

Supporting information

Descriptor-based Rational Design of Two-Dimensional Self-Assembled Nano-architectures Stabilized by Hydrogen Bonds

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Synthesis of phenyleneethynylene derivatives

The synthesis of molecule **PE4A** was carried out by following a previously reported procedure.¹ Synthesis and characterization of carboxylic acid derivatives of phenyleneethynylene, **PE4B** and **PE3A** were carried out by following the reactions shown in Figure S1. The key step for the synthesis of phenyleneethynylene derivatives **PE3A** and **PE4B** is Sonogashira cross coupling reaction, using Pd(II) as catalyst.²⁻³ The intermediate compound **3** was prepared by the dimerization of compound **2** by following Glaser reaction using CuCl as a catalyst, which on further reaction with 1 equivalent of phenylacetylene, using Pd(II) as the catalyst, gave the intermediate **4**. Compounds **1** and **4** on reaction with dimethyl 5-ethynylisophthalate (note: 2 equivalents for **1** and 1 equivalent for **4**) yielded products **6** and **5**, respectively. The precursors of carboxylic acid derivatives (**5** and **6**) were purified using recycling HPLC to maintain a high purity of the final products. The purified products **5** and **6** were hydrolyzed using KOH to yield the corresponding acid derivatives **PE4B** and **PE3A**, respectively. All the compounds were characterized using various analytical and spectroscopic techniques. ¹H NMR and ¹³C NMR spectra were recorded on Bruker 500 MHz spectrometers. Elemental analysis was carried out using Elementar Vario Micro Cube. All melting points are uncorrected and were determined with a Stuart SMP30 melting point apparatus.

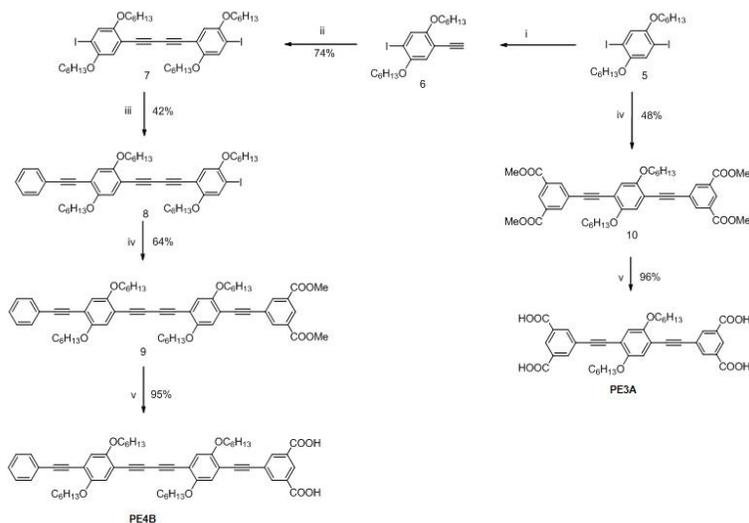


Figure S1. i) Pd(PPh₃)₂Cl₂, CuI, PPh₃, diisopropylamine, TMSA; KOH, MeOH. ii) CuCl, diazabicyclo[5.4.0]undec-7-ene; N,N-diisopropylethylamine; iii) Pd(PPh₃)₂Cl₂, CuI, PPh₃, diisopropylamine, phenylacetylene; iv) Pd(PPh₃)₂Cl₂, CuI, PPh₃, diisopropylamine, dimethyl 5-ethynylisophthalate; v) KOH, MeOH.

Synthesis of 4. Compound **3** (1.5 g, 1.75 mmol) was kept in a two necked flask and deoxygenated by purging with argon for 15 min, to which Pd(PPh₃)₂Cl₂ (24.5 mg, 0.035mmol), CuI (6.6 mg, 0.035 mmol), PPh₃ (18 mg, 0.07 mmol), phenyl acetylene (0.178 g, 1.75 mmol), diisopropylamine (10 mL), and THF (30 mL) were added. The mixture was stirred for 12 h at room temperature. After removal of the solvent under reduced pressure, the crude product was purified by column chromatography over silica gel (100-200 mesh) using a mixture (1:10) of CHCl₃ and hexane as eluent to yield 0.6 g (42 %) of **4** as a yellow solid. mp 123-124 °C; ¹H NMR (500 MHz, CDCl₃, TMS): δ 7.56-7.54 (m, 2H), 7.37-7.35 (d, 3H), 7.31 (s, 1H), 7.00 (d, 2H), 6.89 (s, 1H), 4.01-3.94 (m, 8H), 1.86-1.81 (m, 8H), 1.59-1.52 (m, 7H), 1.38-1.37 (m, 16H), 0.94-0.91 (m, 12H). ¹³C NMR (500 MHz, CDCl₃, TMS): δ 155.59, 155.01, 153.49, 151.79, 131.61, 128..33, 123.99, 123.34, 117.73, 116.95, 116.62, 115.21, 112.48, 112.31, 95.53, 87.09, 85.81, 70. 11, 70.09, 69.80, 69.65, 31.60, 31.53, 31.51, 31.49, 30.92, 29.70, 29.10, 29.08, 25.73, 25.61, 25.57, 22.59, 14.03. Anal. Calcd for C₄₈H₆₁IO₄ : C, 69.55; H, 7.42; I, 15.31; O, 7.72 Found: C, 69.62; H, 7.34.

Synthesis of 5. Compound **4** (525 mg, 0.64 mmol) was kept in a two necked flask and deoxygenated by purging with argon for 15 min, to which Pd(PPh₃)₂Cl₂ (21 mg, 0.03 mmol), CuI (6 mg, 0.03 mmol), PPh₃ (15 mg, 0.06 mmol), dimethyl 5-ethynylisophthalate (140 mg, 0.64 mmol), diisopropylamine (10 mL), and THF (30 mL) were added. The mixture was stirred for 12 h at room temperature. After removal of the solvent under reduced pressure, the crude product was purified by column chromatography over silica gel (100-200 mesh) using a mixture (1:10) of CHCl₃ and hexane as eluent to yield 375 mg (64 %) of **5** as a yellow solid. mp 117-118 °C; ¹H NMR (500 MHz, CDCl₃, TMS): δ 8.55 (s, 1H), 8.28 (s, 2H), 7.47-4.45 (m, 5H), 6.93-6.91 (d, 4H), 3.96-3.89 (m, 14H), 1.78-1.75 (t, 8H), 1.45-1.43 (m, 8H), 1.30-1.27 (m, 8H), 0.85-0.81 (m, 12H). ¹³C NMR (500 MHz, CDCl₃, TMS): δ 206.97, 165.62, 155.04, 153.04, 136.45, 134.16, 131.62, 130.99, 128.39, 128.33, 127.61, 124.48, 123.34, 117.74, 117.04, 116.95, 113.34, 87.81, 85.81, 69.85, 69.82, 69.66, 52.53, 31.60, 31.53, 30.92, 29.28, 29,18, 29.10, 25.73, 25.61, 22.63, 22.61, 14.04, 14.02. Anal. Calcd for C₆₀H₇₀O₈ : C, 78.60; H, 7.68; O, 13.92 Found: C, 78.32; H, 7.39.

Synthesis of 6. Compound **1** (700 mg, 1.30 mmol) was kept in a two necked flask and deoxygenated by purging with argon for 15 min, to which Pd(PPh₃)₂Cl₂ (91 mg, 0.13 mmol), CuI (12 mg, 0.13 mmol), PPh₃ (67 mg, 0.26 mmol), dimethyl 5-ethynylisophthalate (566 mg, 1.60 mmol), diisopropylamine (10 mL), and THF (30 mL) were added. The mixture was stirred for 12 h

at room temperature. After removal of the solvent under reduced pressure, the crude product was purified by column chromatography over silica gel (100-200 mesh) using a mixture (3:2) of CHCl_3 and hexane as eluent to yield 450 mg (48 %) of **6** as a yellow solid. mp 162-163 °C; ^1H NMR (500 MHz, CDCl_3 , TMS): δ 8.64 (s, 2H), 8.38 (s, 4H), 7.06 (s, 2H), 4.09-4.06 (t, 4H), 3.98 (s, 6H), 1.91-1.88 (t, 4H), 1.60-1.56 (m, 4H), 1.43-1.34 (m, 8H), 0.92-0.89 (m, 6H). ^{13}C NMR (500 MHz, CDCl_3 , TMS): δ 165.62, 153.79, 134.36, 136.44, 130.98, 130.08, 124.52, 117.01, 113.78, 92.92, 87.83, 69.71, 52.52, 31.54, 29.23, 25.76, 22.59, 14.00. Anal. Calcd for $\text{C}_{42}\text{H}_{46}\text{O}_{10}$: C, 70.97; H, 6.52; O, 22.51 Found: C, 71.12; H, 6.39.

Synthesis of PE4B and PE3A. The synthesis of compounds **PE4B** and **PE3A** was achieved through hydrolysis of corresponding ester derivatives (**5** and **6**) by adopting a general procedure given below. The ester derivative (100 mg) was dissolved in THF (10 mL) and to this solution KOH (1.0 g) and methanol (20 mL) were added and refluxed for about 5 h. The reaction mixture was cooled to room temperature, neutralized with dilute HCl and water (50 mL) was added. The precipitated product was either filtered or extracted with chloroform. The solvent was evaporated under reduced pressure to yield the corresponding carboxylic acid as solid product. The spectral details of the individual compounds are provided below.

PE4B. 92 mg (95%). mp 216 °C; ^1H NMR (500 MHz, THF-d_8 , TMS): δ 8.50 (s, 1H), 8.20 (s, 2H), 7.39-7.37 (d, 2H), 7.25-7.21 (m, 3H), 7.04-6.96 (m, 4H), 3.95-3.91 (m, 8H), 1.73-1.69 (m, 8H), 1.50-1.42 (m, 8H), 1.29-1.24 (m, 16H), 0.84-0.78 (m, 12H). ^{13}C NMR (500 MHz, THF-d_8 , TMS): δ 165.33, 155.06, 155.03, 153.82, 153.64, 135.75, 131.94, 131.25, 128.22, 128.17, 124.15, 123.60, 117.27, 116.56, 116.47, 114.35, 112.86, 112.25, 85.87, 78.79, 69.25, 69.20, 66.13, 31.58, 31.54, 29.25, 29.10, 29.08, 25.85, 25.73, 22.55, 13.45, 13.41. Anal. Calcd for $\text{C}_{58}\text{H}_{66}\text{O}_8$: C, 78.17; H, 7.47; O, 14.34 Found: C, 78.53; H, 6.65.

PE3A 88 mg (96%). mp above 300 °C; ^1H NMR (500 MHz, THF-d_8 , TMS): δ 8.52 (s, 2H), 8.23 (s, 4H), 7.07 (s, 2H), 4.00-3.98 (t, 4H), 1.78-1.75 (t, 4H), 1.53-1.50 (m, 4H), 1.435-1.30 (m, 8H), 0.92-0.89 (m, 6H). ^{13}C NMR (500 MHz, THF-d_8 , TMS): δ 165.32, 153.95, 135.75, 131.88, 130.08, 124.28, 116.59, 113.75, 92.70, 87.56, 69.20, 31.59, 29.30, 25.86, 22.57, 14.42. Anal. Calcd for $\text{C}_{38}\text{H}_{38}\text{O}_{10}$: C, 69.71; H, 5.85; O, 24.44 Found: C, 70.01; H, 5.39.

STM imaging: The STM imaging was carried out in the constant-height mode under ambient conditions at the liquid-solid interface immediately after drop casting the solution on HOPG surface using a multimode scanning tunneling microscope.

Computational Methods

Density functional theory calculations

Density functional theory (DFT) calculations were performed using the SIESTA package.⁴⁻⁵ We used a double zeta polarized basis set with confinement energy of 0.01 Ry. The mesh cutoff size of the real-space grid was taken to be 250 Ry. Troullier-Martin norm-conserving pseudopotentials were used.⁶ Exchange-correlation interactions were treated within the PBE form of the generalized gradient approximation.⁷ Van der Waals interactions were incorporated by using the DFT-D2 treatment.⁸ A vacuum with a spacing of 16 Å was introduced along non-repeating directions. For the large unit cells used in this study, the Brillouin zone was sampled at the zone center Γ only. Geometry optimization was performed using the conjugate gradient algorithm. In the calculations, highly oriented pyrolytic graphite (HOPG) was replaced by a single atomic layer of graphene. All atomic coordinates were relaxed until the forces on all atoms were smaller than 0.03 eV/Å. Basis set superposition errors were eliminated by applying the counterpoise correction procedure.⁹⁻¹⁰ The simulated STM images were obtained using the Tersoff-Hamann approach in the constant height mode.¹¹

In all cases, the unit cells used corresponded to the primitive unit cell of the corresponding HEX or LIN pattern. These primitive unit cells contain three host molecules in the HEX arrangements of PE4A and PE3A, six host molecules in the HEX arrangement of PE4B, and two host molecules in the LIN arrangements of all three host molecules (see Fig. S2, where the black rhombi and rectangles mark the boundaries of the unit cells for the HEX and LIN, respectively). In all cases, the unit cell contains only one guest molecule (when present).

The initial coordinates of the molecules in the gas phase were obtained using the Gaussview software package. The unit cell parameters were varied stepwise, with the constraint that they be commensurate with the underlying lattice constant of graphene. The coordinates of all molecules were allowed to relax, following a three-step procedure: (i) the coordinates of the isolated molecules

were then relaxed using DFT. Next, for each set of cell parameters: (ii) a monolayer of molecules was assembled in either the HEX or LIN pattern in the gas phase (i.e., no graphene substrate) and the coordinates were further relaxed, with the constraint that they remain in the x-y plane, (iii) this monolayer was then placed on graphene and the coordinates was further relaxed, with no geometric constraints. The optimized cell parameters thus obtained, along with the corresponding values of the adsorption energy E_{ads} are listed in Table S1 below.

Table S1. Results from DFT for geometries and energetics of the HEX and LIN arrangements of the host molecules when adsorbed on graphene. N_{host} is the number of host molecules contained in the 2D primitive unit cell, whose optimal size is given by the cell parameter a (in HEX) or a, b (in LIN). We see that the optimal lattice parameters as obtained from DFT are very close to those determined from the STM experiments. For ease of reference, the corresponding geometries are given in Fig. S2 below. E_{ads} is the adsorption energy of the host assemblies on graphene.

Host molecule →		PE4A	PE4B	PE3A
HEX	N_{host} (molecules/cell)	3	6	3
	a [DFT](nm)	4.23	5.23	3.24
	a [expt](nm)	4.2 ± 0.1	$5.2 \pm 0.1^*$	$3.2 \pm 0.1^*$
	E_{ads} (meV/Å ²)	-13.1	-14.2	-16.8
LIN	N_{host} (molecules/cell)	2	2	2
	a [DFT](nm)	1.49	1.49	1.49
	a [expt](nm)	NA	1.5 ± 0.1	1.5 ± 0.1
	b [DFT] (nm)	6.38	5.17	3.88
	b [expt](nm)	NA	5.2 ± 0.1	3.9 ± 0.1
	E_{ads} (meV/Å ²)	-12.0	-15.2	-17.1

* In presence of coronene

Determination of form of Host Descriptor η

The form of η was determined by demanding that it involve a combination of powers of the three physical parameters L , N_{COOH} and N_{alkoxy} ; this combination should correlate linearly with the difference in ΔG between the HEX and LIN patterns.¹² Here, L is the length of the central backbone of the host molecule, measured between the centers of the terminal phenyl rings, N_{COOH} is the number of terminal carboxyl groups, and N_{alkoxy} is the number of alkoxy side chains. The power of N_{COOH} was set = 1 (based on the physical argument that the main stabilization of the assembly comes from dimeric hydrogen bonds between COOH groups). Thus the trial form chosen was $\eta = N_{\text{COOH}} \times (L)^\alpha \times [1 + (N_{\text{alkoxy}})^\beta]$; this form ensures that η does not become zero or infinite in the absence of alkoxy side chains. The powers α and β were required to be integers or simple fractions and then determined so as to minimize the error obtained by a linear fit; this led to $\alpha = 1$ and $\beta = 1/8 = 0.125$.

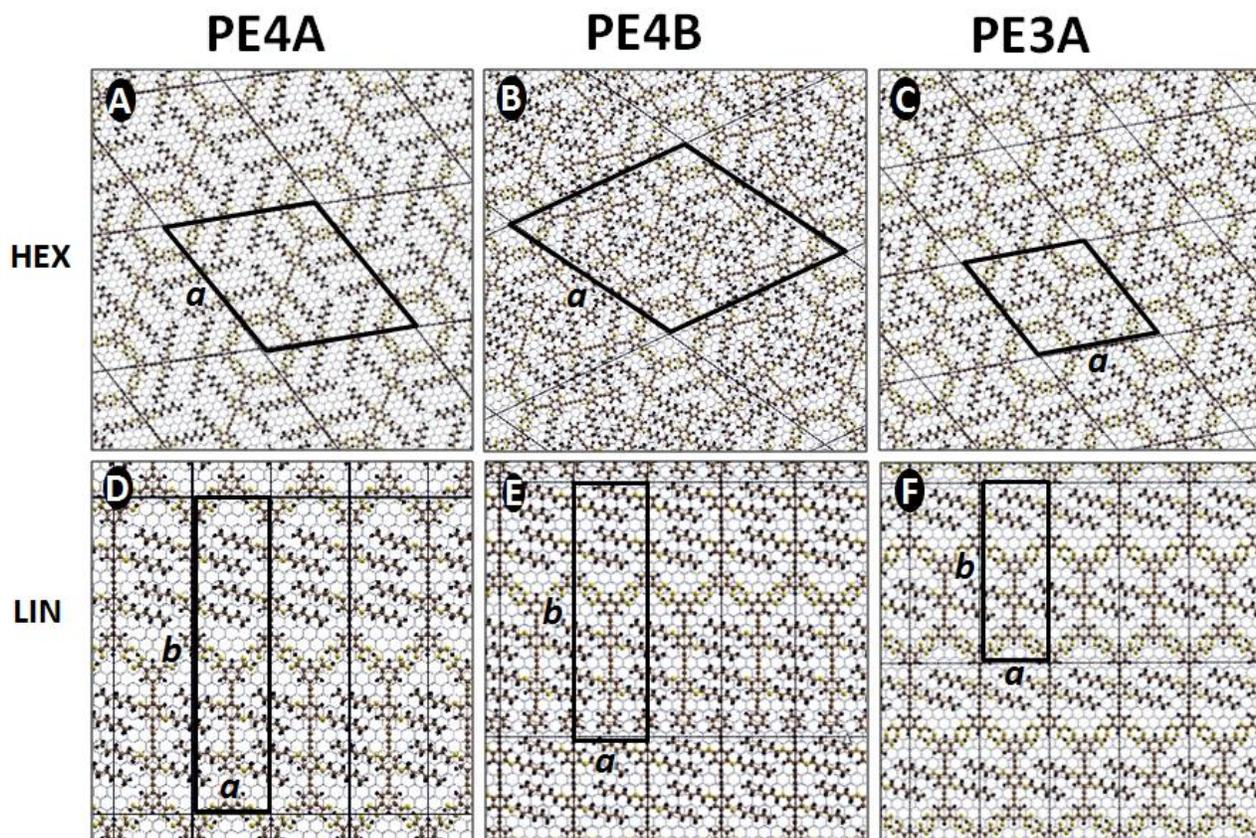


Figure S2: Atomistic structures of host assemblies on graphene. All coordinates have been relaxed using DFT calculations. Black rhombi and rectangles mark the boundaries of the unit cells for the HEX and LIN patterns, respectively. Atomic colors: C of host: brown, H: black, O: yellow, C of graphene: gray, C of guest: green.

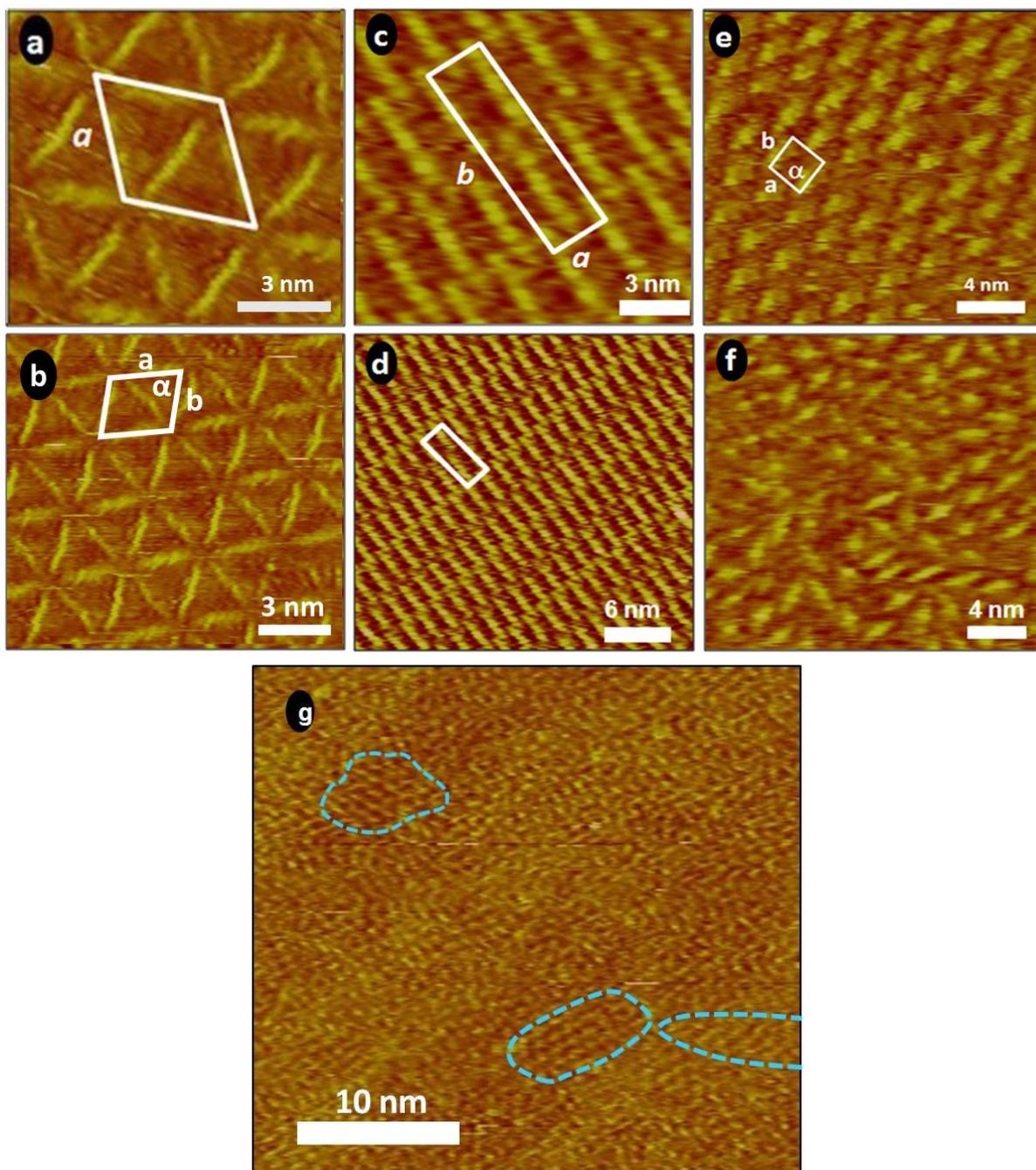


Figure S3. Experimental STM images of host assemblies. PE4A in HEX pattern: (a) zoomed in view (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1500 \text{ mV}$; $I_t = 60 \text{ pA}$) and (b) larger area STM image (scan size: $30 \times 30 \text{ nm}^2$; $V_{\text{bias}} = -850 \text{ mV}$; $I_t = 50 \text{ pA}$). PE4B in LIN pattern: (c) zoomed in view (scan size: $15 \times 15 \text{ nm}^2$; $V_{\text{bias}} = -1450 \text{ mV}$; $I_t = 100 \text{ pA}$) and (d) larger area STM image (scan size: $40 \times 40 \text{ nm}^2$; $V_{\text{bias}} = -1450 \text{ mV}$; $I_t = 100 \text{ pA}$). PE3A: (e) local LIN domains (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1925 \text{ mV}$; $I_t = 230 \text{ pA}$), (f) random 2D glass arrangement (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1750 \text{ mV}$; $I_t = 230 \text{ pA}$) of PE3A and (g) larger area, showing 2D glass but with some domains of ordered LIN (outlined by blue dashed lines).

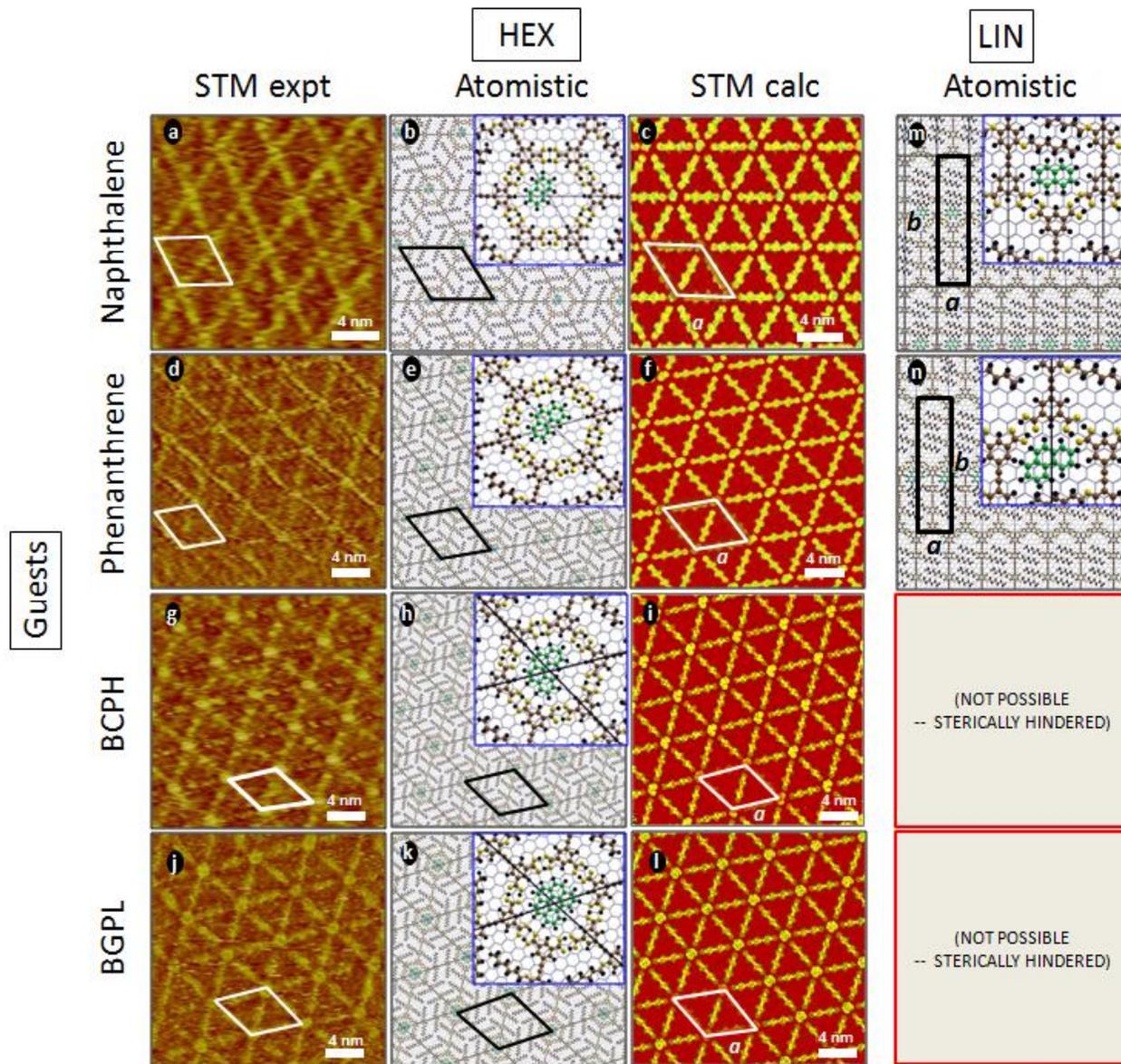


Figure S4. STM images and atomistic structures of PE4A with guest molecules on HOPG or graphene in HEX (a)-(l) and LIN (m-n) pattern. Experimental STM images of PE4A with (a) naphthalene (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -600 \text{ mV}$; $I_t = 120 \text{ pA}$), (d) phenanthrene (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1150 \text{ mV}$; $I_t = 153 \text{ pA}$), (g) BCPH (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -539 \text{ mV}$; $I_t = 60 \text{ pA}$) and (j) BGPL (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1400 \text{ mV}$; $I_t = 100 \text{ pA}$) as guest in HEX pattern. The corresponding atomistic structures of PE4A with (b) naphthalene, (e) phenanthrene, (h) BCPH and (k) BGPL as guest. The insets are zoomed in views. Simulated STM images of PE4A with (c) naphthalene ($V_{\text{bias}} = -2720 \text{ mV}$), (f) phenanthrene ($V_{\text{bias}} = -2720 \text{ mV}$), (i) BCPH ($V_{\text{bias}} = -2720 \text{ mV}$) and (l) BGPL ($V_{\text{bias}} = -2720 \text{ mV}$). Atomistic structures of PE4A with (m) naphthalene and (n) phenanthrene in LIN pattern. The black/white lines indicate the unit cell; with a and b are the unit cell parameter. Atomic colors: C of host: brown, H: black, O: yellow, C of graphene: gray, C of guest: green.

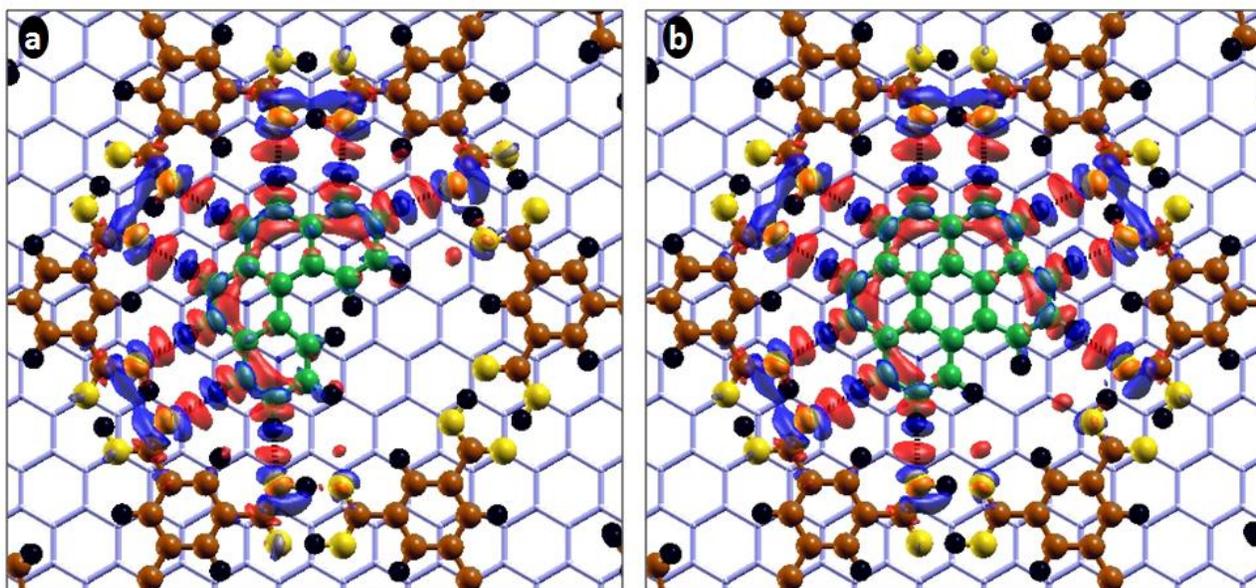


Figure S5. Theoretically computed charge redistribution plots between PE4A and different guest molecules. PE4A (a) with benzo-c-phenanthrene (BCPH) and (b) benzo-ghi-perylene (BGPL) on graphene. Red and blue lobes indicate electron accumulation and depletion respectively, plotted at an isosurface value of 0.0002 e/bohr³. Note the clear signatures of eight and ten host-guest hydrogen bonds in (a) and (b), respectively. Color scheme: brown: C of PE4A; yellow: O, black: H; green: C of guest. Gray lines indicate bonds between C atoms of graphene.

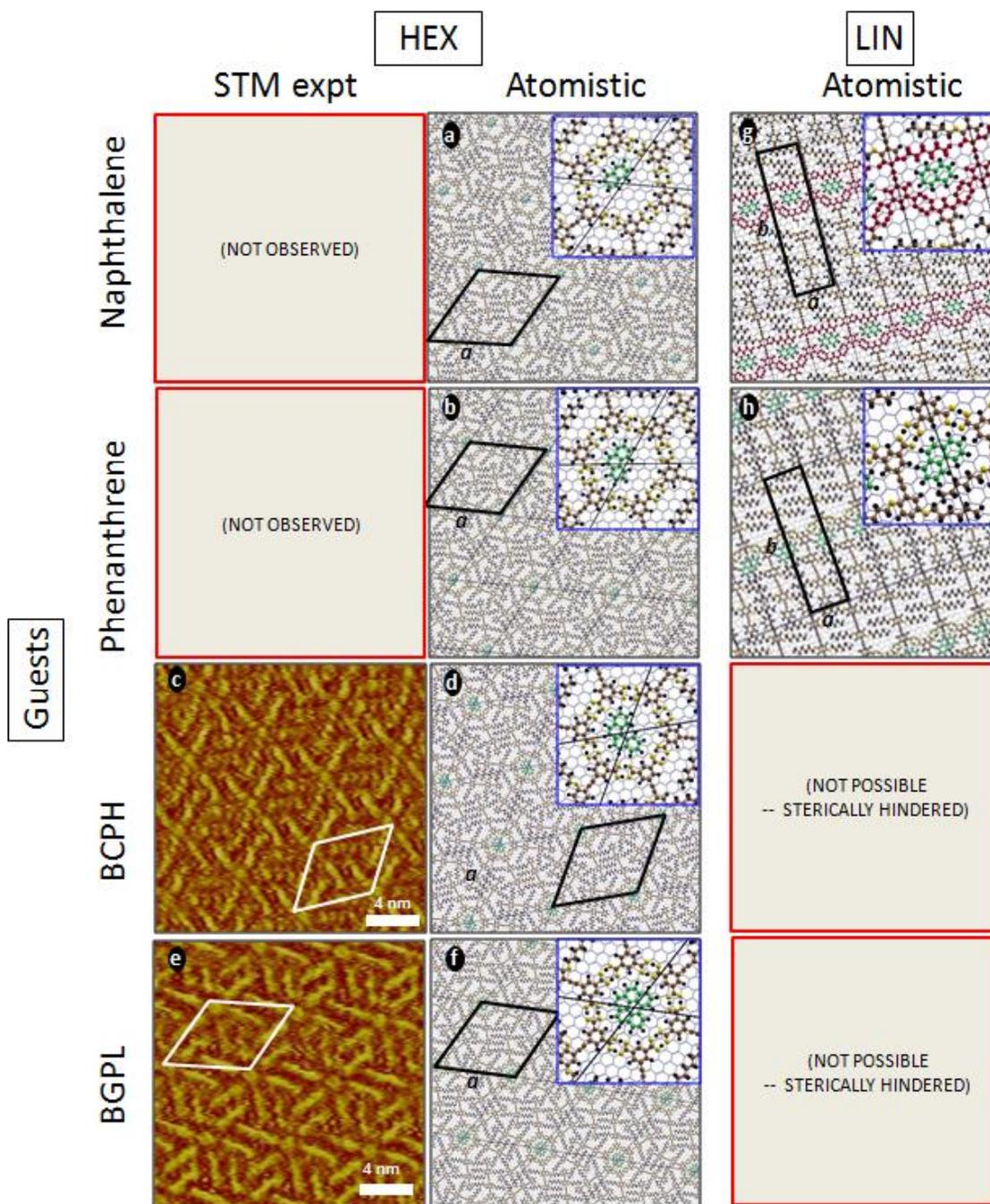


Figure S6. Atomistic structures from DFT and experimental STM images of PE4B with various guests in (a)-(f) HEX and (g)-(h) LIN patterns. Atomistic structures of PE4B with (a) naphthalene (b) phenanthrene, (d) BCPH and (f) BGPL (PE4B in HEX); (g) naphthalene and (h) phenanthrene (PE4B in LIN). Experimental STM images of PE4B with (c) BCPH (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1018 \text{ mV}$; $I_t = 120 \text{ pA}$) and (e) BGPL (scan size: $20 \times 20 \text{ nm}^2$; $V_{\text{bias}} = -1500 \text{ mV}$; $I_t = 150 \text{ pA}$). The black/white lines indicate the unit cell; with a and b are the unit cell parameter. Atomic colors: C of host: brown, H: black, O: yellow, C of graphene: gray, C of guest: green.

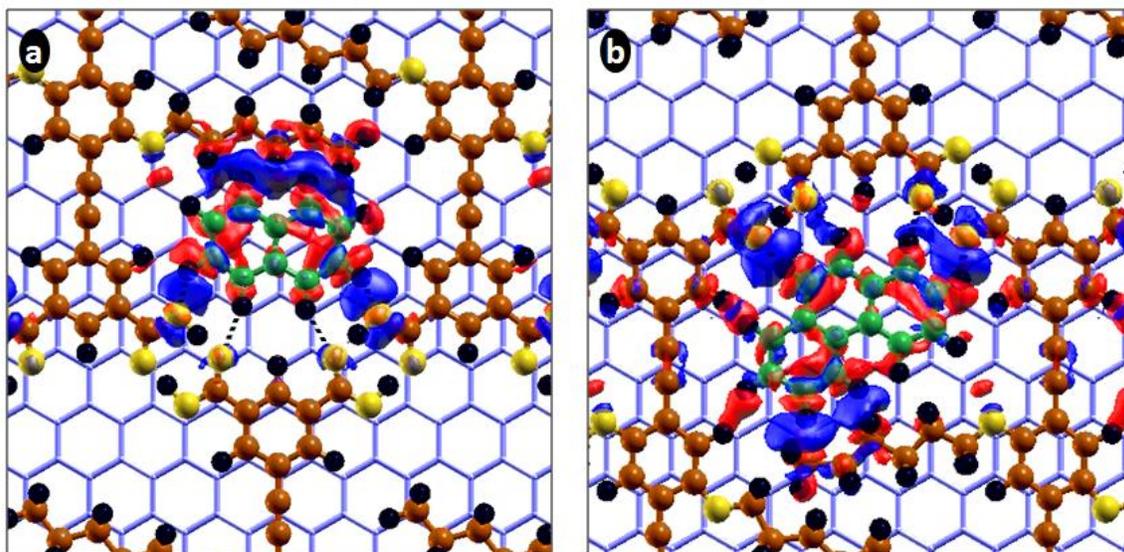


Figure S7. Theoretically computed charge redistribution plots between PE4B and different guest molecules. PE4B with (a) naphthalene and (b) phenanthrene on graphene in LIN pattern. Red and blue lobes indicate electron accumulation and depletion respectively, plotted at an isosurface value of 0.00045 e/bohr^3 . Note the total number of host-guest bonds are hard to define precisely in both (a) and (b), due to lower symmetry of LIN pattern. Color scheme: brown: C of PE4B; yellow: O, black: H; green: C of guests. Gray lines indicate bonds between C atoms of graphene.

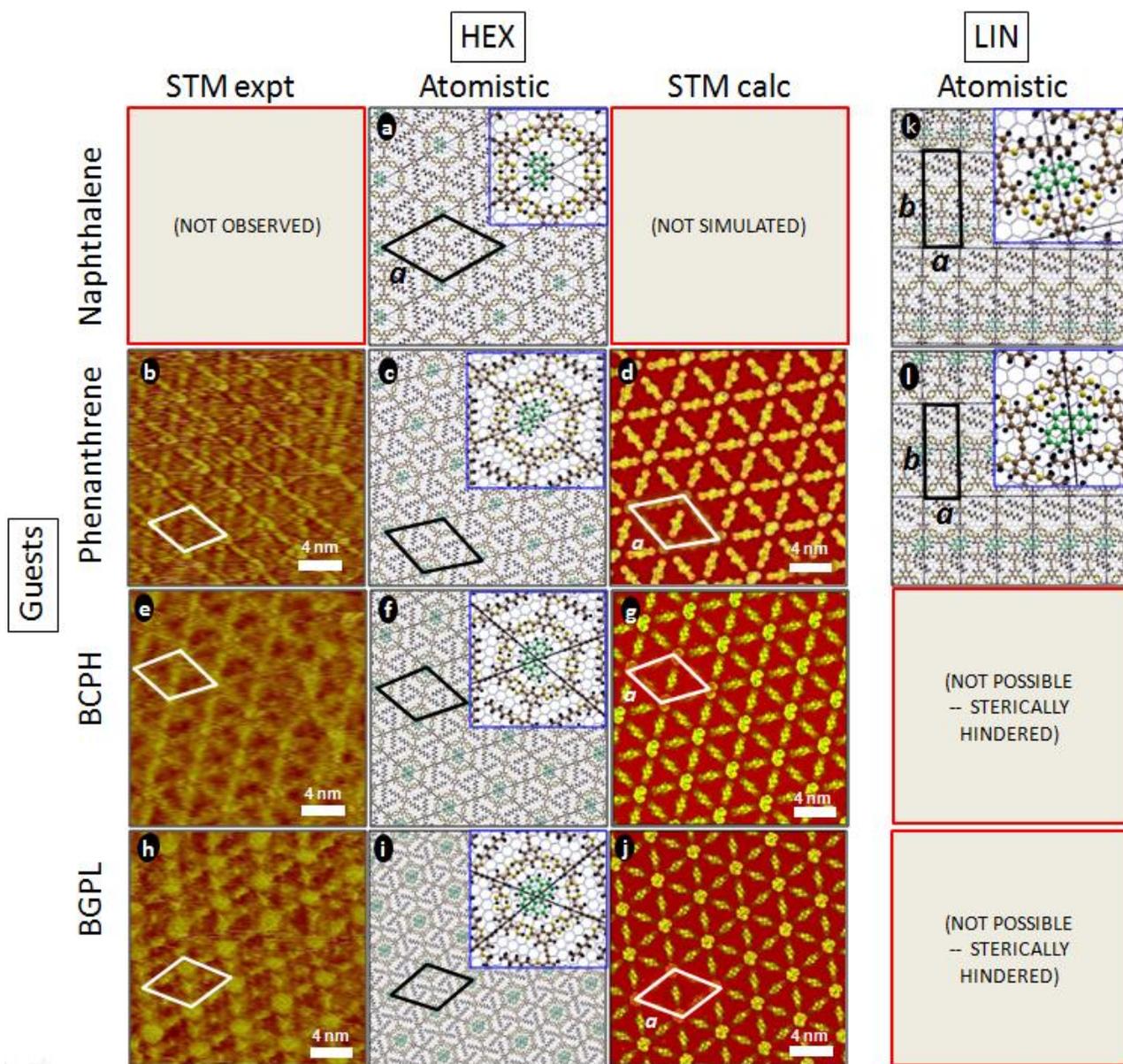


Figure S8. STM images and atomistic structures of PE3A with various guests in HEX (a)-(j) and LIN (k)-(l) patterns. Experimental STM images of PE3A with (b) phenanthrene (scan size: 20×20 nm²; $V_{\text{bias}} = -553$ mV; $I_t = 229$ pA), (e) BCPH (scan size: 20×20 nm²; $V_{\text{bias}} = -600$ mV; $I_t = 450$ pA) and (h) BGPL (scan size: 20×20 nm²; $V_{\text{bias}} = -1214$ mV; $I_t = 122$ pA) as guests in HEX pattern. Atomistic structures from DFT of PE3A with (a) naphthalene, (c) phenanthrene, (f) BCPH and (i) BGPL in HEX pattern; (k) naphthalene and (l) phenanthrene in LIN pattern. Simulated STM images of PE3A with (d) phenanthrene, (g) BCPH and (j) BGPL as guests. Simulated STM images are obtained at $V_{\text{bias}} = -2720$ mV. The black/white lines indicate the unit cell; with a and b are the unit cell parameters. Atomic colors: C of host: brown, H: black, O: yellow, C of graphene: gray, C of guest: green.

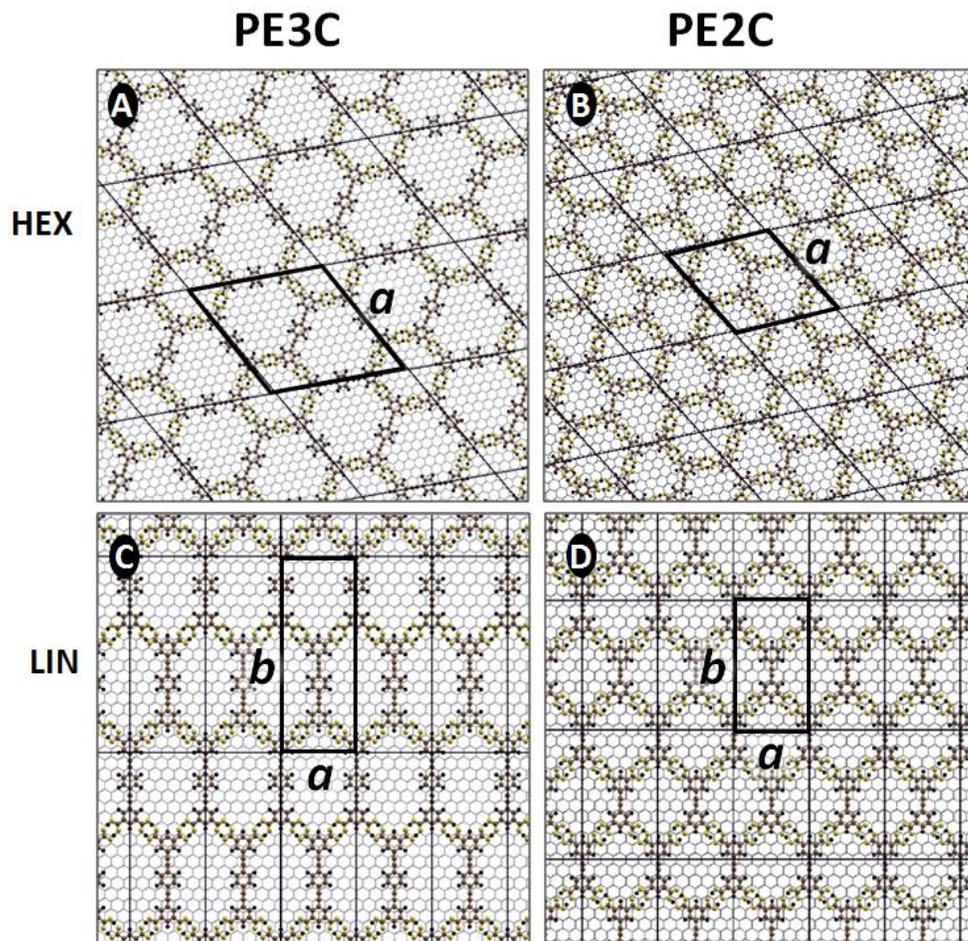


Figure S9: Atomistic structures of host assemblies of the test molecules on graphene. All coordinates have been relaxed using DFT calculations. Black rhombi and rectangles mark the boundaries of the unit cells for the HEX and LIN patterns, respectively. Atomic colors: C of host: brown, H: black, O: yellow, C of graphene: gray, C of guest: green.

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