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## **Supporting Information**

## Coronene Guest Molecule Selectivity in the Host Templates Formed by Hydrogen Bond and Van der Waals Force at Liquid/Solid Interface

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#### S1. Acronyms table

Table S1 The definition of Acronyms in the main text.		
Acronyms	Full name	
H <sub>6</sub> PDB	1,3,5-tris(4-(3,5-dicarboxyphenyl) diphenyl) benzene	
H <sub>6</sub> PAB	1,3,5-tris (4-(3,5-dicarboxyphenylacetylene) phenyl) benzene	
COR	1,3,5-tris (4-(3,5-dicarboxyphenylacetylene) phenyl) benzene	
HPB	Hexaphenylbenzene derivatives	
H <sub>4</sub> ETTC	4',4''',4'''',4''''''''''(ethene-1,1,2,2-tetrayl) tetrakis-([1,1-biphenyl]-3-carboxylic acid)	
H <sub>8</sub> ETTB	4',4"",4""",4"""'-(ethene-1,1,2,2-tetrayl) tetrakis([1,1'-biphenyl]-3,5-dicarboxylic acid)	

### **S2.** Sample Preparation

Table S2 The information of all the samples used in the experiment.			
Sample	Obtained from	Purity	
H <sub>6</sub> PDB	Jilin Chinese Academy of Sciences-Yanshen Technology Co., Ltd.	98%	
H <sub>6</sub> PAB	Jilin Chinese Academy of Sciences-Yanshen Technology Co., Ltd.	98%	
HPB	synthesized by the previous reference	98%	
COR	Beijing HWRK Chemical Corporation	98%	
1-heptanoic acid (HA)	J&K Scientific	98%	

All the samples were at 98% purity and used directly without further purification. All the samples were dissolved in HA solvent. The concentrations of H<sub>6</sub>PDB, H<sub>6</sub>PAB and HPB solution were 10% of their saturated concentrations. COR/HA solution was prepared in C1 =  $10^{-5}$  mol·L<sup>-1</sup>, C2 = 8 × C1, and saturated concentration (C3) respectively. H<sub>6</sub>PAB and HPB solution was premixed in 1:1 ratio. The volume of COR added in-suit was the same as the premixed solution.

S3. Large scale STM image of H<sub>6</sub>PDB and H<sub>6</sub>PAB co-assembled with saturated COR/HA solution(C3)



Figure S1. (a) Large scale STM image of  $H_6PDB/COR(C3)$  co-assembled structure ( $I_{set} = 259.4$  pA,  $V_{bias} = 809.9$  mV, 100 nm × 100 nm); (b) Large scale STM image of  $H_6PAB/COR(C3)$  co-assembled structure ( $I_{set} = 268.6$  pA,  $V_{bias} = 806.3$  mV, 100 nm × 100 nm).

# S4. High resolution STM image and molecular model of HPB self-assembled structure and HPB/COR co-assembled structure



Figure S2. (a) High resolution STM image of HPB self-assembled structure ( $I_{set} = 271 \text{ pA}$ ,  $V_{bias} = 799 \text{ mV}$ , 25 nm × 25 nm); (b) High resolution STM image of HPB/COR(C3) co-assembled structure ( $I_{set} = 198.4 \text{ pA}$ ,  $V_{bias} = 739.7 \text{ mV}$ , 25 nm × 25 nm); (c) (d) Proposed models of the structures in (a) and (b) respectively.