

Optimal fold symmetry of LH2 rings on a photosynthetic membrane

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An intriguing observation of photosynthetic light-harvesting systems is the N -fold symmetry of light-harvesting complex 2 (LH2) of purple bacteria. We calculate the optimal rotational configuration of N -fold rings on a hexagonal lattice and establish two related mechanisms for the promotion of maximum excitation energy transfer (EET). (i) For certain fold numbers, there exist optimal basis cells with rotational symmetry, extendable to the entire lattice for the global optimization of the EET network. (ii) The type of basis cell can reduce or remove the frustration of EET rates across the photosynthetic network. We find that the existence of a basis cell and its type are directly related to the number of matching points S between the fold symmetry and the hexagonal lattice. The two complementary mechanisms provide selection criteria for the fold number and identify groups of consecutive numbers. Remarkably, one such group consists of the naturally occurring 8-, 9-, and 10-fold rings. By considering the inter-ring distance and EET rate, we demonstrate that this group can achieve minimal rotational sensitivity in addition to an optimal packing density, achieving robust and efficient EET. This corroborates our findings *i* and *ii* and, through their direct relation to S , suggests the design principle of matching the internal symmetry with the lattice order.

Photosynthesis lies at the heart of all life on the planet. Excitation energy created in a peripheral light-harvesting antenna complex, upon absorption of a photon, is efficiently transferred via an exciton mechanism, sometimes over hundreds of angstroms, to a reaction center where charge separation takes place, creating a transmembrane electrochemical potential difference that ultimately drives the production of ATP. Light-harvesting antenna complexes, whose primary function is to increase the effective cross-section for photon absorption, while maintaining an efficient energy transfer network, consist of well-ordered membrane-associated arrays of light-absorbing pigments, namely chlorophyll or bacteriochlorophyll (BChl), embedded in a protein environment (1–4).

Perhaps the most intriguing aspect of light-harvesting systems is the N -fold symmetry displayed by light-harvesting complex 2 (LH2) of purple nonsulfur bacteria. Fold symmetry occurs in systems as diverse as carbon nanotubes, tubular dye aggregates, and the mosaic virus protein, as well as in snow flakes, tori, and nonperiodic tilings (5). In LH2, the N -fold symmetry arises from the N repetitions of a basic heterodimer subunit consisting of two transmembrane polypeptides, α and β , arranged in a circular structure (6–8). Each $\alpha\beta$ -heterodimer subunit noncovalently binds one B800 BChl and two B850 BChls (Fig. 1A), so named according to their respective room-temperature absorption bands in the infrared region of the spectrum. In most bacterial species, the $\alpha\beta$ -heterodimer repeats eight or nine times to form an octameric or nonameric ring structure (in *Rhodospseudomonas paulisterium* 10-fold rings have also been observed). Since the breakthrough of the first crystallizations of LH2 (9, 10), X-ray crystallography has determined the nine- and eightfold structures of *Rhodospseudomonas acidophila* and *Rhodospirillum molischianum*, respectively, to within a resolution of ~ 2 Å. Subsequent, equilibrium and nonequilibrium molecular dynamics simulations of their subunit heterodimers have shown that the mean values of separations between heterodimers are nearly fixed (11), so that the heterodimer subunit is independent of fold number.

Consequently, a simple variation of the angle attained during self-assembly between heterodimers can achieve a complete ring of specific fold number N . This raises an important question, as to the role of specific fold symmetry in the highly efficient excitation energy transfer (EET) observed in light-harvesting complexes.

However, another intriguing aspect, in addition to the fold symmetry of the individual LH2 complexes, is the lattice structure evident in their supramolecular organization (12–14). Recent progress in atomic force microscopy (AFM), which allows direct observation of the bacterial photosynthetic complexes on their native membrane, has revealed that LH2 complexes occur in well-ordered hexagonal domains (Fig. 1B) (15–17). This is especially evident in bacterial membranes grown under low light intensity, where large LH2 domains, some hundreds of angstroms wide, form a hexagonal lattice to increase absorption. Not surprisingly, hexagonal order is the closest packing structure on a two-dimensional surface. An important observation is that well-ordered hexagonal domains occur for both octameric and nonameric structures, suggesting that a close packing order is preferable to a crystalline order. Understanding the symmetry properties on the level of both the individual complex and the supramolecular complex is necessary to form a complete picture of efficient excitation energy transfer in the light-harvesting antenna (18, 19) and has direct implications for the design of efficient synthetic devices (20–26).

In this work, we investigate the role of fold symmetry in promoting excitation energy transfer in LH2 hexagonal domains, using an N -fold model geometry of a B850 ring based on the $\alpha\beta$ -heterodimer subunit of *Rps. acidophila* (outlined in *SI Materials and Methods*, Fig. S1). Considering the effective Hamiltonian of the N -fold geometry, and noting an important property of N -fold symmetric ring structures, namely the existence of $N - 2$ and $N - 1$ pairs of degenerate eigenstates for even and odd N (*SI Materials and Methods*), we arrive at a simple metric for determining the maximum excitation energy transfer rate between two B850 rings, based on the multichromophoric Förster theory (Eq. 6). We first discuss the issues of symmetry and frustration on a hexagonal lattice and then explore the optimal rotational configuration of N -fold rings and establish two related fold symmetry mechanisms for the promotion of maximum excitation energy transfer. Using these mechanisms as selection criteria, we can identify groups of consecutive fold numbers. However, to understand why one consecutive group may be preferable to another we require additional considerations (the fold symmetry mechanisms effectively provide a prescreening of fold numbers). In particular, supposing an optimal packing density of heterodimers on the bacterial membrane,

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Table 1. Matching points and basis cells for fold numbers 2–18

N	S points	Basis cell	Optimization
2	2	Rhombus	Yes
3	3	Triangle	Yes
4	2	Rhombus	Yes
5	1	None	No
6	6	Single ring	Yes
7	1	None	No
8	2	Rhombus	Yes
9	3	Triangle	Yes
10	2	Rhombus	Yes
11	1	None	No
12	6	Single ring	Yes
13	1	None	No
14	2	Rhombus	Yes
15	3	Triangle	Yes
16	2	Rhombus	Yes
17	1	None	No
18	6	Single ring	Yes

illustrate how the intermediate group of consecutive fold numbers 8, 9, and 10 may be considered preferable.

Packing Density and Rotational Sensitivity

To elucidate why 8-, 9-, and 10-fold symmetries appear preferable in nature, we extend our considerations beyond EET (relevant to the antenna network efficiency) to include the packing density ρ (relevant to the absorption cross section). For LH2 rings, the packing density can be defined as $\rho = N_H/A$, where N_H is the number of heterodimers contained in the primitive cell and A is the area of the cell. For the hexagonal lattice the primitive cell is a hexagon containing $N_H = 3N$ heterodimers with area $A = 3\sqrt{3}d_c^2/2$. Thus, we have

$$\rho = \frac{N_H}{A} = \frac{2N}{\sqrt{3}d_c^2}. \quad [1]$$

In Fig. 5 we plot Eq. 1 (right axis, dashed line) as a function of the fold number N for the fixed inter-ring distance of the 9-fold rings $d = d_c - 2R = 21.8$ Å, where the N -dependent radius R is

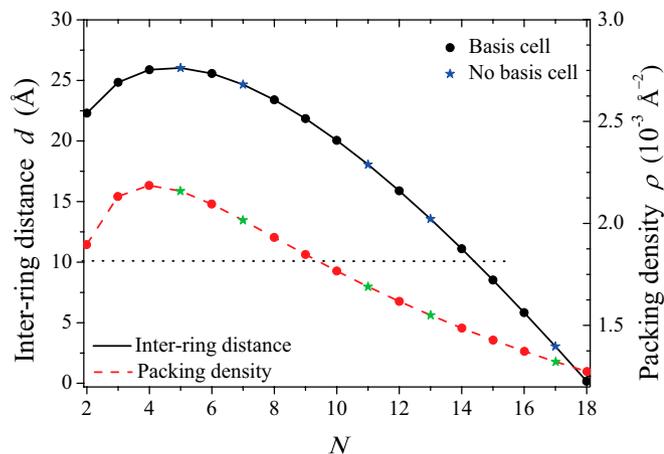


Fig. 5. Inter-ring distance d for constant density $\rho = \rho_9$ (left axis, solid line) and the packing density ρ for constant inter-ring distance $d = 21.8$ Å (right axis, dashed line) as a function of fold number N . The fold numbers for which a or no basis cell exists are indicated with a solid circle or star. The minimum physical inter-ring distance $d_{\min} = 10$ Å is indicated by the dotted line.

given in *SI Materials and Methods* (Eq. S1). Maintaining a fixed inter-ring distance results in a diminishing density for large fold symmetry. Alternatively, we can suppose the existence of an optimal packing density of heterodimers on the native bacterial membrane that achieves maximum absorption. To maintain this optimal packing density, the fixed-structure heterodimer subunits must self-assemble into a homogeneous N -fold ring system with a varying inter-ring distance. We choose the optimal density to be $\rho_9 \approx 1.8 \times 10^{-3} \text{ Å}^{-2}$, whereon the center-to-center distance d_c in terms of N becomes

$$d_c = \sqrt{\frac{2N}{\sqrt{3}\rho_9}} = 25 \sqrt{N}. \quad [2]$$

The inter-ring distance $d = d_c - 2R$ (left axis, solid line), where d_c is given by Eq. 2, is shown in Fig. 5. Maintaining the criterion of an optimal packing density disallows large N -fold rings, because LH2 rings of fold number greater than 14 result in an inter-ring distance less than the physical minimum (Fig. 5, dashed line) and exceed steric contact with neighboring rings. In particular, the 14-, 15-, and 16-fold consecutive group is not possible under optimal packing density. Note an initial rise occurs in ρ and d because, for $N \leq 4$, $R \sim 21.8/2 \sim 10$ Å, respectively.

The excitation energy transfer rates and diffusion constants of the remaining consecutive groups 2-, 3-, and 4-fold and 8-, 9-, and 10-fold differ significantly. To demonstrate this we calculate the generalized Förster rate (Eq. 5), where we must now include the spectral overlap integral (the spectral overlap differs for each fold number so that comparison of $|J|^2$ alone would be invalid). In Fig. 6 we plot the rotationally averaged transfer rate k and the corresponding diffusion constant kd_c^2 as a function of the fold number N for the cases of fixed d and ρ . In the latter (solid lines), both the transfer rate and the diffusion constant increase as N increases. In the former (dashed lines), the transfer rate k increases and then plateaus for large N (due to absence of shortening inter-ring distance), whereas the diffusion constant increases for all N . In both cases, it is preferable to have rings of larger fold number, simply to increase the absolute transfer rate and/or diffusion constant. Thus, the 8-, 9-, and 10-fold consecutive group appears preferable to the 2-, 3-, and 4-fold group.

Finally, we consider the rotational sensitivity of the energy transfer between the LH2 rings (the rotational sensitivity is indicative of the roughness of the energy landscape seen by the migrating exciton). To this end we define the percentage difference ratio, $\delta = (|J|_{12\max}^2 - |J|_{12\min}^2) / |J|_{\text{av}}^2$, where $|J|_{\text{av}}^2 = (|J|_{12\max}^2 + |J|_{12\min}^2) / 2$. In Fig. 7, the percentage difference ratio for EET between two rings is shown as a function of the fold number N for fixed density (solid line). As the fold number increases, the sensitivity decreases to a minimum at $N = 10$, due to the increased aggregation of the heterodimers (this is complementary to reducing the degree of frustration). For larger fold numbers $N \geq 11$, however, the effect of shortening inter-ring distance (solid line in Fig. 5) dominates increasing aggregation and the rotational sensitivity increases so that the dependence of the EET rate on dipole orientation is minimized around the 8-, 9-, and 10-fold consecutive group. The absence of this effect in the case of fixed inter-ring distance (dashed line) removes this minimum. Hence the requirements of maintaining an optimal packing density and minimal rotational sensitivity are met exclusively by the 8-, 9-, and 10-fold consecutive group.

Discussion

We have investigated the role of the fold symmetry of LH2 in promoting maximum excitation energy transfer on the hexagonal lattice. Considering the related issues of symmetry and

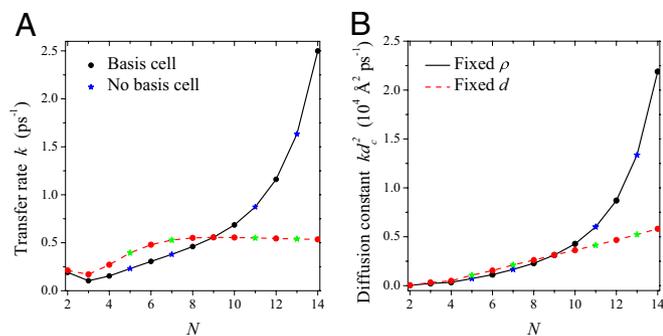


Fig. 6. (A) Excitation energy transfer rate k between two rings as a function of the fold number N for fixed packing density (solid line) and inter-ring distance (dashed line). (B) Corresponding diffusion constants kd_c^2 .

frustration, we conclude that the fold symmetry can promote maximum EET via two complementary mechanisms. First, the existence of an optimal basis cell allows optimization of the entire hexagonal domain, thus enhancing the network efficiency. Second, the type of basis cell can significantly reduce or remove the degree of EET frustration between basis cells across the lattice, thus smoothing the EET landscape and enhancing robustness. EET efficiency and robustness are precisely the design criteria of a functioning light-harvesting organism, e.g., the distance over which an exciton can migrate before relaxation has a direct consequence for the ratio of LH2 to LH1/reaction center complexes (required to capture the exciton for further use) in natural systems (1–4).

A key result is that the existence of an optimal basis cell and its type are directly related to the number of matching points S . Additionally, basis cells exist for groups of consecutive fold numbers, particularly the group of naturally occurring 8-, 9-, and 10-fold rings. Considering the inter-ring distance, EET rate, and diffusion constant, we have demonstrated how this group achieves minimal rotational sensitivity in addition to an optimal packing density. Minimal rotational sensitivity and optimal packing density facilitate the biological function of a pigment-protein complex in a light-harvesting role, where essentially isotropic, delocalized behavior is desired (26). In other words, minimization of the roughness of the energy landscape allows for robust EET, whereas an optimal packing density enhances the transfer efficiency. This essentially corroborates our finding that the existence and type of an optimal basis cell achieve global optimization and minimal frustration across the hexagonal lattice and, through their direct relation to S , suggests the design principle of matching the internal symmetry of the LH complex to the underlying lattice order.

All of the above considerations provide theoretical insight into the role of fold symmetry, arising at the molecular level, in promoting efficient and robust EET at the supramolecular level of the light-harvesting antenna. Recent experiments in synthetic light harvesting (21, 22), for example, have found that hexagonal arrays of 9-fold LH2 achieve diffusion constants greatly enhanced over those observed in LH2 domains in their natural membrane due to increased coupling of the complexes. Our design principle predicts that this enhancement should be relatively reduced for 8- and 10-fold rings due to lattice mismatch and increased EET frustration. It is hoped this work will inform photosynthetic, self-assembled materials, and lithographic communities striving to understand solar light harvesting, its limits, and its design principles.

Finally, we note that increasingly sophisticated techniques continue to yield exciting discoveries at the molecular level of LH2 (27). Although we have here considered only the EET

between individual LH2 complexes, thus ignoring the dynamics that occur within the complex, our qualitative conclusions should remain intact. For example, we know that neither static nor thermal disorder in an LH2 ring is strong enough to remove its excitonic character, so that the main contribution to the average transfer rate between two rings still comes from the lowest (formerly) degenerate eigenstates (even with disorder a multichromophoric FRET theory between rings must be used). Furthermore, although disorder quantitatively changes the average transfer rate between two rings, the average transfer rate will still be $2\pi/N$ periodic with respect to rotation and have a mirror symmetry phase of $\phi_N = \pi/N$ (N odd) and $\phi_N = 0$ (N even) (Eq. S7). We thus expect our conclusions regarding the fold symmetry mechanisms to hold in the disordered scenario. The extension of our understanding of both the structure and the dynamics of individual molecular complexes to their supramolecular function presents a significant challenge for the future.

Materials and Methods

The system Hamiltonian of two B850 rings may be written

$$H = \begin{pmatrix} H_D & H_c \\ H_c^\dagger & H_A \end{pmatrix}, \quad [3]$$

where H_D and H_A are the donor and acceptor Hamiltonians of the two individual rings as discussed in *SI Materials and Methods* (Eq. S2), H_c is the coupling Hamiltonian between the two rings $H_c = \sum_{m=1}^{N_D} \sum_{n=1}^{N_A} V_{m,n} \langle m|n \rangle$, and the inter-ring electronic coupling elements are simply $V_{m,n} = \langle m|H_c|n \rangle$ and are calculated using the dipole-dipole approximation.

Now, due to the weak electronic coupling between LH2 rings, the excitation energy transfer is incoherent. Thus, the hopping rate is adequately described by the multichromophoric Förster theory

$$k = \sum_{m,m'}^{N_D} \sum_{n,n'}^{N_A} \frac{V_{m,n} V_{m',n'}}{2\pi\hbar^2} \int_{-\infty}^{\infty} E_{m',m}^D(\omega) I_{n,n'}^A(\omega) d\omega, \quad [4]$$

where $E_{m',m}^D(\omega)$ and $I_{n,n'}^A(\omega)$ are the site basis elements of the donor emission and acceptor absorption spectra; i.e., $E_{m',m}^D(\omega) = \langle m'|E_D(\omega)|m \rangle$ and $I_{n,n'}^A(\omega) = \langle n|I_A(\omega)|n' \rangle$. The diffusion constant of excitons between rings is simply kd_c^2 , where d_c is the ring center-to-center distance. Now if we use the diagonal (secular) approximation in the eigenstate basis of the donor and acceptor

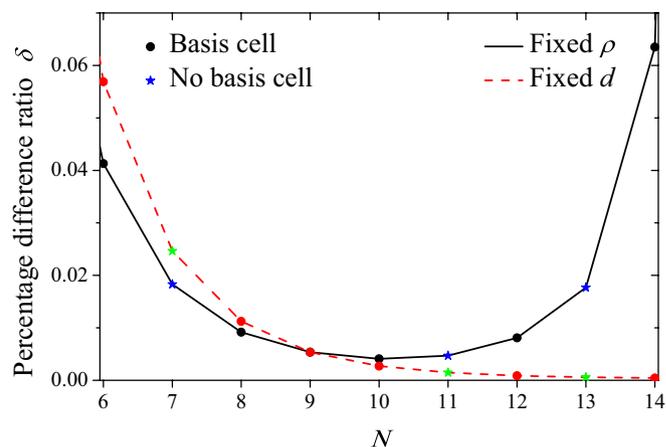


Fig. 7. Percentage difference ratio δ as a function of the fold number N for fixed density ρ (solid line) and fixed inter-ring distance d (dashed line). As the fold number increases, the rotational sensitivity obtains a minimum for 10-fold symmetry.

systems so that $\langle \varepsilon_\mu^D | E_D(\omega) | \varepsilon_\nu^D \rangle = E_\nu^D(\omega) \delta_{\mu,\nu}$ and $\langle \varepsilon_\mu^A | I^A(\omega) | \varepsilon_\nu^A \rangle = I_\nu^A(\omega) \delta_{\mu,\nu}$, we have the diagonal multichromophoric Förster rate

$$k = \frac{1}{2\pi\hbar^2} \sum_{\mu} \sum_{\nu} \left| \sum_{m=1}^{N_D} \sum_{n=1}^{N_A} V_{m,n} C_\mu^m C_\nu^{n*} \right|^2 \times \int_{-\infty}^{\infty} E_\mu^D(\omega) I_\nu^A(\omega) d\omega, \quad [5]$$

often referred to as the generalized Förster rate. Clearly excitation energy transfer can occur between two rings via any pair of eigenstates $|\varepsilon_\mu^D\rangle$ and $|\varepsilon_\nu^A\rangle$. However, noting that most of the total oscillator strength of a single ring is contained in its lowest degenerate eigenstates $|\varepsilon_2\rangle$ and $|\varepsilon_3\rangle$, we simplify our calculation by considering only excitation energy transfer between these two eigenstates so that we need only count the four terms given by $\mu=2,3$ and $\nu=2,3$ in Eq. 5. Furthermore, because these eigenstates are degenerate, we have $E_2^D(\omega) = E_3^D(\omega)$ and $I_2^A(\omega) = I_3^A(\omega)$ so that the overlap integral may be factorized out. Upon translation or rotation of a ring, the spectrum does not change, and hence the contribution from the spectral

overlap remains constant. Therefore, to find the optimal rotational positions of two B850 rings, we need only calculate the effective dipolar coupling term

$$|J|^2 \equiv \sum_{\mu=2}^3 \sum_{\nu=2}^3 \left| \sum_{m=1}^{N_D} \sum_{n=1}^{N_A} V_{m,n} C_\mu^m C_\nu^{n*} \right|^2 = |\langle \varepsilon_2^A | H | \varepsilon_2^D \rangle|^2 + |\langle \varepsilon_2^A | H | \varepsilon_3^D \rangle|^2 + |\langle \varepsilon_3^A | H | \varepsilon_2^D \rangle|^2 + |\langle \varepsilon_3^A | H | \varepsilon_3^D \rangle|^2, \quad [6]$$

where the second relation in terms of H from Eq. 3 is obtained by extending the dimension of $|\varepsilon_\mu^D\rangle$ and $|\varepsilon_\nu^A\rangle$ from $2N$ to $4N$ by adding zeros at the end and beginning of the eigenvectors, respectively.

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