**Electronic Supplementary Information** 

## The self-assembly and pyridine regulation of a hydrogen-bonded dimeric

## building block formed by low-symmetric aromatic carboxylic acid

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## Table of Contents:

1	DFT simulated single molecular models	S3
2	Supplementary examples of pyridine molecules co-assembled with H <sub>5</sub> BHB	S5
3	Supplementary large scale (120 nm × 120 nm) STM images	S6

1 DFT simulated single molecular models.



**Fig. S1** DFT simulated molecular models in the gas phase. (a-d) Top view of  $H_5BHB$ , BPD, BPDYB, TPDYB respectively. (e-h) Side view of  $H_5BHB$ , BPD, BPDYB, TPDYB, respectively.



**Fig. S2** DFT simulated molecular models on the graphite surface. (a-d) Top view of  $H_5BHB$ , BPD,  $H_5BHB$  dimer,  $H_5BHB$ –BPD unit cell, respectively. (e-h) Side view of  $H_5BHB$ , BPD,  $H_5BHB$  dimer,  $H_5BHB$ –BPD unit cell, respectively.

2 Supplementary examples of pyridine molecules co-assembled with H<sub>5</sub>BHB.



**Fig. S3** Supplementary failure and success examples of pyridine molecules that can form stable ordered co-assembled structure with  $H_5BHB$  molecule. Failure examples: (a) 1,2-Di(4-pyridyl)ethylene (DPE). (b) 3,3',5,5'-Tetrakis(pyridin-4-ylethynyl)-1,1'-biphenyl (TPYB). Success example: (c) 1,4-Di(pyridin-4-yl)benzene (DPYB).



**Fig. S4** STM images of supplementary experiment:  $H_5BHB$  co-assembled with DPYB at heptanoic acid–HOPG liquid–solid interface. (a) Large scale (120 nm × 120 nm) STM image,  $I_{set} = 311.3$  pA,  $V_{bias} = 686.6$  mV. (b) Large scale (60 nm × 60 nm) STM image,  $I_{set} = 152.6$  pA,  $V_{bias} = 1024$  mV. (c) High-resolution STM image (30 nm × 30 nm),  $I_{set} = 158.7$  pA,  $V_{bias} = 904.8$  mV. Two different co-assembled structures were observed, labeled with stru1 and stru2, respectively. The blue triangle shape represents for a H<sub>5</sub>BHB molecule, while the white rod shape represents for a DPYB molecule. The co-assembly behavior of H<sub>5</sub>BHB and DPYB was similar to that of H<sub>5</sub>BHB and BPDYB. 3 Supplementary large scale (120 nm × 120 nm) STM images.



**Fig. S5** Supplementary large scale (120 nm × 120 nm) STM images at the heptanoic acid–HOPG liquid–solid interface of (a)  $H_5BHB_self$ -assembled,  $I_{set} = 231.9$  pA,  $V_{bias} = 815.4$  mV. (b)  $H_5BHB-BPD$  co-assembled structure,  $I_{set} = 146.5$  pA,  $V_{bias} = 932.0$  mV. (c)  $H_5BHB-BPDYB$  co-assembled structure,  $I_{set} = 177.9$  pA,  $V_{bias} = 932.3$  mV. (d)  $H_5BHB-TPDYB$  co-assembled structure,  $I_{set} = 164.8$  pA,  $V_{bias} = 941.8$  mV.