

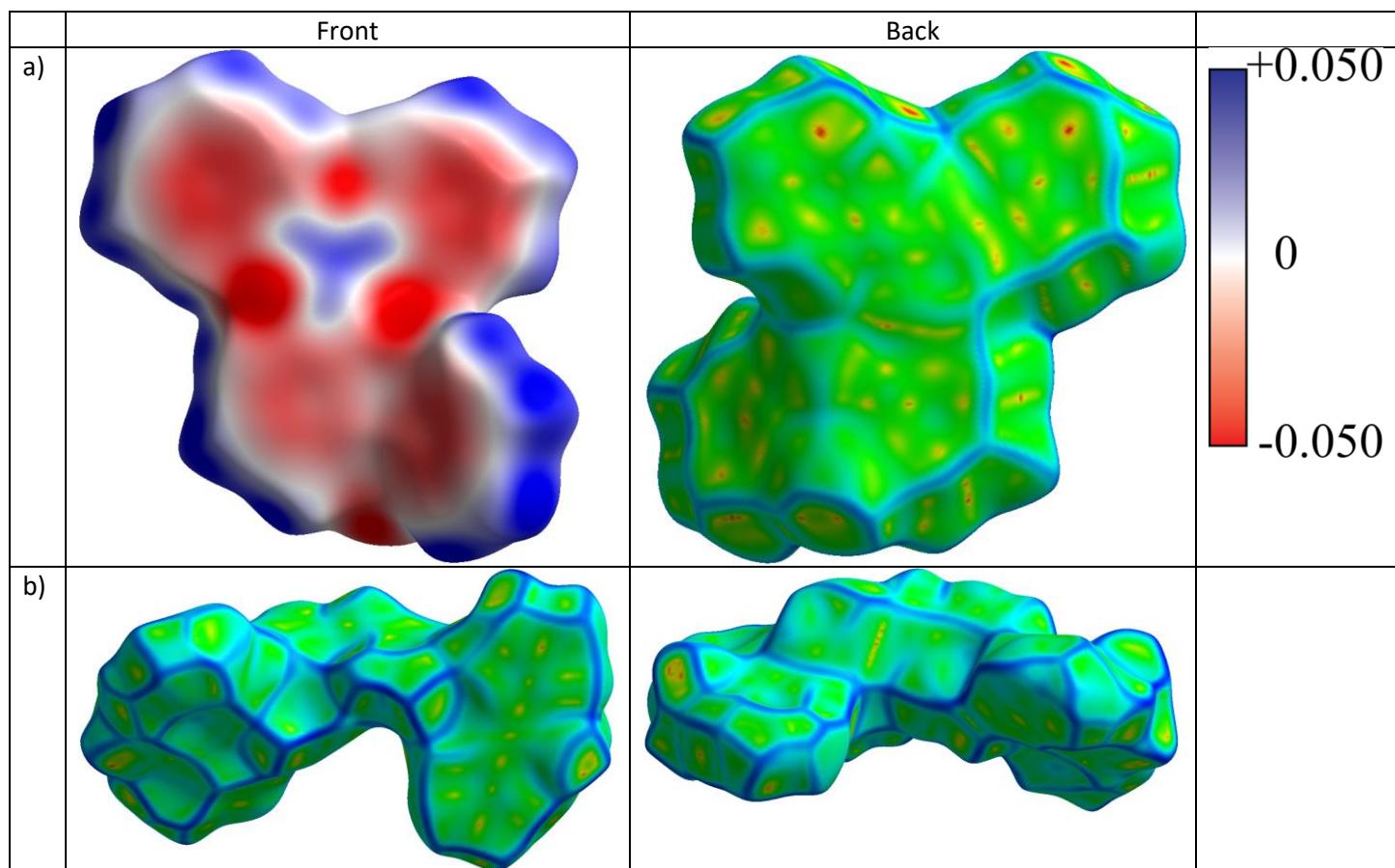
Supplementary Materials

Charge transport in organic semiconducting crystals exhibiting TADF: insight from quantum-chemical calculations

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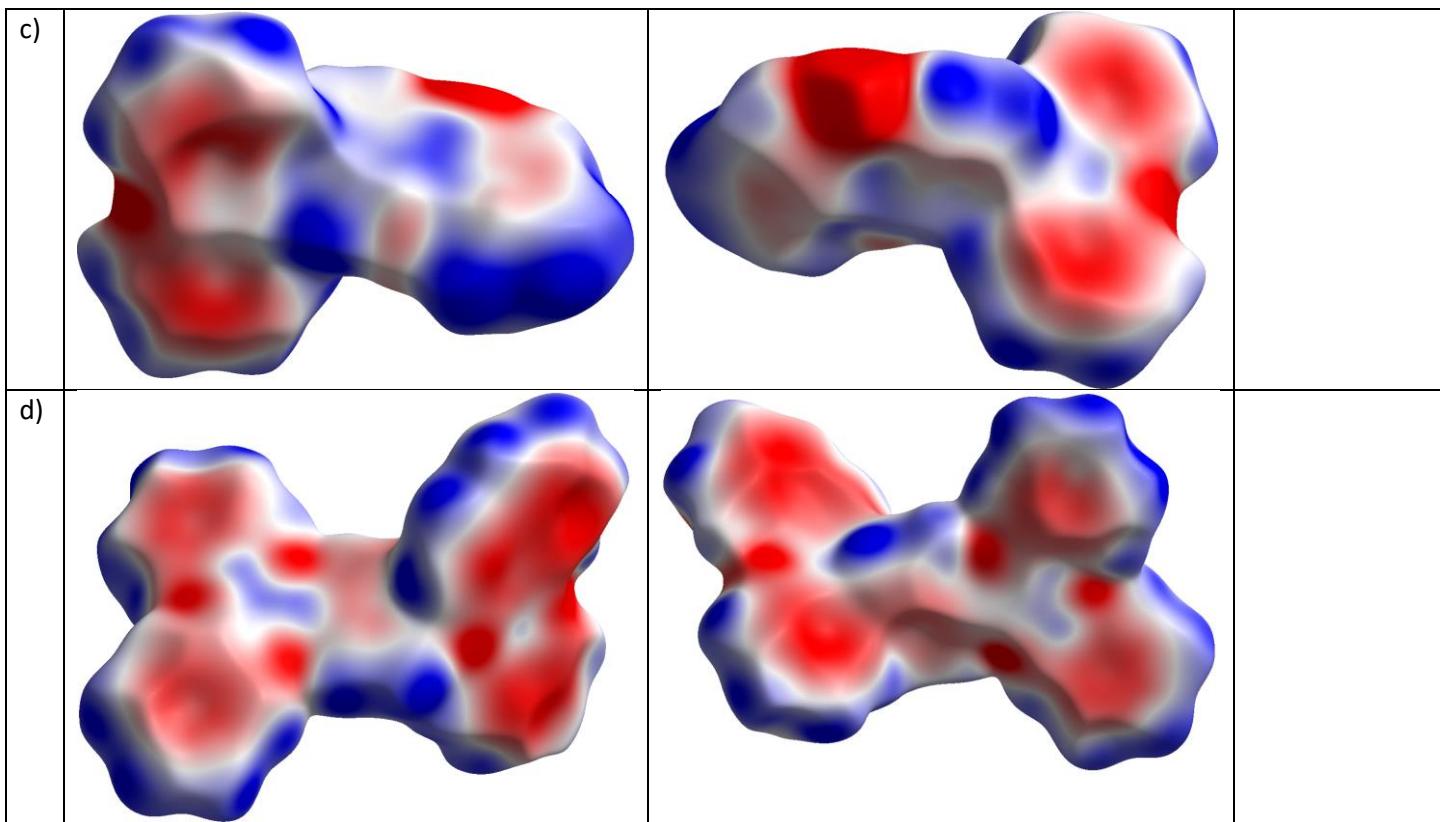


Figure S1. Hirshfeld surfaces (front and back of the molecules) mapped with electrostatic potential for crystalline oTE-DRZ (a), CPPD (b), PXZ-XO (c) and TRZ-c-BPXZ (d).

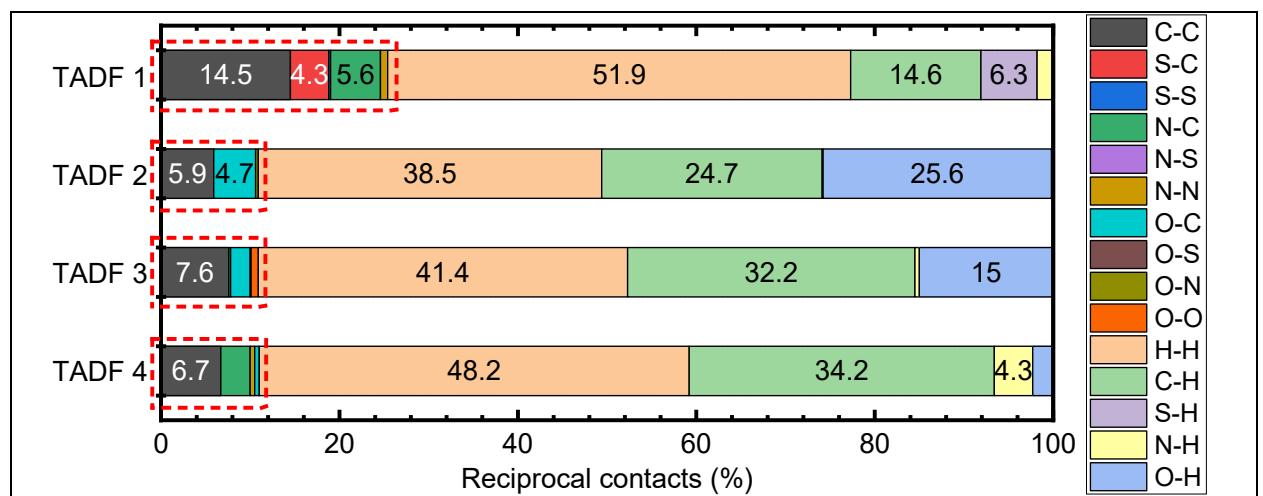


Figure S2. Distribution of reciprocal intermolecular contacts for oTE-DRZ, CPPD, PXZ-XO and TRZ-c-BPXZ crystals. “Conducting” contacts (*i.e.*, the contacts between the atoms bearing large HOMO and/or LUMO electron density) are highlighted with the red dashed frames.

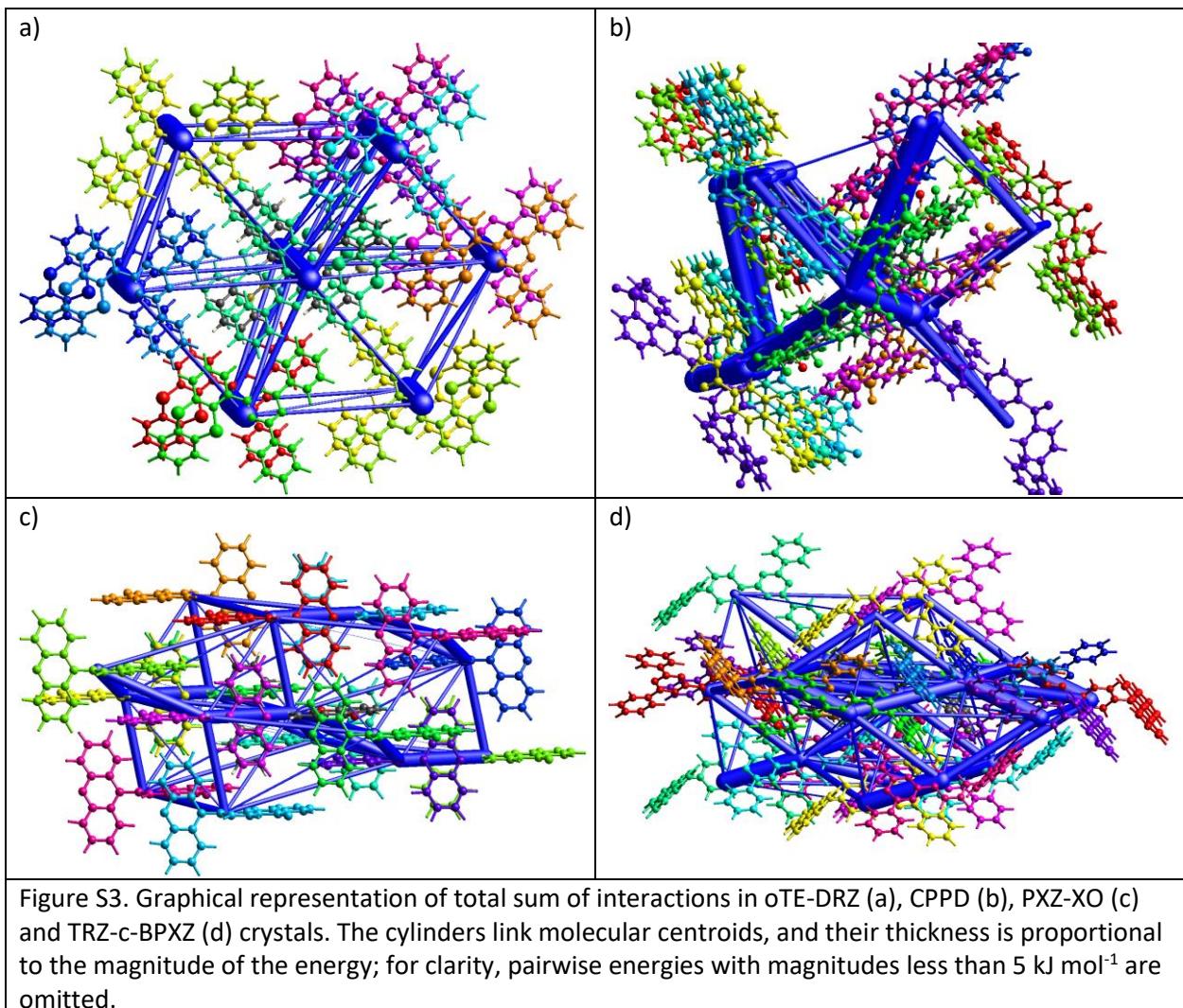


Figure S3. Graphical representation of total sum of interactions in oTE-DRZ (a), CPPD (b), PXZ-XO (c) and TRZ-c-BPXZ (d) crystals. The cylinders link molecular centroids, and their thickness is proportional to the magnitude of the energy; for clarity, pairwise energies with magnitudes less than 5 kJ mol^{-1} are omitted.

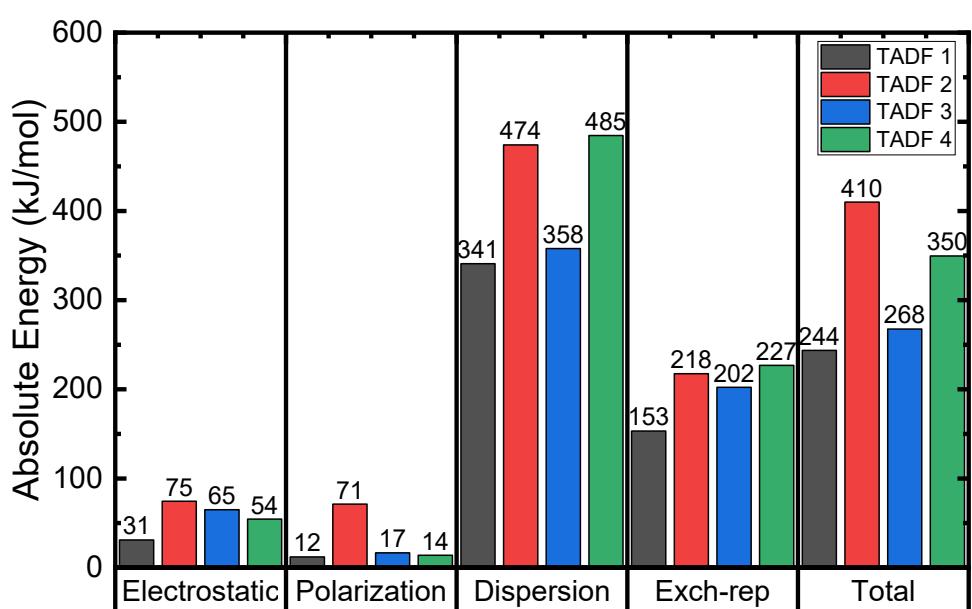


Figure S4. Distribution of sum of interaction energies (absolute values) in oTE-DRZ, CPPD, PXZ-CMO and TRZ-c-BPXZ for 3.8\AA cluster.

Table S1. Different interaction energies of the molecular pairs for oTE-DRZ in kJ mol⁻¹: N is the number of pairs, R is the distance between the molecule centroids, E_ele is the classical electrostatic energy of interaction between monomer charge distributions, E_pol is the polarization energy estimated as a sum over atoms with terms of the kind $-\frac{1}{2}\alpha|\mathbf{F}|^2$, where the electric field F is computed at each atomic nucleus from the charge distribution of the other monomer and α are isotropic atomic polarizabilities, E_dis is Grimme's D2 dispersion correction summed over all intermolecular atom pairs, E_rep is the exchange–repulsion energy, obtained from the antisymmetric product of the monomer spin orbitals, and E_tot is the total energy.

| | N | R (Å) | E_ele | E_pol | E_dis | E_rep | E_tot |
|---|-------|-------|-------|--------|-------|-------|-------|
| 1 | 11.31 | -6.8 | -0.9 | -32.0 | 17.1 | -25.1 | |
| 0 | 13.81 | 1.4 | -1.0 | -17.2 | 0.0 | -14.2 | |
| 1 | 12.44 | -1.1 | -0.4 | -12.1 | 0.0 | -11.9 | |
| 0 | 13.81 | 0.8 | -0.3 | -7.1 | 0.0 | -5.6 | |
| 1 | 12.38 | -0.5 | -0.8 | -21.6 | 0.0 | -20.0 | |
| 1 | 3.98 | -13.0 | -5.3 | -151.4 | 96.9 | -89.7 | |
| 0 | 11.88 | -1.5 | -0.4 | -10.5 | 5.6 | -7.5 | |
| 0 | 14.10 | -0.6 | -0.2 | -9.3 | 0.0 | -8.9 | |
| 1 | 13.91 | 1.2 | -0.6 | -17.6 | 0.0 | -14.5 | |
| 0 | 10.99 | -6.5 | -0.8 | -30.1 | 19.9 | -21.4 | |
| 0 | 13.50 | -0.4 | -0.6 | -11.8 | 0.0 | -11.1 | |
| 0 | 11.50 | -4.0 | -0.7 | -20.2 | 13.7 | -13.8 | |

Table S2. Different interaction energies of the molecular pairs for CPPD in kJ mol⁻¹.

| | N | R (Å) | E_ele | E_pol | E_dis | E_rep | E_tot |
|---|-------|-------|-------|--------|-------|-------|-------|
| 2 | 15.95 | 1.7 | -0.3 | -3.3 | 0 | -1.3 | |
| 1 | 6.80 | -19.1 | -13 | -88.5 | 54 | -73.5 | |
| 2 | 11.14 | -5.5 | -3.4 | -29.1 | 21.2 | -20.5 | |
| 2 | 12.99 | 18.3 | -4 | -27.1 | 0 | -7.1 | |
| 2 | 8.62 | -6.4 | -4.6 | -35.6 | 15 | -31.8 | |
| 2 | 11.36 | -25.5 | -9.4 | -40.1 | 33 | -48.4 | |
| 2 | 16.81 | -1 | -0.5 | -4.9 | 0 | -5.8 | |
| 1 | 14.18 | -3.9 | -10.4 | -66.5 | 0 | -69.8 | |
| 2 | 14.46 | -13.6 | -3.2 | -11.5 | 0 | -26.8 | |
| 1 | 5.47 | -7.6 | -17.6 | -151.9 | 94.4 | -95.1 | |
| 1 | 18.53 | -11.9 | -4.9 | -15.6 | 0 | -29.8 | |

Table S3. Different interaction energies of the molecular pairs for PXZ-XO in kJ mol⁻¹.

| | N | R (Å) | E_ele | E_pol | E_dis | E_rep | E_tot |
|---|-------|-------|-------|-------|-------|-------|-------|
| 1 | 7.27 | -9.7 | -1 | -46.8 | 29 | -33.8 | |
| 1 | 11.91 | -1.6 | -0.2 | -8.1 | 2.8 | -7.2 | |
| 1 | 12.61 | 0.3 | -0.7 | -22.3 | 0 | -19.6 | |

| | | | | | | | |
|--|---|-------|-------|------|-------|------|-------|
| | 2 | 16.27 | 0.2 | -0.1 | -2 | 0 | -1.6 |
| | 2 | 7.69 | -11.5 | -3.6 | -30.4 | 21.2 | -28.1 |
| | 1 | 6.51 | -15.9 | -3 | -61.9 | 43.7 | -45.9 |
| | 2 | 9.84 | -2.7 | -0.3 | -20.6 | 11.6 | -13.9 |
| | 1 | 14.58 | -3.3 | -0.4 | -15.3 | 0 | -17.1 |
| | 1 | 8.74 | -7.4 | -2.7 | -66.9 | 46.6 | -39.4 |
| | 1 | 8.52 | -13.4 | -3.5 | -74.9 | 47.3 | -52.7 |
| | 2 | 12.38 | 0 | -1.1 | -8.7 | 0 | -8.3 |

Table S4. Different interaction energies of the molecular pairs for TRZ-c-BPXZ in kJ mol⁻¹.

| | N | R (Å) | E_ele | E_pol | E_dis | E_rep | E_tot |
|--|---|-------|-------|-------|--------|-------|-------|
| | 2 | 16.52 | 0.2 | -0.7 | -19.1 | 0 | -16.9 |
| | 1 | 14.46 | 3.3 | -1.4 | -58.7 | 0 | -48.6 |
| | 2 | 11.76 | -3.5 | -0.7 | -17.8 | 10.3 | -13.4 |
| | 1 | 7.12 | -12.1 | -2.1 | -119.2 | 63.5 | -79 |
| | 2 | 11.28 | -7.9 | -0.9 | -34.9 | 20.8 | -26.6 |
| | 2 | 16.85 | 0.8 | -0.2 | -5.9 | 0 | -4.5 |
| | 2 | 10.02 | -11.1 | -1.6 | -57.4 | 37.7 | -39.6 |
| | 1 | 9.12 | -7 | -1.4 | -44.5 | 28.1 | -29.7 |
| | 1 | 10.71 | -11.6 | -3.6 | -86.1 | 45.9 | -61.5 |
| | 2 | 14.51 | 1.2 | -0.3 | -9.9 | 0 | -7.5 |
| | 2 | 8.65 | -4.4 | -0.6 | -21.9 | 16.7 | -13.9 |
| | 2 | 11.52 | -2.3 | -0.4 | -9.3 | 3.9 | -8.4 |

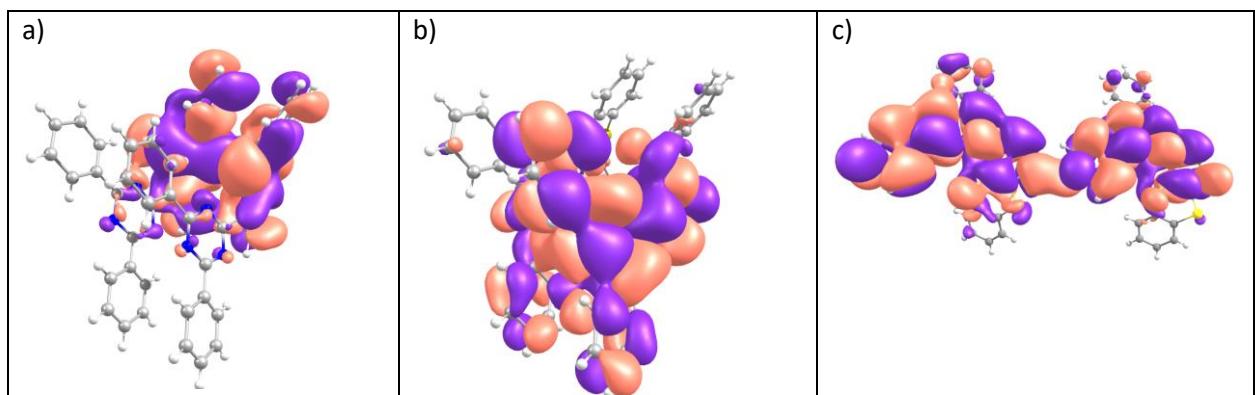


Figure S5. (a) HOMO and (b) LUMO of the oTE-DRZ π -stacking dimer. (c) LUMO of the oTE-DRZ dimer showing the largest electron-transfer integral.

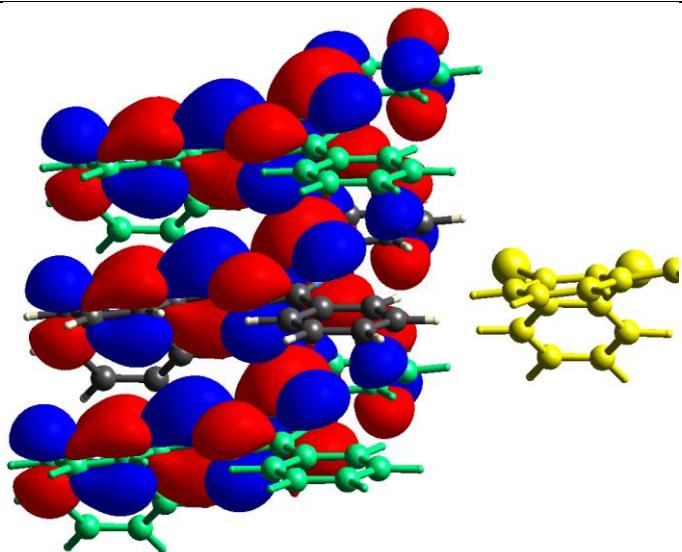


Figure S6. LUMO of the oTE-DRZ π -stack. Wavefunction phase mismatch between adjacent molecules is clearly observed.