

# **Efficient Energy Transfer in Light-Harvesting Systems, III: The Influence of the Eighth Bacteriochlorophyll on the Dynamics and Efficiency in FMO**

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As mentioned in the main text, the Hamiltonian for FMO is constructed from the crystal structure recently deposited in the protein data bank (pdb code: 3eoj).<sup>1</sup> The site energies are taken from those computed in Ref. 4 and the coupling element between sites  $n$  and  $m$  is calculated from the dipole-dipole approximation,

$$V_{nm} = C \left( \frac{\mathbf{d}_n \cdot \mathbf{d}_m}{|\mathbf{r}_{nm}|^3} - 3 \frac{(\mathbf{d}_n \cdot \mathbf{r}_{nm})(\mathbf{d}_m \cdot \mathbf{r}_{nm})}{|\mathbf{r}_{nm}|^5} \right). \quad (1)$$

Following the prescriptions used previously for constructing the dipole-dipole interactions,<sup>2-4</sup> the unit vectors,  $\mathbf{d}_n$ , in Eq. (1) point along the axis connecting the  $N_b$  and  $N_d$  atoms of the  $n$ -th Bchl and  $\mathbf{r}_{nm}$  is the vector connecting the Mg atoms of Bchl  $n$  and  $m$ . Setting the constant  $C = 155000 \text{ cm}^{-1} \text{ \AA}^3$  leads to an effective dipole strength of  $30 \text{ D}^2$ . With these specifications the system Hamiltonian (in  $\text{cm}^{-1}$ ) for the eight site model is

$$H_{\text{FMO}} = \begin{pmatrix} 310.0 & -97.9 & 5.5 & -5.8 & 6.7 & -12.1 & -10.3 & 37.5 \\ -97.9 & 230.0 & 30.1 & 7.3 & 2.0 & 11.5 & 4.8 & 7.9 \\ 5.5 & 30.1 & 0.0 & -58.8 & -1.5 & -9.6 & 4.7 & 1.5 \\ -5.8 & 7.3 & -58.8 & 180.0 & -64.9 & -17.4 & -64.4 & -1.7 \\ 6.7 & 2.0 & -1.5 & -64.9 & 405.0 & 89.0 & -6.4 & 4.5 \\ -12.1 & 11.5 & -9.6 & -17.4 & 89.0 & 320.0 & 31.7 & -9.7 \\ -10.3 & 4.8 & 4.7 & -64.4 & -6.4 & 31.7 & 270.0 & -11.4 \\ 37.5 & 7.9 & 1.5 & -1.7 & 4.5 & -9.7 & -11.4 & 505.0 \end{pmatrix}, \quad (2)$$

where the zero of energy is  $12195 \text{ cm}^{-1}$ . Note that there is an error in the sign of the coupling between sites 1 and 2 in the table provided in Ref. 4. Aside from this, these values reproduce all of the couplings listed therein to within  $3 \text{ cm}^{-1}$ .

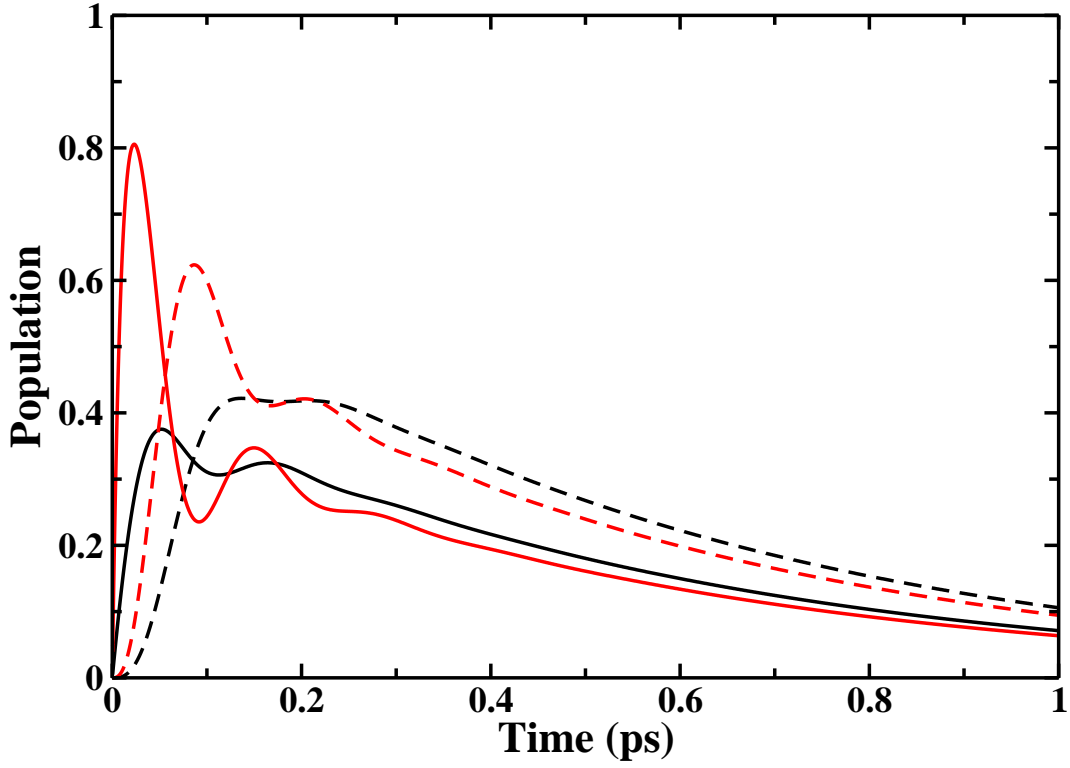


Figure 1: The site populations of the dimer calculated using the NIBA discussed in the main text. The solid and dashed lines correspond to the populations of site 1 and 2, respectively. The black lines are calculated with a decay rate from site 8 of  $\gamma = 15 \text{ ps}^{-1}$  and the red lines with  $\gamma = 100 \text{ ps}^{-1}$ . In both cases, the temperature is 300 K with a reorganization energy of  $35 \text{ cm}^{-1}$  and cutoff frequency of  $\omega_c^{-1} = 50 \text{ fs}$ . Note in particular, the onset of oscillations in the dynamics of site 2 when  $\gamma = 15 \text{ ps}^{-1}$  (dashed black) and the recovery of the dynamics in Figure 2(c) of the main text when  $\gamma = 100 \text{ ps}^{-1}$  (red).

## References

- (1) Tronrud, D. E.; Wen, J.; Gay, L.; Blankenship, R. E. The Structural Basis for the Difference in Absorbance Spectra for the FMO Antenna Protein from Various Green Sulfur Bacteria. *Photosynth. Res.* **2009**, *100*, 79–87.
- (2) Hu, X.; Ritz, T.; Damjanović, A.; Schulten, K. Pigment Organization and Transfer of Electronic Excitation in the Photosynthetic Unit of Purple Bacteria. *J. Phys. Chem. B* **1997**, *101*, 3854–3871.
- (3) Müh, F.; Madjet, M. E.; Adolphs, J.; Abdurahman, A.; Rabenstein, B.; Ishikita, H.; Knapp, E.; Renger, T.  $\alpha$ -Helices Direct Excitation Energy Flow in the Fenna-Matthews-Olson Protein. *Proc. Natl. Acad. Sci. USA* **2007**, *104*, 16862–16867.
- (4) Schmidt am Busch, M.; Müh, F.; Madjet, M. E.; Renger, T. The Eighth Bacteriochlorophyll Completes the Excitation Energy Funnel in the FMO Protein. *J. Phys. Chem. Lett.* **2011**, *2*, 93–98.