## **Supporting information**

## Unravelling the Conformations of di-(perylene bisimide acrylate) by combining Time-Resolved Fluorescence-Anisotropy experiments and Molecular Modelling

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#### I. <u>Time rescaling in MD simulations</u>

Since the focus of our work is comparing the photophysical properties of monomeric PerAcr with those of dimeric (PerAcr)<sub>2</sub> we choose the long-time anisotropy decay of the former as a reference time scale. From simulations on a single monomer we obtain the anisotropy decay depicted in figure SI\_1. It features a clear mono-exponential tail with a time constant  $\theta_{sim}$ =1140ps. Comparing with the experimental value  $\theta_{exp}$ =460ps yields a scaling factor of 2.5. All times obtained from MD simulations were divided by this constant factor without any further adjustments.



Fig.SI\_1: Anisotropy decay of a single PerAcr monomer (solid line) with a fit to the monoexponential tail (red dashed line).

#### II. Förster-distance for PBI-PBI homo-ET in (PerAcr)2

Förster resonance energy transfer (FRET) refers to the transfer of electronic excitation energy between a donor (D) and an acceptor (A) molecule, that are only weakly coupled. This allows to take advantage of time-dependent perturbation theory and to derive under several further approximations (restriction of the electrostatic interaction to the leading dipole-dipole term) an expression from Fermi's golden rule for the energy transfer rate from D to A. This yields that the rate depends on the D-A distance as  $R^{-6}$ , the reciprocal lifetime of the excited state of the unperturbed donor, and further scaling factors that can be lumped together in a constant termed the Förster radius  $R_0$ , which is given as:<sup>1</sup>

$$R_0^{\ 6} = \frac{9(\ln 10)\kappa^2 Q_D}{128\pi^5 Nn^4} \int_0^\infty F_D(\lambda) \varepsilon_A(\lambda) \lambda^4 d\lambda$$
<sup>(1)</sup>

Here N denotes Avogadro's constant, n refers to the refractive index of the solvent;  $F_D$  is the (normalised) fluorescence spectrum of the donor,  $Q_D$  is the fluorescence quantum yield of the unperturbed donor,  $\varepsilon_A$  denotes the extinction coefficient of the acceptor, and  $\kappa^2$  is the orientational factor of the two interacting transition-dipole moments. Under the assumption of dynamic random averaging of the donor/acceptor orientation this yields 2/3. The Förster radius corresponds to that distance between the donor and the acceptor where the probability is 50% for energy transfer from D to A during the lifetime of the donor excited state. Using the fluorescence and absorption spectra reported for PerAcr earlier,<sup>2</sup> we find a value of  $R_0 = 5$  nm for the Förster radius between the two PBIs in (PerAcr)<sub>2</sub>.

## III. Derivation of Eq. 9 of the main text

Let  $p_a(t')$  denote the probability that the excitation is located on monomer *a* at time *t'*. With the reaction rate *k*, its rate of change is given by

$$\mathbf{p}_{a}^{k}(t') = -p_{a}(t')\mathbf{k} + (1 - p_{a}(t'))\mathbf{k}$$
  
=  $-2p_{a}(t')\mathbf{k} + \mathbf{k}$  (2)

which can be solved by the ansatz

$$p_a(t') = \frac{1}{2} + C_0 e^{-2kt'}$$
(3)

where  $C_0$  is a constant. Assuming that  $p_a(t'=0) = 1$  fixes  $C_0 = \frac{1}{2}$ .

Thus, the probability that the excitation is located on monomer *a* at time  $\Delta t$ , given that it was on *a* at *t*'=0, is

$$p_a(\Delta t) = \frac{1}{2} + \frac{1}{2}e^{-2k\Delta t}.$$
(4)

This is the resting probability used in the main text.

# IV. <u>Calculation of the hopping constant $k_0$ and the</u> <u>orientation factor $\kappa$ in Eq. 10 of the main text</u>

The reaction rate k(t) is given by <sup>3</sup>

$$k(t) = \frac{3}{2} \frac{\kappa(t)^2}{\tau} \left(\frac{R_0}{R}\right)^6.$$
(5)

The fluorescence life time  $\tau$  for the ISO state has been determined to be 3.7ns <sup>2</sup> and the Förster radius  $R_0$  =5nm which leads to  $k_0$ =6.33 nm<sup>6</sup>/ps.

The orientation factor  $\kappa(t)$  is calculated from <sup>3</sup>

$$\kappa(t) = \left(\cos\alpha(t) - 3\cos\theta_D(t)\cos\theta_A(t)\right)^2$$
(6)

where  $\alpha$  is the angle between the two dipole vectors as in Fig. 3 of the main text and  $\theta_D$  ( $\theta_A$ ) is the angle between the acceptor (donor) dipole with the vector connecting the centres-of-mass of the two monomers.

#### References

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