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Disentanglement of excited-state dynamics with implications for FRET measurements: two-dimensional electronic spectroscopy of a BODIPY-functionalized cavitand†

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Förster Resonance Energy Transfer (FRET) is the incoherent transfer of an electronic excitation from a donor fluorophore to a nearby acceptor. FRET has been applied as a probe of local chromophore environments and distances on the nanoscale by extrapolating transfer efficiencies from standard experimental parameters, such as fluorescence intensities or lifetimes. Competition from nonradiative relaxation processes is often assumed to be constant in these extrapolations, but in actuality, this competition depends on the donor and acceptor environments and can, therefore, be affected by conformational changes. To study the effects of nonradiative relaxation on FRET dynamics, we perform two-dimensional electronic spectroscopy (2DES) on a pair of azaboraindacene (BODIPY) dyes, attached to opposite arms of a resorcin [4]arene cavitand. Temperature-induced switching between two equilibrium conformations, *vase* at 294 K to *kite* at 193 K, increases the donor–acceptor distance from 0.5 nm to 3 nm, affecting both FRET efficiency and nonradiative relaxation. By disentangling different dynamics based on lifetimes extracted from a series of 2D spectra, we independently observe nonradiative relaxation, FRET, and residual fluorescence from the donor in both *vase* to *kite* conformations. We observe changes in both FRET rate and nonradiative relaxation when the molecule switches from *vase* to *kite*, and measure a significantly greater difference in transfer efficiency between conformations than would be determined by standard lifetime-based measurements. These observations show that changes in competing nonradiative processes must be taken into account when highly accurate measurements of FRET efficiency are desired.

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Introduction

Förster Resonance Energy Transfer (FRET) moves an excitation nonradiatively from a donor fluorophore to a nearby acceptor *via* dipole–dipole interactions.¹ FRET can be elegantly described by readily accessible experimental parameters, such as the distance between chromophores (R), the fluorescence spectrum of the donor, and the absorption spectrum of the acceptor.² The

mechanism's well known R^{-6} distance dependence has led to widespread application^{3,4} of FRET as a spectroscopic measure of distances on the single nanometer scale.^{5,6} In particular, FRET is often used to probe interactions between biomolecules such as proteins^{6–10} and DNA^{11–14} and to image those interactions within cells.^{15–21} An extensive array of donor–acceptor pairs has been developed for these purposes.^{8,12,22–28}

In Förster's model, the coupling between the donor and acceptor electronic states is treated as a second-order perturbation in the electronic coupling. The interaction is governed by the transition dipole moments of the excited electronic states. The coupling is sufficiently weak (and the transfer rate therefore sufficiently slow) that the donor relaxes to thermal vibrational equilibrium in its electronic excited state before transfer can occur. Although no absorption or emission occurs in the transfer itself (FRET is nonradiative), the same transition dipoles that are responsible for donor emission and acceptor absorption also drive the transfer, so resonance can be determined by absorption and emission spectra, and transfer competes with donor fluorescence and nonradiative relaxation.

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Experimentally, FRET efficiency can be determined by photon counts of donor or acceptor fluorescence or by changes in the donor's decay lifetime.^{17,29} The latter approach is often preferred due to relative immunity to cross-talk and concentration artifacts,^{7,16,22} and is based on kinetic competition of decay processes. In this scenario, efficiency can be expressed as

$$E = 1 - \tau_{DA}/\tau_D = \frac{1}{1 + (R/R_0)^6} \quad (1)$$

where τ_D and τ_{DA} are measured lifetimes of the donor alone and, in the presence of the acceptor, R is the donor-acceptor distance, and R_0 is the distance at which the transfer rate is equal to τ_D , corresponding to 50% transfer.^{9,18} The donor lifetime depends on the relative rates of competing processes as

$$1/\tau_D = k_D = k_{D,f} + k_{D,nr} \quad (2a)$$

$$1/\tau_{DA} = k_{DA} = k_{D,f} + k_{D,nr} + k_{tr} \quad (2b)$$

where $k_{D,f}$, $k_{D,nr}$, and k_{tr} correspond to rates of fluorescence, nonradiative relaxation, and energy transfer. Lifetime-based FRET determinations of this nature rely on assumptions that all processes involved are in kinetic competition, and that processes other than FRET, *i.e.* nonradiative relaxation, behave similarly for the donor regardless of the donor-acceptor distance.³⁰ However, local solvation conditions and system-bath coupling can change as molecular conformation changes. Such changes break the assumptions above and frustrate accurate measurement of FRET efficiency. We demonstrate this scenario using 2D electronic spectroscopy and show how conformation-dependent nonradiative processes affect FRET measurements.

To investigate the interactions between nonradiative relaxation and FRET, we sought a pair of chromophores that undergo both excited-state processes. Moreover, we sought a means of exerting control over both the distance separating the chromophore pair and the vibrational bath that defines the local environment of each chromophore. Quinoxaline-bridged resorcin[4]arene cavitands emerged as ideal molecular scaffolds for fulfilling the latter requirement, as these conformational switches can be toggled between a pursed *vase* state and a ringent, *kite* state using various external stimuli, including temperature.^{31,32} Due to solvation-related entropic effects, the cavitand exists in the *vase* conformation at room temperature and above; conformational switching commences at *ca.* 223 K, and at 193 K, only the *kite* conformation is observed.³³ To adapt this scaffold for our studies, we sought to append each of a pair of chromophores to opposing walls of the cavitand.^{34–36} We selected a pair of arylethynyl-functionalized azaborindacene (BODIPY) chromophores for this purpose because they exhibit the requisite photophysical properties,^{37–39} and because both a donor and an acceptor chromophore are conveniently accessible from a single, divergent synthetic route.⁴⁰

Given the ability of 2DES measurements to elucidate ultrafast transfer and relaxation dynamics in both biological^{41–43} and synthetic systems,^{44–46} we sought to study the BODIPY-functionalized cavitand using this technique. By separating the signal based on time dynamics, we disentangle three

relaxation pathways (nonradiative relaxation, energy transfer, and residual fluorescence) and isolate their lineshapes from the donor in both conformations (*kite* at 193 K and *vase* at 294 K). Based on our signal isolation analysis, we observe that the contribution of nonradiative relaxation to the overall donor decay changes with conformation. We show that, due to this change in nonradiative relaxation, fluorescence lifetime measurements underreport the transfer efficiency difference between *kite* and *vase* conformations, highlighting the need to test assumptions about nonradiative processes when measuring FRET in complex environments.

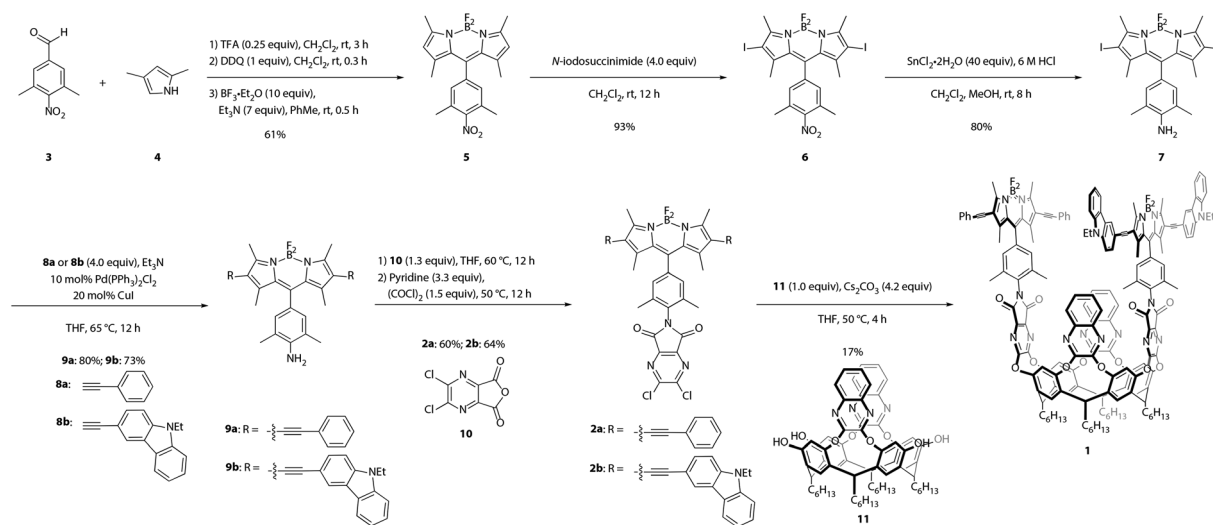
Results and discussion

Synthesis, variable-temperature characterization, and computed structures of the BODIPY-functionalized resorcin[4]arene cavitand

Synthesis of the target cavitand **1**, a new compound, was based on various precedents in the BODIPY and resorcin[4]arene literatures as indicated below. The process commenced with preparation of BODIPY-functionalized walls **2a** and **2b** (Scheme 1). Condensation of commercially available nitrobenzaldehyde **3** with 2,4-dimethylpyrrole (**4**) provided the nitro-BODIPY core **5**.⁴⁷ Treatment of nitro-BODIPY **5** with 4 equiv. *N*-iodosuccinimide yielded diiodo-BODIPY **6**, which was subsequently converted to anilino-BODIPY **7** using stannous chloride to selectively reduce the nitro moiety without affecting the iodides.⁴⁸ Sonogashira cross-coupling of anilino-BODIPY **7** with phenylacetylene (**8a**) yielded donor dye **9a**; acceptor dye **9b** was prepared analogously using *N*-ethylcarbazoyl alkyne **8b**.⁴⁰ Condensation of donor dye **9a** and acceptor dye **9b** with anhydride **10** afforded walls **2a** and **2b**, respectively.³⁴ Final assembly of cavitand **1**, which involves four nucleophilic substitutions, was achieved in a one-pot procedure with 17% yield, wherein a mixture of tetrol **11** and both walls, **2a** and **2b**, were treated with cesium carbonate. Target cavitand **1** was purified from the resultant mixture using flash column chromatography, followed by recycling gel permeation chromatography. The successful installation of two different walls on tetrol **11** in one pot to access cavitand **1** represents an alternative to previously reported methods for synthesizing non-symmetrically-substituted resorcin[4]arene cavitands. The one-pot reaction obviates the need for multiple purification steps which reduce overall yield.³⁴

Temperature-dependent conformational behavior of cavitand **1** was characterized by variable-temperature ¹H NMR,³³ absorption, and fluorescence spectroscopies.³⁴ As shown in Fig. 1, the ¹H NMR spectrum of cavitand **1** at 323 K exhibits two sets of sharp multiplets between 5.7 and 5.8 ppm, corresponding to the resorcin[4]arene methine protons (★) and indicative of the *vase* conformation. Cooling the sample to 223 K resulted in a diagnostic upfield shift of these methine proton signals to *ca.* 3.8 ppm due to increased shielding from the proximity of the aromatic wall flaps in the *kite* conformation. Consistent with previous reports, the *vase-to-kite* transition was also observed upon addition of acid (see Fig. S27 in the ESI†).³⁴



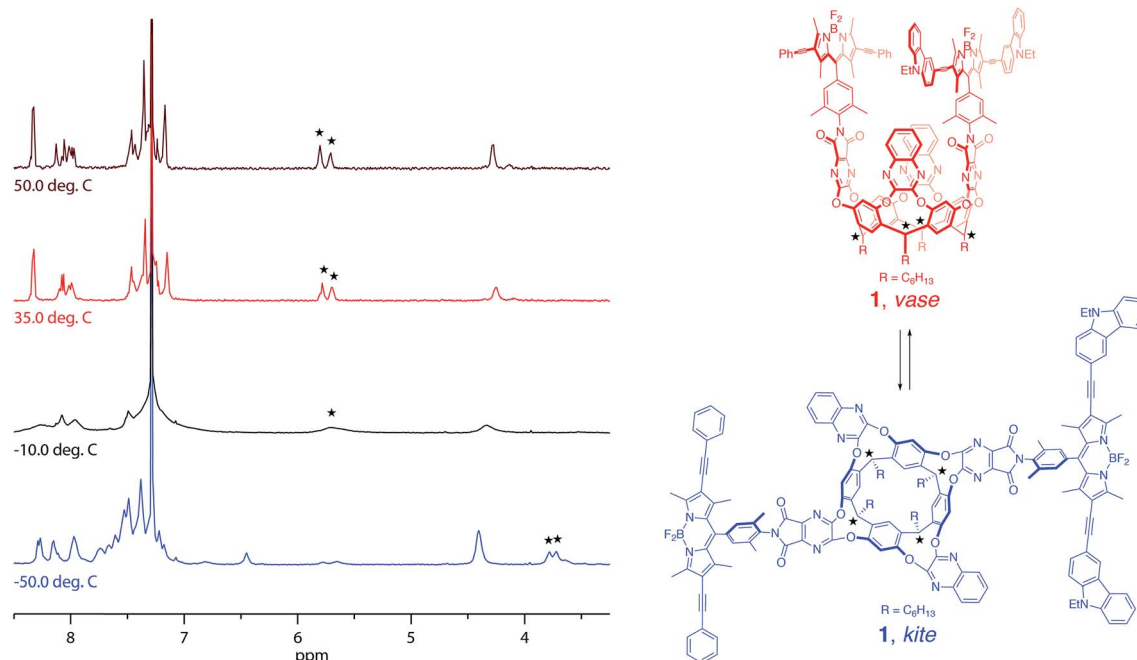


Scheme 1 Synthesis of cavitaand 1.

The temperature-dependent conformational switching of cavitaand 1 was corroborated by fluorescence and absorption spectroscopies, as depicted in Fig. 2. The *vase* conformation of cavitaand 1 predominates at 294 K (Fig. 2a), evidenced by a blue-shift of the absorption maximum of cavitaand 1 relative to that of the dye mixture; interactions between the donor and acceptor dyes in the *vase* conformation of cavitaand 1 result in shifts of their ground- and excited-state energies. At 193 K, the absorption spectrum of cavitaand 1 resembles that of the dye mixture, which is consistent with the weak interaction between the donor and acceptor dyes in the *kite* conformation (Fig. 2b). Fluorescence spectra of cavitaand 1 at 294 K and 193 K (excitation at 525 nm for both conformations) show a clear increase in

donor emission at 610 nm with a concurrent decrease in acceptor emission at 670 nm as the temperature is lowered. These fluorescence changes indicate a decrease in FRET efficiency and an increase in the distance between the donor and acceptor dye at lower temperature, which is consistent with conversion of *vase* to *kite* with decreasing temperature.

Computational studies provided further structural insights into the two conformations of cavitaand 1. The geometries of the *vase* and *kite* conformations were optimized using density functional theory calculations (B3LYP/3-21G(d)) in the Gaussian 09 software package,⁴⁹ and are depicted in Fig. 2c and d, respectively. In both conformations, terminal C₆H₁₃ groups on the cavitaand base have been replaced with CH₃. In the *vase*

Fig. 1 ¹H NMR spectra of cavitaand 1 in CDCl₃ at various temperatures.

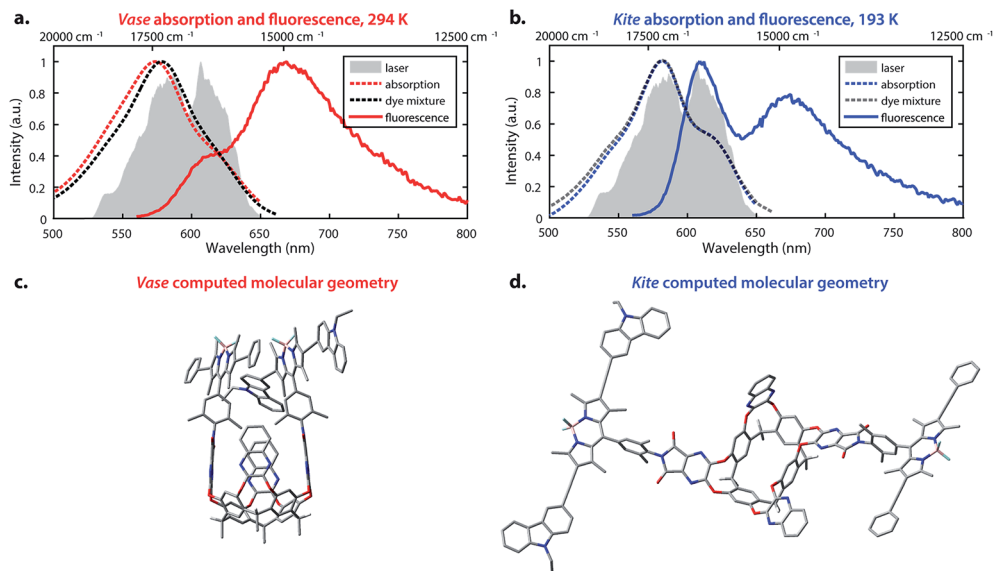


Fig. 2 Optical properties and corresponding calculated structures of the vase and kite conformations of cavitand 1. (a) Absorption (red dashed line) and fluorescence (solid line, excitation at 525 nm) of cavitand 1 in the vase conformation at 294 K; absorption of donor–acceptor mixture, 1 : 1 molar ratio at 294 K (black dashed line); laser spectrum (filled grey area). (b) Absorption (blue dashed line) and fluorescence (solid line, excitation at 525 nm) of cavitand 1 in the kite conformation at 193 K; absorption of donor–acceptor mixture, 1 : 1 molar ratio at 193 K (dark grey dashed line); laser spectrum (filled grey area). (c) Optimized geometry of the vase conformation calculated at the B3LYP/3-21G(d) level of theory. (d) Optimized geometry of the kite conformation calculated at the B3LYP/3-21G(d) level of theory. In (c) and (d), terminal C₆H₁₃ groups have been replaced with CH₃.

conformation, the donor and acceptor are sterically constrained to an orientation that promotes dipole–dipole coupling. The donor–acceptor distance in the vase conformation is *ca.* 0.5 nm, which is too close for independent solvation of the two chromophores, suggesting that they interact vibrationally, as well as electronically. In contrast, the donor and acceptor are separated by *ca.* 3 nm in the kite conformation, providing the chromophores with relative orientational freedom. Although the donor and acceptor are separated by less than R_0 , they interact weakly enough that the approximations made by FRET theory accurately describe their interaction. Having successfully synthesized cavitand 1 and confirmed that its conformation and transfer dynamics could be externally controlled by temperature, we next sought to probe the excited-state dynamics of cavitand 1, focusing on the interactions between nonradiative relaxation and FRET processes.

Two-dimensional electronic spectroscopy

Two-dimensional electronic spectroscopy (2DES) is a four-wave mixing technique that uses three ultrafast pulses at controlled time delays to interact with a sample, generating a third-order nonlinear signal. Two frequency axes, ω_τ and ω_t , are produced *via* Fourier transformation over the first and third time delays (coherence time, τ , and rephasing time, t). These axes represent the frequencies of excitation and detection, respectively, and correspond to the horizontal (ω_τ) and vertical (ω_t) axes on all 2D spectra shown in this paper. Plotted as the change in transmitted light (ΔT), positive features can be attributed to stimulated emission (SE) or to ground state bleach (GSB; a decrease in absorption caused by depletion of the ground state population).

Negative features are due to excited state absorption (ESA). Signal recorded at a series of waiting times (T) can reveal the time dynamics of the interrogated sample over 5–6 orders of magnitude, ranging from femtoseconds to nanoseconds.

2D spectra are frequency–frequency correlation maps, describing the optical response of a system (recorded on ω_t axis) at a series of waiting times, T , after initial excitation at ω_τ . Signals along the diagonal indicate features detected at the same energy at a series of waiting times, T , after initial excitation, while signals above or below the diagonal indicate coupling or population transfer between excited states. Additionally, the lineshapes of 2D signals provide information on the mechanisms that lead to broad spectral signals.⁵⁰ Inhomogeneous broadening (extension along the diagonal) gives insight into static variation in chemical environments, which perturb the excitation energies of individual chromophores.⁵¹ In contrast, homogeneous broadening (which determines antidiagonal linewidth) occurs when frequency fluctuations due to bath interactions occur quickly compared with the experimental timescale, such that an individual chromophore experiences all configurations of its local environment during a given measurement and thus represents the entire ensemble.⁵² In the homogeneous limit, one can estimate the dephasing time of a transition (the time required to lose phase correlation in the ensemble of sample molecules) by its homogeneous linewidth.⁵³

Two-dimensional electronic spectra of the BODIPY-functionalized cavitand

We performed 2DES measurements on the BODIPY-cavitand dimer at both 193 K and 294 K in two sets of waiting times—one from –20 to 1000 fs with a 10 fs step size, and the other



from -5 to 800 ps with a 5 ps step size. Representative, real-valued 2D spectra for the *kite* and *vase* conformations of the dimer are shown in Fig. 3. Four logarithmically-spaced time series and highlighted time points are illustrated for comparison. The 2D spectra of each conformation in Fig. 3c are normalized to the maximum of 2D spectrum at $T = 0$ fs in both datasets, and the frames in the highlighted spectra (Fig. 3a, b, d, and e) are normalized individually.



Fig. 3 Representative real-valued phased 2DES spectra for *vase/kite* conformation. (a) 2DES spectra of *vase* at 20 fs waiting time. (b) 2DES spectra of *vase* at 260 ps waiting time. Highlighted points are $17\,100\text{ cm}^{-1}$ (585 nm) diagonal (white star) and crosspeak (black star) at the same ω_{τ} ($17\,100\text{ cm}^{-1}$, 585 nm) and ω_{ϵ} of $16\,400\text{ cm}^{-1}$ (610 nm). (c) Time series of 2D spectra ranging 0 – 1000 fs and 5 – 800 ps for both *vase* and *kite*, normalized to $T = 0$. (d) 2DES spectra of *kite* at 20 fs waiting time. (e) 2DES spectra of *kite* at 260 ps waiting time, with same highlights as in (b). Spectra in (a) and (d) are normalized independently for the two conformations at 20 fs, while spectra (b) and (e) are normalized independently for the two conformations at 260 ps. The growth of the cross peak (black star, $17\,100\text{ cm}^{-1}$ in ω_{τ} and $16\,400\text{ cm}^{-1}$ in ω_{ϵ}) corresponds to donor relaxation to donor fluorescence as well as FRET transfer to acceptor absorption during waiting time.

At $T = 20$ fs, 2D spectra of both *vase* (Fig. 3a) and *kite* (Fig. 3d) show an inhomogeneously broadened peak along the diagonal at roughly $17\,100\text{ cm}^{-1}$. This signal corresponds to the initial absorption of the donor; at 20 fs the system has not yet relaxed past this starting point. Vibrational relaxation (aka Stokes shift) from this initial absorption occurs within the first 100 fs, leading to the below-diagonal cross peak in all subsequent spectra (see ESI† for more detail).

At $T = 260$ ps (Fig. 3b and e), two dominant features are present in both 2D spectra—a diagonal peak at $17\,100\text{ cm}^{-1}$ (white star) and a crosspeak at $17\,100\text{ cm}^{-1}$ (585 nm) in ω_{τ} and $16\,400$ (610 nm) cm^{-1} in ω_{ϵ} (black star). The diagonal feature at $T = 260$ ps is due to ground state bleach on the donor, because Stokes shift from the initial excitation has completed by this time. The below-diagonal crosspeak corresponds primarily to stimulated emission from the Stokes-shifted excited state of the donor. Along the anti-diagonal direction, all features are very broad, indicating the homogeneity of excited donor molecules. The width of these homogeneously broadened features is caused by rapid dephasing on the scale of tens of fs (Fig. S30†). A red shift in early time dynamics of the complex occurs on the same timescale, (see Fig. S31†) and the processes driving the red shift may also be the cause of this rapid dephasing. Further discussion of dephasing lifetimes and ultrafast dynamics can be found in the ESI.†

Decay-associated spectra

Decay-associated spectra (DAS) are generated from global fitting of the decay dynamics of each pixel in a 2D spectrum, and quantifying the contributions of various pathways to the overall signal. This and similar approaches have previously been applied to 2D spectra by the Engel group⁵⁴ and others.^{55–58} We performed DAS analysis on the long waiting time (5 ps resolved) 2D data, specifically from 5 ps to 800 ps.

The waiting time decay dynamics were fit to the following equation

$$S(T) = A_1 e^{-T/\tau_1} + A_2 e^{-T/\tau_2} + A_3 \quad (3)$$

where τ_1 and τ_2 are globally optimized decay constants for different transfer or relaxation processes (ESI Table TS3†). The A_i values correspond to the contribution of each process with A_3 corresponding to the contribution of signals with decay constant $\tau_3 \gg 800$ ps (our experimental timescale). A detailed description of how the maps are generated is included in the ESI.†

The left (grey) column in Fig. 4 shows a fast (~ 25 ps) decay for cavitand **1** in the *vase* (294 K , top) and *kite* (193 K , bottom) conformations. These lifetimes are consistent with reports of a decay in the 15 – 30 ps range for analogous *meso*-aryl substituted boron dipyrroin structures.^{37,38} We assign this fast decay to a nonradiative, vibrationally driven relaxation arising from a combination of flexing in the dipyrroin backbone and rotation of the aryl group, as suggested by Li *et al.*³⁸ and Kee *et al.*³⁷ The multi-exponential nature of the decay means that this process (corresponding to the fastest exponential decay) does not compete kinetically with the processes observed in the



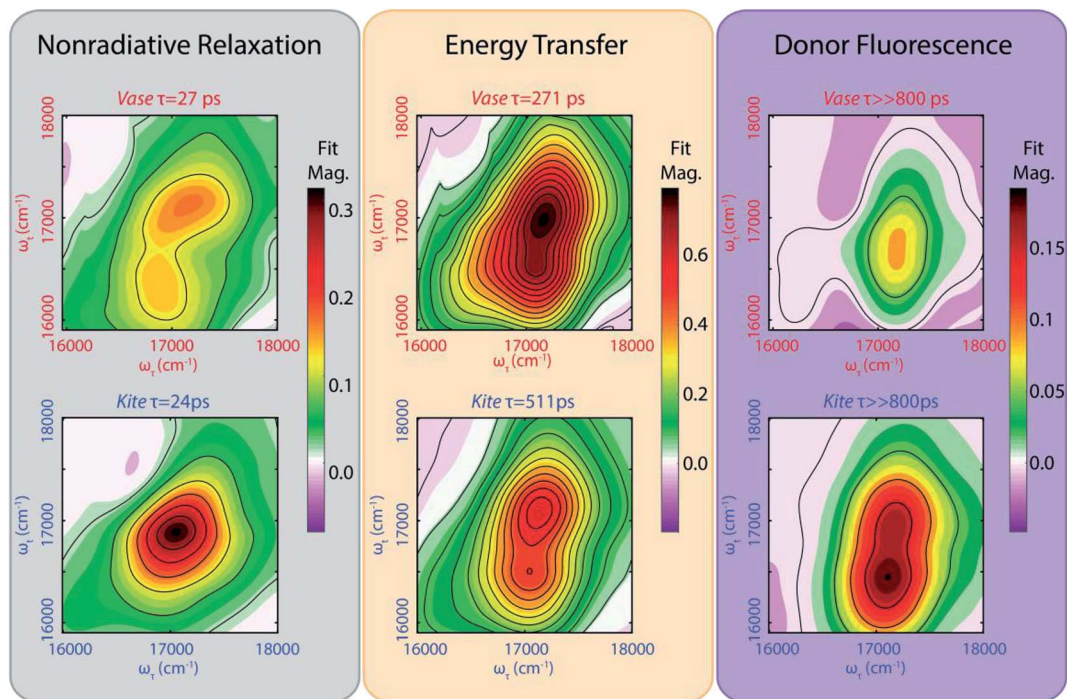


Fig. 4 Decay-associated spectra (DAS) reveal lineshapes of decay dynamics. DAS demonstrate relative distribution and strength (indicated by colorbar) of features decaying at the labeled rate. Sum of all DAS for a given conformation equals the normalized real-valued 2D spectra at $T = 5$ ps. The grey, yellow, and purple columns correspond to nonradiative relaxation, energy transfer (FRET), and donor fluorescence processes respectively.

other two exponential decays. This observation may be explained by a multi-well potential energy surface for the excited donor, as illustrated in Fig. 4 and observed by Li *et al.*³⁸

While the lifetimes of this nonradiative relaxation pathway are very similar for both conformations, the corresponding lineshapes in the 2D lifetime map differ notably. In the *kite* spectrum (Fig. 4, lower left plot, grey), the signal is primarily due to one homogeneously broadened feature at 17 000 cm⁻¹. The antidiagonal full width at half maximum (FWHM) of this feature corresponds to a homogeneous process with a dephasing lifetime of 48 fs, coincident with the timescale of relaxation within the electronic excited state as seen *via* Stokes shift in fs-resolved 2DES (Fig. S31†). By comparison, the *vase* spectrum (Fig. 3, upper left plot) shows an inhomogeneously broadened diagonal feature at 17 100 cm⁻¹, as well as a crosspeak feature at an excitation frequency of 17 000 cm⁻¹ and detection frequency of 16 400 cm⁻¹. Inhomogeneous broadening indicates that subsections of the sample population have differing energetic environments, such as those arising from conformational variation, that persist on the timescale of the experiment.

The middle (orange) column in Fig. 4 represents the strength of signal that decays with a lifetime of several hundred ps. This decay lifetime is not observed in fluorescence lifetime measurements of the donor solution. We assign this signal to energy transfer between donor and acceptor (FRET). The energy of the diagonal feature matches with GSB from the donor, while the energy of the below-diagonal feature matches SE from the

Stokes-shifted excited donor, as well as the absorption of the acceptor. Both features decay due to FRET transfer to the acceptor. Because the initial vibrational relaxation in the donor excited state is complete within 100 fs (Fig. S31†), all signal at the diagonal observed on this timescale is due to GSB of the donor, meaning that only the loss of donor excitation is observed. The below-diagonal features may contain signal from the acceptor and have longer lifetimes than features on the diagonal (Table TS4 in ESI†). Hence, the diagonal feature is used for donor lifetime measurements. This FRET transfer pathway contributes to the majority of the signal, particularly in the *vase* conformation where the chromophores are in close proximity to one another.

The right (purple) column in Fig. 4 shows the portion of the 2D signal that does not decay on the timescale of 800 ps. We assign this signal to residual donor fluorescence based on the measured fluorescence lifetime of the donor (2–3 ns, ESI Table TS2†). The double peak lineshape of this signal can be described by a combination of GSB at the diagonal (corresponding to the original absorption) and SE at the crosspeak (corresponding to Stokes-shifted emission). Simulated spectra generated from the absorption and fluorescence of the donor alone (Fig. S32†) qualitatively match the lineshape of this long-lived signal for both conformations, further supporting our assignment. Previous work on FRET between orthogonal dyes has highlighted that low frequency vibrations (analogous here to the difference in orientational freedom between *vase* and *kite*) can alter transfer efficiency dramatically.⁵⁹



Conformation-dependent changes in excited-state dynamics and effects on FRET

The excited-state dynamic processes in cavitand **1** elucidated by 2DES and DAS analyses are summarized in Fig. 5. After absorption in the donor from ground state D_0 to initial excited state D' , rapid vibrational relaxation on the excited state surface occurs within the first 100 fs, as shown in inset 1 of Fig. 5 (brown arrows). Excitations relax into (at least) two local minima, with one containing 20–30% of the original excited state population leading to nonradiative relaxation on a ~ 25 ps timescale. All remaining excitations relax into the local minimum corresponding to dipole-driven relaxation, either as FRET or fluorescence. At this juncture, the available population of FRET donors is limited again, but by a distribution of dipole orientations between donor and acceptor. Inset 2 of Fig. 5 shows two example chromophore orientations in the *kite* conformation, one in which FRET occurs readily (bottom) and one in which FRET is prohibited as dipole coupling approaches zero (top). This distribution of dipole orientations leads to the two remaining decays illustrated in Fig. 5, one due to energy transfer (orange in Fig. 4, dashed orange arrows in Fig. 5), and one due to residual fluorescence (purple in Fig. 4, dark purple arrow in Fig. 5). With a rate of a few hundred ps, FRET is the dominant dipole-driven relaxation pathway for the donor, regardless of conformation. The acceptor then quickly relaxes out of our laser bandwidth and is not observed. While FRET remains the dominant pathway, conformational change between *vase* and *kite* affects both the transfer rate and the relative fractions of excited donor molecules decaying along all three paths.

While we identify the second of the three lifetimes as corresponding to energy transfer, it should be noted that the measured lifetime is distinct from the time actually required for transfer. FRET relies upon the same dipole coupling between ground and excited states that leads to donor fluorescence, meaning that fluorescence always occurs along with FRET and

that the branching ratio between them is determined by their relative rates. As such, the measured decay in excited state population will have a rate k_{2D} such that

$$k_{2D} = k_{tr} + k_{d,0} \quad (4)$$

where k_{tr} is the transfer rate, k_{2D} is the observed decay rate, and $k_{d,0}$ is the fluorescence decay rate of the unperturbed donor.⁵² As before, any rate k is defined as the inverse of the respective lifetime τ . Using this approach with k_{2D} as the rate labeled “energy transfer,” we estimate the FRET lifetimes to be 303 ± 10 ps for the *vase* conformation and 602 ± 58 ps for the *kite* conformation.

In addition to determining transfer lifetimes, we can calculate the branching ratio of FRET in this system using eqn (1), where τ_{DA} is the lifetime we assign to energy transfer in DAS and τ_D is the fluorescence decay lifetime of the donor in solution, measured *via* time-correlated single photon counting (TCSPC). This method results in a calculated transfer efficiency of $92.1 \pm 1.3\%$ transfer for the *vase* conformation and $78.0 \pm 2.9\%$ for the *kite* conformation. This approach only accounts for kinetic competition, however, and therefore only determines the efficiency of the subpopulation measured in DAS. If we use the peak diagonal intensities from DAS to approximate branching between the three subpopulations, we obtain overall FRET efficiencies of $68.7 \pm 1.1\%$ for the *vase* conformation and $44.3 \pm 1.8\%$ for the *kite* conformation, a significant reduction in overall values and a dramatically greater difference between conformations than the difference measured *via* eqn (1).

The results above highlight how conformation-dependent variations in nonradiative relaxation processes affect the accuracy of lifetime-based FRET measurements. These effects should be considered in situations where the chromophore structure(s) or their relative orientations result in either: (1) breakdown of the ideal dipole approximation (IDA); or (2) appreciable competition in the excited state population between FRET and nonradiative relaxation. The former is

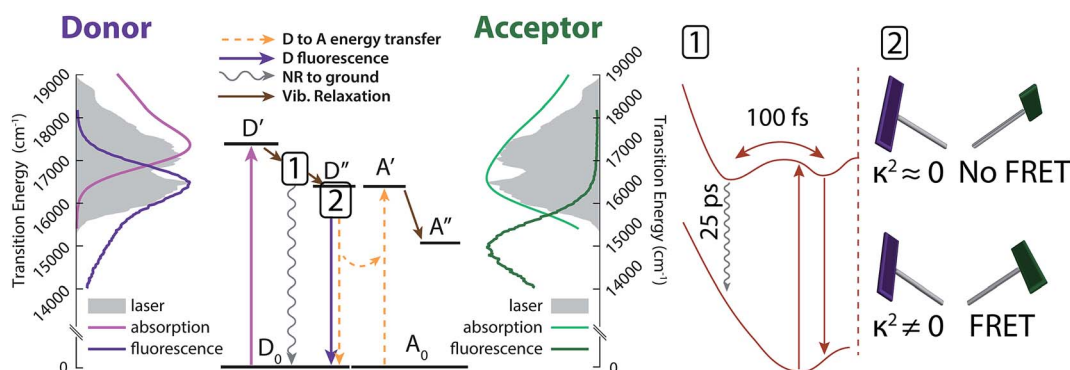


Fig. 5 Energy diagram interpretation of the observed dynamics. Initial excitation to D' (light purple) quickly relaxes vibrationally into one of two wells (see inset 1). Excited donors then relax back to the ground state either nonradiatively (wavy grey arrow) or *via* dipole coupling from well D'' (straight dark purple arrow). Part of the decay from D'' transfers energy to the acceptor *via* FRET (dashed orange arrows), and part decays without transfer, as shown in inset 2. Side panels show donor (purple) and acceptor (green) absorption and emission spectra (light and dark lines) overlaid with the laser spectrum (solid grey). Inset 1 shows relaxation into multiple wells in the excited state (top) leading to nonradiative relaxation (wavy grey arrow) and dipole relaxation (straight brown arrow). Inset 2 shows example dipole orientations in the *kite* conformation. A distribution of dipole orientations is responsible for multiple lifetimes of dipole-based donor relaxation.



relevant to chromophore structures that exhibit delocalized transition densities^{60,61} or to systems in which chromophore orientations are restricted such that isotropic dipole distributions that might mitigate IDA breakdown cannot be achieved.^{62,63} The latter situation might arise in donor chromophore structures that exhibit either modest quantum yields or conformational freedom that exacerbates nonradiative relaxation processes.³⁷ In addition to considering the effect of variations in nonradiative relaxation on FRET measurements, these variations might also be exploited productively as probes for changing viscosity in cell environments *via* motion-induced change in emission measurements (MICE).⁶⁴

Conclusion

When excited-state populations undergo competing FRET and nonradiative relaxation processes that change with donor-acceptor distance, the nonradiative relaxation processes must be accounted for in order to obtain accurate FRET efficiencies. To illustrate how nonradiative processes can affect FRET efficiency determinations, we designed a BODIPY-functionalized resorcin [4]arene cavitand that undergoes well-defined, temperature-induced conformational changes, allowing external control over donor-acceptor distance, as well as interactions between the chromophores and vibrational bath. The *vase* and *kite* conformations of the cavitand were subjected to 2DES studies, in which transitions could be separated with respect to both frequency and time, thereby enabling direct observation of changes in non-radiative relaxation that occur upon switching the cavitand conformation. Accurate determination of FRET efficiency required disambiguating overlapping transitions arising from nonradiative relaxation, FRET, and residual donor fluorescence; efficiencies extrapolated using this method differ significantly from those determined from lifetime measurements that assume that nonradiative relaxation is independent of conformation. Thus, in systems where nonradiative pathways compete with FRET, understanding how these pathways change with variations in donor and acceptor environments is vital for accurate determination of transfer efficiency. This understanding will enable improved FRET efficiency measurements in the complex environments inherent to biological systems and advanced materials.

Experimental section

General methods

Detailed synthesis procedures and spectra are provided in the ESI.†

UV/vis absorption spectra were recorded using a Cary 5000 dual-beam UV/VIS/NIR spectrophotometer. Steady state fluorescence was measured *via* a Horiba Fluorolog 3, and fluorescence quantum yields were recorded using the Horiba QuantaPhi attachment. Time-domain lifetimes were measured on a ChronosBH lifetime fluorometer (ISS, Inc.) using Time-Correlated Single Photon Counting (TCSPC) methods. The fluorometer contained Becker-Hickl SPC-130 detection electronics and an HPM-100-40 Hybrid PMT detector. Tunable picosecond pulsed excitation at 560 nm was provided by

a Fianium SC400-2 supercontinuum laser source with integrated pulse picker and AOTF. Emission wavelengths were selected with bandpass filters (Chroma HQ615/85 nm). The Instrument Response Function (IRF) was measured to be approximately 120 ps FWHM in a 1% scattering solution of Ludox LS colloidal silica. Multi-component exponential decay lifetimes were fit *via* a forward convolution method in the Vinci control and analysis software.

Two-dimensional electronic spectra were acquired using a homebuilt instrument that has been described previously.⁴⁵ A Ti:sapphire oscillator and regenerative amplifier (Micra and Legend Elite, Coherent) generate a 5 kHz, 100 fs pulse train that is directed into a noncollinear optical parametric amplifier (NOPA, TOPAS). The output of the NOPA is then compressed using a prism pair followed by a spatial light modulator array (FemtoJock, Biophotonics). The resulting pulse (12 fs, 530–650 nm FWHM) is then split into four *via* a beamsplitter and a transmissive grating. The four beams are directed to the sample with a boxcar geometry. The time delays are controlled by paired glass wedges (coherence time, τ) and a translational stage (waiting time, T). Beams 1, 2 and 3 interact with the sample to generate the third-order signal. Beam 4 is attenuated by three orders of magnitude and serves as local oscillator for heterodyne detection. Signal, along with the local oscillator, is directed to a spectrometer and the resulting interferogram is recorded on a CCD array (Andor). All 2D data in the main text were collected in the same 24 h period in the same cryostat, varying only the temperature between the two measurements.

Conflicts of interest

There are no conflicts to declare.

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References

- 1 T. Förster, in *Istanbul Lectures, Part III: Action of Light and Organic Crystals*, Academic Press, New York, 1965, pp. 93–137.
- 2 R. M. Clegg, *Curr. Opin. Biotechnol.*, 1995, **6**, 103–110.
- 3 P. R. Selvin, *Nat. Struct. Biol.*, 2000, **7**, 730–734.
- 4 P. G. Wu and L. Brand, *Anal. Biochem.*, 1994, **218**, 1–13.
- 5 L. Stryer and R. P. Haugland, *Proc. Natl. Acad. Sci. U. S. A.*, 1967, **58**, 719–726.
- 6 C. G. dos Remedios and P. D. Moens, *J. Struct. Biol.*, 1995, **115**, 175–185.
- 7 R. N. Day, A. Periasamy and F. Schaufele, *Methods*, 2001, **25**, 4–18.
- 8 B. N. G. Giepmans, S. R. Adams, M. H. Ellisman and R. Y. Tsien, *Science*, 2006, **312**, 217–224.



- 9 M. Tramier, I. Gautier, T. Pilot, S. Ravalet, K. Kemnitz, J. Coppey, C. Durieux, V. Mignotte and M. Coppey-Moisan, *Biophys. J.*, 2002, **83**, 3570–3577.
- 10 S. S. Vogel, C. Thaler and S. V. Koushik, *Sci. STKE*, 2006, DOI: 10.1126/stke.3312006re2.
- 11 V. V. Didenko, *BioTechniques*, 2001, **31**, 1106–1121.
- 12 A. Hillisch, M. Lorenz and S. Diekmann, *Curr. Opin. Struct. Biol.*, 2001, **11**, 201–207.
- 13 C.-Y. Zhang, H.-C. Yeh, M. T. Kuroki and T.-H. Wang, *Nat. Mater.*, 2005, **4**, 826–831.
- 14 S. Zadrán, S. Standley, K. Wong, E. Otiniano, A. Amighi and M. Baudry, *Appl. Microbiol. Biotechnol.*, 2012, **96**, 895–902.
- 15 P. Bastiaens, *Trends Cell Biol.*, 1999, **9**, 48–52.
- 16 W. Becker, *J. Microsc.*, 2012, **247**, 119–136.
- 17 E. A. Jares-Erijman and T. M. Jovin, *Nat. Biotechnol.*, 2003, **21**, 1387–1395.
- 18 E. A. Jares-Erijman and T. M. Jovin, *Curr. Opin. Chem. Biol.*, 2006, **10**, 409–416.
- 19 R. B. Sekar and A. Periasamy, *J. Cell Biol.*, 2003, **160**, 629–633.
- 20 K. Truong and M. Ikura, *Curr. Opin. Struct. Biol.*, 2001, **11**, 573–578.
- 21 H. Wallrabe and A. Periasamy, *Curr. Opin. Biotechnol.*, 2005, **16**, 19–27.
- 22 D. W. Piston and G. J. Kremers, *Trends Biochem. Sci.*, 2007, **32**, 407–414.
- 23 A. P. Alivisatos, W. Gu and C. Larabell, *Annu. Rev. Biomed. Eng.*, 2005, **7**, 55–76.
- 24 X. Chi, D. Huang, Z. Zhao, Z. Zhou, Z. Yin and J. Gao, *Biomaterials*, 2012, **33**, 189–206.
- 25 A. R. Clapp, I. L. Medintz and H. Mattoussi, *ChemPhysChem*, 2006, **7**, 47–57.
- 26 P. D. Howes, R. Chandrawati and M. M. Stevens, *Science*, 2014, **346**, 1247390.
- 27 K. E. Sapsford, L. Berti and I. L. Medintz, *Angew. Chem., Int. Ed.*, 2006, **45**, 4562–4588.
- 28 L. Yuan, W. Lin, K. Zheng and S. Zhu, *Acc. Chem. Res.*, 2013, **46**, 1462–1473.
- 29 C. Berney and G. Danuser, *Biophys. J.*, 2003, **84**, 3992–4010.
- 30 U. Noomnarm and R. M. Clegg, *Photosynth. Res.*, 2009, **101**, 181–194.
- 31 V. A. Azov, A. Schlegel and F. Diederich, *Angew. Chem., Int. Ed.*, 2005, **44**, 4635–4638.
- 32 V. A. Azov, A. Beeby, M. Cacciarini, A. G. Cheetham, F. Diederich, M. Frie, J. K. Gimzewski, V. Gramlich, B. Hecht, B. Jaun, T. Latychevskaia, A. Lieb, Y. Lill, F. Marotti, A. Schlegel, R. R. Schlittler, P. J. Skinner, P. Seiler and Y. Yamakoshi, *Adv. Funct. Mater.*, 2006, **16**, 147–156.
- 33 J. R. Moran, S. Karbach and D. J. Cram, *J. Am. Chem. Soc.*, 1982, **104**, 5826–5828.
- 34 I. Pochorovski, B. Breiten, W. B. Schweizer and F. Diederich, *Chem.–Eur. J.*, 2010, **16**, 12590–12602.
- 35 I. Pochorovski, T. Knehans, D. Nettels, A. M. Müller, W. B. Schweizer, A. Caflisch, B. Schuler and F. Diederich, *J. Am. Chem. Soc.*, 2014, **136**, 2441–2449.
- 36 I. Pochorovski and F. Diederich, *Isr. J. Chem.*, 2012, **52**, 20–29.
- 37 H. L. Kee, C. Kirmaier, L. Yu, P. Thamyongkit, W. J. Youngblood, M. E. Calder, L. Ramos, B. C. Noll, D. F. Bocian, W. R. Scheidt, R. R. Birge, J. S. Lindsey and D. Holten, *J. Phys. Chem. B*, 2005, **109**, 20433–20443.
- 38 F. Li, S. I. Yang, Y. Ciringh, J. Seth, C. H. Martin, D. L. Singh, D. Kim, R. R. Birge, D. F. Bocian, D. Holten and J. S. Lindsey, *J. Am. Chem. Soc.*, 1998, **120**, 10001–10017.
- 39 L. Wang, I. S. Tamgho, L. A. Crandall, J. J. Rack and C. J. Ziegler, *Phys. Chem. Chem. Phys.*, 2015, **17**, 2349–2351.
- 40 Y. Wang, D. Zhang, H. Zhou, J. Ding, Q. Chen, Y. Xiao and S. Qian, *J. Appl. Phys.*, 2010, **108**, 033520.
- 41 E. Collini, C. Y. Wong, K. E. Wilk, P. M. G. Curmi, P. Brumer and G. D. Scholes, *Nature*, 2010, **463**, 644–647.
- 42 P. D. Dahlberg, A. F. Fidler, J. R. Caram, P. D. Long and G. S. Engel, *J. Phys. Chem. Lett.*, 2013, **4**, 3636–3640.
- 43 G. S. Schlau-Cohen, A. Ishizaki and G. R. Fleming, *Chem. Phys.*, 2011, **386**, 1–22.
- 44 A. A. Bakulin, S. E. Morgan, T. B. Kehoe, M. W. B. Wilson, A. W. Chin, D. Zigmantas, D. Egorova and A. Rao, *Nat. Chem.*, 2016, **8**, 16–23.
- 45 D. Hayes, G. B. Griffin and G. S. Engel, *Science*, 2013, **340**, 1431–1434.
- 46 C. C. Jumper, J. M. Anna, A. Stradomska, J. Schins, M. Myahkostupov, V. Prusakova, D. G. Oblinsky, F. N. Castellano, J. Knoester and G. D. Scholes, *Chem. Phys. Lett.*, 2014, **599**, 23–33.
- 47 V. A. Azov, B. Jaun and F. Diederich, *Helv. Chim. Acta*, 2004, **87**, 449–462.
- 48 J.-B. Wang, Q.-Q. Wu, Y.-Z. Min, Y.-Z. Liu and Q.-H. Song, *Chem. Commun.*, 2012, **48**, 744–746.
- 49 M. J. Frisch, G. W. Trucks, H. B. Schlegel, G. E. Scuseria, M. A. Robb, J. R. Cheeseman, G. Scalmani, V. Barone, B. Mennucci, G. A. Petersson, H. Nakatsuji, M. Caricato, X. Li, H. P. Hratchian, A. F. Izmaylov, J. Bloino, G. Zheng, J. L. Sonnenberg, M. Hada, M. Ehara, K. Toyota, R. Fukuda, J. Hasegawa, M. Ishida, T. Nakajima, Y. Honda, O. Kitao, H. Nakai, T. Vreven, J. A. Montgomery Jr, J. E. Peralta, F. Ogliaro, M. J. Bearpark, J. Heyd, E. N. Brothers, K. N. Kudin, V. N. Staroverov, R. Kobayashi, J. Normand, K. Raghavachari, A. P. Rendell, J. C. Burant, S. S. Iyengar, J. Tomasi, M. Cossi, N. Rega, N. J. Millam, M. Klene, J. E. Knox, J. B. Cross, V. Bakken, C. Adamo, J. Jaramillo, R. Gomperts, R. E. Stratmann, O. Yazyev, A. J. Austin, R. Cammi, C. Pomelli, J. W. Ochterski, R. L. Martin, K. Morokuma, V. G. Zakrzewski, G. A. Voth, P. Salvador, J. J. Dannenberg, S. Dapprich, A. D. Daniels, Ö. Farkas, J. B. Foresman, J. V. Ortiz, J. Cioslowski and D. J. Fox, *Gaussian 09*, Gaussian Inc., Wallingford, CT, 2009.
- 50 M. Cho, *Chem. Rev.*, 2008, **108**, 1331–1418.
- 51 I. Stiofkin, T. Brixner, M. Yang and G. R. Fleming, *J. Phys. Chem. B*, 2006, **110**, 20032–20037.
- 52 A. Nitzan, *Chemical dynamics in Condensed Phases: Relaxation, Transfer and Reactions in Condensed Molecular Systems*, Oxford University Press, Oxford, New York, 2006.
- 53 P. Hamm and M. T. Zanni, *Concepts and Methods of 2D Infrared Spectroscopy*, Cambridge University Press, Cambridge, New York, 2011.



- 54 C. Wang, M. L. Flanagan, R. D. McGillicuddy, H. Zheng, A. R. Ginzburg, X. Yang, K. Moffat and G. S. Engel, *Biophys. J.*, 2016, **111**, 2125–2134.
- 55 E. E. Ostroumov, R. M. Mulvaney, R. J. Cogdell and G. D. Scholes, *Science*, 2013, **340**(6128), 52–56.
- 56 J. A. Myers, K. L. M. Lewis, F. D. Fuller, P. F. Tekavec, C. F. Yocum and J. P. Ogilvie, *J. Phys. Chem. Lett.*, 2010, **1**, 2774–2780.
- 57 J. Dostal, B. Benesova and T. Brixner, *J. Chem. Phys.*, 2016, **145**, 124312.
- 58 E. Thyryhaug, K. Zidek, J. Dostal, D. Bina and D. Zigmantas, *J. Phys. Chem. Lett.*, 2016, **7**, 1653–1660.
- 59 H. Langhals, A. J. Esterbauer, A. Walter, E. Riedle and I. Pugliesi, *J. Am. Chem. Soc.*, 2010, **132**, 16777–16782.
- 60 Y. R. Khan, T. E. Dykstra and G. D. Scholes, *Chem. Phys. Lett.*, 2008, **461**, 305–309.
- 61 H. Sahoo, D. Roccatano, M. Zacharias and W. M. Nau, *J. Am. Chem. Soc.*, 2006, **128**, 8118–8119.
- 62 A. Muñoz-Losa, C. Curutchet, B. P. Krueger, L. R. Hartsell and B. Mennucci, *Biophys. J.*, 2009, **96**, 4779–4788.
- 63 J. D. Spiegel, S. Fulle, M. Kleinschmidt, H. Gohlke and C. M. Marian, *J. Phys. Chem. B*, 2016, **120**, 8845–8862.
- 64 D. Su, C. L. Teoh, L. Wang, X. Liu and Y. T. Chang, *Chem. Soc. Rev.*, 2017, **46**, 4833–4844.

