input parameters cannot be deduced from independent experimental work. The information regarding the energetic parameters, the medium reorganization energies, the high-frequency modes, and electronic coupling terms can be achieved with quantum mechanical computations. However, until now, these parameters characterizing the RCs have not been available. We thus use the set of parameters that fit the experiments. Several sets of parameters were used to describe a charge transfer in the RC [15–17]. The method for determining the input parameters from theory is by comparison with observed kinetics for different mutated RCs. The problem is that the impact of mutation on the input parameters is not always clear. In this paper, we focus on the electron transfer in two RCs. The first one is the RC of the green bacterium *Chloroflexus (C.) aurantiacus*. The second is the *H(M182)L* mutant from *Rhodobacter (R.) sphaeroides*. Generally, it is believed that RCs have similar structures, where high efficiency of solar energy conversion to chemical energy is based on common electronic properties. The peculiarity of *C. aurantiacus* RC is that it contains the Bacteriopheophytin (BPh) in the *M*-branch, where the Bacteriochlorophyll (BChl) molecule is placed in the *R. sphaeroides* RC [18]. The *M* branch is active in RCs where the BChl molecule is replaced by the BPh

molecule. It is thought that this replacement lowers the energy of accessory molecules, which causes branch *M* to be active. This is not the case for the *C. aurantiacus* RC.

We adapt in this work the set of parameters that characterizes the observed L-side experimental kinetics of wild-type (WT) RCs of *R. sphaeroides* very well [16, 17]. The *C. aurantiacus* RCs and the *H(M182)L* RC mutant from *R. sphaeroides* have a structural similarity, but their charge separation kinetics are different. Both these RCs contain BPh pigment in the *M*-branch at the position where BChl monomer is placed in WT RCs of *R. sphaeroides*. In contrast with this structural similarity, the *H(M182)L* mutant reveals the electron transfer through the *M* branch; in *C. aurantiacus* RCs, the *M*-branch is inactive.

2 Model of reaction center

The RC is an open system. If we describe the RC as a closed system, there ought to be, after some time, a Boltzmann distribution of electron localization on cofactors, and consequently, both branches must be active. To describe the leak of electrons from RC, we have to impose sink parameters into the model. It can be assumed that the system of interest interacts with another part of the overall system with quasicontinuum spectrum, and when an electron is transferred to this subsystem, backward transfer is practically impossible (in the duration of the experiment). There is, for instance, also the possibility of the deactivation of the RC to its ground state. In other words, the system of interest interacts with another part of the whole system, which is not investigated. This part can be assumed to have a continuum spectrum from which the backward electron transfer to the system of interest is not possible in real time. We describe this interaction (channel) by the imaginary part of the effective Hamiltonian, and this imaginary part cannot be neglected in the memory kernel function or in the free electron propagator [19].

Using the standard projection operator techniques [20–23], we can derive a generalized master equation for the populations,

$$\begin{aligned} \partial_t P_j(t) = & -\frac{2\Gamma_j}{\hbar} P_j(t) - \sum_{k=1}^N \int_0^t W_{jk}(t-\tau) P_j(\tau) d\tau \\ & + \sum_{k=1}^N \int_0^t W_{kj}(t-\tau) P_k(\tau) d\tau, \quad j = 1, \dots, N, \quad j \neq k, \end{aligned} \tag{1}$$

where

$$\begin{aligned} W_{jk}(t) = & 2 \frac{|V_{jk}|^2}{\hbar^2} \operatorname{Re} \left\{ \exp \left[-\frac{\Gamma_j + \Gamma_k}{\hbar} t \right] \exp \left[\frac{i(\varepsilon_j - \varepsilon_k)}{\hbar} t \right] \right. \\ & \left. \times \exp \left\{ \sum_{\alpha} \frac{E_{jk}^{\alpha}}{\hbar \omega_{\alpha}} [(\bar{n}_{\alpha} + 1) e^{-i\omega_{\alpha} t} + \bar{n}_{\alpha} e^{i\omega_{\alpha} t} - (2\bar{n}_{\alpha} + 1)] \right\} \right\}. \end{aligned} \tag{2}$$

Here, ε_j is the site energy, V_{ij} are electronic coupling parameters, $\hbar/2\Gamma_j$ has the meaning of the life time of the electron at site j in the limit of zero electronic coupling parameters, ω_{α} is the frequency of the α th mode, $\bar{n}_{\alpha} = [\exp(\hbar\omega_{\alpha}/k_B T) - 1]^{-1}$ is the thermal population of the α th mode, and

$$E_{jk}^{\alpha} = \frac{1}{2} m_{\alpha} \omega_{\alpha}^2 (d_{j\alpha} - d_{k\alpha})^2 \tag{3}$$

is the reorganization energy of the α th mode when the system transfers from state $|j\rangle$ to state $|k\rangle$. Here, m_α is the mass of the α th oscillator and $d_{j\alpha}$ is the equilibrium configuration of the α th oscillator when the system is in the electronic state j .

To describe the first step of the electron transfer processes in the RCs, we have used the five-site kinetic model of RC. We designate the special pair P as site 1, sites 2 and 3 represent the molecules $BChl_M$ and $BChl_L$, and sites 4 and 5 then represent the molecules BPh_M and BPh_L (Fig. 1). We assume that we can neglect the backward electron transfer from quinone molecules, and so we use the complex energies of molecules at sites 4 and 5. Based on experimental observations of ET in RC, it is expected that bacteriochlorophyll plays a crucial role in ET. In this five-site model, we have assumed that ET in RC is sequential. It means that the P^+BChl^- state on both the L and M branch is a real chemical intermediate of electron transfer and not only a virtual state, which is used in the superexchange electron transfer [24]. The imaginary part of energy level 1 describes the probability of electron deactivation to the ground state.

2.1 *Chloroflexus aurantiacus* reaction center

We describe the ET in *C. aurantiacus* RCs by the following kinetic model:

$$\begin{aligned} \partial_t P_1(t) = & - \left(\frac{2\Gamma_1}{\hbar} + k_{12} + k_{13} \right) P_1(t) \\ & + k_{21} P_2(t) + k_{31} P_3(t), \end{aligned} \quad (4a)$$

$$\partial_t P_2(t) = - \left(\frac{2\Gamma_2}{\hbar} + k_{21} + k_{24} \right) P_2(t) + k_{12} P_1(t) + k_{42} P_4(t), \quad (4b)$$

$$\partial_t P_3(t) = - (k_{35} + k_{31}) P_3(t) + k_{13} P_1(t) + k_{53} P_5(t), \quad (4c)$$

$$\partial_t P_4(t) = - \left(\frac{2\Gamma_M}{\hbar} + k_{42} \right) P_4(t) + k_{24} P_2(t), \quad (4d)$$

$$\partial_t P_5(t) = - \left(\frac{2\Gamma_L}{\hbar} + k_{53} \right) P_5(t) + k_{35} P_3(t). \quad (4e)$$

Here, we denote $k_{ij}(s \rightarrow 0^+) = k_{ij}$ and $k_{ij}(s \rightarrow 0^+) = \int_0^\infty W_{ij}(t) dt$.

We assume that the rate constant that characterizes ET can be described by both a low-frequency medium vibrational mode ω_m and a high-frequency intramolecular vibrational mode ω_c . We will work in the limit where the molecular modes are frozen, $\hbar\omega_c \gg k_B T$. In this regime, the constant k_{ij} is in the form [16, 25]:

$$\begin{aligned} k_{ij} = & \int_0^\infty W_{ij}(t) dt = \frac{2\pi V_{ij}^2}{\hbar^2 \omega_m} \exp(-S_{cij} - S_m(2\bar{n}_m + 1)) \\ & \times \sum_{n=0}^\infty \frac{S_{cij}^n}{n!} \left(\frac{\bar{n}_m + 1}{\bar{n}_m} \right)^{p(n)/2} I_{|p(n)|} (2S_m[\bar{n}_m(\bar{n}_m + 1)]^{1/2}). \end{aligned} \quad (5)$$

Here, $p(n) = (G_{ij} - n\hbar\omega_c)/\hbar\omega_m$, $G_{ij} = \epsilon_i - \epsilon_j$, and $S_{cij} = \frac{1}{2\hbar} m_{cij} \omega_{cij} (d_{ci} - d_{cj})^2$ is the scaled reorganization constant of the high-frequency ij th mode when electron is transferring from the state $|i\rangle$ to the state $|j\rangle$, and $S_{mij} = \frac{1}{2\hbar} m_{mij} \omega_{mij} (d_{mi} - d_{mj})^2$ is the scaled

reorganization constant of the low-frequency mode. $I_p(z)$ is the modified Bessel function of order p and $\bar{n}_m = [\exp(\frac{\hbar\omega_m}{k_B T}) - 1]^{-1}$. The back electron transfer reaction rate constant can be calculated by using the detailed balance relation and can be expressed in the form $k_{ji} = k_{ij} \exp(-G_{ij}/k_B T)$.

The quantum yields Φ_L , Φ_M of electronic escape via branch L , M , the quantum yields Φ_G of direct ground state recombination, and $\Phi_2 = 2\Gamma_2/\hbar$ deactivation to the ground state from the site 2 cofactor can be characterized for a five-site kinetic model by the expressions

$$\phi_G = \frac{2\Gamma_1}{\hbar} P_1(s \rightarrow 0^+) + \frac{2\Gamma_2}{\hbar} P_2(s \rightarrow 0^+), \quad (6a)$$

$$\phi_L = \frac{2\Gamma_L}{\hbar} P_5(s \rightarrow 0^+), \quad (6b)$$

$$\phi_M = \frac{2\Gamma_M}{\hbar} P_4(s \rightarrow 0^+), \quad (6c)$$

where the condition $\Phi_L + \Phi_M + \Phi_G = 1$ has to be fulfilled. We start with the assumption that $\Gamma_2 = 0$; this means that we consider no direct decay of the $P^+ BPh_M^-$ state into the ground state. The expressions for electron transfer are given by the inverse Laplace transformation. Therefore, we firstly apply the Laplace transformation to $P(t)$ in the system of Eqs. 4, where the Laplace transformation is defined as

$$P(s) = \int_0^\infty e^{-st} P(t) dt. \quad (7)$$

Next, we apply the inverse Laplace transformation of $P(s)$, where the inverse Laplace transformation is represented by a set of simple poles of $P(s)$. Evaluating it, we obtain

$$P(t) = \sum_{j=1}^5 a_j e^{k_j t}, \quad (8)$$

where a_j are the amplitudes and k_j are the rate kinetic constants describing the electron transfer.

Using the model described above, we would like to find the electron transfer kinetics for the RCs of *C. aurantiacus* [18], where, on the M branch, $BChl_M$ is replaced by BPh_M in the corresponding position. Thus, *C. aurantiacus* RCs contain altogether three BPh molecules and only one BChl monomer. There exists an agreement that the primary charge separation step in purple bacterial RCs occurs with a lifetime of approx. 3 ps at room temperature. This process is slower in RCs of *C. aurantiacus*. We weakly decrease the electronic coupling parameters in comparison with *R. sphaeroides* RCs because the kinetics in this RC is slower than in *R. sphaeroides* RCs. To characterize *C. aurantiacus*, we start with the set of parameters that characterize the kinetics of WT RCs of *R. sphaeroides*. We assume that the free energy difference of the $P^+ BChl_M^-$ state in *R. sphaeroides* RCs and the $P^+ BPh_M^-$ state of *C. aurantiacus* RCs is about $1,000 \text{ cm}^{-1}$, similar to the L -branch of *R. sphaeroides* RCs. The values $\varepsilon_2 = -50 \text{ cm}^{-1}$ and $\varepsilon_2 = -650 \text{ cm}^{-1}$ were used in our computations. The value $\varepsilon_2 = -50 \text{ cm}^{-1}$ is less favorable for electron transfer to the M -branch when

the high-frequency modes $\hbar\omega_c = 1,500 \text{ cm}^{-1}$ are used. The following values of input parameters were used: scaled reorganization constants $S_{mij} = 10$, $S_{cij} = 0.5$, high-frequency modes $\hbar\omega_{cij} = 1,500 \text{ cm}^{-1}$, and low-frequency mode $\hbar\omega_{mij} = 80 \text{ cm}^{-1}$, where $i, j = 1, 3, 5$ for the *L*-side and $i, j = 1, 2, 4$ for the *M*-side of the RC. The values for electronic coupling parameters $V_{24} = V_{35} = 32 \text{ cm}^{-1}$ and $V_{12} = V_{13} = 15 \text{ cm}^{-1}$ were used. The sink parameters $2\Gamma_M/\hbar = 2\Gamma_L/\hbar = (200 \text{ ps})^{-1}$, $2\Gamma_1/\hbar = (170 \text{ ps})^{-1}$ were used in accordance with experimental observation, which characterizes the ET to quinone molecules and decay to the ground state. The value $\hbar\omega_{mij} = 80 \text{ cm}^{-1}$ for low-frequency medium mode was chosen in accordance with results of the work [26]. The value $S_{mij} = 10$ of the scaled reorganization energy constant was used to obtain the reorganization energy of medium mode $\lambda_m = \hbar\omega_m S_m = 800 \text{ cm}^{-1}$. Since BChl_M is replaced by BPh_M in the corresponding position, we decrease the free energy in site 2. The calculated rate constants and quantum yields for the concrete energy levels are collected in Table 1, upper part.

We can see that, in this case, we get electron transfer through the *M* branch, which is not in accordance with experimental observations. To avoid this discrepancy, we must assume that asymmetry in the electronic coupling exists. To describe the experimental kinetics for the *C. aurantiacus* RC, we used the following asymmetry in electronic couplings: $V_{12} = 10 \text{ cm}^{-1}$ and $V_{13} = 15 \text{ cm}^{-1}$. We used the energy $\varepsilon_2 = -50 \text{ cm}^{-1}$ in the calculation because this case is less favorable for electron transfer through the *M*-branch, and so needs less asymmetry in the electronic coupling to achieve agreement with experimental data.

The calculated rate constants and quantum yields for the concrete energy levels are shown in Table 1, bottom part. The solution of Eq. 4 can be expressed in the form ($T = 295\text{K}$)

$$P_1(t) = 0.002e^{-0.53t} + 0.018e^{-0.34t} + 0.98e^{-0.12t}, \quad (9a)$$

$$P_2(t) = -0.05e^{-0.53t} + 0.005e^{-0.34t} + 0.04e^{-0.12t} + 0.005e^{-0.01t}, \quad (9b)$$

$$P_3(t) = -0.4e^{-0.34t} + 0.4e^{-0.12t}, \quad (9c)$$

$$P_4(t) = 0.053e^{-0.53t} - 0.003e^{-0.34t} - 0.22e^{-0.12t} + 0.13e^{-0.01t} + 0.04e^{-0.009t}, \quad (9d)$$

$$P_5(t) = 0.4e^{-0.34t} - 1.3e^{-0.12t} - 0.5e^{-0.01t} + 1.4e^{-0.009t}. \quad (9e)$$

The behavior of the occupation probabilities $P_i(t)$ at 295K is shown in Fig. 2. Similar to the work [18], we describe the kinetics with five exponential components. In our case, it is because the five-site model was used.

2.2 *Rhodobacter sphaeroides* *H(M182)L* reaction center

In this mutant, BChl_M is replaced with BPh_M . The new cofactor is referred to as ϕ_B . The new state $P^+\phi_B^-$ is formed during the decay of P^* and recombines to the ground state with a lifetime of 200 ps. The yield of $P^+\phi_B^-$ is about 0.35 at room temperature and the yield of $P^+\text{BChl}_L^-$ is about 0.65. There does not appear to be any further electron transfer from ϕ_B to BPh_M . The yield of the $P^+\phi_B^-$ state decreases to 0.12 at 77 and 9K. It is thought that, in the *H(M182)L* mutant, the $P^+\phi_B^-$ state is lower in energy than $P^+\text{BPh}_M^-$ [27, 28].

To elucidate the electron kinetics in this RC, we test two models. The first model is a coherent model where we assume that the free energies of the $P^+\phi_B^-$ and $P^+\text{BPh}_M^-$ states

Table 1 Computed constants $1/k_j$ and quantum yield dependence on temperature for *C. aurantiacus* RCs

Sample	T K	ϵ_2 cm^{-1}	ϵ_3 cm^{-1}	ϵ_4 cm^{-1}	ϵ_5 cm^{-1}	$1/k_{12}$ ps	$1/k_{21}$ ps	$1/k_{13}$ ps	$1/k_{31}$ ps	$1/k_{24}$ ps	$1/k_{42}$ ps	Φ_G	Φ_M	Φ_L
<i>C. auranti.</i>	295	-50	-450	-1,000	-2,000	12	15	6	52	2	188	0.02	0.31	0.67
$V_{12} = V_{13}$	200					15	21	5	138	2	1,559	0.02	0.25	0.73
<i>C. auranti.</i>	295	-650	-450	-1,000	-2,000	9.1	217	11	96	2.7	14.6	0.03	0.49	0.48
$V_{12} = V_{13}$	200					7.7	826	9.6	246	2.5	31	0.02	0.54	0.44
<i>C. auranti.</i>	295	-50	-450	-1,000	-2,000	47	60	11	93	2	188	0.05	0.16	0.79
$V_{12} \neq V_{13}$	200					58	84	10	246	2	1,559	0.05	0.12	0.83

The constants $1/k_{35} = 3(3.8)$ ps and $1/k_{53} = 5,894(264,544)$ ps for $T = 295(200)$ K

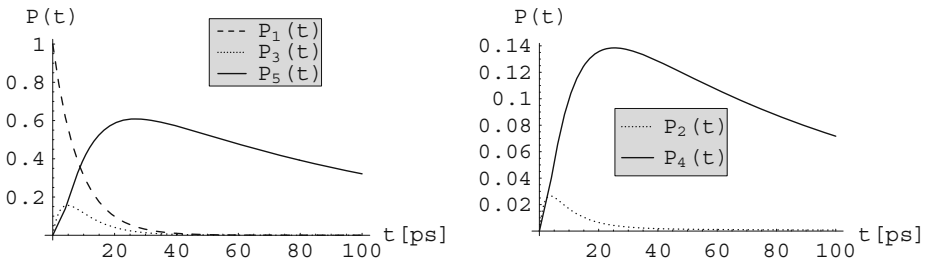


Fig. 2 The occupation probabilities $P(t)$ for *C. aurantiacus* RC in the asymmetric case $V_{12} = 10 \text{ cm}^{-1}$ and $V_{13} = 15 \text{ cm}^{-1}$

are closely spaced and the reorganization energy is practically zero when an electron is transferred between these two states. The second model is an incoherent model where the free energy of the $P^+\phi_B^-$ state is substantially below $P^+BPh_M^-$. Both models predict a small probability of finding electron at site 4. The following values of input parameters were used: scaled reorganization constants $S_{mij} = 10$, $S_{cij} = 0.5$, high-frequency mode $\hbar\omega_{cij} = 1,500 \text{ cm}^{-1}$, low-frequency mode $\hbar\omega_{mij} = 80 \text{ cm}^{-1}$ where $i, j = 1, 3, 5$ for the *L*-side and $i, j = 1, 2, 4$ for the *M*-side of RC. The values for electronic coupling parameters $V_{24} = V_{35} = 59 \text{ cm}^{-1}, V_{12} = V_{13} = 32 \text{ cm}^{-1}$ were used. The sink parameters $2\Gamma_M/\hbar = 2\Gamma_L/\hbar = (200 \text{ ps})^{-1}$, $2\Gamma_1/\hbar = (170 \text{ ps})^{-1}$ were used in accordance with experimental observation, which characterizes the ET to quinone molecules and decay to the ground state.

To explain the electron transfer in this mutant, we started with the coherent model. This means that we assume that the reorganization energy for ET from the $P^+\phi_B^-$ state to $P^+BPh_M^-$ is practically zero. The electron kinetics has to be described by the following system of equations:

$$\begin{aligned} \partial_t P_1(t) = & - \left(\frac{2\Gamma_1}{\hbar} + k_{12} + k_{13} \right) P_1(t) \\ & + k_{21} P_2(t) + k_{31} P_3(t), \end{aligned} \tag{10a}$$

$$\begin{aligned} \partial_t P_2(t) = & - \left(\frac{2\Gamma_2}{\hbar} + k_{21} \right) P_2(t) - \int_0^t W_{24}(t - \tau) P_2(\tau) d\tau + k_{12} P_1(t) \\ & + \int_0^t W_{42}(t - \tau) P_4(\tau) d\tau \end{aligned} \tag{10b}$$

$$\partial_t P_3(t) = - (k_{35} + k_{31}) P_3(t) + k_{13} P_1(t) + k_{53} P_5(t), \tag{10c}$$

$$\partial_t P_4(t) = - \frac{2\Gamma_M}{\hbar} P_4(t) - \int_0^t W_{42}(t - \tau) P_4(\tau) d\tau + \int_0^t W_{24}(t - \tau) P_2(\tau) d\tau, \tag{10d}$$

$$\partial_t P_5(t) = - \left(\frac{2\Gamma_L}{\hbar} + k_{53} \right) P_5(t) + k_{35} P_3(t). \tag{10e}$$

In this case, the memory function $W_{24} = W_{42}$ can be expressed in the form: $W_{24}(t) = 2\pi \frac{|V_{24}|^2}{\hbar^2} \text{Re} \left\{ \exp \left[- \frac{\Gamma_M + \Gamma_2}{\hbar} t \right] \exp \left[\frac{i(\epsilon_2 - \epsilon_4)}{\hbar} t \right] \right\}$, we begin with the assumption that $\Gamma_2 = 0$. This means that recombination to the ground state from site 2 does not exist. The calculated

Table 2 Computed constants $1/k_j$ and quantum yield dependence on temperature for *R. sphaeroides H(M182)L* mutant RCs

Coherent sample	T K	ϵ_2 cm^{-1}	ϵ_3 cm^{-1}	ϵ_4 cm^{-1}	ϵ_5 cm^{-1}	$1/k_{12}$ ps	$1/k_{21}$ ps	$1/k_{13}$ ps	$1/k_{31}$ ps	$1/k_{24}$ ps	$1/k_{42}$ ps	Φ_G	Φ_M	Φ_L
<i>H(M182)L</i> $\Gamma_2 = 0$	295	-850	-450	-1,000	-2,000	1.9	121	2.4	21	1,293	1,293	0.01	0.09	0.90
	200					1.6	729	2.1	54	1,293	1,293	0.01	0.30	0.69
<i>H(M182)L</i> $\Gamma_2 \neq 0$	77					1.1	10^6	1.9	8,479	1,293	1,293	0.004	0.628	0.368
	295	-850	-450	-1,000	-2,000	1.9	121	2.4	21	646	646	0.29	0.07	0.64
	200					1.6	729	2.1	54	646	646	0.43	0.10	0.47
	77					1.1	10^6	1.9	8,479	646	646	0.51	0.12	0.37

The constants $1/k_{35} = 0.9(1.1)(3.3)$ ps and $1/k_{53} = 1734(77820)(10^{13})$ ps for $T = 295(200)(77)$ K, $2\Gamma_2/\hbar = (200 \text{ ps})^{-1}$

values of parameters for the coherent model, which describes the $H(M182)L$ mutant RCs, are collected in Table 2, upper part. As can be seen, there is very small decay to the ground state. To fit the experimental data of the $H(M182)L$ mutant RCs, we need to introduce the possibility of a recombination of the $P^+\phi_B^-$ state directly into the ground state. So we also include sink $\Gamma_2 \neq 0$ into Eq. 10, where $2\Gamma_2/\hbar = (200 \text{ ps})^{-1}$. The computed rate constants and quantum yields after this assumption are shown in the bottom part of Table 2. In this case, the occupation probabilities $P_i(t)$ at 295K can be expressed in the form:

$$P_1(t) = 0.212e^{-1.23t} + 0.784e^{-0.88t} + 0.004e^{-0.008t}, \tag{11a}$$

$$P_2(t) = -0.073e^{-1.23t} - 0.379e^{-0.88t} + 0.448e^{-0.008t} + 0.004e^{-0.005t} + 0.002e^{-0.006t} \sin(36t), \tag{11b}$$

$$P_3(t) = -1.224e^{-1.23t} + 1.222e^{-0.88t} + 0.001e^{-0.008t} + 0.001e^{-0.005t}, \tag{11c}$$

$$P_4(t) = -0.017e^{-1.23t} - 0.09e^{-0.88t} + 0.106e^{-0.008t} + 0.001e^{-0.005t} - 0.002e^{-0.006t} \sin(36t), \tag{11d}$$

$$P_5(t) = 1.108e^{-1.23t} - 1.544e^{-0.88t} - 0.56e^{-0.008t} + 0.996e^{-0.005t}. \tag{11e}$$

We can see in this case that occupation probabilities $P_3(t)$ and $P_4(t)$ have an oscillating character. The frequency of oscillations depends on the free energy difference and electronic coupling parameter between these two sites [29]. The electron transfer kinetics for coherent models are shown in Fig. 3.

Now we intend to elucidate the observed ET kinetics with incoherent models. The value of the free energies used to calculate the rate constant are listed in Table 3. It was assumed that the free energy of $P^+\phi_B^-$ is significantly below $P^+BPh_M^-$ because of the electron transfer stops at ϕ_B [27, 30]. In the case $\Gamma_2 = 0$, we get a very small probability of finding an electron on the BPh_M molecule, but the quantum yield through the branch M is substantial (Table 3, upper part). So we have to impose, as in the previous case, the possibility that the $P^+\phi_B^-$ state can recombine directly into a ground state. The value $2\Gamma_2/\hbar = (200 \text{ ps})^{-1}$ was used. The computed rate constants and quantum yields after this assumption are collected in the

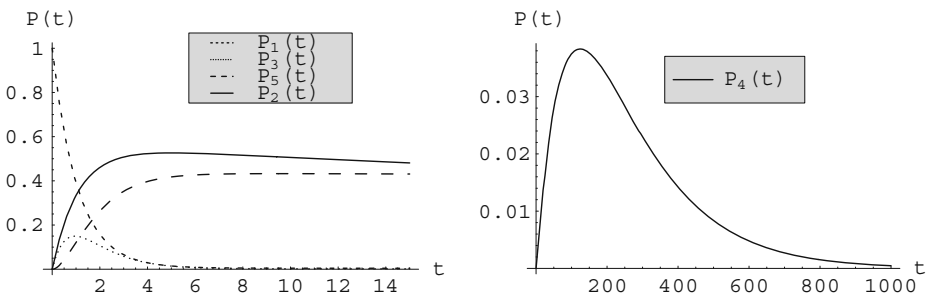


Fig. 3 The occupation probabilities $P(t)$ for the mutant $H(M182)L$ in the case of the coherent model of electron transfer where $2\Gamma_2/\hbar = (200 \text{ ps})^{-1}$, $T = 295\text{K}$

Table 3 Computed constants $1/k_{ij}$ and quantum yield dependence on temperature for *R. sphaeroides* *H(M182)L* mutant RCs

Incoherent sample	T K	ϵ_2 cm^{-1}	ϵ_3 cm^{-1}	ϵ_4 cm^{-1}	ϵ_5 cm^{-1}	$1/k_{12}$ ps	$1/k_{21}$ ps	$1/k_{13}$ ps	$1/k_{31}$ ps	$1/k_{24}$ ps	$1/k_{42}$ ps	Φ_G	Φ_M	Φ_L
<i>H(M182)L</i> $\Gamma_2 = 0$	295	-1,600	-450	-1,000	-2,000	3.2	7,797	2.4	21	11	0.6	0.01	0.34	0.65
	200					3.9	398,011	2.1	54	39	0.5	0.01	0.33	0.66
<i>H(M182)L</i> $\Gamma_2 \neq 0$	77					12	10^{14}	1.9	8,479	27,477	0.37	0.01	0.13	0.86
	295	-1,600	-450	-1,000	-2,000	3.2	7,797	2.4	21	11	0.6	0.42	0.02	0.56
	200					3.9	398,011	2.1	54	39	0.5	0.35	0.01	0.64
	77					12	10^{14}	1.9	8,479	27,477	0.37	0.14	0	0.86
	9					4.9	10^{111}	0.68	10^{31}	10^{41}	0.38	0.125	0	0.875

The rate constants $1/k_{35} = 0.9(1.1)(3.3)(3.11)$ ps and $1/k_{53} = 1734(77820)(10^{13})(10^{108})$ ps for $T = 295(200)(77)(9)K$; $2\Gamma_2/\hbar = (200 \text{ ps})^{-1}$

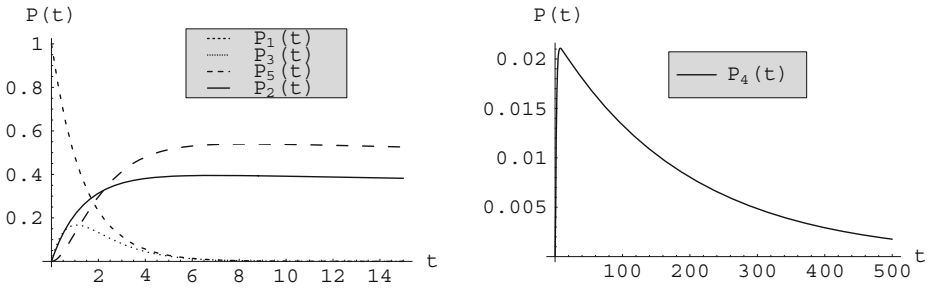


Fig. 4 The occupation probabilities $P(t)$ for the $H(M182)L$ mutant in the case of the incoherent model of electron transfer where $2\Gamma_2/\hbar = (200 \text{ ps})^{-1}$, $T = 295\text{K}$

bottom part of Table 3. We found the following expressions for the occupation probabilities $P_i(t)$ for an incoherent model at 295K in this case:

$$P_1(t) = 0.1e^{-1.2t} + 0.9e^{-0.7t}, \tag{12a}$$

$$P_2(t) = -0.02e^{-1.7t} - 0.02e^{-1.2t} - 0.36e^{-0.7t} + 0.4e^{-0.005t}, \tag{12b}$$

$$P_3(t) = -0.85e^{-1.2t} + 0.85e^{-0.7t}, \tag{12c}$$

$$P_4(t) = 0.02e^{-1.7t} - 0.04e^{-0.7t} + 0.02e^{-0.005t}, \tag{12d}$$

$$P_5(t) = 0.8e^{-1.2t} - 1.37e^{-0.7t} + 0.57e^{-0.005t}. \tag{12e}$$

The time dependence of the site-occupation probabilities $P_i(t)$ for an incoherent model is presented in Fig. 4.

The system deactivates to the ground state mainly through the $P^+\phi_B^-$ state. To get the quantum yields in accordance with experimental data, we have to use the value $S_{mij} = 5$ at $T = 9\text{K}$.

3 Conclusion

We have dealt with electron transfer in *C. aurantiacus* and *R. sphaeroides H(M182)L* mutant RCs. In spite of their structural similarity, their functionality is very different. The *H(M182)L* mutant reveals the *M*-branch active in electron transfer.

Discussions about the main factors determining electron transfer directionality can be found in the papers [26, 31–34]. Previously, it was suggested that the difference in the electronic coupling parameters in the two branches is the dominant factor. Later, experimental work cast doubt on the dominance of electronic coupling as a mechanism that causes the asymmetry in ET through branches [27, 30]. Our position is that the parameters describing electron transfer from the bacteriochlorophyll dimer to bacteriochlorophyll monomers are crucial for unidirectionality. Obviously, mutations in RCs with a goal to change these parameters can identify what the most important parameter for unidirectionality is.

In RCs, electron transfer is sensitive to the free energy difference between the charge-separated states. In the case of *C. aurantiacus* RCs, we assume that the free energy of the $P^+BPh_M^-$ state at site 2 is in the range of $1,000\text{ cm}^{-1}$ below the $P^+BChl_M^-$ state at site 2 of *R. sphaeroides* RCs. It is estimated that this energy is about 800 cm^{-1} above P^* in *R. sphaeroides* RCs [35]. We have examined two possible values of the free energy of the $P^+BPh_M^-$ state at site 2, $\varepsilon_2 = -50\text{ cm}^{-1}$ and $\varepsilon_2 = -650\text{ cm}^{-1}$. The value $\varepsilon_2 = -50\text{ cm}^{-1}$ is less favorable for electron transfer through the *M*-branch. In both cases, the *M*-branch is active. So we have to introduce coupling integrals asymmetry with to get accordance with the experimental data.

In the case of the *R. sphaeroides H(M182)L* mutant RCs, we have investigated two possibilities. The first one has free energy of the $P^+\phi_B^-$ and $P^+BPh_M^-$ states closely spaced. The second one has free energy of the $P^+\phi_B^-$ state 600 cm^{-1} below free energy of $P^+BPh_M^-$. In the first case, we used the model where, between the sites 2 and 4, electron transfer has a coherent character. This means that the electron is delocalized between these two sites. It was assumed that reorganization energy is small when electron is transferred from ϕ_B to the BPh_M molecule. The coherent model was used because is not clear why this mutation can decrease the free energy of the BPh (refereed as ϕ_B) molecule so substantially below the P^* state. The incoherent model in this case gives a significant probability to find the system at the $P^+BPh_M^-$ state. Without assuming the possibility that the $P^+\phi_B^-$ state can decay directly to the ground state, the coherent model gives more realistic results then the incoherent model. But in both models, there is the outlet through the *M* branch, which is not in accordance with experimental observations. The problem of both the coherent and incoherent models is that they do not predict enough decay to the ground state without the assumption that there is a possibility of $P^+\phi_B^-$ state recombination to the ground state. If we take this possibility into account, both models give results in good agreement with experimental observation at 295K. The coherent model predicts an incorrect dependence of quantum yield on temperature. The coherent model does not produce results that are in accordance with experimental data in this case but can be used, for instance, when we want to investigate how the *BChl* dimer can contribute to electron transfer asymmetry [36]. A similar model ought to be used to characterize electron transfer in the PSII (PSI) system, where dimer molecules of *BChl* are not so close as in the bacterial RCs.

The incoherent model can elucidate electron transfer in *R. sphaeroides H(M182)L* mutant RCs without asymmetry in electronic coupling parameters using the set of parameters that characterizes the observed *L*-side experimental kinetics very well [16, 17, 23]. To characterize the *H(M182)L* mutant, we used the value $\varepsilon_2 = -1,600\text{ cm}^{-1}$ for the free energy of the $P^+\phi_B^-$ state. At 9K, the value $S_{mij} = 5$ has to be used to fit the experimental data. To get appropriate results at $T = 9\text{K}$, we have to change the reorganization energy of the low-frequency mode from the value 800 cm^{-1} to the value 400 cm^{-1} . Using the value 800 cm^{-1} , the yield of the $P^+\phi_B^-$ state is 0.04, which is not in accordance with experimental data. The decrease of the reorganization energy at low temperature can be due to the fact that friction with the surrounding medium is greater. The result is that the configuration displacement $d_{mi} - d_{mj}$ at low temperature is smaller than that at room temperature.

One of the unsolved problems of electron transfer in the RC is unidirectionality. The electron used only one of two possible branches. Much experimental work has been performed to solve this problem [9, 10, 27, 28, 37, 38]. Generally, it is assumed that RCs have common features and the unidirectionality is caused mainly by asymmetry in the coupling integral or asymmetry in the free energy, or that both these asymmetries contribute

to unidirectionality. This paper shows that there is a possibility that different RCs can have different grounds for unidirectionality.

To derive the kinetic equations, the projection operator technique has been used. The physics depends on the concrete projection operator, which was used. If we assume that, after excitation, the electron transfer is so fast that relevant vibrational modes have no time to relax, we can average over the surrounding medium, which is in equilibrium with the dimer in the ground state and not in the excited state. The models used in the presented paper can describe such quick electron transfer.

To also describe the bath dynamics, additional free parameters are needed [39]. The theoretical model where the bath relaxation is imposed also has to fit the experimental data. The data determine a model in which phonon relaxation may or may not play an important role. If the relaxation is important, there has to be another set of parameters in the model, in which this process is taken into account in comparison with the presented model. For instance, both models give different predictions for electron transfer in experiments where the electric fields are applied. In the optimal case, a comparison with experimental data can guide the selection of a correct model, and establish a role of phonon relaxation in the electron transfer in the RCs.

To describe both the unidirectionality and the results of the experiments with mutated RCs, we used the five-site model. For this complex system, the parameters, which describe the *L*-side electron transfer very well, were applied. Despite a large discussion about these parameters, the key problem is, however, to determine the *M*-side parameters. Because of the two-fold symmetry of RCs, the *L*-side parameters were also used to describe the *M*-side electron transfer. But there must be asymmetry in some parameter to get unidirectionality of electron transfer. Our goal was to determine which parameters are critical and what are the values of these parameters. Previously, it was presumed that the asymmetry in coupling integrals results in the asymmetry of electron transfer. Several studies with mutated RCs showed that it is not a sufficient explanation. It cannot be only the asymmetry of coupling integrals, but rather the asymmetry of the free energy of bacteriochlorophyll monomers located on different branches that appears to be the key factor of unidirectionality. We believe that the presented model can identify the parameters that are changed in the mutated RC in comparison to WT and also determine the impact of mutation on the RCs. We also believe that we can identify the parameters that play a key role in the unidirectionality in the RC of *C. aurantiacus*. The values of these parameters are reasonable. Without their change in comparison to *R. sphaeroides* mutant RC, we have *L*-side electron transfer, which is not observable in the RC of *C. aurantiacus*. Generally, it was believed that asymmetry in coupling integrals or asymmetry in free energy or both these asymmetries together are the main reason for unidirectionality in bacterial RCs. We show here that there is some diversification of mechanisms among different RCs. In some RCs, the main impact on the unidirectionality comes from the asymmetry in coupling integrals (*C. aurantiacus*), and in others, the key factor can be the asymmetry in free energies (*R. sphaeroides*). Until now, such a possibility has not been taken into account, and it can be important if artificial RCs with similar performance are constructed. We utilized the rate constant, which allows us to make a prediction for electron transfer at very low temperature (9K).

It is important to compute the temperature dependence of quantum yields and also to compute the electron transfer kinetics. It can help to estimate more correctly the values of model parameters and subsequently to contribute to better understanding of the primary processes of photosynthesis.

Acknowledgements The work was supported by the Slovak Academy of Sciences under the CEX NANOFLUID, APVV 0509-07, Centrum of Excellence No 26220120021 and VEGA grant 2/7056/27.

References

1. Deisenhofer, J., Epp, O., Miki, K., Huber, R., Michel, H.: Structure of the protein subunits in the photosynthetic reaction centre of *Rhodospseudomonas viridis* at 3 Å resolution. *Nature* **318**, 618–624 (1985)
2. Allen, J.P., Willams, J.C.: Photosynthetic reaction centers. *FEBS Lett.* **438**, 5–9 (1998)
3. Ames, J.: The heliobacteria, a new group of photosynthetic bacteria. *J. Photochem. Photobiol. B Biol.* **30**, 89–96 (1995)
4. Sakurai, H., Kusumoto, N., Inoue, K.: Function of the reaction center of green sulfur bacteria. *Photochem. Photobiol.* **64**, 5–13 (1996)
5. Golbeck, J.H.: Shared thematic elements in photochemical reaction centers. *Proc. Natl. Acad. Sci. U.S.A.* **90**, 1642–1646 (1993)
6. Krauß, N., Schubert, W.-D., Klukas, O., Fromme, P., Witt, H.T., Saenger, W.: Photosystem I at 4 Å resolution represents the first structural model of a joint photosynthetic reaction center and core antenna system. *Nat. Struct. Biol.* **3**, 965–973 (1996)
7. Van Brederode, M.E., Jones, M.R., Van Mourik, F., Van Stokkum, I.H.M., Van Grondelle, R.: A new pathway for transmembrane electron transfer in photosynthetic reaction centers of *Rhodobacter sphaeroides* not involving the excited special pair. *Biochemistry* **36**, 6855–6861 (1997)
8. Heller, B.A., Holten, D., Kirmaier, C.: Control of electron transfer between the *L*- and *M*-sides of photosynthetic reaction centers. *Science* **269**, 940–945 (1995)
9. Chuang, J.I., Boxer, S.G., Holten, D., Kirmaier, C.: Temperature dependence of electron transfer to the *M*-side bacteriopheophytin in *Rhodobacter capsulatus* reaction centers. *J. Phys. Chem. B* **112**, 5487–5499 (2008)
10. Kirmaier, Ch., Holten, D.: Evidence that a distribution of bacterial reaction centers underlies the temperature and detection-wavelength dependence of the rates of the primary electron-transfer reactions. *Proc. Natl. Acad. Sci. U.S.A.* **87**, 3552–3556 (1990)
11. Takahashi, E., Wraight, C.A.: Proton and electron-transfer in the acceptor quinone complex of *Rhodobacter sphaeroides* reaction centers – characterization of site-directed mutants of the 2 ionizable residues, GLUL212 and ASPL213, in the QB binding-site. *Biochemistry* **31**, 855–866 (1992)
12. Shuvalov, V.A., Duysens, L.N.M.: Primary electron transfer reactions in modified reaction centers from *Rhodospseudomonas sphaeroides*. *Proc. Natl. Acad. Sci. U.S.A.* **83**, 1690–1694 (1986)
13. Gehlen, J.N., Marchi, M., Chandler, D.: Dynamics affecting the primary charge transfer in photosynthesis. *Science* **263**, 499–502 (1994)
14. Müller, M.G., Drews, G., Holzwarth, A.: Primary charge separation processes in reaction centers of an antenna-free mutant of *Rhodobacter capsulatus*. *Chem. Phys. Lett.* **258**, 194–202 (1996)
15. Marchi, M., Gehlen, J.N., Chandler, D., Newton, M.: Diabatic surfaces and the pathway for primary electron transfer in a photosynthetic reaction center. *J. Am. Chem. Soc.* **115**, 4178–4190 (1993)
16. Tanaka, S., Marcus, R.A.: Electron transfer model for the electric field effect on quantum yield of charge separation in bacterial photosynthetic reaction centers. *J. Phys. Chem. B* **101**, 5031–5045 (1997)
17. Bixon, M., Jortner, J., Michel-Beyerle, M.E.: A kinetic analysis of the primary charge separation in bacterial photosynthesis. Energy gaps and static heterogeneity. *Chem. Phys.* **197**, 389–404 (1995)
18. Müller, M.G., Griebenow, K., Holzwarth, A.R.: Primary processes in isolated photosynthetic bacterial reaction centers from *Chloroflexus aurantiacus* studied by picosecond fluorescence spectroscopy. *Biochim. Biophys. Acta* **1098**, 1–12 (1991)
19. Capek, V., Szocs, V.: Is the sink model of exciton trapping in molecular condensates satisfactory? *Phys. Status Solidi, B* **125**, K137–K142 (1984)
20. Spargaglione, M., Mukamel, S.: Dielectric friction and the transition from adiabatic to nonadiabatic electron transfer. I. Solvation dynamics in Liouville space. *J. Chem. Phys.* **88**, 3263–3280 (1988)
21. Zwanzig, R.: On the identity of three generalized master equations. *Physica* **30**, 1109–1123 (1964)
22. Shibata, F., Takashi, Y., Hashitsume, N.: A generalized Stochastic Liouville equation. Non-Markovian versus memoryless master equations. *J. Stat. Phys.* **17**, 171–187 (1977)
23. Pudlak, M.: Primary charge separation in the bacterial reaction center: validity of incoherent sequential model. *J. Chem. Phys.* **118**, 1876–1882 (2003)
24. Marcus, R.A.: An internal consistency test and its implications for the initial steps in bacterial photosynthesis. *Chem. Phys. Lett.* **146**, 13–22 (1988)

25. Jortner, J.: Temperature dependent activation energy for electron transfer between biological molecules. *J. Chem. Phys.* **64**, 4860–4867 (1976)
26. Pudlak, M., Pincak, R.: Modeling charge transfer in the photosynthetic reaction center. *Phys. Rev. E* **68**, 061901–7 (2003)
27. Katilius, E., Turanchik, T., Lin, S., Taguchi, A.K.W., Woodbury, N.W.: B-side electron transfer in a *Rhodobacter sphaeroides* reaction center mutant in which the B-side monomer bacteriochlorophyll is replaced with bacteriopheophytin. *J. Phys. Chem. B* **103**, 7386–7389 (1999)
28. Katilius, E., Katilene, Z., Lin, S., Taguchi, A.K.W., Woodbury, N.W.: B-side electron transfer in a *Rhodobacter sphaeroides* reaction center mutant in which the B-side monomer bacteriochlorophyll is replaced with bacteriopheophytin: low-temperature study and energetics of charge-separated states. *J. Phys. Chem. B* **106**, 1471–1475 (2002)
29. Pudlak, M.: Electron transfer driven by conformational variations. *J. Chem. Phys.* **108**, 5621–5625 (1998)
30. Kirmaier, Ch., He, Ch., Holten, D.: Manipulating the direction of electron transfer in the bacterial photosynthetic reaction center by swapping Phe for Tyr near BChlM (L181) and Tyr for Phe near BChlL (M208). *Biochemistry* **40**, 12132–12139 (2001)
31. Michel-Beyerle, M.E., Plato, M., Deisenhofer, J., Michel, H., Bixon, M., Jortner, J.: Unidirectionality of charge separation in reaction centers of photosynthetic bacteria. *Biochim. Biophys. Acta* **932**, 52–70 (1988)
32. Pincak, R., Pudlak, M.: Noise breaking the twofold symmetry of photosynthetic reaction centers: electron transfer. *Phys. Rev. E* **64**, 031906–10 (2001)
33. Plato, M., Möbius, K., Michel-Beyerle, M.E., Bixon, M.: Intermolecular electronic interactions in the primary charge separation in bacterial photosynthesis. *J. Am. Chem. Soc.* **110**, 7279–7285 (1988)
34. Pudlak, M., Pincak, R.: The role of accessory bacteriochlorophylls in the primary charge transfer in the photosynthetic reaction center. *Chem. Phys. Lett.* **342**, 587–592 (2001)
35. Parson, W.W., Chu, Z.T., Warshel, A.: Electrostatic control of charge separation in bacterial photosynthesis. *Biochim. Biophys. Acta* **1017**, 251–272 (1990)
36. Yamasaki, H., Nakamura, H., Takano, Y.: Theoretical analysis of the electronic asymmetry of the special pair in the photosynthetic reaction center: effect of structural asymmetry and protein environment. *Chem. Phys. Lett.* **447**, 324–329 (2007)
37. Kirmaier, C., Weems, D., Holten, D.: M-side electron transfer in reaction center mutants with a lysine near the non-photoactive bacteriochlorophyll. *Biochemistry* **38**, 11516–11530 (1999)
38. Haffa, A.L.M., Lin, S., Williams, J.A.C., Bowen, B., Taguchi, A.K.W., Allen, J.P., Woodbury, N.: Controlling the pathway of photosynthetic charge separation in bacterial reaction centers. *J. Phys. Chem. B* **108**, 4–7 (2004)
39. Parson, W.W., Warshel, A.: A density matrix model of photosynthetic electron transfer with microscopically based estimated vibrational relaxation times. *Chem. Phys.* **296**, 201–216 (2004)