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Controllable molecular aggregation and fluorescence properties of 1,3,4-oxadiazole derivatives†

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The molecular self-assembly behaviour of 2,2'-bis-(4-hexyloxyphenyl)-bi-1,3,4-oxadiazole (BOXD-6) in solution, on surfaces and in bulk crystals, and its photo-physical properties were studied via a combination of experimental techniques and theoretical calculations. It is found that BOXD-6 molecules self-assemble into both H- and J-aggregates at moderate concentration ($\sim 10^{-4}$ M) and then transit to exclusive J-aggregates at higher concentration ($\sim 10^{-3}$ M) in tetrahydrofuran. In H-aggregation (α polymorph), BOXD-6 adopts a linear conformation and forms a one-dimensional layered structure; in J-aggregation (β polymorph), it adopts a Z-shaped conformation and forms a more ordered two-dimensional layered structure. A π -stacking structure is observed in both cases, and adjacent molecules in J-aggregation show larger displacement along the molecular long axis direction than that in H-aggregation. Although J-aggregates are almost the only component in concentrated solutions (10^{-3} M), both H- and J-aggregates can be obtained if concentrated solution is transformed onto substrates through a simple drop-casting method. Such a phase transition during film formation can be easily avoided by adding water as a precipitator; a film with pure J-aggregates is then obtained. In order to get more information on molecular self-assembly, intermolecular interaction potential energy surfaces (PES) were evaluated via theoretical calculations at the DFT level (M062x/6-31G**). The PES not only confirm the molecular stacking structures found in crystals but also predict some other likely structures, which will be the target of future experiments.

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1 Introduction

π -Conjugated organic semiconductors have attracted much attention due to their applications in low-cost and flexible organic electronic devices, *e.g.*, organic light-emitting diodes (LEDs),^{1–7} field-effect transistors (FETs),^{8–12} and solar cells (SCs).^{13–18} It has been demonstrated that not only the molecular structure but also the molecular packing in the solid state plays an important

role in the light-emitting and charge-transporting properties.^{19–25} Although great progress has been made in understanding the effect of the molecular structure on the optoelectronic properties, a challenge still remains in developing the transition from some well-established ‘molecular’ characteristics to some desirable ‘material’ (collective) properties.²¹ For example, molecules can show high light-emission efficiency in dilute solutions; however, quenching occurs in concentrated solutions and solid states because of the aggregation of chromophores.²⁶ Therefore, it is useful to develop organic solid-state films with high light-emission efficiency.

Molecular crystals, where the molecular conformational and packing structures are clear, provide ideal models for studying the relationship between the molecular arrangements and the semi-conducting properties in the solid state. Thus, they have attracted many efforts both in academia and in industry.^{23–25,27–30} However, many of these crystals grow spontaneously and it is highly desirable to gain control over the molecular self-assembled structures. With this goal in mind, here we present the characterization of the molecular self-assembled structures of 2,2'-Bis-(4-hexyloxyphenyl)-bi-1,3,4-oxadiazole (BOXD-6, Scheme 1)

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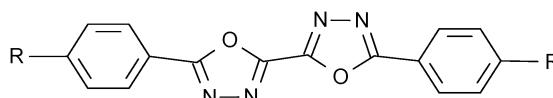
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BOXD-1: R = -OCH₃, BOXD-6: R = -OC₆H₁₃Scheme 1 The molecular structure of BOXD-*n* (*n* = 1, 6).

in solutions, on surfaces, and in bulk crystals. An effective way was discovered to control the molecular self-assembly on substrates. In order to get an in-depth understanding of the molecular self-assembly, the intermolecular interaction potential energy surfaces were also evaluated through theoretical calculations at the Density Functional Theory (DFT) level.

2 Experimental and computational details

2.1 Materials and experimental techniques

The synthesis of BOXD-6 was reported previously; the purity was verified by FT-IR, ¹H NMR spectroscopy, and elemental analysis.³¹ All the solvents for spectral measurements were of spectroscopic grade and used as received. UV-vis absorption spectra were obtained using a Shimadzu UV-2550 spectrometer. Photoluminescence spectra were collected using a Perkin-Elmer LS55 spectrophotometer. The room-temperature luminescence quantum yields in solutions were determined relative to quinine sulfate in sulfuric acid aqueous solution (0.546), and calculated according to the following equation: $\Phi_{\text{unk}} = \Phi_{\text{std}}(I_{\text{unk}}/A_{\text{unk}})(A_{\text{std}}/I_{\text{std}})(\eta_{\text{unk}}/\eta_{\text{std}})^2$, where Φ_{unk} is the radiative quantum yield of the sample; Φ_{std} is the radiative quantum yield of the standard; I_{unk} and I_{std} are the integrated emission intensities of the sample and the standard, respectively; A_{unk} and A_{std} are the absorptions of the sample and the standard at the excitation wavelength, respectively; and η_{unk} and η_{std} are the indexes of refraction of the sample and standard solutions (pure solvents were assumed), respectively. Powder X-ray diffraction (XRD) measurements were carried out using a Bruker Avance D8 X-ray diffractometer (Cu K α radiation, $\lambda = 1.54$ Å). The single crystal data were collected on a Rigaku R-AXIS RAPID IP diffractometer using a graphite monochromator for Mo K α radiation ($\lambda = 0.71073$ Å) at room temperature (293 K) using the ω -scan technique. Absorption corrections were applied. The structures were solved by direct methods and refined by full matrix least-square techniques using the SHELXTL package. Anisotropic thermal parameters were assigned to all non-hydrogen atoms. The hydrogens were included in the structure factor calculation at idealized positions by using a riding model and refined isotropically. The crystallographic data of the derivatives are listed in Table 1. SEM measurements were taken using a JEOL JSM-6700F apparatus. All the samples were coated with gold before observation.

2.2 Computational details

In the calculation, the hexyloxy group (-OC₆H₁₃) of BOXD-6 was replaced by a methoxy group (BOXD-1, Scheme 1) to save

Table 1 Summary of crystallographic data for the two polymorphs of BOXD-6

	α	β
Empirical formula	C ₂₈ H ₃₄ N ₄ O ₄	C ₂₈ H ₃₄ N ₄ O ₄
Molecular weight	490.59	490.59
Crystal system	Triclinic	Triclinic
Space group	$P\bar{1}$	$P\bar{1}$
<i>Z</i>	2	1
<i>a</i> , Å	8.166(6)	8.134(6)
<i>b</i> , Å	13.066(9)	8.142(7)
<i>c</i> , Å	14.357(14)	10.657(12)
α , °	65.13(3)	110.13(4)
β , °	73.59(3)	93.00(4)
γ , °	86.14(3)	98.55(3)
<i>V</i> , Å ³	1330.84	651.219
ρ	1.224	1.251
Total reflections	10 326	6429
Unique reflections	4582	2936
<i>R</i> _{int}	0.0456	0.0282
<i>R</i> ₁ , <i>wR</i> ₂	0.0699, 0.1677	0.0507, 0.1281
<i>R</i> ₁ , <i>wR</i> ₂ (all data)	0.1521, 0.2043	0.0883, 0.1432

computational time. The monomer structure of BOXD-1 in the ground state was optimized at the M062x/6-311+G** level.³² The intermolecular interaction potential energy surfaces (PESS) were constructed first by calculating single-point energies with a face-to-face dimer model, where the non-covalent interactions are considered with the M062x functional; this functional has a good track record in investigations of main-group thermochemistry, kinetics, non-covalent interactions, and electronic structures.³²⁻³⁴ The monomer geometry obtained at the M062x/6-311+G** level was used for further PES scans of BOXD-1 dimers. The energy scans start from an ideal face-to-face π -stacked structure with an intermolecular separation of 3.33 Å (taken from crystal data),³¹ and then move over the displacements along both the molecular long axis (*y*-displacement, $0 \leq y \leq 11.0$ Å) and the short axis (*x*-displacement, $0 \leq x \leq 2.0$ Å) in steps of 0.2 Å. All the calculations for molecular dimers were done at the M062x/6-31G** level; the counterpoise procedure was utilized to correct for basis set superposition error (BSSE). The Gaussian 09 software package (version A.02) was used for all the calculations.³⁵

The energy decomposition analysis was carried out using the open-source *ab initio* Electronic Structure Package PSI 4.0.0-beta5 driver (SAPT0, paired with the jun-cc-pVDZ basis set).³⁶

3 Results and discussion

3.1 Concentration-dependent photophysical properties of BOXD-6 in THF

As shown in Fig. 1a and b and reported previously, BOXD-6 shows an intense absorption band at around 320 nm and strong emission at 386 nm in dilute tetrahydrofuran (THF) solution ($\sim 10^{-6}$ M).³⁷ The position of the main absorption band is almost independent of the concentration (Fig. 1a); only relative absorption intensity of short wavelength (under 300 nm, Fig. S1a, ESI†) and the peak width increase (more obvious from 2×10^{-4} M, Fig. S1b, ESI†) with the increasing concentration, indicative of the



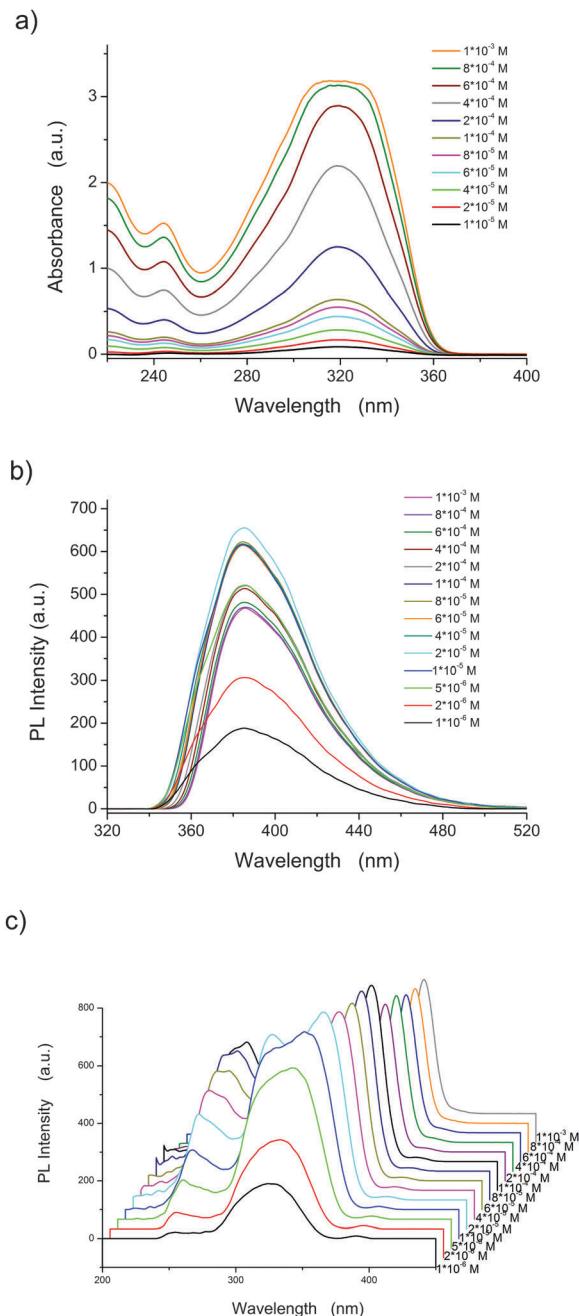


Fig. 1 UV-vis absorption (a), photoluminescence (PL) emission (b), and excitation (c, $\lambda_{\text{em}} = 386$ nm) spectra of BOXD-6 in tetrahydrofuran at different concentrations.

formation of molecular aggregates. These molecular aggregates can be further confirmed by the photoluminescence excitation and emission spectra (more sensitive to molecular aggregates, Fig. 1b and c). It can be seen that the excitation spectrum at the lower concentration ($\sim 10^{-6}$ M) in THF presents a main peak at 320 nm (Fig. 1c and Fig. S2, ESI[†]), and is consistent with the shape of the absorption spectrum (Fig. 1a). These observations strongly suggest that BOXD-6 is molecularly dissolved in THF at about 10^{-6} M or below. With the increase in concentration, a significant broadening of the excitation peak is observed

(Fig. S2a, ESI[†]), indicating the formation of aggregates. Two new bands, *i.e.* a blue-shifted one (~ 300 nm) and another red-shifted one (~ 350 nm), were observed at 2×10^{-5} M (Fig. S2b, ESI[†]), indicative of the simultaneous formation of H- and J-aggregates. With a further increase in concentration, the two bands split more obviously due to their opposite-directional shift, and the intensity of the peak at 300 nm (H-aggregates) decreases compared to the one at 355 nm; the peak at 355 nm (J-aggregates) red-shifts gradually to 370 nm and is the only one prominent at higher concentrations (Fig. S2b, ESI[†]), implying that J-aggregates are the main components in concentrated solutions ($\sim 10^{-3}$ M). It should be noted that the intensity of the band at 250 nm, which should be assigned to those transitions to the higher excited states,³⁷ also increased when the concentrations are increasing from 1×10^{-6} to 1×10^{-4} M, but decrease on further concentration increase (Fig. 1c and Fig. S2b, ESI[†]), indicating that the intensity change should be related with the formation of H-aggregates. However, a detailed mechanism is still beyond our understanding. The emission spectra of BOXD-6 show obvious changes in the peak shape rather than in band shifts (Fig. 1b and Fig. S3, ESI[†]): upon increasing concentration, the intensity of the higher energy vibrational band decreases gradually, this might be attributed to the re-absorption of molecular aggregates. These molecular aggregates can be observed using a fluorescence microscope with the mean size of ~ 4 μm and ~ 6 μm from 1×10^{-4} and 1×10^{-3} M THF solutions, respectively (Fig. S4, ESI[†]). These observations indicated that photoluminescence emission and excitation, especially excitation spectra, are more sensitive to molecular aggregates than UV-vis absorption spectra. Only peak broadening was observed in absorption spectra might be due to the simultaneous formation of H- and J-aggregates, and the nature that absorption spectra are less sensitive to molecular aggregates.

The interesting trends in quantum yields with increasing concentration (Fig. 2) can be explained by the observed phase transitions of molecular aggregation in solution. The initial decrease in quantum yield (from 1×10^{-5} to 6×10^{-5} M) can be attributed to concentration-quenching *via* the formation of H-aggregates; consequent increase (from 8×10^{-5} to 4×10^{-4} M) is probably caused by the increasing population of J-aggregates due

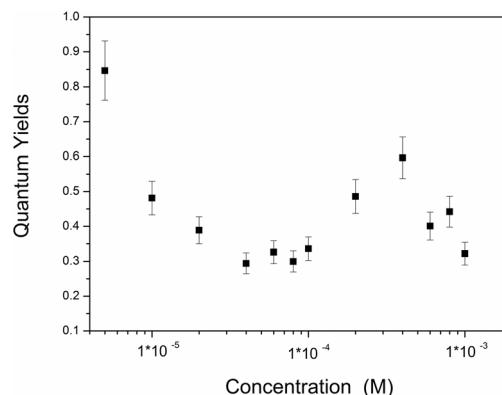


Fig. 2 Concentration-dependent photoluminescence quantum yields of BOXD-6 in tetrahydrofuran (THF).

to the transition from H-aggregation to J-aggregation; and the final decrease (from 4×10^{-4} to 1×10^{-3} M) might be due to serious re-absorption of these molecular aggregates observed in the thick solution. H-aggregates are generally non-emissive; however, in this case, excitations at both 300 (H-aggregates) and 355 nm (J-aggregates) could lead to emission at 386 nm, which might suggest that energy transfer from H-aggregates to J-aggregates is involved, however, unfortunately, there is no further evidence.

3.2 Crystal structures of BOXD-6

Single crystals of BOXD-6 suitable for X-ray analysis were grown by slow evaporation from dilute acetonitrile (ACN), chloroform (CHL), tetrahydrofuran (THF), or ethanol (ETO) solutions ($\sim 10^{-4}$ M). Two polymorphs of BOXD-6 crystals were obtained; one polymorph (referred to as α , see below) was grown under various conditions (in different solvents), while the other one (β) was sometimes found to grow from ETO. Table 1 shows the crystallographic data of BOXD-6 from X-ray structural analysis. The ORTEP representations of the molecular structures are displayed in Fig. 3.

The molecules adopt different conformations and have very different packing structures in the two polymorphs (Fig. 3 and 4). In α polymorph, BOXD-6 adopts a linear conformation (Fig. 3a) and crystallizes in the triclinic space group $P\bar{1}$. Such linear molecules stack up along the crystal a axis and form molecular layers parallel to the ab plane (Fig. 4a). To geometrically analyze the extent of π -stacking of neighboring molecules (dimers), the long molecular axis is defined as the line through the ether O atom and the molecular center, the short molecular axis is in the molecular plane and perpendicular to it. The nearest neighbors of BOXD-6 are separated by about 3.58 Å in the stacking direction, and are anti-parallel to each other along the long molecular axis with a rotation of 21.4° (Fig. 4c). Neglecting this rotation, adjacent molecules slip from a 'co-facial' π -stacking by 6.46 Å along the long molecular axis direction (D_1) and 1.08 Å in the short molecular axis direction (D_2).

The β polymorph also shows triclinic symmetry and a π -stacked structure; here, the molecules adopt a Z-shaped conformation (Fig. 3b). These Z-shaped molecules stack up along the crystal a axis and form a more ordered two-dimensional layered structure; the molecular layer parallel to the ac plane is of

single molecular layer type, while the molecular layer parallel to the ab plane had the alkoxy chains intercalated, termed as the intercalated layer (Fig. 4b).^{38,39} The molecules in the stacking direction are arranged in a parallel fashion and separated by 3.37 Å (Fig. 4d). The molecules in the crystals of β polymorph are packed more tightly than that in α polymorph, which is confirmed by the calculated density. The density of the crystal of β polymorph is 1.251 g cm⁻³, which is slightly larger than that of α polymorph, 1.224 g cm⁻³ (Table 1). Adjacent molecules show larger displacements along the long molecular axis direction, calculated to be 7.43 Å (D_1), but almost no displacement along the short molecular axis direction (D_2). In comparison with that of BOXD-1,³¹ where $D_1 = 5.50$ Å and D_2 is small, it appears that the introduction of the long terminal alkoxy chains enlarged the intermolecular displacements and result in J-aggregation.

3.3 Fine tuning the molecular aggregation structure

Even though the J-aggregates are almost the only component at a concentration of 10^{-3} M, a solid film with complex layered structures (Fig. 5a) is found when we transfer these molecular aggregates onto a substrate (silicon or glass) using a drop-casting method. The XRD pattern of this solid film shows several sharp peaks in the low-angle region with a d spacing of 12.80, 10.10, 7.53 and 6.40 Å. The diffraction peak at $d = 12.80$ Å and its harmonic peak at $d = 6.40$ Å are very similar to the diffraction peak found in the crystal of α polymorph (Fig. 5c); the other two peaks (at $d = 10.10$ and 7.53 Å) are similar to the diffraction pattern of crystals of β polymorph (Fig. 5d). This implies that both H- and J-aggregation modes form in the solid film, and J-aggregates can transform into H-aggregates during film formation. A question arises: how can we get only J-aggregation in the film? By examining the whole procedure of film formation, we think that several factors, for example, the interaction between the molecular aggregates and substrates or the evaporation of the solvents, can have strong influence on the film structure. So, in order to avoid these effects, we caused the J-aggregates first to precipitate from solution by adding water into the THF solution; then, these suspended precipitates in solution were transferred onto the substrates. Films with only J-aggregates are successfully obtained *via* this method (see the XRD curve in Fig. 5b). This observation can be further demonstrated by spectroscopic study. As shown in Fig. S5 (ESI[†]), increasing the water ratio can decrease the intensity of the peak at 295 nm, indicating that more J-aggregates can be stabilized by adding water into THF solution.

SEM was used to observe the morphology of the films (Fig. 6). As shown in Fig. 6a, the SEM images suggest that thick plates with sharp crystal faces are formed in drop-cast films. These plates are about 100 nm in thickness and several micrometers in other two dimensions. The BOXD-6 aggregates precipitated by adding water into THF solutions assemble into relatively thinner sheets (about 10 nm, Fig. 6b), which are several micrometers in length and 0.5 μ m in width; this indicates that molecules essentially assemble along one dimension.

The possible mechanism for the phase transition from J-aggregates to H-aggregates during film formation can be

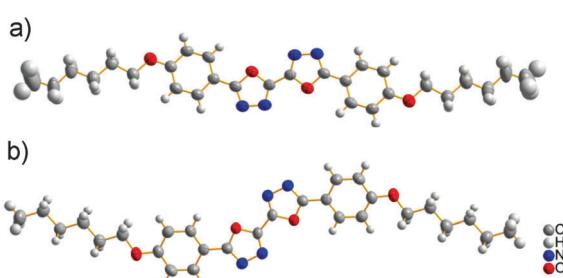


Fig. 3 The molecular structures of BOXD-6 with displacement ellipsoids drawn at the 50% probability level: (a) from α polymorph, linear shape and (b) from β polymorph, Z-shape.



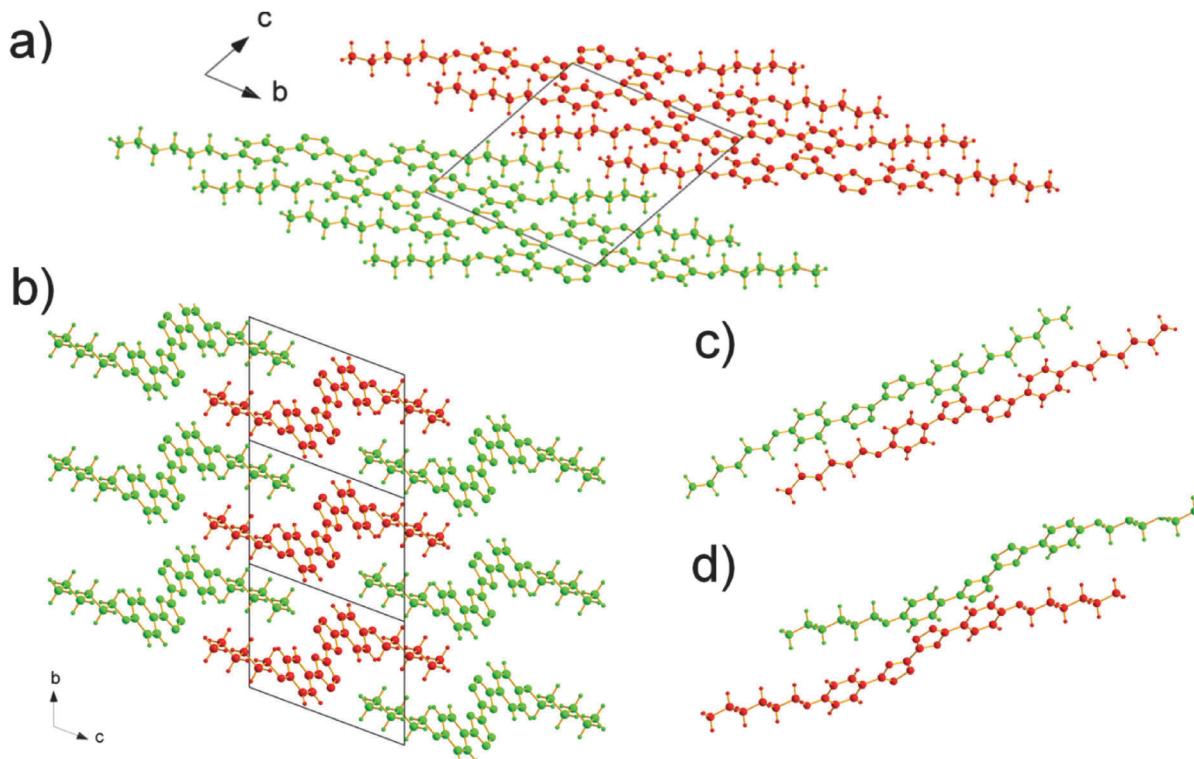


Fig. 4 The molecular stacking structures of BOXD-6 in both α and β polymorphs, where molecules are colored green or red to make the layer structure or molecular stacking structure more distinguishable. (a) α polymorph, viewed from the crystal a axis: molecules assemble into a layered structure. (b) β polymorph, viewed from the crystal a axis: molecules assemble into a layered structure with two-dimensional order, and the molecular layer can be observed in parallel with both ab and ac planes; (c) in α polymorph, the nearest neighbors in the stack are antiparallel to each other, and are separated by about 3.58 Å; and (d) in β polymorph, the nearest neighbors in a stack are parallel to each other, and are separated by about 3.37 Å.

attributed to the ease of the formation of α polymorph (H-aggregation), as has been confirmed in single crystal growth experiments. The crystallization of α polymorph disturbed the equilibrium of the molecular conformation and aggregation, making molecules in J-aggregates re-dissolved into solvents, change their conformation, and re-crystallize as H-aggregates. Adding water into the solution can make J-aggregates precipitate out and prevent the following re-dissolving process.

3.4 Potential energy surface (PES)

In order to get further information on molecular aggregation, the intermolecular interaction potential energy surfaces (PESs) were evaluated by calculating the single-point energies using a dimer model. As shown in Fig. 7, the dimer structures were built up in a π -stacked face-to-face orientation with an intermolecular separation of 3.33 Å. In order to simulate the molecular packing structures in the crystals, both parallel and anti-parallel dimer models were taken into account in this study. The potential energy was computed as a function of molecular shifts along both the molecular long axis (y -displacement) and short axis (x -displacement) at the M062x level.³²

In both configurations, the intermolecular interaction potential energy surface points to several energy minimum states (Fig. 8 and Fig. S6, ESI[†]). For the anti-parallel stacking mode, the global minimum is found at $x = 0.4$ Å, $y = 4.0$ Å (denoted as $M_{AP}1(0.4, 4.0)$, and the subscript AP indicates

anti-parallel packing). The binding energy is calculated to be $E_{bin} = -10.98$ kcal mol⁻¹. Besides, two local energy minimum points, $M_{AP}2(0, 1.6)$ and $M_{AP}3(0.2, 6.2)$, are also predicted. The energy difference for these energy minima is under 0.5 kcal mol⁻¹. For the parallel stacking mode, the global minimum is found at $x = 1.0$ Å, $y = 5.2$ Å (denoted as $M_P1(1.0, 5.2)$, and the subscript P indicates parallel packing). The binding energy is calculated to be $E_{bin} = -12.37$ kcal mol⁻¹. Besides, several local energy minimum points, $M_P2(0.6, 7.4)$, $M_P3(1.4, 2.8)$ and $M_P4(0, 1.4)$, are also obtained. The energy difference among these energy minima is smaller than 3 kcal mol⁻¹. The molecular packing structures corresponding to $M_{AP}3(0.2, 6.2)$, $M_P1(1.0, 5.2)$ and $M_P2(0.6, 7.4)$ are very similar to that found in the crystals. If we use the same definition of the molecular long axis and short axis as in the crystal structure, the displacement (D_1 and D_2) of the neighboring molecules in these theoretical predicted structures are very close to that found in corresponding crystal structures (Table 2). The difference in D_1 is less than 0.3 Å; a relatively larger deviation could be found for D_2 (less than 0.6 Å), which might be due to the small energy differences involved in x displacements.

In addition, to get better insight into the origins of the binding in terms of the various fundamental intermolecular forces, we carried out a Symmetry Adapted Perturbation Theory (SAPT) analysis on these energy-minimum molecular stacking structures, as well as the crystal packing and the ideal face-to-face stacking (referred to as $H(0,0)$ structures. As shown in

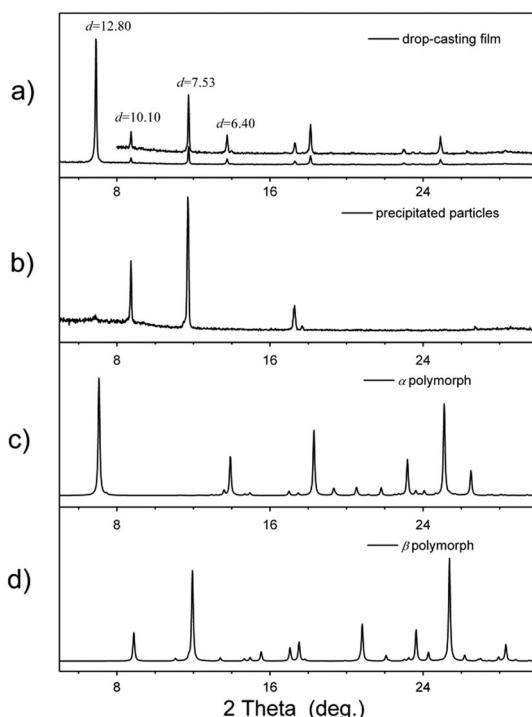


Fig. 5 XRD patterns of BOXD-6 in different states: (a) drop-casting films by directly dropping thick tetrahydrofuran solution of BOXD-6 ($\sim 10^{-3}$ M) onto the substrates; (b) precipitated particles by adding water into a thick tetrahydrofuran solution of BOXD-6; (c) calculated powder XRD curve from the single crystal structure of BOXD-6 in α polymorph; and (d) calculated powder XRD curve from the single crystal structure of BOXD-6 in β polymorph.

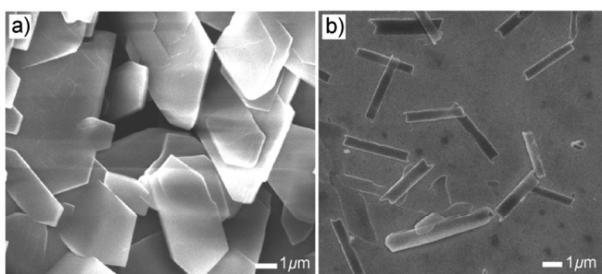


Fig. 6 SEM images of BOXD-6 in different states: (a) drop-cast films by directly dropping a thick tetrahydrofuran solution of BOXD-6 ($\sim 10^{-3}$ M) onto the substrates and (b) precipitated particles by adding water into a thick tetrahydrofuran solution of BOXD-6.

Fig. 9, London dispersion forces are the strongest attractive components in the binding (~ -25 to -45 kcal mol $^{-1}$). The electrostatic stabilization (~ -8 to -13 kcal mol $^{-1}$) is about 1/3–1/2 of the dispersion term. Induction, at about -1.5 to -5 kcal mol $^{-1}$, provides the remaining stabilization. The dominance of dispersion is consistent with the results of many other π -stacked systems.^{34,40,41} Compared to electrostatics and induction interactions, exchange and dispersion interactions are more sensitive to the molecular packing structures. The most favorable dispersion energy is for ideal face-to-face configurations ($H_{AP}(0,0)$ and $H_P(0,0)$); this is consistent with the fact that

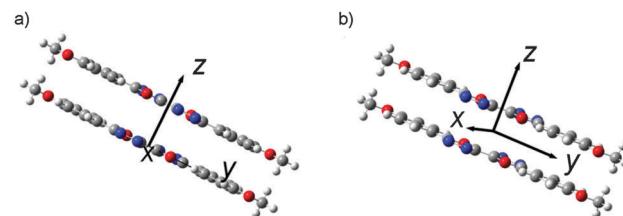


Fig. 7 Intermolecular coordinates (dimer model) used for scanning the intermolecular interaction potential energy surface: (a) the molecules stack antiparallel to each other (anti-parallel model) and (b) the molecules stack parallel to each other (parallel model).

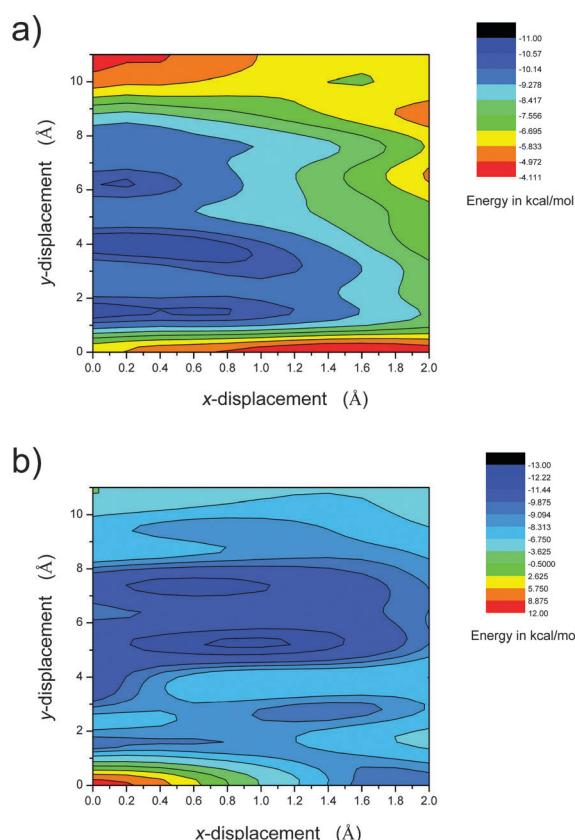


Fig. 8 Contour images of the intermolecular interaction potential energy surface of BOXD-6. The potential energies were computed as a function of molecular shifts along both the molecular long axis (y-displacement) and the short axis (x-displacement) using the DFT M062x/6-31G** method based on a dimer model. (a) For the anti-parallel model, the molecules stack antiparallel to each other and (b) for the parallel model, the molecules stack parallel to each other.

in these configurations, the two molecules are showing the closest contact. However, the greatest exchange repulsive forces are, as expected, also observed in these configurations, especially in the parallel mode (63.35 kcal mol $^{-1}$). This is the reason why such face-to-face configurations are not favored. Other configurations with moderate dispersion and exchange terms are more favored (~ -16 kcal mol $^{-1}$). The interaction energy obtained in SAPT analysis is slightly larger than that from the M062x/6-31G** method (Fig. S7, ESI †), due to the difference in the methodologies.



Table 2 Comparing the intermolecular displacements (D_1 and D_2) in the theoretically predicted and crystal structures

No.	Stacking structure	D_1 (Å)	D_2^a (Å)
1	$M_{AP3}(0.2, 6.2)$	6.18	0.45
	BOXD-6- α	6.46	1.08
2	$M_P1(1.0, 5.2)$	5.27	0.44
	BOXD-1	5.50	0
3	$M_P2(0.6, 7.4)$	7.42	0.11
	BOXD-6- β	7.43	0

^a The values for D_2 in BOXD-1 and BOXD-6- β crystal are negligible.

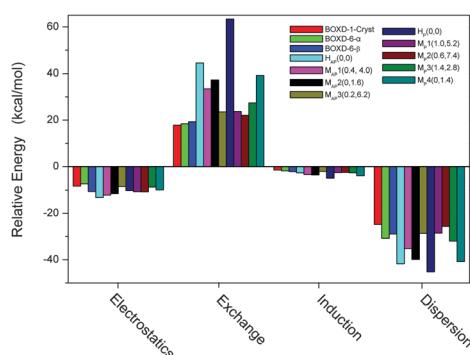


Fig. 9 Intermolecular interaction energy components of BOXD- n dimers. BOXD-1-Cryst, BOXD-6- α , and BOXD-6- β are the molecular stacking structures from the single-crystal X-ray analysis. M_{AP1} – M_{AP3} and M_P1 – M_P4 (offset π -stacking structures) are the energy-minimum molecular stacking structures of BOXD-1 found in the PES scanning. $H_{AP}(0, 0)$ and $H_P(0, 0)$ refer to the ideal face-to-face π -stacking configurations.

However, two methods show very similar trends for these structural changes, which confirm their reliability. Analysis of those crystal packings indicates that the two types of molecular packings found in different crystalline structures of BOXD-6 have similar intermolecular interaction energies (~ -22 kcal mol $^{-1}$). Although BOXD-6 shows a larger offset in crystalline packing than BOXD-1, the dispersion term for BOXD-6 is somewhat larger than that of BOXD-1; thus, the introduction of the long terminal alkoxy chains results in an increase in dispersive attraction.

4 Conclusions

BOXD-6 forms both H- and J-aggregates in THF at moderate concentration ($\sim 10^{-4}$ M), and exclusively J-aggregates at higher concentration ($\sim 10^{-3}$ M). The molecular self-assembled structures were determined by powder and single-crystal X-ray diffraction analysis. In H-aggregation (α polymorph), BOXD-6 adopts a linear conformation and forms a one-dimensional layered structure; in J-aggregation (β polymorph), the molecules adopt a Z-shaped conformation and form a more ordered two-dimensional layered structure. BOXD-6 exhibits a π -stacking structure in either H or J-aggregation. Compared to that of H-aggregation, the displacement along the molecular long-axis direction in J-aggregation is larger. Moreover, an effective method was developed to control the molecular self-assembled structures

deposited on substrates. Films with pure J-aggregates of BOXD-6 were successfully obtained by adding water as a precipitator into THF solution. The calculated intermolecular interaction potential energy surfaces (the M062x/6-31G** method) are fully consistent with the molecular stacking structures. London dispersion forces are found to play an important role in stabilizing the molecular aggregation. Several additional stable structures are also predicted by these PESs, which will be the target of future experiments.

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