



# Local protein solvation drives direct down-conversion in phycobiliprotein PC645 via incoherent vibronic transport

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The mechanisms controlling excitation energy transport (EET) in light-harvesting complexes remain controversial. Following the observation of long-lived beats in 2D electronic spectroscopy of PC645, vibronic coherence, the delocalization of excited states between pigments supported by a resonant vibration, has been proposed to enable direct excitation transport from the highest-energy to the lowest-energy pigments, bypassing a collection of intermediate states. Here, we instead show that for phycobiliprotein PC645 an incoherent vibronic transport mechanism is at play. We quantify the solvation dynamics of individual pigments using *ab initio* quantum mechanics/molecular mechanics (QM/MM) nuclear dynamics. Our atomistic spectral densities reproduce experimental observations ranging from absorption and fluorescence spectra to the timescales and selectivity of down-conversion observed in transient absorption measurements. We construct a general model for vibronic dimers and establish the parameter regimes of coherent and incoherent vibronic transport. We demonstrate that direct down-conversion in PC645 proceeds incoherently, enhanced by large reorganization energies and a broad collection of high-frequency vibrations. We suggest that a similar incoherent mechanism is appropriate across phycobiliproteins and represents a potential design principle for nanoscale control of EET.

photosynthesis | light harvesting | excitation energy transfer | quantum coherence | molecular dynamics

Understanding the solvation dynamics of multiple pigments in a heterogeneous medium is essential for being able to control excited-state dynamics in artificial materials. Natural photosynthesis demonstrates this capability in light-harvesting complexes (LHCs), which exert nanoscale control over excitation transport between pigments via the protein environment. Currently, there is no spectroscopic technique that can disentangle excitation dynamics in pigment aggregates from the many timescales of the vibrational bath, which include intramolecular pigment vibrations, fluctuations of neighboring amino acids, reorientation of proximate biological water, and reorganization of the bulk water solvent. Thus, quantitatively connecting the atomic-scale motions of the pigment–protein environment to the dynamics of excitation transport in LHCs remains a grand challenge at the interface of materials science and chemical physics.

Are there simple design principles that connect the complex atomic structure of pigment environments to the regulation of excitation energy transfer (EET) pathways in LHCs (1–5)? As demonstrated by the environmentally assisted quantum transport (ENAQT) mechanism (6), thermally occupied low-frequency vibrations can enhance transport between energetically detuned pigments and represent the simplest design principle connecting atomistic dynamics and excitation transport. The presence of a high-frequency vibration resonant with the excitation energy gap between pigments can also enhance transport. If the vibration supports coherence between the pigment excited states, *i.e.*, vibronic coherence, then excitation transport

occurs by transient delocalization; otherwise incoherent excitation transport occurs by discrete hopping. Persistent beat signals in nonlinear spectroscopic measurements of multiple LHCs have been interpreted as evidence for coherent vibronic transport (7–16). Phycobiliprotein PC645, in particular, has been studied extensively and has been suggested to leverage a coherent vibronic enhancement in the down-conversion process central to its physiological function (10). Here, we instead show that down-conversion in PC645 occurs via an incoherent vibronic mechanism.

*In vivo*, PC645 absorbs high-energy photons and down-converts the excitation to be resonant with the reaction center, where charge separation occurs. Transient absorption measurements on isolated PC645 reveal the process of down-conversion: Photoexcitation of the high-energy core dihydrobiliverdins (DBVs) (Fig. 1A, blue and green) is followed by transfer to the low-energy phycocyanobilin 82s (PCB82s) (Fig. 1A, red and cyan), skipping the energetically intermediate mesobiliverdins (MBVs) (Fig. 1A, brown and magenta) (17). Direct down-conversion cannot be explained by interpigment electronic couplings (18), suggesting that the vibrational dynamics arising from the local protein environments determine the pathways of EET.

## Significance

To harness sunlight for growth, plants and algae rapidly convey absorbed excitation energy from antennae pigments to a reaction center, where the excitations convert to chemical energy. In deep water environments, cryptophyte algae survive by using the molecular motion of phycobiliprotein antennae to funnel excitations to low-energy pigments. The mechanism of down-conversion in phycobiliproteins remains controversial: Could specific vibrations resonant with pigment energy gaps support coherence between excited states and thereby enhance the rate of transport by transient delocalization? Here, we demonstrate that down-conversion in a specific phycobiliprotein, PC645, is both incoherent and enhanced by a broad range of high-frequency vibrations. We suggest that a similar incoherent mechanism is appropriate across phycobiliproteins.

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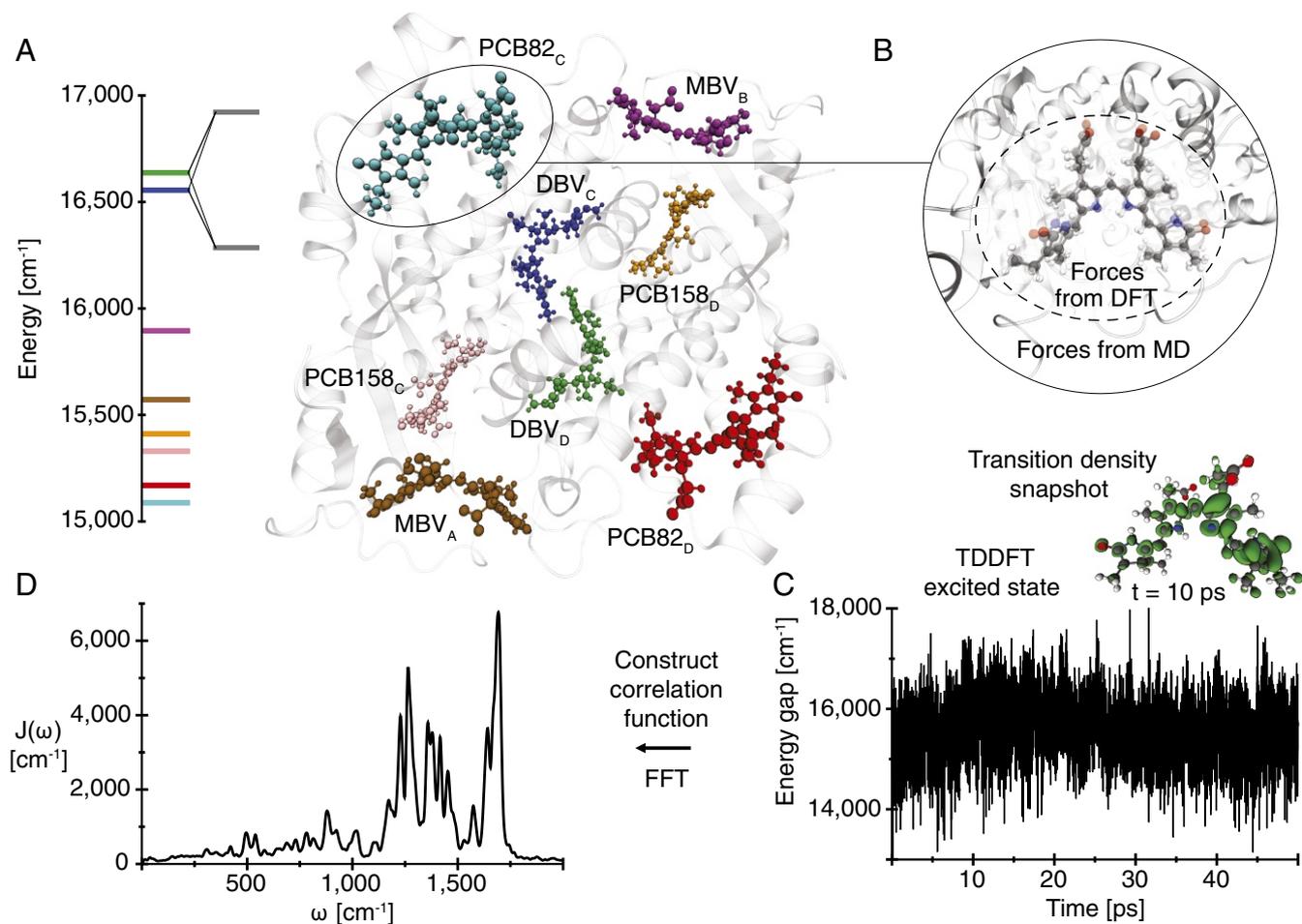
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**Fig. 1.** PC645 and computational methodology for spectral density construction. (A) Structure and reorganized energy levels of PC645, where we have defined a bilin color scheme that remains consistent throughout. Gray lines in the energy-level diagram are drawn schematically to represent the exciton eigenstates of the core DBVs (blue and green) due to the significant excitonic coupling of  $319.4 \text{ cm}^{-1}$ , more than three times larger than any other coupling in PC645. (B) For each bilin, we simulate 50 ps of mixed QM/MM nuclear dynamics in which that bilin is treated quantum mechanically, while the remaining protein is treated with a classical force field. (C) We construct energy-gap trajectories by calculating full TDDFT excited states for each bilin on geometries extracted at 2-fs intervals from our QM/MM trajectories. (D) We construct an energy-gap correlation function for each bilin and Fourier transform to obtain eight unique bilin spectral densities.

Quantifying the ultrafast vibrational dynamics of specific chromophores in LHCs remains a challenge for both theory and experiment. Using a combination of linear and nonlinear spectroscopic data, one can parameterize a model that describes the local vibrational environments of each pigment (19). However, multiple distinct vibrational environments quickly lead to an intractable number of free parameters. Atomistic nuclear dynamics simulations have the potential to bypass the inverse problem (20–23) but have not previously managed to quantitatively reproduce spectroscopic signals, thus restricting their ability to inform on biological function.

Here, we successfully connect unique pigment–protein vibrational environments to spectroscopic signals by incorporating pigment forces calculated with ground-state density functional theory (DFT). We extract bilin reorganization energies with a wider spread and larger average value than previously expected (10, 18, 24), allowing for quantitative reproduction of linear spectra in the absence of free parameters. Additionally, our EET simulations yield transfer pathways and timescales in good agreement with experimental transient absorption measurements. To address the underlying mechanism, we construct a representative vibronic dimer and establish the regimes of coherent and incoherent vibronic transport. We find that PC645 is far into the

incoherent parameter regime and support this conclusion by demonstrating that down-conversion in PC645 does not depend on the frequency, or even the presence, of the resonant vibrational mode previously suggested to support vibronic coherence. Our results demonstrate that transport between bilins in PC645 occurs incoherently and that Förster spectral overlaps tuned by local bilin–protein environments control EET and enable direct down-conversion. We further suggest that vibronic transport in phycobiliproteins, generally, is incoherent and enhanced by the presence of a broad collection of high-frequency, intramolecular vibrations.

## Results and Discussion

### Ab Initio Simulations of Protein Solvation Reproduce Linear Spectra.

To understand the role of the local pigment environments in controlling EET, we first characterize the atomistic origin of the bilin solvation dynamics. The excitation energy of a pigment bound in a protein pocket, given as the energy difference between the ground and first excited state, fluctuates due to the nuclear motions of the chromophore, the surrounding protein residues, and the proximate water. Simulating excitation energy fluctuations requires performing nuclear dynamics that sample the ground-state potential energy surface (PES) and calculating



lengths and either one or two covalent protein linkages. The variability of the reorganization energies between chemically identical bilins bound in distinct protein environments (e.g., MBV<sub>A</sub> and MBV<sub>B</sub>) demonstrates the importance of the protein scaffold in determining the local solvation dynamics. On the other hand, all eight bilin spectral densities show a peak near 1,650 cm<sup>-1</sup> which is consistent with the assignment of a long-lived 1,580-cm<sup>-1</sup> mode as intramolecular C = C or C = N vibrations in broad-band transient absorption measurements (10).

Incorporating the ab initio QM/MM spectral densities into absorption and fluorescence simulations (solid black lines) results in excellent agreement with experimental spectra (open gray circles), as shown in Fig. 2. We note that the negative features observed in simulated spectra are unphysical but do not impact the quality of the lineshapes (*SI Appendix*, Fig. S4). To account for both the large reorganization energies and the multiple timescales of vibrational relaxation encoded in the structure of our spectral densities, we use numerically exact hierarchical equations of motion (HEOM) (32, 33), as implemented in QMaster (34). HEOM has been shown to yield realistic simulations of exciton dynamics in LHCs (35–37), but its computational complexity limits our spectral densities to including a total of 24 Drude–Lorentz peaks for full-system simulations, each representing a distributions of bath modes. We construct four classes of abridged spectral densities for each bilin, starting with class 1 (eight or nine peaks per bilin) and successively coarse graining to incorporate fewer peaks until we reach class 4 (two peaks per bilin). Peak parameters for all spectral densities are given in *SI Appendix*, Tables S1–S4. Abridged spectral densities used for spectroscopic calculations are shown with our consistent bilin color scheme in the subpanels of Fig. 2. We justify our abridged spectral densities by comparing simulated monomer absorption and fluorescence spectra, as described in *SI Appendix*, section 2. *SI Appendix*, section 2 discusses the system Hamiltonian while *SI Appendix*, section 3 covers ab initio transition dipole moments, details of the fluorescence simulations, and inhomogeneous broadening.

We validate our spectral densities by comparing them to two features of the experimental spectra: the Stokes shift, defined as the frequency difference between the highest-energy fluorescence peak and the lowest-energy absorption peak, and the overall width of the absorption spectrum. Together, these observables are quite sensitive to the distribution of reorganization energies between the different bilins of PC645. Using the lowest-energy bilin spectral density for all pigments, such as would be extracted from a fluorescence line-narrowing experiment, results in an absorption spectrum with a full-width at half maximum (FWHM) that is 25% too small (*SI Appendix*, Fig. S5). The absorption spectrum contracts in this case because we underestimate the reorganization energies of the PCB158s and MBVs. If we instead use the average of the eight bilin spectral densities for all pigments, we obtain an absorption spectrum with a Stokes shift that is 55% too large (*SI Appendix*, Fig. S6). The Stokes shift expands because we substantially overestimate the reorganization energy of the PCB82s that dominate the fluorescence and low-energy absorption features. Thus, 18% error between the experimental and simulated Stokes shift and very good agreement with the overall absorption linewidth in the absence of free parameters represent a strong validation of the solvation dynamics encoded in our atomistic spectral densities.

**Protein Solvation Drives Down-Conversion.** The spectral density connects the atomistic dynamics of the bilin vibrational environment to both spectroscopic lineshapes and EET pertinent to light harvesting. Having validated our atomistic QM/MM spectral densities against absorption and fluorescence spectra, we now confirm that they also describe the experimental observables associated with direct down-conversion in PC645. Experimen-

tal transient absorption measurements, combined with global kinetic analysis, indicate that initial excitation of the DBV core is followed by direct transport to the low-energy PCB82 pigments with a rate of 1.7 ps<sup>-1</sup> (10, 17). Neither the rates nor selectivity of direct down-conversion have been successfully reproduced using previously estimated unstructured spectral densities (24). Here, we show that numerically exact HEOM simulations combined with our spectral densities predict pathways and rates of down-conversion that reproduce experimental results.

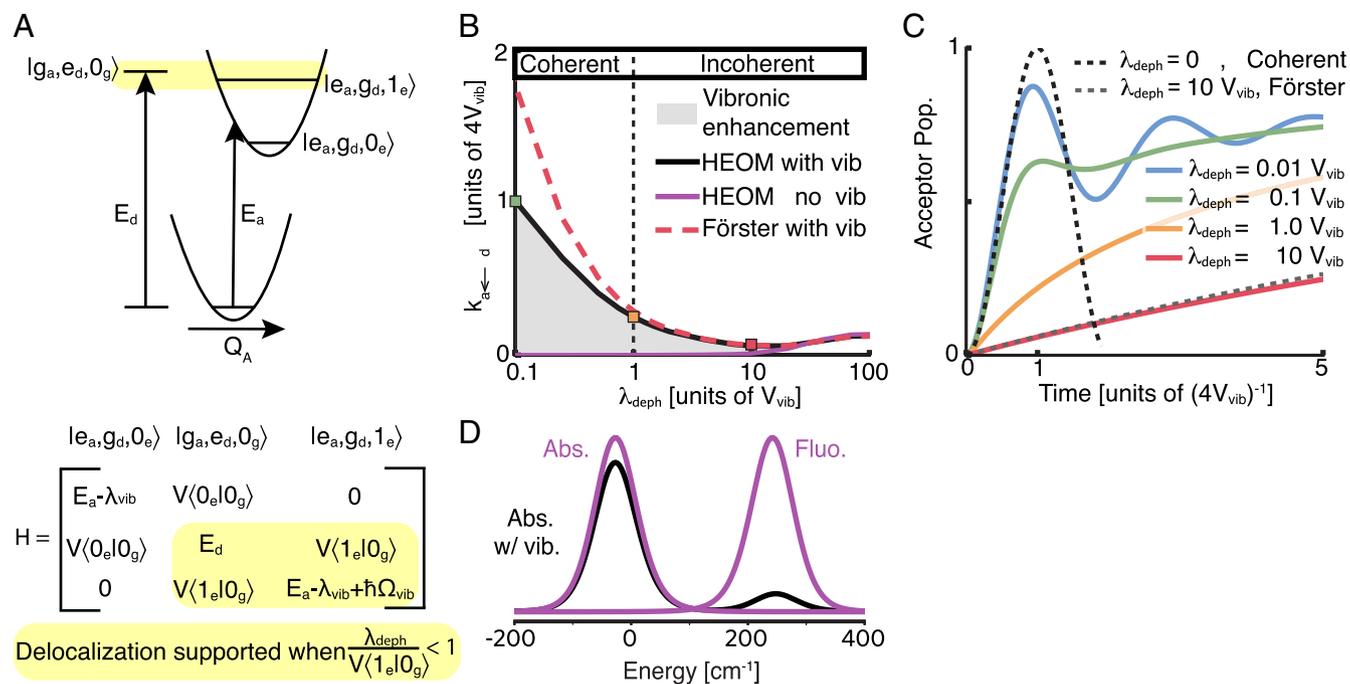
The large reorganization energies of our QM/MM spectral densities are essential for correctly predicting the selectivity of transport from the DBV core to the low-energy PCB82s. Simulated EET pathways following photoexcitation of the DBV core are shown in Fig. 3*A* and *B*. Arrow thickness in Fig. 3 is proportional to exciton flux, the net amount of excitation that is transferred from the DBVs to an acceptor pigment in a 1.0-ps interval following photoexcitation. We perform exciton flux calculations using class 2 spectral densities for the MBVs and class 4 spectral densities for the remaining pigments (*SI Appendix*, section 2). In Fig. 3*A* we rescale the spectral densities to match the average magnitude of reorganization energy used in previous simulations ( $\langle \lambda \rangle = 260$  cm<sup>-1</sup>, labeled 30%). In the presence of a smaller average reorganization energy, the more weakly solvated DBV core primarily transports excitation to the energetically adjacent MBV<sub>B</sub>, qualitatively reproducing previous results (24). EET simulations using the full reorganization energy (Fig. 3*B*, labeled 100%), however, show enhanced direct transfer to the low-energy PCB82s, in reasonable agreement with global kinetic analysis of transient absorption measurements.

In addition to influencing selectivity, the large reorganization energies also increase the rate of direct down-conversion. HEOM is a non-Markovian theory, and as a result, the rates of transport vary as a function of time in response to changes in the vibrational energy distribution. We extract a best-fit rate of down-conversion from HEOM simulations using class 1 spectral densities with a four-site model containing only the core DBVs and the low-energy PCB82s (*SI Appendix*, section 4). Class 1 spectral densities (e.g., *SI Appendix*, Fig. S1*A*) explicitly incorporate the high-frequency mode ( $\sim 1,650$  cm<sup>-1</sup>) previously assigned to an intramolecular bilin vibration. The rescaled spectral densities (Fig. 3*A*, 30%) result in a transport rate of 0.8 ps<sup>-1</sup>, substantially slower than the experimentally observed 1.7 ps<sup>-1</sup>. However, using the full reorganization energy of our spectral densities (Fig. 3*B*, 100%), we find the simulated rate of transport to be 1.6 ps<sup>-1</sup>, within 8% of the experimental value.

#### Down-Conversion Occurs via an Incoherent Vibronic Mechanism.

We have demonstrated that HEOM simulations using our atomistic spectral densities reproduce both the linear spectra and the rate of direct down-conversion in PC645. However, the question remains: What are the mechanistic principles connecting bilin vibrational environments to regulation of EET pathways? In particular, the relative importance of short-lived, low-frequency, intermolecular motions vs. long-lived, high-frequency, intramolecular vibrations remains controversial (1, 15). In this section, we assess how the mechanism of EET in PC645 arises from the interplay of electronic couplings between pigments with structured spectral densities. First, we introduce the simplest representative model system that captures the essential features of vibronic transport in the presence of a long-lived high-frequency vibration and distinguish between the coherent and incoherent transport regimes. Second, we demonstrate that PC645 experiences an incoherent vibronic mechanism and that the long-lived near-resonant vibration assigned to support vibronic coherence is not essential for efficient transport. Finally, we show that generalized Förster theory (38–41) explains the mechanism of transport in PC645, consistent with recent work on other PPC aggregates (34, 42–45), and that local protein





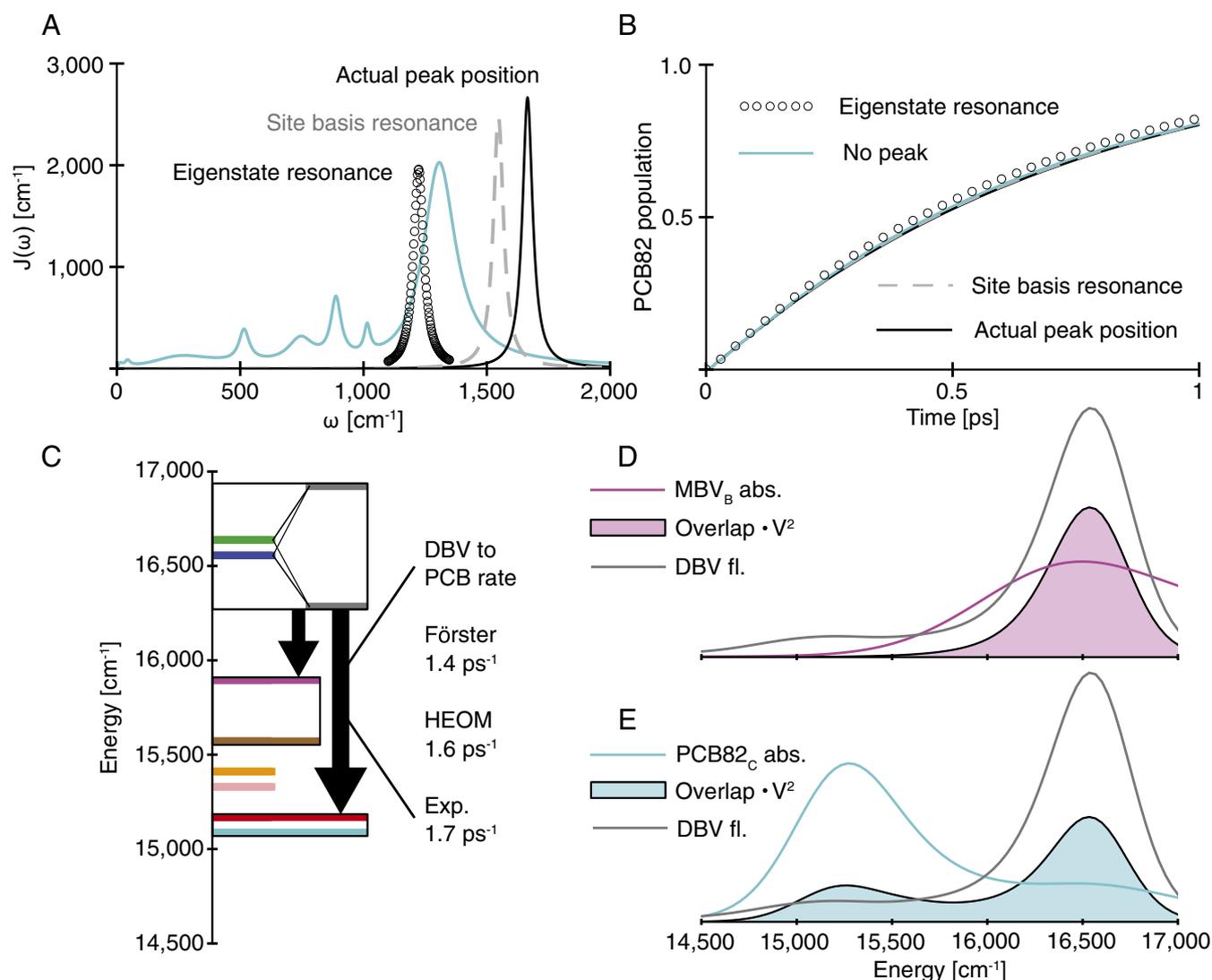
**Fig. 4.** Regimes of coherent and incoherent vibronic transport. (A) Energy levels of a detuned model dimer with a vibration on the acceptor pigment that is resonant with the site energy gap. The yellow shading represents the possible delocalization between the donor and the vibrationally excited acceptor states. (B) Rate of transport from donor to acceptor as a function of the reorganization energy of the low-frequency background ( $\lambda_{\text{deph}}$ ), where  $E_D - E_A = 250 \text{ cm}^{-1}$ ,  $V = 2.75 \text{ cm}^{-1}$ ,  $\lambda_{\text{vib}} = 24.7 \text{ cm}^{-1}$ ,  $\Omega_{\text{vib}} = 274.7 \text{ cm}^{-1}$ , and  $\gamma_{\text{vib}} = 0.125 \text{ cm}^{-1}$ . For the model dimer, our calculations use a vibrational frequency much lower than that found in PC645 to ensure that no negative populations are observed in the coherent regime (we check the validity of PC645 calculations in *SI Appendix, Fig. S4*). Calculated rates of transport are shown for HEOM with a high-frequency vibration (HEOM with vib, solid black line), HEOM without vibration (HEOM no vib, solid purple line), and Förster with vibration (Förster with vib, dashed red line). The solid gray area represents the vibronic enhancement arising from the presence of the high-frequency vibration. The green, orange, and red boxes represent the population dynamics shown in C. (C) The acceptor population dynamics following the donor excitation are plotted for  $\lambda_{\text{deph}} = 0.01, 0.1, 1,$  and  $10$  in blue, green, orange, and red, respectively. The first oscillation of purely coherent dynamics between the donor and vibrationally excited acceptor is plotted as a dashed black line. The population dynamics calculated using Förster theory for  $\lambda_{\text{deph}} = 10$  are plotted as a dashed gray line. (D) Mechanism of incoherent vibronic enhancement of transport, in which a nearly resonant vibration can generate a vibronic sideband in the absorption (black) that can enhance overlap with the fluorescence. Absorption and fluorescence lineshapes in the absence of a high-frequency vibration are shown in purple.

relevant vibronic coherence in PC645 (10). However, recent work has demonstrated that oscillatory vibronic signatures in 2D electronic spectra do not necessarily imply a role of vibronic coherence in excitation energy transfer (50). We directly probe the role of vibronic coherence in PC645 by manipulating the high-frequency vibration previously assigned to support vibronic signatures and demonstrate that it is not required for efficient transport. Coherent vibronic transport is known to exhibit a sharp resonance condition as a function of the vibrational frequency (7, 10). The resonance condition arises because vibronic delocalization between the donor and the vibrationally excited acceptor occurs only when the energy gap is smaller than, or the same order of magnitude as, the vibronic coupling (i.e.,  $V_{\text{vib}} \sim 12 \text{ cm}^{-1}$ ). In the case of PC645, partial delocalization in the DBVs means that there are two possible resonance conditions to consider, depending on whether the DBV core eigenstates or pigment states act as the donor. To explore the possible resonance conditions, we examine population dynamics calculated in a four-site model containing only the DBVs and PCB82s. We use class 1 spectral densities with four possible positions of the high-frequency mode (Fig. 5A): (i) the original positions ( $\sim 1,650 \text{ cm}^{-1}$ ), (ii) the energy difference between the DBV<sub>D</sub> and PCB82<sub>C</sub> sites, (iii) the energy difference between the low energy DBV exciton and the PCB82<sub>C</sub> site, and (iv) the complete removal of the high-frequency peak. We observe minimal differences in the resulting population dynamics (Fig. 5B). An additional exhaustive scan of the peak position between  $1,100 \text{ cm}^{-1}$  and  $1,800 \text{ cm}^{-1}$  shows no greater variations in popula-

tion dynamics. We have therefore demonstrated the absence of a sharp resonance condition, further supporting our assignment of incoherent vibronic transport.

Consistent with the incoherent transport regime, Förster theory captures the dominant contributions to the rate of down-conversion in PC645. To describe incoherent transport in the presence of strong coupling between DBVs, we use a generalization of Förster theory (38) that treats excitations within the DBV core as delocalized but assumes transport out of the core can be described as an incoherent hop. Due to rapid excitation relaxation within the DBV core, the rate of transport out of the core is determined by the overlap between the lowest-energy DBV exciton fluorescence and the absorption spectra of the remaining pigments. We simulate absorption spectra for the non-DBV pigments using Kubo lineshapes, which are exact for local excitations. We simulate the low-energy DBV exciton fluorescence using an extension of Kubo theory (38, 42, 43), which neglects the localization of excitons resulting from vibrational fluctuations (*SI Appendix, Fig. S10*). Generalized Förster theory predicts a rate of  $1.4 \text{ ps}^{-1}$ , within 15% of the HEOM result (Fig. 5C).

Using generalized Förster theory, we can explain the relative mechanistic role of low-frequency, intermolecular vibrations and high-frequency, intramolecular vibrations in controlling direct down-conversion in PC645. To understand the impact of bilin vibrational environments on transport rates and pathways, we examine the absorption/fluorescence overlap between the DBV core (Fig. 5D and E, gray lines) and either PCB82<sub>C</sub> (Fig. 5E,



**Fig. 5.** Direct down-conversion in PC645 is driven by incoherent vibronic transport. (A) Four different PCB82<sub>c</sub> spectral densities that we use to examine the impact of the high-frequency vibration on direct down-conversion. While cyan represents the case where the high-frequency vibration has been removed entirely, it is also implicitly added to the following three cases in which the high-frequency vibration is shown by itself for clarity: the vibration in its original position in our QM/MM spectral densities (black), the vibration shifted to be in resonance with the DBV<sub>B</sub> – PCB82<sub>c</sub> site energy difference (1,550 cm<sup>-1</sup>, dashed gray line), and the vibration shifted to be in resonance with the low-energy DBV exciton – PCB82<sub>c</sub> energy difference (1,220 cm<sup>-1</sup>, black open circles). Note that all four sites present in our population dynamics simulations use their individual spectral densities, but the changes to the high-frequency peak are equivalently applied to the other three sites. (B) Population dynamics of a four-site system containing the DBVs and PCB82s. Lines representing the sum PCB82 population are labeled by their representation in A. (C) An energetic depiction of down-conversion where site energies are fully reorganized and arrows represent the combined flux from the DBV core to the MBVs or PCB82s. The flux arrow from DBVs to PCB82s is labeled with the effective transport rate obtained from theory and experiment. Note that excitons are drawn schematically to represent DBV delocalization. (D) Kubo monomer absorption for MBV<sub>B</sub> (magenta), Kubo DBV core fluorescence (gray), and absorption/fluorescence overlap weighted by the coupling to the lowest-energy DBV exciton (outlined magenta). (E) Kubo monomer absorption for PCB82<sub>c</sub> (cyan), Kubo DBV core fluorescence (gray), and absorption/fluorescence overlap weighted by the coupling to the lowest-energy DBV exciton (outlined cyan).

cyan) or MBV<sub>B</sub> (Fig. 5D, magenta). The Stokes shift of the DBV core ( $\sim 280$  cm<sup>-1</sup>) enhances the energetic overlap with the lower-energy pigments and dramatically increases the overall rates of transport. Despite almost perfect alignment of the DBV fluorescence with the MBV absorption spectra, excitation is preferentially transported to PCB82s due to the presence of broad vibronic sidebands. In contrast to the narrow resonance condition associated with coherent vibronic transport (width on the order of  $|V\langle 1_c | 0_g \rangle|$ ), the existence of wide absorption and fluorescence features allows for incoherent vibronic enhancement with a broad resonance condition (width on the order of the homogeneous linewidth). Finally, we note that given the

importance of the Stokes shift in determining the rate and pathways of excitation transport, the demonstration of modified local solvation dynamics for chemically identical bilins within PC645 represents a potential design principle for nanoscale control of EET.

### Concluding Remarks

Here, we have demonstrated how specific features of the pigment vibrational environments in PC645 exert nanoscale control of EET pathways and enable direct down-conversion from the DBV core to the low-energy PCB82s. We accessed the atomistic origin of local solvation for each bilin, using ab initio QM/MM nuclear

dynamics. In contrast with the previous assumption of identical baths, our simulations identified substantial disparities between bilin vibrational environments which were essential to reproducing the experimental absorption and fluorescence spectra. Further, we have determined that down-conversion in PC645 proceeds via an incoherent vibronic transport mechanism where (i) excitations are localized on individual bilins (except for the DBV core) and transport occurs via incoherent hops and (ii) direct down-conversion is enhanced by large reorganization energies and the presence of a wide collection of high-frequency vibrations that induce broad vibronic sidebands. Given similar bilin flexibility and weak electronic couplings generally reported for biliproteins, our findings suggest that vibronic transport in cryptophyte algal antenna complexes proceeds incoherently.

The incoherent vibronic mechanism assigned here to PC645 is far more robust to imperfections than its coherent counterpart and could act as a blueprint for the design of artificial excitonic materials. Recent advances in biomimetic light-harvesting technologies have demonstrated novel architectures, e.g., metal-organic frameworks (MOFs) (51) and DNA origami (52), that can precisely place pigments within a hierarchically organized assembly. To fully realize nanoscale control of EET, we must iteratively simulate and design the local vibrational environments of pigment-scaffold architectures. However, the complexity of our current procedure is infeasible for high-throughput application, pointing toward the need for new and more efficient computational strategies and approximations.

## Materials and Methods

We perform eight QM/MM nuclear dynamics simulations where the forces on one bilin are constructed from ground-state DFT calculations for each trajectory. We begin each QM/MM trajectory with 10 ps of equilibration

followed by 40 ps of production run. We construct energy gap trajectories for each bilin, using TDDFT calculations on geometries sampled every 2 fs, for a total of 20,000 geometries per bilin. We use the energy gap trajectories to construct two-time correlation functions and then Fourier transform to obtain the spectral density for each bilin. Overall, the eight 50-ps QM/MM trajectories took over 9 mo to run and cost more than 2 million central processing unit (CPU) hours. The details of the initial geometry preparation, separation of the quantum from classical regions, excited-state energy calculations, and the construction of the spectral densities are described in [SI Appendix, section 1](#).

Exciton dynamics simulations were performed with the QMaster software package (34) that provides a high-performance implementation of HEOM, which runs flexibly on both graphics processing unit (GPU) and CPU architectures (13, 34, 53). HEOM fluorescence calculations used a development version of QMaster. All HEOM results presented were run at a hierarchy depth of six except for rate calculations, which were run at a hierarchy depth of five ([SI Appendix, section 4](#)).

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