

Supporting Information for

Photoinduced energy and electron transfers at graphene quantum dot /azobenzene interfaces

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Excitation energies

Table S1 Detailed description of excited states for isolated AZO. Energies of each excited states are reported in nm. Oscillator strength is reported in bracket.

Excited state	<i>t</i> AZO		<i>c</i> AZO	
	Fold	Open	Fold	Open
1	434 (0.00) HOMO-3 → LUMO	435 (0.00) HOMO-3 → LUMO	459 (0.06) HOMO-1 → LUMO+1	454 (0.05) HOMO-1 → LUMO+1
2	332 (0.57) HOMO-1 → LUMO	328 (1.15) HOMO-1 → LUMO	317 (0.23) HOMO → LUMO	309 (0.33) HOMO → LUMO
3	317 (0.07) HOMO → LUMO	308 (0.17) HOMO → LUMO+1	309 (0.03) HOMO → LUMO+2	307 (0.08) HOMO → LUMO+2
4	314 (0.46) HOMO → LUMO+1	306 (0.19) HOMO → LUMO+1	288 (0.30) HOMO-1 → LUMO+1	285 (0.29) HOMO-3 → LUMO+1
5	308 (0.08) HOMO → LUMO+2	258 (0.01) HOMO-1 → LUMO+4	282 (0.03) HOMO → LUMO+1	263 (0.08) HOMO-1 → LUMO+3
6	272 (0.02) HOMO-1 → LUMO+1	253 (0.00) HOMO → LUMO+3	274 (0.03) HOMO-1 → LUMO	259 (0.05) HOMO-4 → LUMO+1
7	263 (0.02) HOMO-6 → LUMO	252 (0.00) HOMO-5 → LUMO	266 (0.06) HOMO-3 → LUMO+1	252 (0.00) HOMO → LUMO+5
8	257 (0.01) HOMO-1 → LUMO+3	244 (0.01) HOMO → LUMO	262 (0.01) HOMO-1 → LUMO+3	247 (0.00) HOMO-1 → LUMO+3
9	252 (0.01) HOMO → LUMO+4	242 (0.32) HOMO-4 → LUMO	253 (0.00) HOMO → LUMO+5	241 (0.39) HOMO-2 → LUMO
10	252 (0.03) HOMO-6 → LUMO	241 (0.15) HOMO-4 → LUMO	249 (0.01) HOMO-1 → LUMO+6	241 (0.03) HOMO-5 → LUMO

11	247 (0.24) HOMO-2 → LUMO+1	240 (0.00) HOMO-7 → LUMO+1	246 (0.22) HOMO-2 → LUMO	229 (0.00) HOMO → LUMO+1
12	246 (0.00) HOMO-6 → LUMO+1	225 (0.00) HOMO-1 → LUMO+1	244 (0.02) HOMO-5 → LUMO	228 (0.00) HOMO-1 → LUMO
13	242 (0.01) HOMO-4 → LUMO	220 (0.15) HOMO-6 → LUMO	233 (0.03) HOMO-2 → LUMO+1	224 (0.20) HOMO-6 → LUMO+1
14	236 (0.00) HOMO-3 → LUMO+1	217 (0.00) HOMO-2 → LUMO+3	228 (0.03) HOMO → LUMO+4	222 (0.10) HOMO-1 → LUMO+4
15	225 (0.20) HOMO-1 → LUMO+2	215 (0.21) HOMO-9 → LUMO+6	227 (0.19) HOMO-5 → LUMO+1	217 (0.01) HOMO-2 → LUMO+5
16	224 (0.01) HOMO-9 → LUMO+6	215 (0.74) HOMO-2 → LUMO+2	226 (0.06) HOMO → LUMO+4	215 (0.94) HOMO-2 → LUMO+2
17	222 (0.04) HOMO → LUMO+4	208 (0.19) HOMO-1 → LUMO+4	224 (0.01) HOMO-3 → LUMO	214 (0.05) HOMO-9 → LUMO+8
18	220 (0.13) HOMO-2 → LUMO+2	206 (0.01) HOMO → LUMO+7	223 (0.00) HOMO → LUMO+4	213 (0.01) HOMO-7 → LUMO+2
19	218 (0.13) HOMO-7 → LUMO	204 (0.01) HOMO-2 → LUMO	221 (0.01) HOMO → LUMO+3	206 (0.02) HOMO → LUMO+7
20	217 (0.07) HOMO-4 → LUMO+1	204 (0.01) HOMO-8 → LUMO+1	220 (0.03) HOMO → LUMO	205 (0.05) HOMO-1 → LUMO+9

Internal reorganization energy calculation for interfaces

When considering the interface consisting of donor and acceptor fragments, one can separately calculate the internal reorganization energy as an average of the backward (λ_{Bw}) and forward transitions (λ_{Fw}) as:

- PET

$$\lambda_{Bw} = E(D^* @ D^+) - E(D^* @ D^*) + E(A @ A^-) - E(A @ A) \quad (S1)$$

$$\lambda_{Fw} = E(D^+ @ D^*) - E(D^+ @ D^+) + E(A^- @ A) - E(A^- @ A^-)$$

- PHT

$$\lambda_{Bw} = E(D @ D^+) - E(D @ D) + E(A^* @ A^-) - E(A^* @ A^*) \quad (S2)$$

$$\lambda_{Fw} = E(D^+ @ D) - E(D^+ @ D^+) + E(A^- @ A^*) - E(A^- @ A^-)$$

- EET

$$\lambda_{Bw} = E(D^* @ D) - E(D^* @ D^*) + E(A @ A^*) - E(A @ A) \quad (S3)$$

$$\lambda_{Fw} = E(D @ D^*) - E(D @ D) + E(A^* @ A) - E(A^* @ A^*)$$

where D indicates the donor fragment and A the acceptor fragment in their excited states (i.e., anionic, cation or excitonic states).

Table S2. Reorganization energy, coupling, driving force and Marcus rates for the three processes considered in this study, for the AZO molecule isolated.

	Process	λ [eV]	Coupling [eV]	Involved transitions
<i>t</i>AZO	EET	0.14	0.021	$D_{PYR}^*(S_3, S_4), A_{AZO}^*(S_2)$
	PET	0.75	0.009	$D_{PYR}^*(S_3, S_4), D_{PYR \rightarrow AZO}(S_8)$
	PHT	0.64	0.003	$A_{AZO}^*(S_2), A_{AZO \rightarrow PYR}(S_8)$
<i>c</i>AZO	EET	0.24	0.010	$D_{PYR}^*(S_2), A_{AZO}^*(S_1)$
	PET	1.26	0.010	$D_{PYR}^*(S_2), D_{PYR \rightarrow AZO}(S_5)$
	PHT	1.24	0.003	$A_{AZO}^*(S_1), A_{AZO \rightarrow PYR}(S_5)$

Photo-induced energy and electron transfer couplings

The photoinduced interaction between donor and acceptor moieties can essentially be written as the sum of two components: a long-range Coulomb interaction between the transition density of the two moieties/fragments and a short range Dexter-like exchange interaction [1]:

$$V = V_{Coul} + V_{short}$$

The first term resembles a classical interaction between two density distributions. The first term of the dipolar expansion of the density distribution is the dipole moment. Therefore, the Coulomb term can be written as a dipole-dipole interaction which scales as $1/r^3$ as a function of the distance and it is mostly related to the reciprocal orientation of the interacting transition dipoles (leading to the famous Forster-like equation) and the magnitude of the interacting dipoles. As such, the distance dependence is not so strong.

The second term is a Dexter-like exchange and decays exponentially with the distance between the two moieties (thus, much faster than the Coulomb interaction). This second term is more sensitive to the actual orbital shape and specific overlap between the wavefunction two moieties. Usually, it is negligible compared to the first Coulomb term. However, it might be important in the case of the interaction of almost forbidden transition with a small oscillator strength (which nullify the Coulomb interaction).

Importantly, since the diabaticization scheme used in this work starts from the adiabatic wavefunction of the full system, it contains both Coulomb and Dexter-like exchange included in the final coupling values. We refer to Ref 1-4 for a deep analysis of the diabaticization method used.

Optimized structure and absorption spectra of the azobenzene moiety

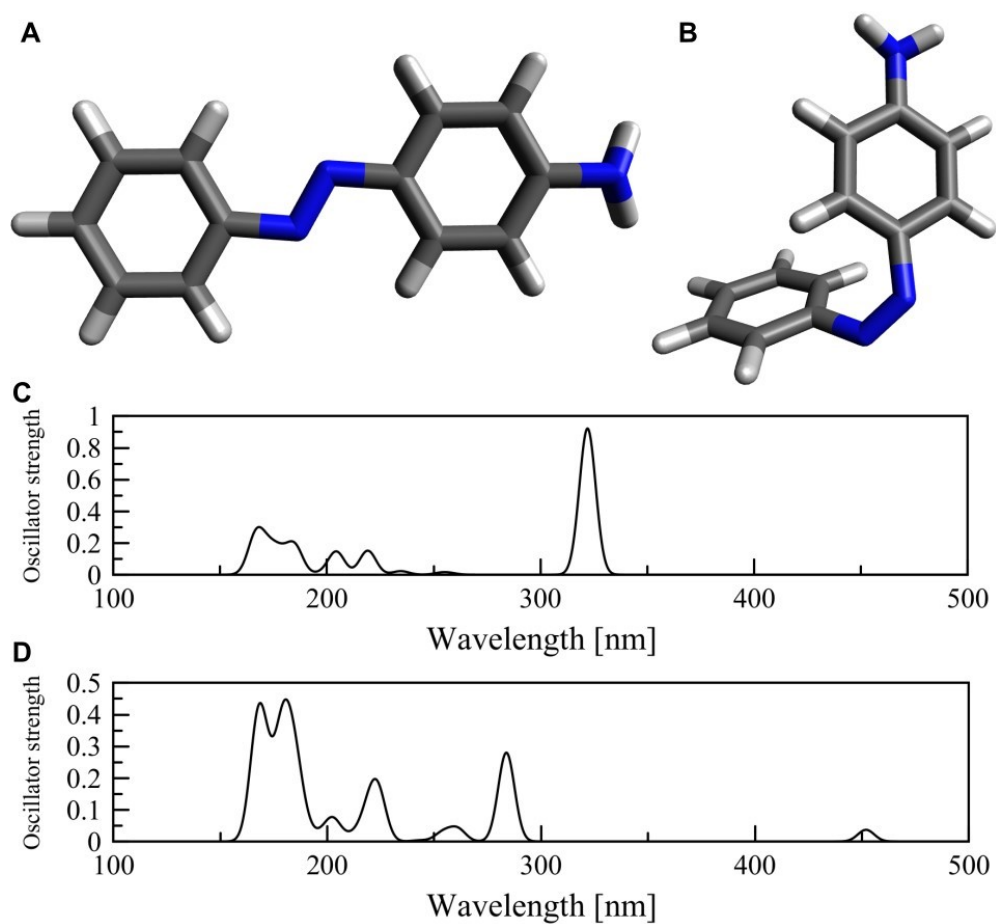


Figure S1 The structure of NH_2 -functionalized azobenzene molecules (A-*trans*, B-*cis*), C - UV-Vis spectra for isomer *trans*, D - UV-Vis spectra for isomer *cis*

Discussion on the open-AZO conformation

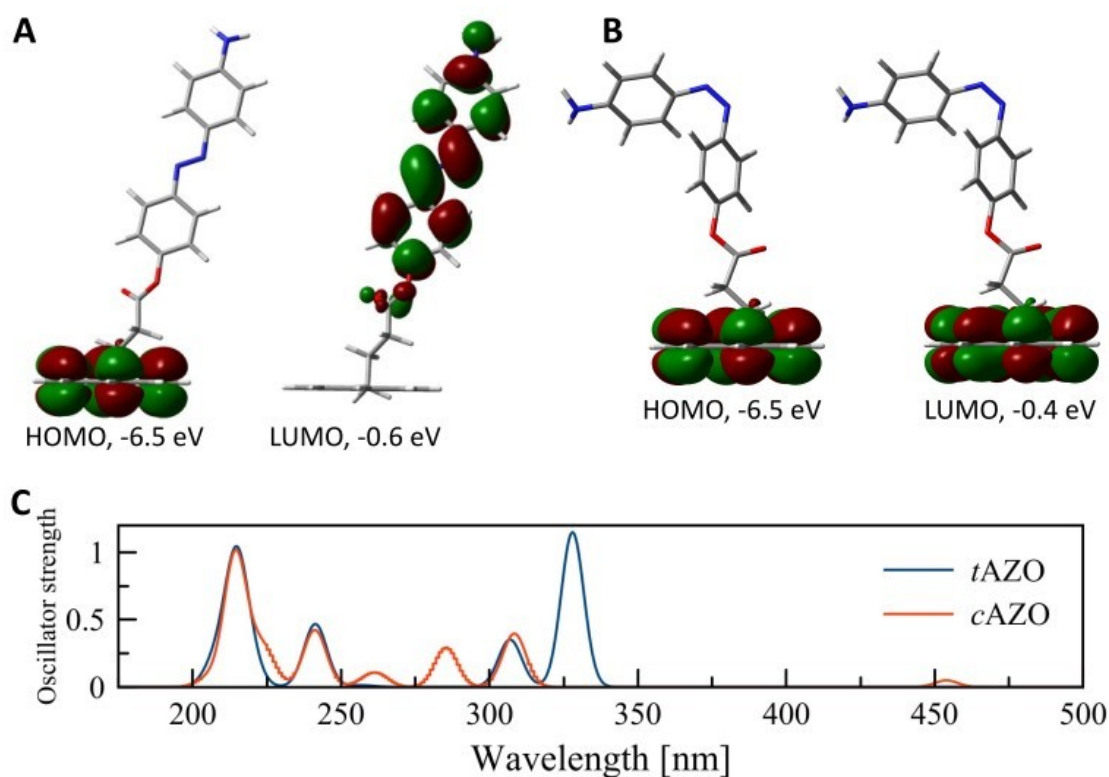


Figure S2 Frontier orbitals of isolated AZO molecules (A – *trans*, B – *cis*), C - UV-Vis spectra for isolated AZO molecules.

The opening of the conformation lowers the energies of the frontier orbitals, with HOMO at -6.5 eV and LUMO at -0.6 eV for *tAZO* and -0.4 eV for *cAZO*, which leads to energy gap of 5.9 eV and 6.1 eV, respectively. For *tAZO* HOMO is localized on pyrene, whereas LUMO is located on AZO, which suggests a possible intramolecular charge transfer upon excitation (Figure S1A). Both frontier orbitals for *cAZO* in the open conformation are localized on the pyrene moiety, which suggest limitations in charge transfer ability for this isomer (Figure S1B). For *tAZO* the first excited state is a forbidden $n \rightarrow \pi^*$ transition (HOMO-3 to LUMO), therefore is not visible on the spectra (see Table S1). The first bright peak is at 328 nm, which is related to the second excited state (S_2) and corresponds to a HOMO-1 to LUMO transition ($\pi \rightarrow \pi^*$). The next peak at 306 nm corresponds to HOMO to LUMO+1 transition. For the *cAZO* isomer, similar absorption peaks are obtained compared to the *tAZO*, albeit blue shifted by 20 nm. In addition, now the first excited state is weakly allowed and it shows as a peak at 454 nm, which corresponds to $n \rightarrow \pi^*$ (HOMO-1 to LUMO+1) transition. The next peak at 309 nm corresponds to a HOMO to LUMO transition.

From a geometrical point of view, all three processes require less energy for *t*AZO than for *c*AZO. For PET and PHT the amount of energy required for geometrical changes is similar and nearly twice as high for *c*AZO compared to *t*AZO. Considerably lower λ is observed for EET, with the value of only 0.14 eV for *t*AZO and 0.24 eV for *c*AZO, suggesting that (similar to fold conformation) this process might be favored.

The coupling for *t*AZO in open conformation favors the EET process between almost degenerated pyrene excitations at 308 nm and 306 nm and AZO excitation at 328 nm. The EET coupling for this molecule (21.3 meV) is one magnitude higher than the coupling for both charge transfers. Quite low couplings for all three processes are calculated for *c*AZO in the open conformation and it seems that for this molecule EET (between D_{PYR}^* at 309 nm and A_{AZO}^* at 454 nm) and PET (between D_{PYR}^* at 309 nm and $D_{PYR \rightarrow AZO}$ at 263 nm) can be competitive processes.

Dominant transitions for the isolated molecule

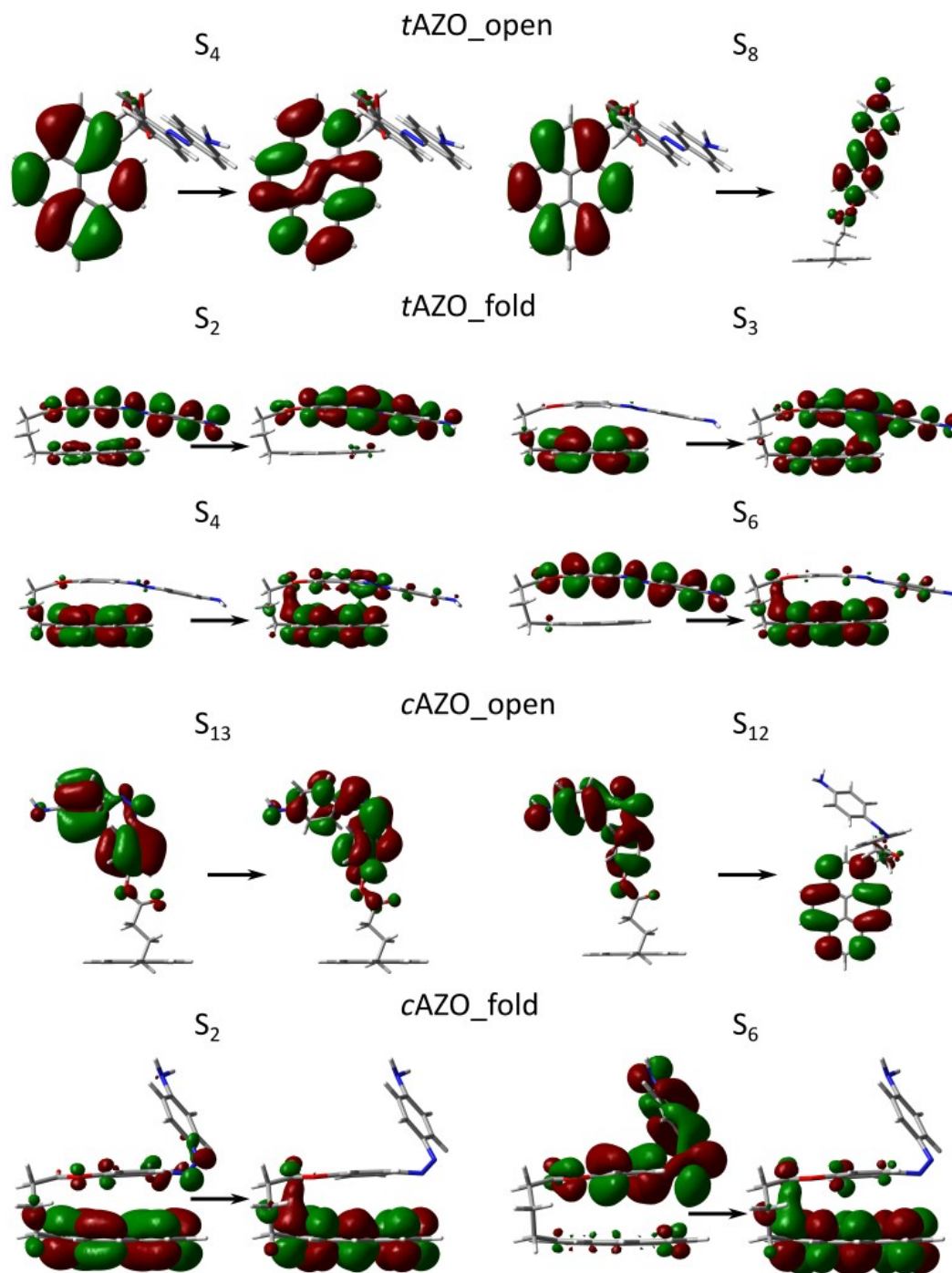


Figure S3 NTOs of AZO's excited states.

Geometrical displacements

	tAZO	tAZO (top)	cAZO	cAZO (top)
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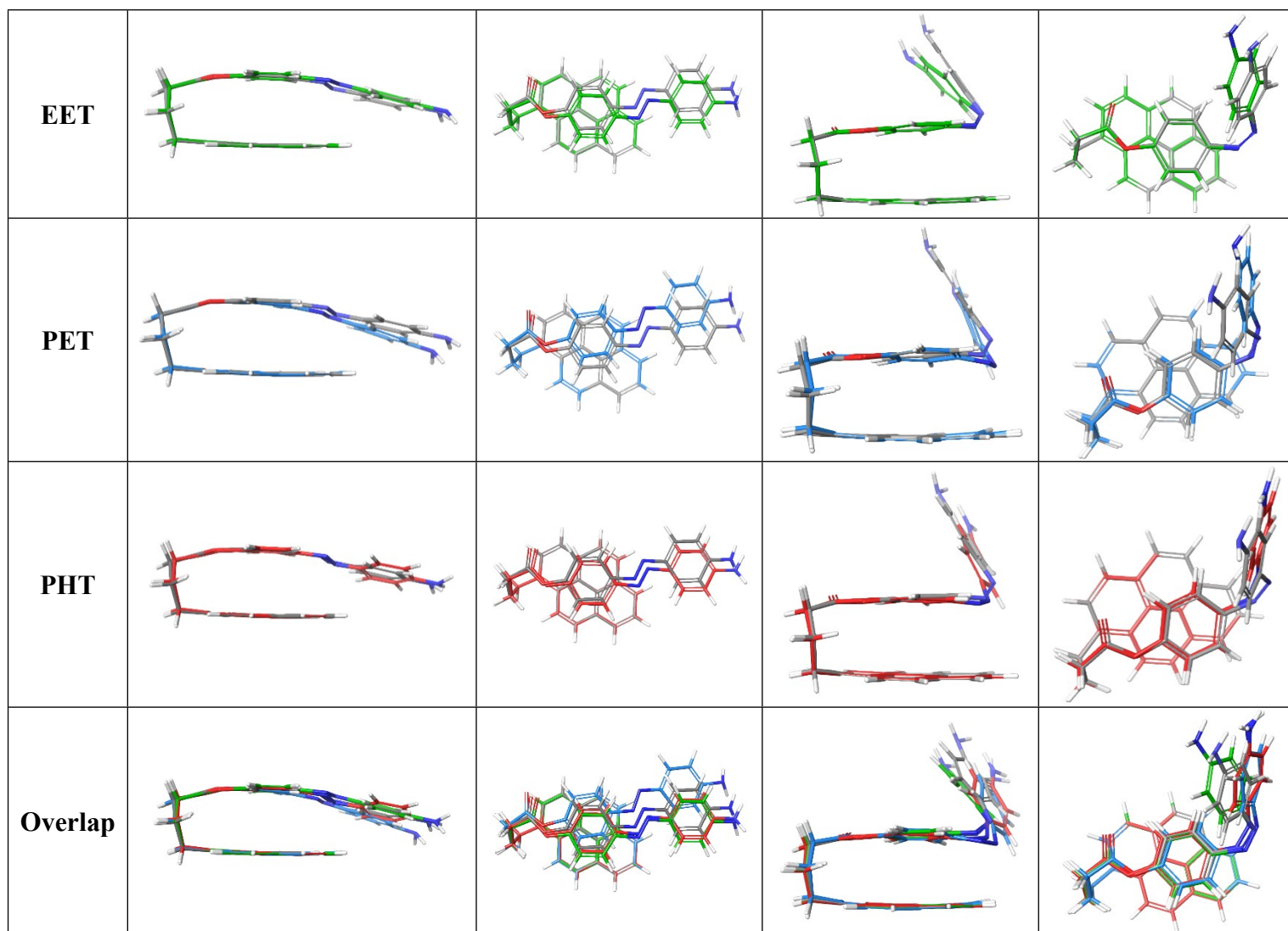


Figure S4 Geometrical differences for the optimized tAZO and cAZO structures for the three considered processes. Gray: structure in the ground state geometry; green: structure in the excited state, blue: structure with one additional electron; red: structure with one additional hole. The overlap among PHT, PET and EET is also reported.

Density of states analysis for the GQD-AZO interfaces

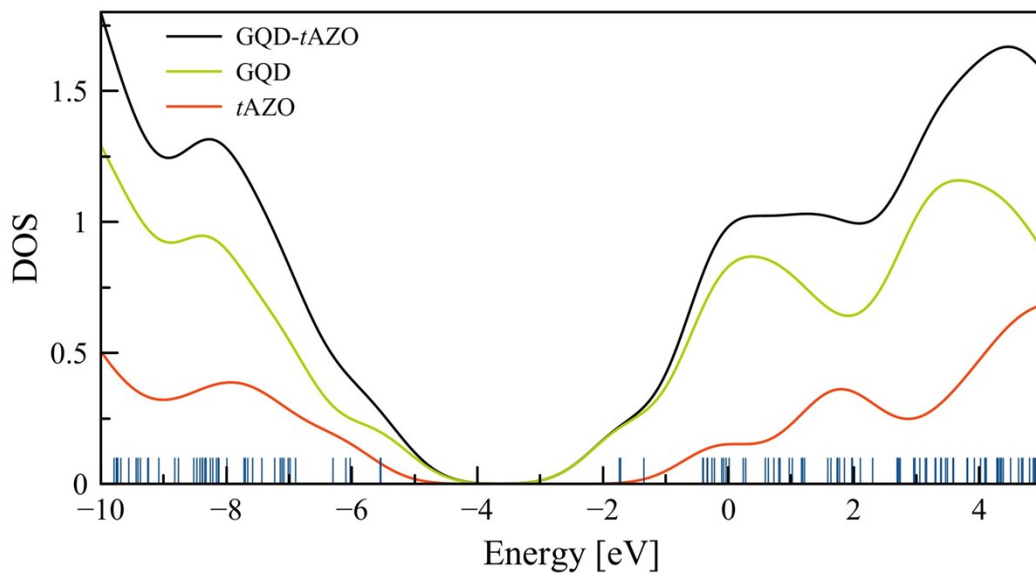


Figure S5 Density of states for GQD-tAZO as well as partial DOS for GQD and tAZO.

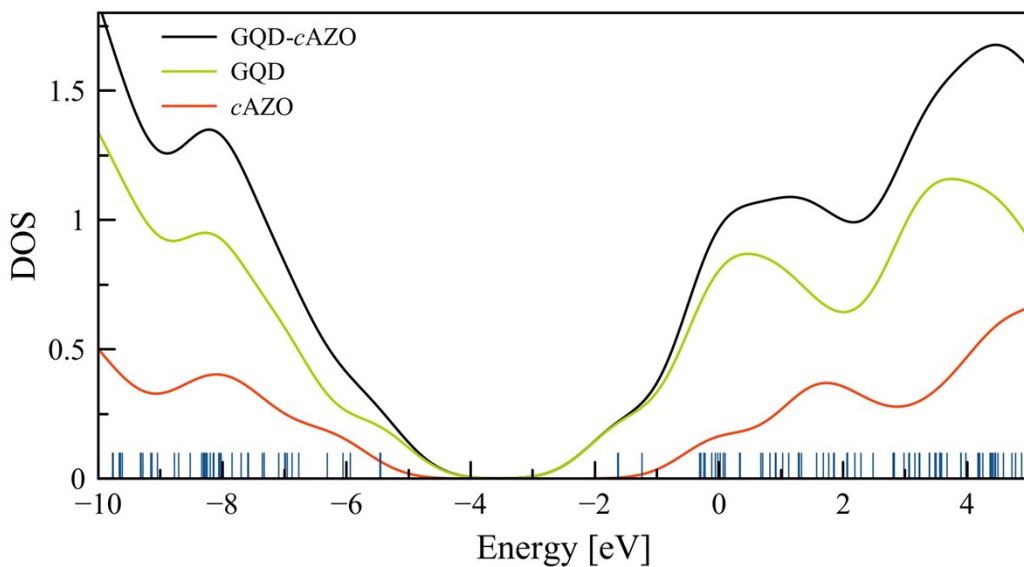


Figure S6 Density of states for GQD-cAZO as well as partial DOS for GQD and cAZO.

Energy level alignment at the GQD-AZO interfaces

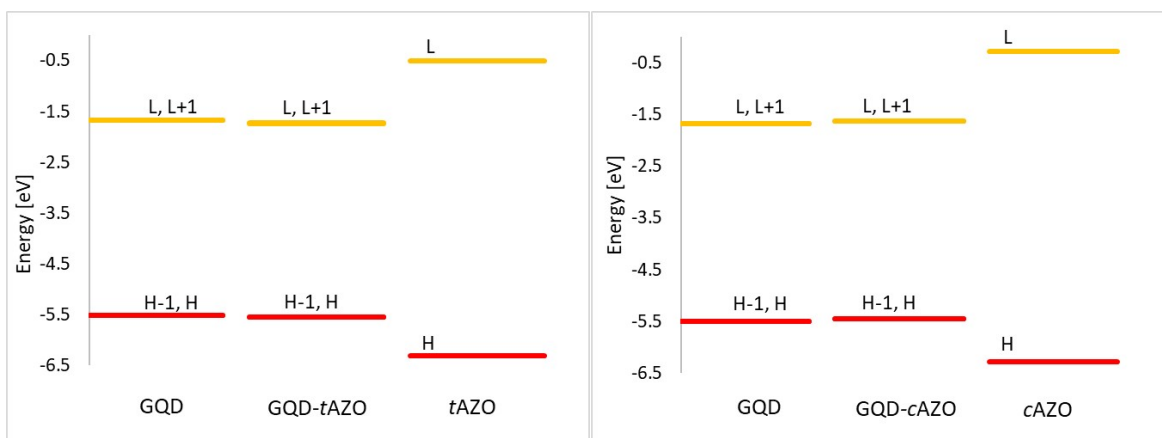


Figure S7 The alignment of frontier orbitals' energies for GQD, AZO and GQD-AZO for GQD-*t*AZO (left) and GQD-*c*AZO (right).

Dominant transitions for the interfaces

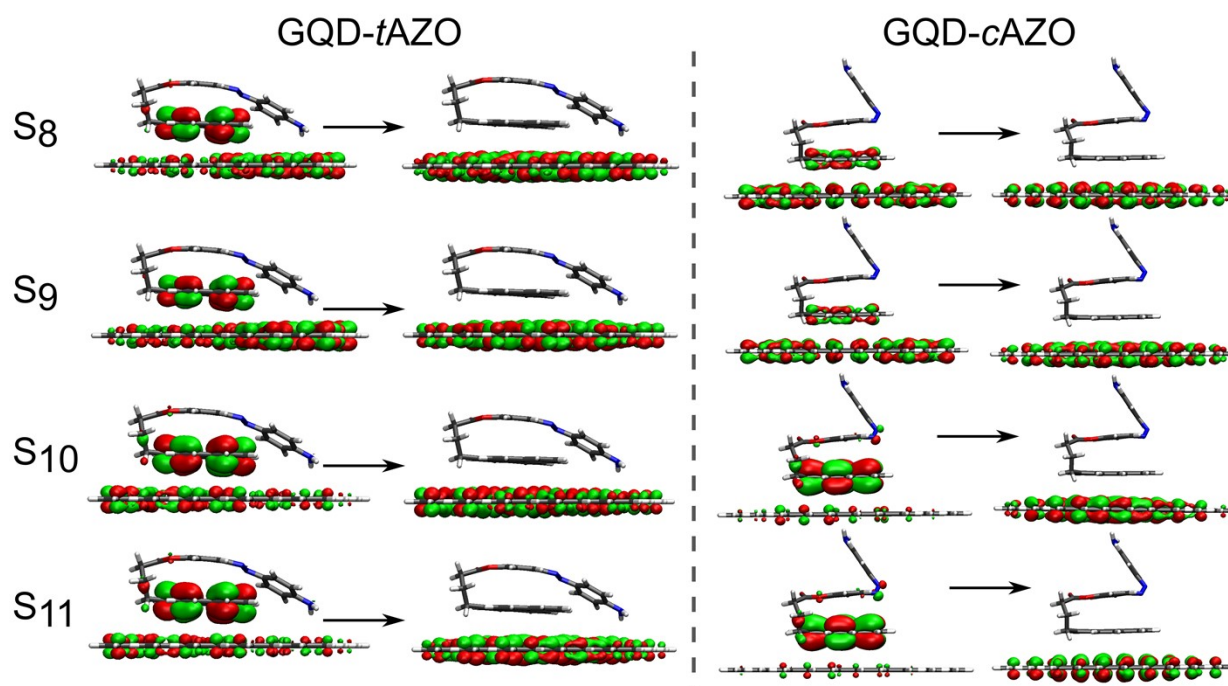


Figure S8 Natural transition orbitals for excited states that take part in PHT for GQD-*t*AZO (left) and GQD-*c*AZO (right).

Geometry of the assemblies

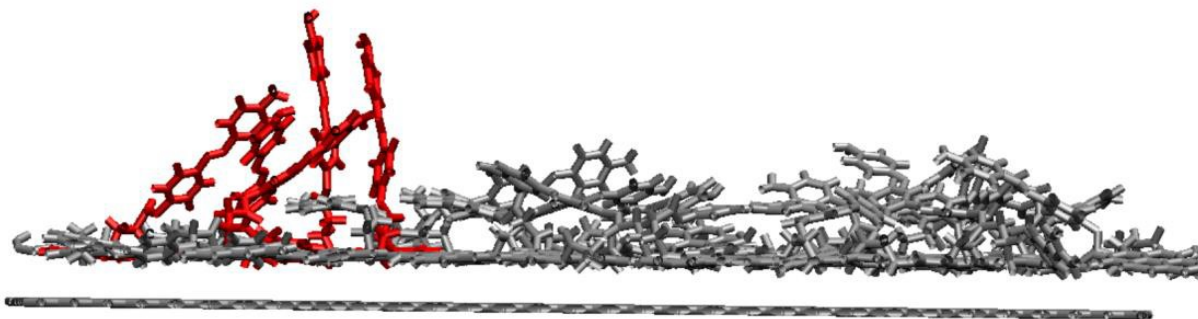


Figure S9 The final geometry of the interface composed of graphene and 42 *c*AZO molecules. Red molecules resemble open conformation of *c*AZO.

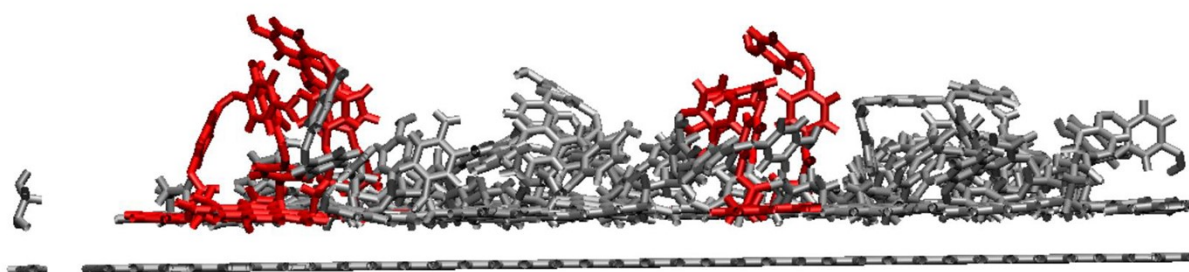


Figure S10 The final geometry of the interface composed of graphene and 42 *t*AZO molecules. Red molecules resemble open conformation of *t*AZO.

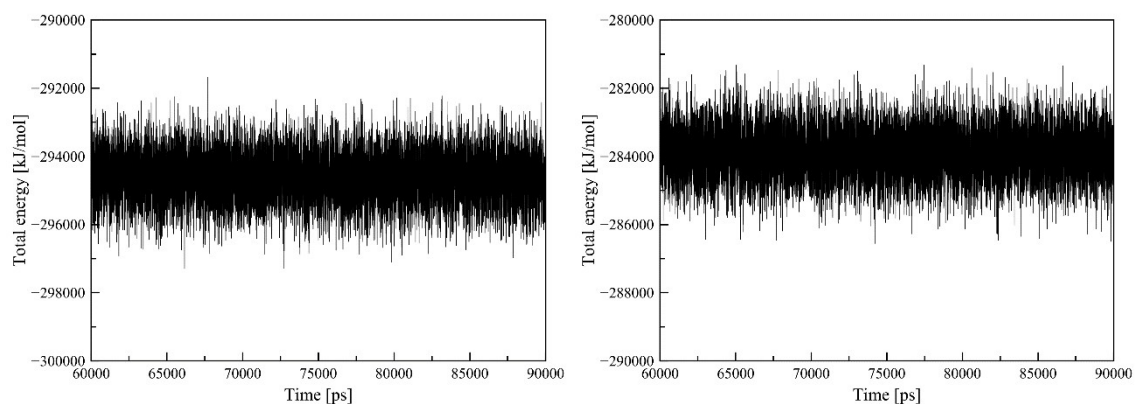


Figure S11 Total energy equilibration during MD for SLG-*t*AZO (left) and SLG-*c*AZO (right).

Photophysical properties for the interfaces from MD simulations

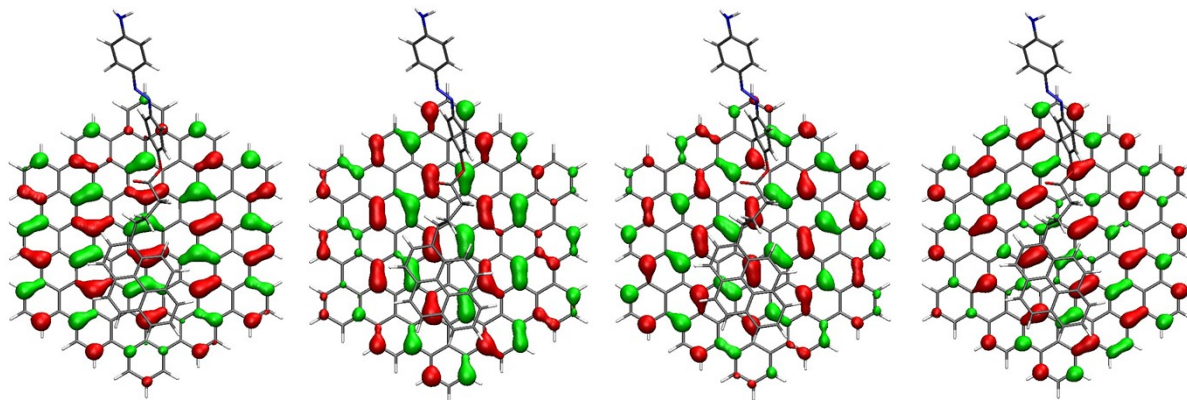


Figure S12 The visualization of FMOs for the first MD frame.

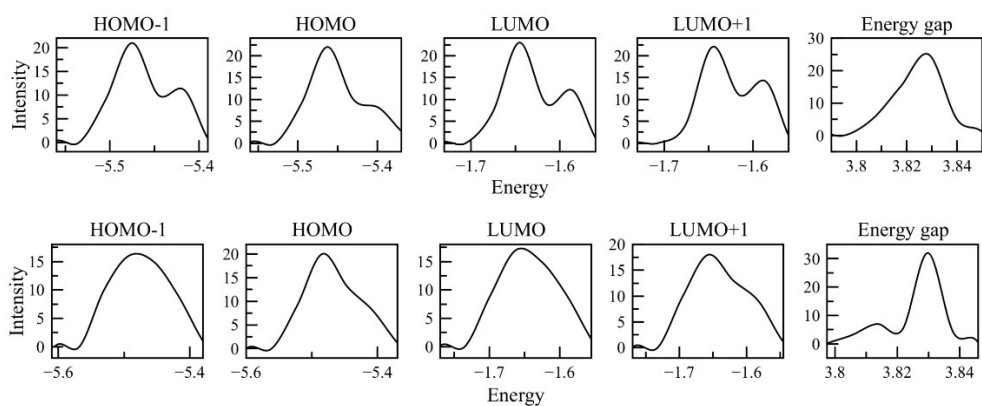


Figure S13 Distribution of FMOs' energy for the extracted frames.

Table S3. Coupling analysis for the different interfaces.

Model	Process's type	Involved transitions	Coupling [meV]
<i>t</i> AZO_fold	EET	$D_{PYR}^* (S_4), A_{AZO}^* (S_2)$	46.4
	PET	$D_{PYR}^* (S_4), D_{PYR \rightarrow AZO} (S_3)$	44.9
	PHT	$A_{AZO}^* (S_2), A_{AZO \rightarrow PYR} (S_3)$	64.3
<i>c</i> AZO_fold	EET	$D_{PYR}^* (S_2, S_3), A_{AZO}^* (S_4)$	9.0
	PET	$D_{PYR}^* (S_2, S_3), D_{PYR \rightarrow AZO} (S_{11})$	8.9
	PHT	$A_{AZO}^* (S_4), A_{AZO \rightarrow PYR} (S_{11})$	105.5
GQD- <i>t</i> AZO	EET	$D_{AZO}^* (S_7), A_{GQD}^* (S_3, S_4)$	5.3
	PET	$D_{AZO}^* (S_7), D_{AZO \rightarrow GQD} (S_{10}, S_{11})$	0.5
	PHT	$A_{GQD}^* (S_8, S_9), A_{GQD \rightarrow AZO} (S_{10}, S_{11})$	51.2
GQD- <i>c</i> AZO	EET	$D_{AZO}^* (S_3), A_{GQD}^* (S_4, S_5)$	10.3
	PET	$D_{AZO}^* (S_3), D_{AZO \rightarrow GQD} (S_{10}, S_{11})$	4.3
	PHT	$A_{GQD}^* (S_8, S_9), A_{GQD \rightarrow AZO} (S_{10}, S_{11})$	70.4
GQD- <i>t</i> AZO_MD	EET	$D_{AZO}^* (S_7), A_{GQD}^* (S_3, S_4)$	2.1
	PET	$D_{AZO}^* (S_7), D_{AZO \rightarrow GQD} (S_{10}, S_{11})$	0.2
	PHT	$A_{GQD}^* (S_8, S_9), A_{GQD \rightarrow AZO} (S_{10}, S_{11})$	41.5
GQD- <i>c</i> AZO_MD	EET	$D_{AZO}^* (S_3), A_{GQD}^* (S_4, S_5)$	6.9
	PET	$D_{AZO}^* (S_3), D_{AZO \rightarrow GQD} (S_{10}, S_{11})$	0.5
	PHT	$A_{GQD}^* (S_8, S_9), A_{GQD \rightarrow AZO} (S_{10}, S_{11})$	24.8

References

- [1] Cupellini, L., Corbella, M., Mennucci, B. & Curutchet, C. Electronic energy transfer in biomacromolecules. *WIREs Comput. Mol. Sci.* **9**, 1–23 (2019)
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