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# Vibronic Coherence in the Charge Separation Process of the *Rhodobacter sphaeroides* Reaction Center

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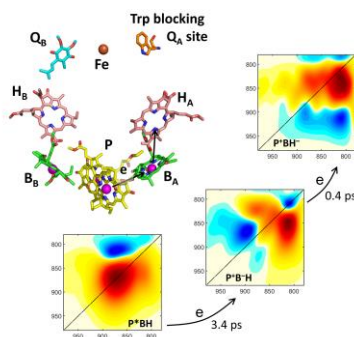
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ABSTRACT. Two-dimensional electronic spectroscopy was applied to a variant of the reaction center (RC) of purple bacterium *Rhodobacter sphaeroides* lacking the primary acceptor ubiquinone in order to understand the ultrafast separation and transfer of charge between the bacteriochlorin cofactors. For the first time characteristic 2D spectra were obtained for the participating excited and charge-transfer states, and the electron transfer cascade (including two different channels, the P\* and B\* channels) was fully mapped. By analyzing quantum beats using 2D frequency maps, excited-state vibrational modes at 153 and 33 cm<sup>-1</sup> were identified. We speculate that these modes couple to the charge separation (CS) process and collectively optimize the CS and is responsible for the super-high efficiency.

## TOC GRAPHICS



**KEYWORDS** bacterial reaction center, photosynthetic charge separation, electron transfer cascade, 2DES, vibronic coherence

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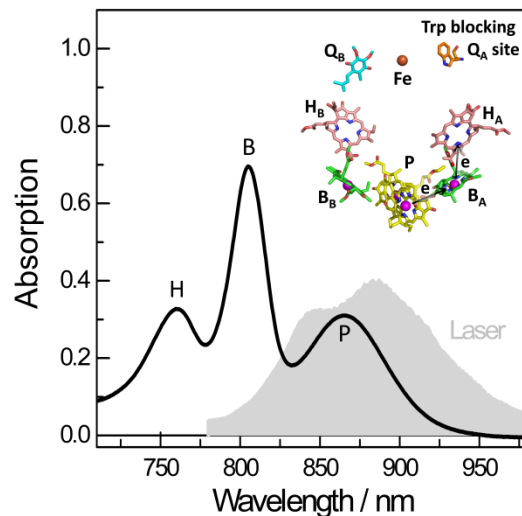
Reaction center (RC) pigment-proteins facilitate the key reaction of photosynthesis in which harvested solar energy is used to power separation of electrical charge across a lipid bilayer membrane.<sup>1,2</sup> Many of the fundamental aspects of ultrafast light-powered charge separation (CS) have been unraveled in the relatively simple RC from the purple bacterium *Rhodobacter (Rba.) sphaeroides*. The bacteriochlorin cofactors of this RC are a pair of bacteriochlorophyll (BChl) *a* (P), two monomeric BChl *a* (B) and two bacteriopheophytin *a* (H) molecules arranged in two branches around an axis of quasi two-fold symmetry (Figure 1).<sup>3</sup> CS is initiated from the first singlet excited state of P ( $P^*$ ), an electron being transferred to a ubiquinone ( $Q_A$ ) via the intervening  $B_A$  and  $H_A$  that make up the “active branch” or “A-branch” of cofactors.<sup>4</sup> At room temperature the  $P^* \rightarrow P^+B_A^-$ ,  $P^+B_A^- \rightarrow P^+H_A^-$  and  $P^+H_A^- \rightarrow P^+Q_A^-$  electron transfer (ET) steps occur with time constants of 3–5 ps, 0.5–1 ps and ~200 ps, respectively.<sup>1,5</sup> CS can also occur via another ET channel,  $B^* \rightarrow P^+B_A^-$ , with a time constants of 0.2–0.5 ps.<sup>6,7</sup> The CS process proceeds with a near-unity quantum efficiency.

Coherent two-dimensional electronic spectroscopy (2DES) has developed into a powerful tool for the study of excitation energy transfer (EET) and ET dynamics in photosynthetic systems.<sup>8–20</sup> The related Photosystem II RC from oxygenic phototrophs has been studied by 2DES,<sup>14,15</sup> but the ET cascades were difficult to resolve because the absorption bands of the chlorin cofactors strongly overlap, producing high signal congestion. In contrast the *Rba. Sphaeroides* RC exhibits near-discrete absorption bands with maxima at 865, 805 and 760 nm attributable to the P pair,  $B_A/B_B$  and  $H_A/H_B$ , respectively. This raises the prospect that 2DES can resolve details of the ET kinetics in bacterial RCs, a possibility supported by modeling.<sup>21</sup>

To date, 2DES studies performed on *Rba. sphaeroides* RCs have exclusively probed EET processes in complexes where P was oxidized to  $P^+$ <sup>11,22,23</sup> or kept reduced<sup>24</sup>, and direct

observation of the ET cascade has not been carried out. A complication in applying 2DES to native *Rba. sphaeroides* RCs is that the final states  $P^+Q_A^-$  and  $P^+Q_B^-$  have recombination lifetimes on the order of milliseconds to seconds, such that P spends significant periods in the photo-oxidized state and is unable to be excited and to perform CS.<sup>25</sup> In this work we overcame this problem by using an engineered RC in which an alanine at residue 260 of the M-polypeptide is replaced by tryptophan (AM260W). This mutation causes the RC to assemble without a  $Q_A$  ubiquinone<sup>26,27</sup> such that the forward  $P^+H_A^- \rightarrow P^+Q_A^-$  ET step is blocked and the majority of  $P^+H_A^-$  decays to the ground state with a lifetime of  $\sim 17$  nanoseconds.<sup>28</sup> As a result, no inactive RCs with an oxidized P persist beyond the repetition time between excitations (1 ms in this work), but the  $P^* \rightarrow P^+B_A^- \rightarrow P^+H_A^-$  ET rates are essentially identical to those in the wild-type RC.<sup>29</sup>

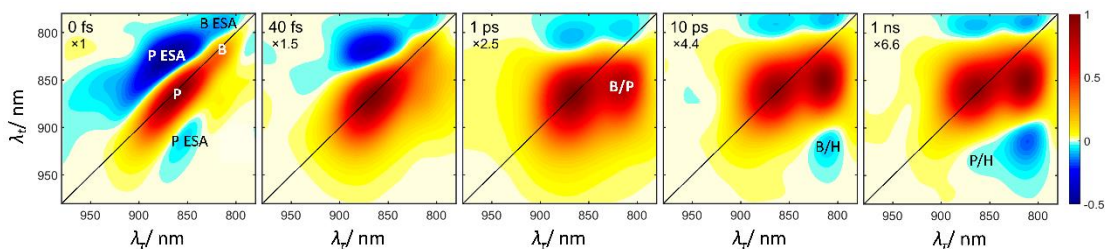
Potentially important aspects of the mechanism of CS in the *Rba. sphaeroides* RC are quantum effects, i.e. quantum beating arising from coupling between electronic, vibrational or mixed (vibronic) states. Previous pump-probe experiments have found a rich structure of oscillations,<sup>30-33</sup> which were thought to play a role in efficient ET.<sup>34</sup> In this work, we used 2DES to distinguish each CT intermediate along the two ET pathways of the neutral AM260W RC, and tried to understand the role of the long-lived quantum beats in CS.



**Figure 1.** Room temperature linear absorption spectrum of the AM260W RC (black) overlaid with the laser spectrum (grey). The inset shows the arrangement of cofactors and the ET pathway in the AM260W RC, which lacks a Q<sub>A</sub> acceptor due to an Ala to Trp replacement.

The excitation pulse predominantly excited P with some minor excitation of B (Figure 1). The absorptive 2D spectra at five population times (T) are shown in Figure 2. Signals associated with the P→P\* transition, the positive ground state bleaching (GSB) and stimulated emission (SE) on the diagonal and the negative excited-state absorption (ESA) off the diagonal, appeared simultaneously following excitation. At the same time small GSB and ESA signals attributable to B\* also appeared, and a small cross peak centered at (815, 870) nm ((excitation wavelength, detection wavelength)) was visible. This cross peak may reflect dynamic coherence between P\* and B\* at T=0 fs, and later its growth may reflect two processes: EET or/and ET from B\* to P. Both processes were fast, the EET was complete in around 200 fs<sup>1,25</sup> and the ET had a time constant of 0.2–0.5 ps.<sup>6,7</sup> The 2D spectrum at T=0 fs allowed an upper estimate of the homogeneous line width of the exciton between the ground and excited state, which was 340 cm<sup>-1</sup> (2γ) determined by the full width at half-maximum of the antidiagonal broadening of the P GSB band. γ is the dephasing rate and is inversely proportional to the coherence lifetime,<sup>19</sup> thus

the electronic coherence lifetime was estimated to be 54 fs. The P GSB decreased significantly during the first 50 fs and its shape became more rounded. This spectral diffusion reflected redistribution of the excitation energy among the exciton state manifold coupled to the protein bath.



**Figure 2.** Absorptive total 2D spectra of AM260W RC at the indicated population time  $T$ . Symbols  $\lambda_\tau$  and  $\lambda_t$  denote the excitation and detection wavelengths, respectively. The spectra are normalized to the maximum of the diagonal P signal; the relative amplitude multiplier is shown below the  $T$ . The identities of particular signals are indicated in each of the spectra: diagonal peaks are labeled with a single symbol designating the corresponding state; cross peaks are labeled using two symbols, with the first representing the locally excited state and the second representing the acceptor state.

Three different cross peaks appeared on a picosecond time scale. The positive one centered at (815, 855) nm arose with a time constant of 430 fs. It is slower than the  $B^* \rightarrow P$  EET (200 fs), so most probably it corresponded to  $B^* \rightarrow P^+ B_A^-$  ET (labeled “B/P”) and thus was a feature of the  $P^+ B_A^-$  CT state. The remaining two cross peaks were negative and centered at (810, 920) and (865, 927) nm. They formed nearly synchronously on the time scale of several picoseconds, although the weak (865, 927) nm cross peak was partially obscured by the P GSB signal and therefore seemed to appear later. According to previous pump-probe experiments, the positive signal at 920~930 nm can be attributed to the absorption of  $H^-$ , with the absorption of  $B^-$  centered at 1025 nm possibly making a minor contribution.<sup>35</sup> Control pump-probe measurements with the same excitation and probe pulses as used for 2DES also showed a positive signal after a

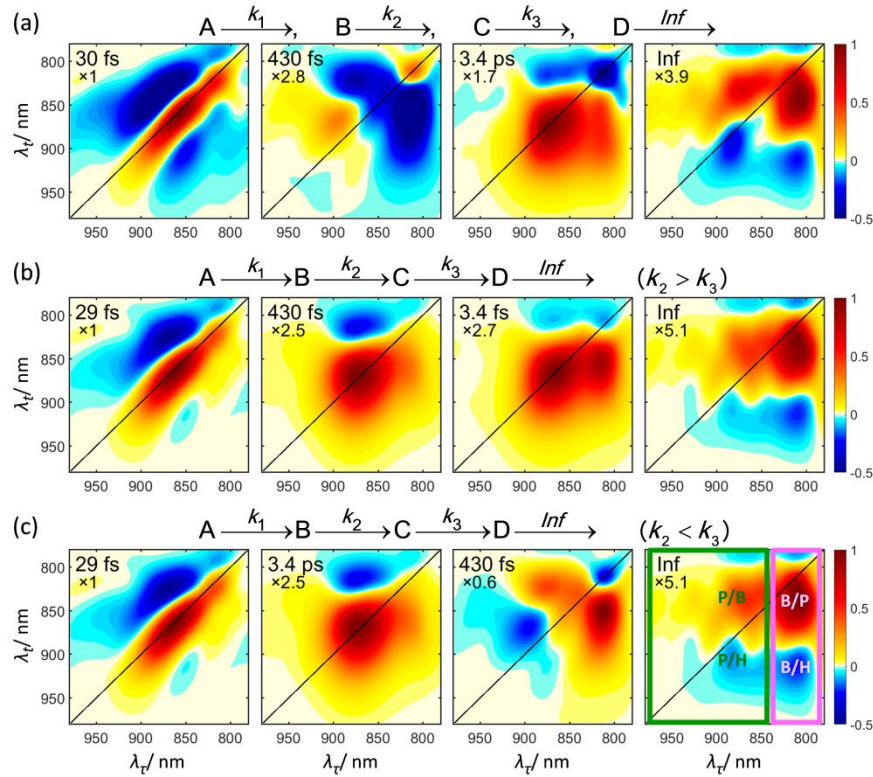
few picoseconds (Figure S1), indicating the formation of  $H^-$ . The (810, 920) and (865, 927) nm cross peaks therefore reflected ET to  $H_A$ , being features of the  $B_A^+H_A^-$  and  $P^+H_A^-$  CT states, respectively. The spectral shapes of all the three cross peaks remained nearly unchanged between 10 ps and 1 ns. This period corresponded to decay of  $P^+H_A^-$  to the ground state by charge recombination or via formation of the triplet state of P.<sup>29</sup>

One should note that the three CT cross peaks belonged to two different ET channels. The (815, 855) and (810, 920) nm ones respectively corresponded to the first and second CT state in the  $B^* \rightarrow P^+B_A^- \rightarrow P^+H_A^-$  channel. The (865, 927) nm one corresponded to the second CT state in the  $P^* \rightarrow P^+B_A^- \rightarrow P^+H_A^-$  channel. The first CT state expected to be at around (855, 810) nm was not observed, mainly because that its decay is faster than its formation, and furthermore it was obscured by the P ESA. It will be revealed by global fitting as discussed below. Comparing the amount of formed  $H^-$  from the two channels, we can conclude that because  $B^* \rightarrow P^+B_A^-$  (0.4 ps) is faster than  $P^* \rightarrow P^+B_A^-$  (3.4 ps) ET, the  $B^*$  channel is more efficient. The results show that 2DES can not only track the two channels, but also provides a way to quasi-quantitatively assess the relative yield of competing channels, which is almost impossible in 1D measurements.

To obtain the spectral features and evolution time constants of each ET species, a global analysis was applied. The fitting was started with a parallel or a sequential model, which respectively give decay associated spectra (DAS) or evolution associated spectra (EAS).<sup>13,17,18,36</sup> Both models gave the same characteristic time constants: 29 fs, 430 fs, 3.4 ps and >1 ns. Both DAS or EAS may vary from the actual spectra of the true species involved if the model deviates from the real dynamics through which species evolve. For example, in the present case the DAS (Figure 3a) of the species B and C did not exhibit the features that each CT species should exhibit. This discrepancy was because the real dynamic processes are principally sequential for each ET



channel (Figure 3b). Furthermore, according to the well-established cascade scheme for ET in the bacterial RC a four-component sequential model with a faster third step than second step (i.e.  $k_2 < k_3$ ) best agrees with the real situation of the P\* ET channel (Fig. 3c). The spectra obtained with this model are referred to as species associated spectra (SAS). Comparison of the EAS (Figure 3b) and SAS (Figure 3c) showed that the biggest difference was for species C, which is the second of the two species involved in the dynamic inversion. Spectra of the other species were not affected by the inversion. Furthermore, this difference was pronounced for the P\* ET channel while had much smaller influence on the B\* channel.



**Figure 3.** 2D DAS (a), EAS (b) and SAS (c) with the respective kinetic model shown on their top. The time constants of each species are shown in the left-top corner of each panel. The spectra are normalized to the maximum of the diagonal P signal, and the relative amplitude multiplier is shown below the time constant. The 2D spectral regions corresponding to the P\*

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and B\* channels are shown within dark green and pink windows, respectively, in the last plot of c.

Each intermediate state along the ET cascade was well resolved in the four SAS. As discussed above, the SAS model agrees with the P\* ET channel, so below we will focus on the 2D spectra region of this channel (the dark green window in Figure 3c). The first SAS corresponded to P\* at the Franck-Condon point. Its lifetime was very short, 29 fs, due to fast spectral diffusion. The second SAS contained also the GSB, SE and ESA signals. The GSB peak was nearly symmetrically broadened along the antidiagonal direction, and its center position, (872, 869) nm, was red-shifted by about 5 nm along the diagonal compared to that of the first SAS, (863, 865) nm. This indicated that a new equilibrium of the excitonic states manifold was established. An electron was then transferred from P\* to B<sub>A</sub>, the 3.4 ps lifetime of P\* being determined by the P\*→P<sup>+</sup>B<sub>A</sub><sup>-</sup> CT rate.

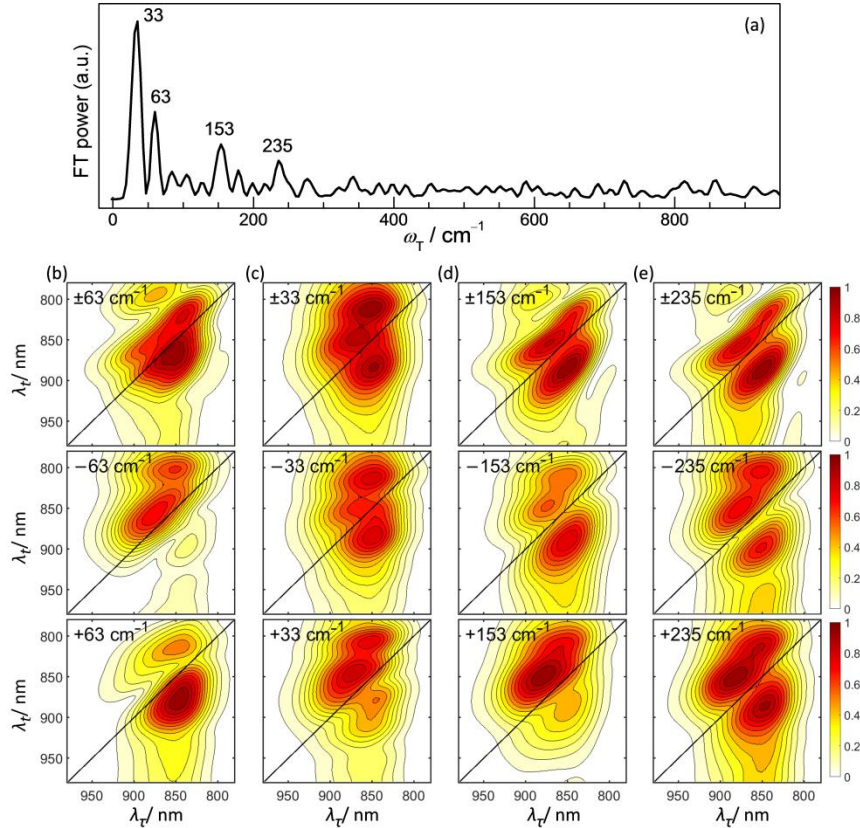
The third SAS contained positive ESA signals for P and B, and two symmetrical cross peaks relative to the diagonal. The above-diagonal cross peak centered at (870, 820) nm corresponded to P\*→P<sup>+</sup>B<sub>A</sub><sup>-</sup> ET, while the below-diagonal cross peak centered at (810, 855) nm to B\*→P<sup>+</sup>B<sub>A</sub><sup>-</sup> ET. They both reflected the properties of the P<sup>+</sup>B<sub>A</sub><sup>-</sup> CT state, but with different origins. We can see that the P<sup>+</sup>B<sub>A</sub><sup>-</sup> CT state originated from P\* was only resolved by fitting with a correct model. The third SAS had a lifetime of 430 fs, which should be determined by the P<sup>+</sup>B<sub>A</sub><sup>-</sup>→P<sup>+</sup>H<sub>A</sub><sup>-</sup> ET rate. However, the time constant of B\*→P<sup>+</sup>B<sub>A</sub><sup>-</sup> CS was also close to this value, 0.2–0.5 ps, so it is difficult to distinguish the two dynamic processes.

The last SAS was more complex, containing weak GSB and ESA signals for both P and B, two positive cross peaks representing P<sup>+</sup>B<sub>A</sub><sup>-</sup> and two negative cross peaks centered at (810, 912) and (882, 905) nm. The latter corresponded to the formation of H<sup>-</sup> from B\* and P\* ET, respectively,

bearing the features of  $B_A^+H_A^-$  and  $P^+H_A^-$  CT states. Thus this SAS not only exhibited features of the  $P^+H_A^-$  state, which is the fourth and final species, but also features of the other CT states that are preceding intermediates in the ET cascade. This SAS decayed only a little until the maximal delay time of 1 ns used in this work, consistent with the measured lifetime of this state in the AM260W RC of 17 ns.<sup>28</sup> To summarize, the sequential ET processes from P to  $H_A$  were fully traced by 2DES to be  $P^* \xrightarrow{29 \text{ fs}} P_A^+P_B^- \xrightarrow{3.4 \text{ ps}} P^+B_A^- \xrightarrow{430 \text{ fs}} P^+H_A^- \xrightarrow{>1 \text{ ns}}$ . In accord with previous observations that  $P^+B_A^-$  decayed much faster than its formation, it was nearly spectrally invisible in raw data and was only revealed by global fitting using the correct model.

In the  $B^*$  ET channel (the pink window in Figure 3c), the rise time constants of the B/P and B/H cross peaks were 430 fs and 3.4 ps. So the dynamic evolution may be better described with EAS than SAS, although the EAS and SAS in this 2D spectral region did not exhibit essential difference. From the above results, we can see that the 2D spectra of each CT intermediate state exhibited more distinguishing characteristics than those can be observed in 1D measurements, being an advantage of exploiting 2DES for complex systems.

The 2D spectra exhibited long-lived (up to 2 ps) oscillations. Figure 4a presents a summary Fourier transform (FT) power spectrum obtained by summing the squared absolute value (Frobenius norm) of the Fourier transforms of the T traces across the 2D spectra. The prominent frequencies were 33, 63, 153 and 235  $\text{cm}^{-1}$ . These frequencies were consistent with those determined previously by pump-probe measurements.<sup>31,32</sup> And the 33, 63 and 153  $\text{cm}^{-1}$  frequencies have their correspondences in resonance Raman spectrum,<sup>37</sup> 33, 70 and 145  $\text{cm}^{-1}$ .



**Figure 4.** (a) Summary FT power spectrum of the oscillations in the real rephasing signal (0–2 ps). (b)–(e) 2D frequency maps of the four  $\omega_T$  frequencies computed from the real-valued rephasing spectra (top) and from the complex-valued rephasing spectra (middle:  $-\omega_T$ , bottom:  $+\omega_T$ ).

To understand the origin of the oscillations, 2D frequency maps were analyzed. The  $\pm\omega_T$  frequency maps were yielded from the real part of the rephasing data while the separate  $-\omega_T$  and  $+\omega_T$  frequency maps were from the full complex part. It has been shown that the characteristic patterns and symmetries in the frequency maps can help to distinguish electronic, vibrational or mixed coherences.<sup>38–42</sup> The  $\pm 63 \text{ cm}^{-1}$  frequency map (Figure 4b for rephasing and Figure S2 for non-rephasing) was dominated by diagonal peak, indicating that the corresponding oscillations originated from vibrational coherence.<sup>39</sup> The features of the  $-63$  and  $+63 \text{ cm}^{-1}$  frequency maps resembled those in a carefully designed experiment for identifying vibrational coherence.<sup>41</sup> The

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below-diagonal peak in  $+\omega_T$  rephasing frequency map reflects ground-state vibrational superpositions while the above-diagonal peak in  $-\omega_T$  rephasing frequency map reflects excited-state vibrational superpositions.<sup>40,41</sup> So the oscillation with  $63\text{ cm}^{-1}$  could be assigned to vibrational coherence in both ground and excited states. It is notable that a peak at about (865, 805) nm appeared in all the frequency maps and was the most pronounced in the  $33\text{ cm}^{-1}$  ones. It could be from the contribution of B. Below our analysis would focus on the main P peak.

The features of the  $33$ ,  $153$  and  $235\text{ cm}^{-1}$  frequency maps (Figure 4c–e) were different. First of all, in the  $\pm\omega_T$  rephasing frequency maps the diagonal peak disappeared, instead, two off-diagonal peaks appeared symmetrically along the diagonal. It indicated a different coherent nature from that for the  $63\text{ cm}^{-1}$  oscillation. Secondly, for  $33$  and  $153\text{ cm}^{-1}$  the dominant peak was below diagonal in the  $-\omega_T$  rephasing frequency map while was above diagonal in the  $+\omega_T$  rephasing frequency map, which was opposite with the  $63\text{ cm}^{-1}$  case. For  $235\text{ cm}^{-1}$  the features were a bit deviated, both above- and below-diagonal peaks with comparable amplitudes appeared in each  $-\omega_T$  and  $+\omega_T$  rephasing frequency maps. The features observed in the  $33$  and  $153\text{ cm}^{-1}$  frequency maps can be explained by strongly coupled excited-state vibronic coherence.<sup>39,42</sup>

Long-lived vibronic coherence was found responsible for increasing the rate of EET<sup>11</sup> and CS<sup>14,15</sup> in RC. It was thought that resonant electronic-vibrational coupling can sustain coherences between electronic states,<sup>43,44</sup> which, however, was doubted later.<sup>45</sup> Very recently the essence of vibronic coherence was proposed to be excited-state vibrational coherences shifted to the ground state as a result of a release of electronic energy during EET, which was concluded from the vibronic coherence coupled with the  $H^* \rightarrow B \rightarrow P$  EET process in *Rba. Sphaeroides* RC.<sup>46</sup> Based on this mechanism, the  $33$  and  $153\text{ cm}^{-1}$  oscillations can be assigned to excited-state vibronic coherence followed with shift into the ground-state vibrational coherence. The assignment of the

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235  $\text{cm}^{-1}$  oscillations was not as definite as the 33 and 153  $\text{cm}^{-1}$  ones. It may be weakly coupled vibronic coherence.

Universal appearance of vibrational modes coupling in the CS process in RCs (both bacterial and high plant) has attracted numerous theoretical interest on how coupling of specific vibrational modes facilitates this process. Our previous simulation of *Rba. Sphaeroides* RC based on Redfield theory revealed that nonequilibrated vibrational modes are involved in CS and determines the dynamics of the excited-state wavepacket.<sup>34</sup> Two modes were identified: the 130  $\text{cm}^{-1}$  one connecting with the intermolecular dynamics within P and the 32  $\text{cm}^{-1}$  one responsible for a stabilization of the primary CT state. Later, ab initio molecular dynamics simulation found that coupling with two vibrational modes, 50 and 100  $\text{cm}^{-1}$ , leads to a unidirectional displacement of electron density to establish the CT state.<sup>47,48</sup> Redfield theory simulation of Photosystem II RC revealed that resonant vibrations can modify the delocalization of the exciton states and thus promote direct CS.<sup>49,50</sup>

Our finding that the 153 and 33  $\text{cm}^{-1}$  modes appeared as excited-state vibronic coherence matches very well with the simulation results. Furthermore, the 153  $\text{cm}^{-1}$  mode, assigned to be associated with the internal CT state within P,<sup>34</sup> did not appear in the 2DES measurement of oxidized *Rba. Sphaeroides* RC<sup>46</sup>, implying that it couples to the CS process. Hence we speculate that excitation of them facilitates the CS process in the *Rba. Sphaeroides* RC. This vibronic coherence mechanism could be responsible for the super-high efficiency of natural photochemical CS, which may offer a blueprint for constructing photochemical devices with high energy conversion efficiencies using abundant materials.

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## ASSOCIATED CONTENT

### **Supporting Information.**

The following files are available free of charge.

Experimental methods. Figure S1. Pump-probe spectra. Figure S2. Non-rephasing 2D frequency maps.

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### **Notes**

The authors declare no competing financial interests.

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Supporting information

# Vibronic Coherence in the Charge Separation Process of the *Rhodobacter sphaeroides* Reaction Center

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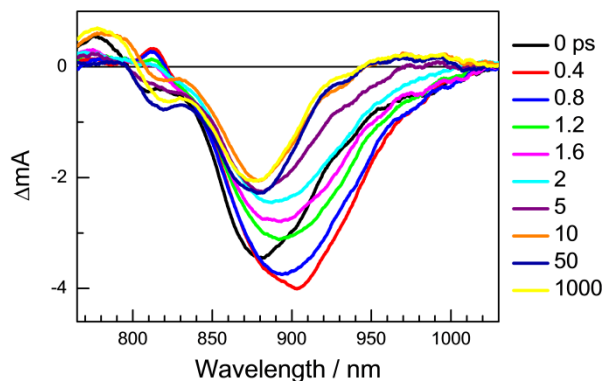
## Experimental methods

AM260W RCs<sup>1</sup> modified with a poly-histidine tag on the PufM polypeptide<sup>2</sup> were isolated from a strain *Rba. Sphaeroides* lacking light harvesting complexes<sup>3</sup> using n-dodecyl-N,N-dimethylamine-N-oxide (LDAO) as the solubilizing detergent. RCs were purified by nickel affinity chromatography followed by size exclusion chromatography, as described in detail previously (S2), and stored as a concentrated solution in 20 mM Tris (pH 8.0)/0.1 % (w/v) LDAO at  $-80$  °C until use. The sample was diluted with a buffer containing 20 mM Tris (pH 8.0)/0.05% LDAO (w/v) to an optical density of  $\sim 0.3$  at the maximum of P band for the 2DES measurement in 0.1 mm quartz cell. The sample was kept static during measurement. After each measurement, the degree of the sample degradation was monitored by UV-Vis spectra and less than 2% degradation was observed.

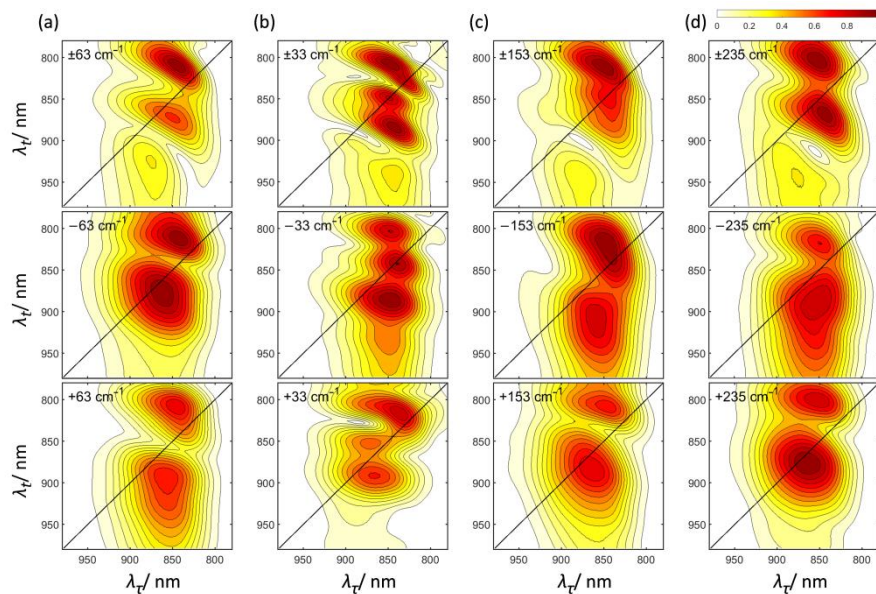
Room temperature 2D electronic spectra were recorded with a diffractive optic-based inherently phase-stabilized four-wave mixing set-up, which was described in detail previously<sup>4</sup>. The repetition rate of the laser system (PHAROS, Light Conversion) was set to 1 kHz. The pulse generated by a home-built non-collinear optical parametric amplifier was centered at 880 nm with a full width half maximum of 105 nm and a duration of 18 fs. The excitation intensity was 2 nJ per pulse, corresponding to a photon density of  $3 \times 10^{13}$  photons/cm<sup>2</sup> at which annihilation was negligible. The polarization scheme was all parallel. The coherence time ( $\tau$ ) was scanned from  $-90$  fs to 100 fs, with 1 fs step by employing movable fused silica wedges. The population time (T) is scanned from  $-1$  ps to 1 ns, with a step of 10 fs in  $-30$  fs to 2 ps region. With this laser repetition frequency and intensity, no before-zero signal was observed, showing that there was no accumulation of long-lived CT components and the RCs were functionally active, which was also proved by the reproducibility of the experimental data (test measurements with less

population time points were employed).

Global fittings were done for the real-valued total (rephasing and non-rephasing) data. Oscillatory components were extracted by subtracting global fits from each  $(\lambda_\tau, \lambda_t)$  data point of the complex rephasing data. Fourier transforming of the oscillating residuals in T and slicing the obtained spectra in  $\omega_T$  resulted in the  $(\lambda_\tau, \lambda_t)$  frequency maps. Fourier transforming the residuals of the real part yielded  $\pm\omega_T$  maps, while transforming the full complex-valued residuals yielded separate positive and negative  $\omega_T$  maps. The resolution of  $\omega_T$  was  $19 \text{ cm}^{-1}$ , determined by the length of the T scan, 0 to 2 ps.



**Figure S1.** Pump-probe spectra at different delay times. The pump and the probe pulses are the same with that used in 2DES experiments.



**Figure S2.** 2D frequency maps of the four  $\omega_T$  frequencies computed from the real-valued non-rephasing spectra (top) and from the complex-valued non-rephasing spectra (middle:  $-\omega_T$ , bottom:  $\omega_T$ ).

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