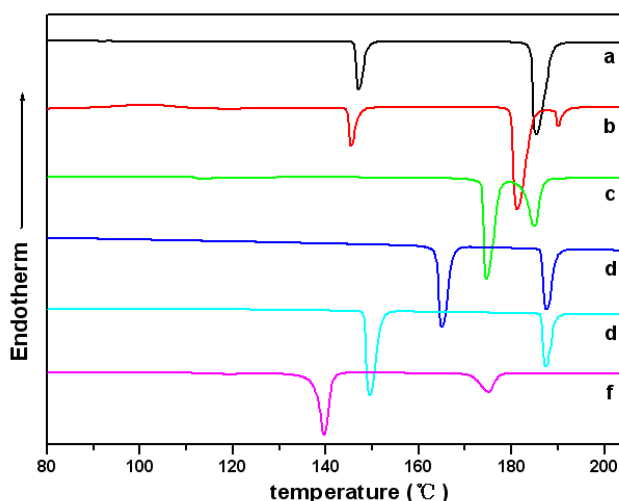
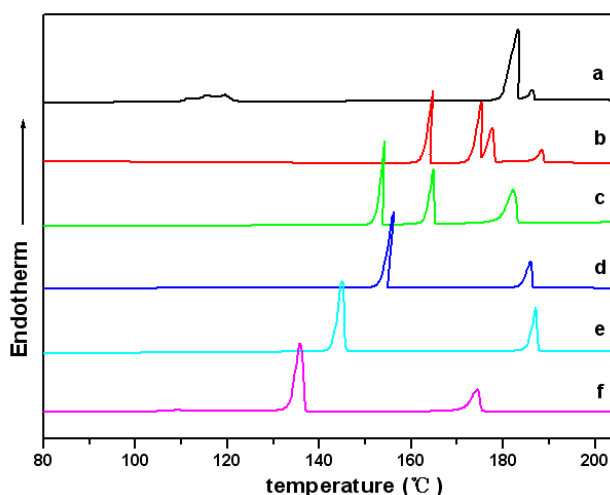


Supporting Information for Self-assemblies of Highly luminescent Bi-1,3,4-oxadiazole Derivatives through Electron Donor-Acceptor Interactions in Three-Dimensional Crystal, Two-Dimensional Layers and Mesophases

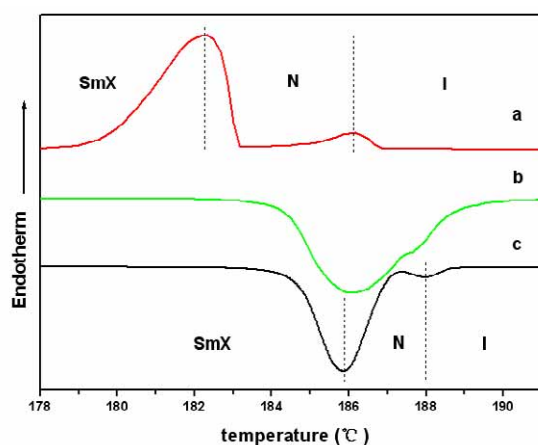
Songnan Qu,^a Xiaofang Chen,^b Xiang Shao,^b Fan Li,^a Hongyu Zhang,^c Haitao Wang,^a Peng Zhang,^a Zhixin Yu,^a Kai Wu,^b Yue Wang^c and Min Li^{*a}



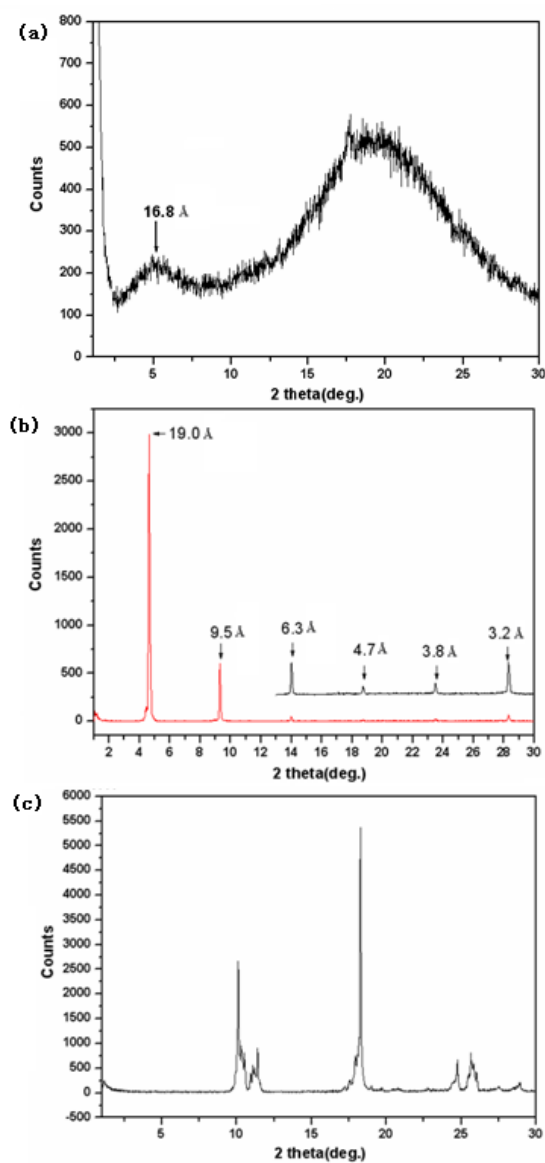
S-Figure 1. DSC curves of (a) BOXD-5, (b) BOXD-6, (c) BOXD-7, (d) BOXD-8, (e) BOXD-10, (f) BOXD-16 on the second heating run (heating rate: 10 °C/min).



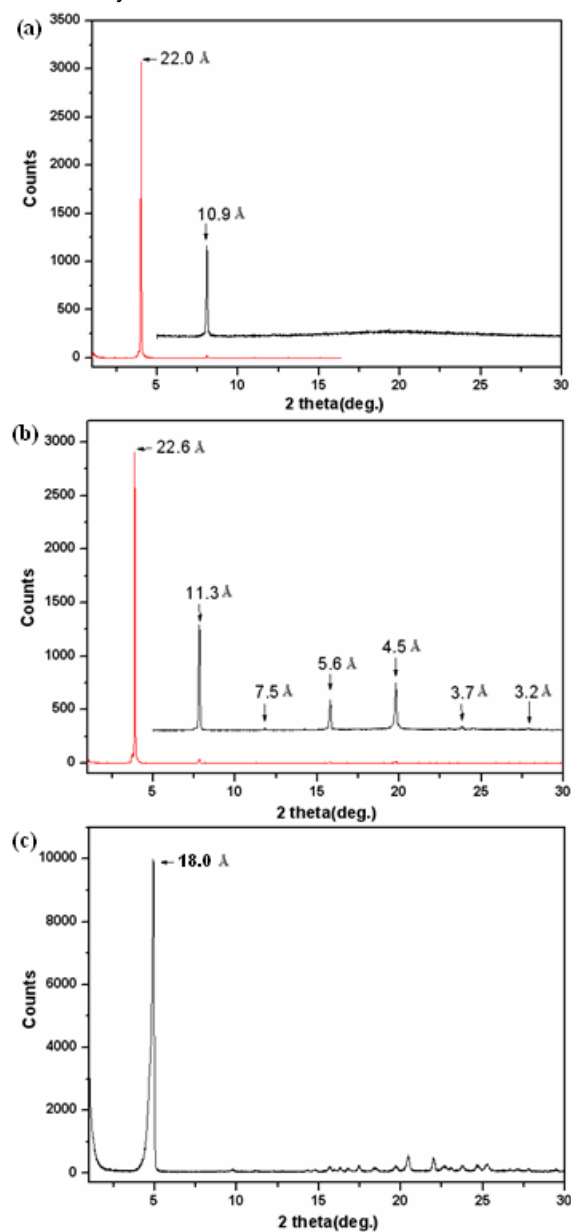
S-Figure 2. DSC curves of (a) BOXD-5, (b) BOXD-6, (c) BOXD-7, (d) BOXD-8, (e) BOXD-10, (f) BOXD-16 on the first cooling run (cooling rate: 10 °C/min).



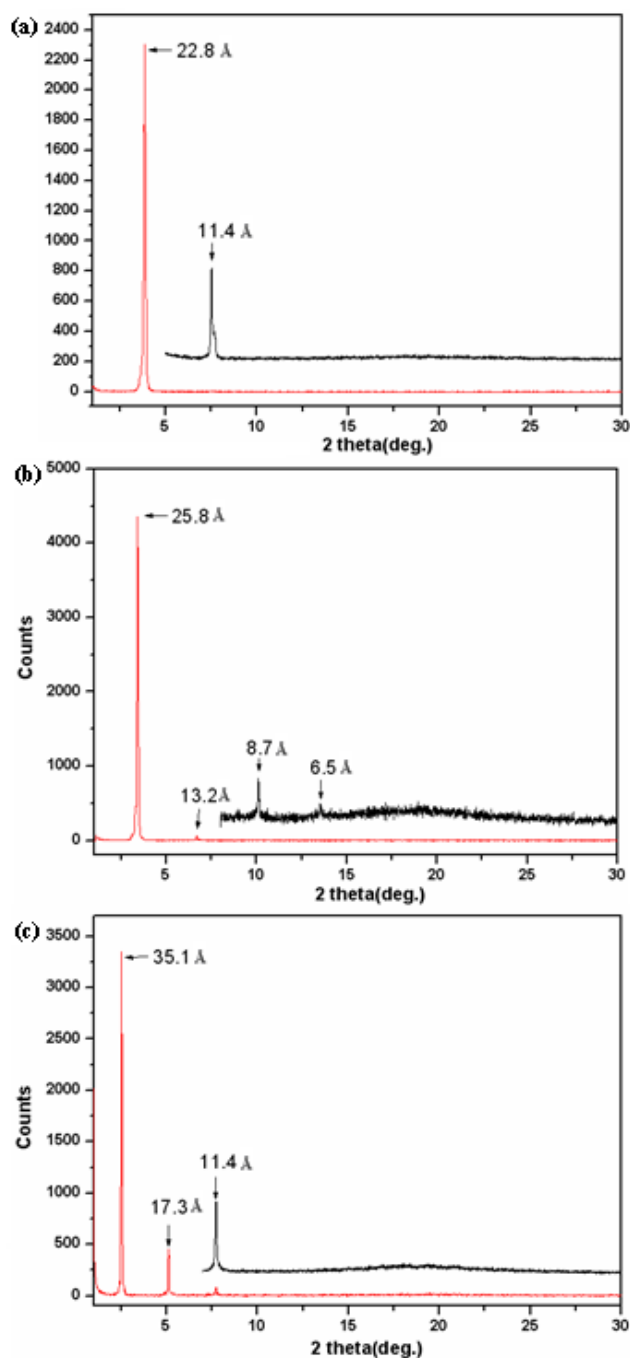
S-Figure 3. DSC curves of BOXD-5 (a) on the first cooling run (10 °C/min), (b) second heating run (10 °C/min) and on the second heating run (2 °C/min).



S-Figure 4. Powder X-ray diffraction patterns of BOXD-5 (a) at 185 °C; (b) 175 °C, (c) 20 °C on the cooling run.



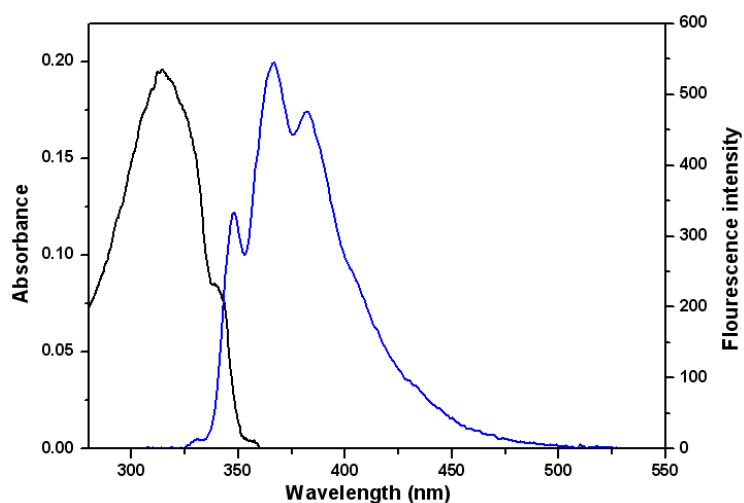
S-Figure 5. Powder X-ray diffraction patterns of BOXD-7 (a) at 175 °C; (b) 160 °C, (c) 20 °C on the cooling run.



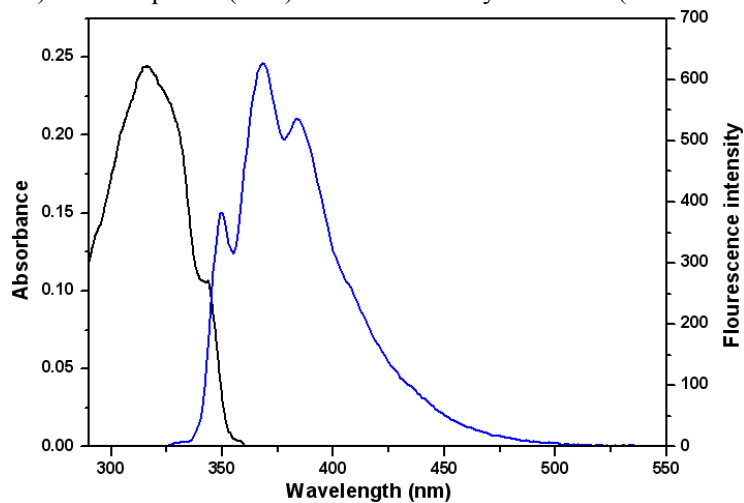
S-Figure 6. Powder X-ray diffraction patterns of (a) BOXD-8 at 180 °C; (b) BOXD-10 at 160 °C, (c) BOXD-16 at 165 °C on the cooling run.

S-Table 1. Photophysical Properties of (BOXD-*n*, *n*=1, 3, 6, 16) in cyclohexane (1×10^{-5} mol/L).

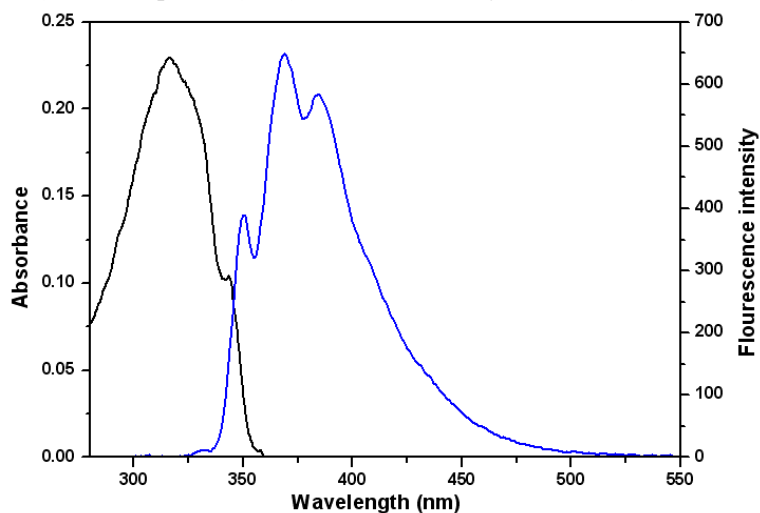
Compound	λ_{max} abs/nm	λ_{max} EM/nm	Φ_{F}
BOXD-1	315	348, 366, 382	0.94
BOXD-3	316	350, 368, 384	0.93
BOXD-6	316	350, 368, 384	0.92
BOXD-16	316	350, 368, 384	0.92



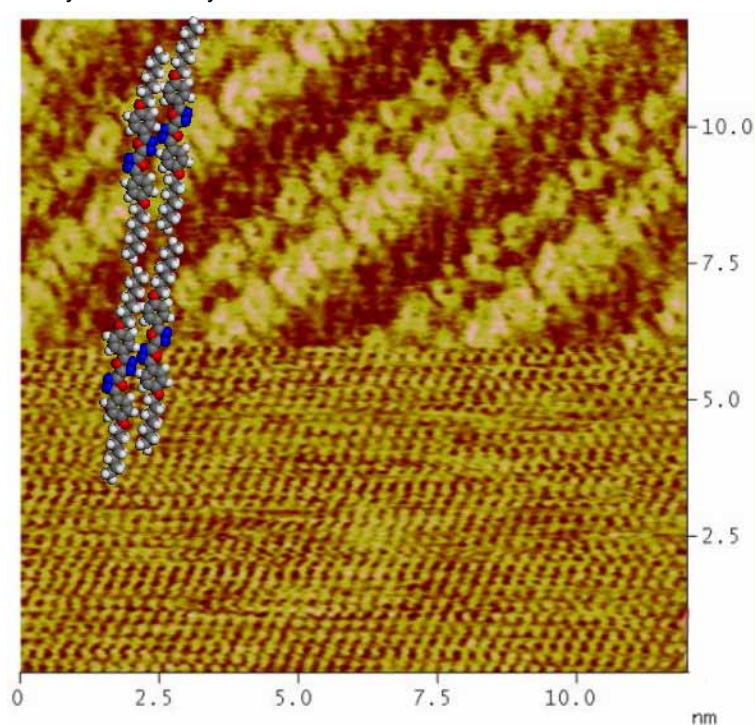
S-Figure 7. UV-vis (black) and PL spectra (blue) of BOXD-1 in cyclohexane (1×10^{-5} mol/L) at room temperature.



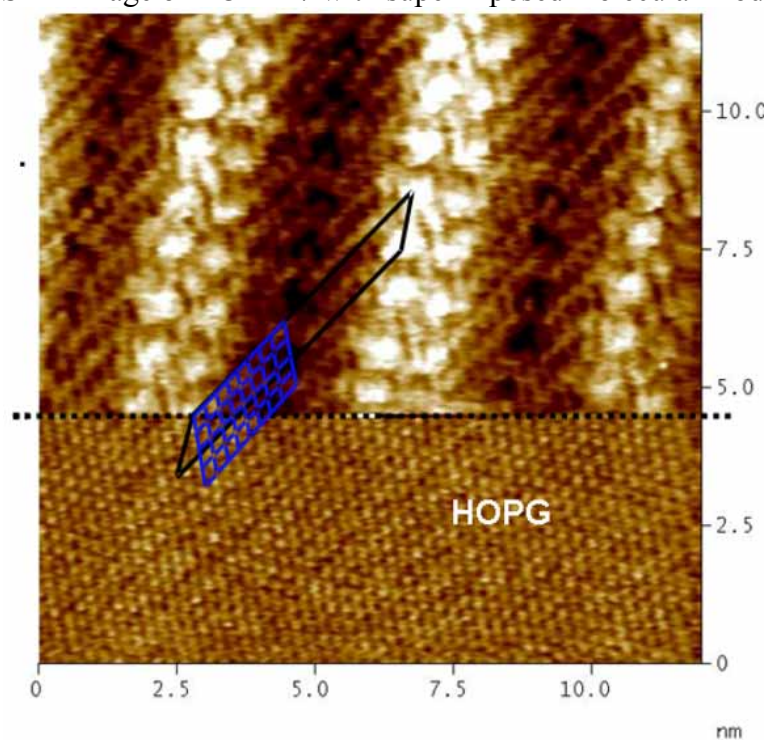
S-Figure 8. UV-vis (black) and PL spectra (blue) of BOXD-3 in cyclohexane (1×10^{-5} mol/L) at room temperature.



S-Figure 9. UV-vis (black) and PL spectra (blue) of BOXD-16 in cyclohexane (1×10^{-5} mol/L) at room temperature.



S-Figure 10. A hybrid STM image of BOXD-7 with superimposed molecular models



S-Figure 11. A hybrid STM image of BOXD-16 showing the mismatch between the adlayer and substrate symmetry

Computational Details

Molecular geometries (BOXD-n) were optimized with the Dmol³ suite of MS Modeling 3.0. The generalized gradient approximation (GGA) method was employed which associated with the PW91

functional for correlation and the 'DNP' basis set. To obtain a high accuracy result, both 'Integration accuracy' and 'SCF tolerance' options were set to "fine".

The face-to-face interaction was estimated by calculating the energy difference between two nearest neighbours of BOXD-1 molecules in the stack which axes is parallel to **a** direction (Figure 2 (c)) and two isolated BOXD-1 molecules. And the estimate of edge-to-edge interaction was obtained by calculating the energy difference between two adjacent BOXD-1 molecules in the (**a** + **b**) direction (Figure 2 (d)) and two isolated BOXD-1 molecules. The face-to-face interaction between the two molecules in this case was about 15.2 kJ mol⁻¹ and the edge-to-edge interaction between the two molecules was about 14.7 kJ mol⁻¹. Both values were obtained by the accurate DFT method mentioned above.

S-Table 2. Crystal data and structure refinement for BOXD-1.

Identification code	BOXD-1
Empirical formula	C ₁₈ H ₁₄ N ₄ O ₄
Formula weight	350.33
Temperature	293(2) K
Wavelength	0.71073 Å
Crystal system, space group	Monoclinic, P2 ₁ /n
Unit cell dimensions	a = 6.419(1) Å α = 90°. b = 8.268(2) Å β = 92.69(3)°. c = 15.213(3) Å γ = 90°.
Volume	806.5(3) Å ³
Z, Calculated density	2, 1.443 Mg/m ³
Absorption coefficient	0.105 mm ⁻¹
F(000)	364
Crystal size	0.36 x 0.16 x 0.13 mm ³
Theta range for data collection	3.39 to 27.47 deg.
Limiting indices	-8 ≤ h ≤ 8, -10 ≤ k ≤ 10, -19 ≤ l ≤ 19
Reflections collected / unique	7435 / 1814 [R(int) = 0.0285]
Completeness to theta = 27.47	97.9 %
Absorption correction	Semi-empirical from equivalents
Max. and min. transmission	0.9869 and 0.9634
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	1814 / 0 / 146
Goodness-of-fit on F ²	1.087
Final R indices [I > 2σ(I)]	R1 = 0.0368, wR2 = 0.0916
R indices (all data)	R1 = 0.0580, wR2 = 0.0983
Largest diff. peak and hole	0.204 and -0.123 e Å ⁻³