Supporting Information

Acid/base responsive pseudo[3]rotaxanes from amine naphthotubes and bis-pyridinium/isoquinolinium guests

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1. Experimental section

1.1 Materials and methods. All reagents were commercially available and used without further purification. Solvents were either employed as purchased or dried prior to use by standard laboratory procedures. ¹H, ¹³C NMR, 2D NMR spectra were recorded on a Bruker Avance-400, 500, or 600 NMR spectrometer. Chemical shifts are reported in ppm with residual solvents or TMS (tetramethylsilane) as the internal standards. High-resolution electrospray-ionization mass spectra (HR-ESI-MS) were recorded on an applied Q EXACTIVE mass spectrometry system. UV-vis absorption spectra were obtained on a Hitachi U-2600 UV-vis spectrophotometer. Compounds $H1^1$ and $G2^2$ were synthesized according to literature procedures.

1.2 Synthetic procedures of G1-G6



Scheme S1. Synthesis of guests G1-G6.

General synthetic procedure: Dibromoalkane (1.0 eq.) in MeCN was dropwise added into the solution of pyridine or isoquinoline (2.5 eq.) in MeCN at 80 °C. The resulting mixture was stirred vigorously overnight. After cooling to room temperature, the solution was concentrated in vacuum. The residue was washed with ether to give corresponding bromide salts. Saturated aqueous NH_4PF_6 was dropwise added into the solution of bromide salts in deionized water. After stirring for 6 h at room temperature, the white precipitate was collected through filtration and washed several times with deionized water. The filter cake was dried to afford corresponding products. The characterization data are shown below:

G1: white solid, yield: 76%. ¹H NMR (400 MHz, DMSO- d_6 , 298 K): δ [ppm] = 9.04 – 8.97 (m, 4H), 8.63 – 8.53 (m, 2H), 8.17 – 8.08 (m, 4H), 4.55 (t, J = 7.4 Hz, 4H), 1.91 – 1.87 (t, J = 7.3 Hz, 4H), 1.34 – 1.25 (m, 4H). ¹³C NMR (101 MHz, DMSO- d_6 , 298 K):

 δ [ppm] = 145.73, 144.79, 128.30, 60.94, 30.53, 24.98. ESI-TOF-HRMS: m/z calcd for [M-PF₆]⁺ C₁₆H₂₂F₆N₂P⁺, 387.1419; found 387.1413 (error = -1.5 ppm).



Figure S1. ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of compound G1.



Figure S2. ¹³C NMR spectrum (101 MHz, DMSO-*d*₆, 298 K) of compound G1. G3: white solid, yield: 48%. ¹H NMR (600 MHz, DMSO-*d*₆, 298 K): δ [ppm] = 9.06 (d, *J* = 5.7 Hz, 4H), 8.60 (t, *J* = 7.7 Hz, 2H), 8.17 – 8.14 (m, 4H), 4.58 (t, *J* = 7.5 Hz, 4H), 1.93 – 1.87 (m, 4H), 1.32 – 1.22 (m, 12H). ¹³C NMR (151 MHz, DMSO-*d*₆, 298 K): δ [ppm] = 145.46, 144.67, 128.07, 60.81, 30.66, 28.66, 28.34, 25.40. ESI-TOF-

HRMS: m/z calcd for $[M-PF_6]^+ C_{20}H_{30}F_6N_2P^+$, 443.2045; found 443.2035 (error = -2.3 ppm).



Figure S3. ¹H NMR spectrum (600 MHz, DMSO-*d*₆, 298 K) of compound G3.



Figure S4. ¹³C NMR spectrum (151 MHz, DMSO-*d*₆, 298 K) of compound G3. G4: white solid, yield: 71%. ¹H NMR (600 MHz, DMSO-*d*₆, 298 K): δ [ppm] =10.05 (s, 2H), 8.78 (d, J = 6.8 Hz, 2H), 8.59 (d, J = 6.7 Hz, 2H), 8.47 (d, J = 8.3 Hz, 2H), 8.35 (d, J = 8.3 Hz, 2H), 8.28 – 8.25 (m, 2H), 8.10 – 8.06 (m, 2H), 4.72 – 4.66 (m, 4H), 2.01 – 1.98 (m, 4H), 1.31 – 1.29 (m, 8H), 1.25 – 1.23 (m, 4H). ¹³C NMR (151 MHz, DMSO-

 d_{6} , 298 K): δ [ppm] =149.88, 136.90, 136.85, 134.86, 131.20, 130.32, 127.25, 127.22, 125.86, 60.80, 30.44, 28.70, 28.42, 25.52. ESI-TOF-HRMS: m/z calcd for [M-PF₆]⁺ C₂₈H₃₄F₆N₂P⁺, 543.2358; found 543.2360 (error = 0.4 ppm).



Figure S5. ¹H NMR spectrum (600 MHz, DMSO-*d*₆, 298 K) of compound G4.



Figure S6. ¹³C NMR spectrum (151 MHz, DMSO-*d*₆, 298 K) of compound G4. G5: white solid, yield: 90%. ¹H NMR (400 MHz, DMSO-*d*₆, 298 K): δ [ppm] = 9.09 – 9.02 (m, 4H), 8.61 – 8.59 (m, 2H), 8.19 – 8.10 (m, 4H), 4.57 (t, *J* = 7.5 Hz, 4H), 1.94 – 1.86 (m, 4H), 1.29 – 1.22 (m, 20H). ¹³C NMR (101 MHz, DMSO-*d*₆, 298 K): δ [ppm]

= 145.51, 144.72, 128.13, 60.91, 30.74, 29.27, 29.08, 29.01, 28.86, 28.45, 25.47. ESI-TOF-HRMS: m/z calcd for $[M-PF_6]^+ C_{24}H_{38}F_6N_2P^+$, 499.2671; found 499.2657 (error = -2.8 ppm).



Figure S7. ¹H NMR spectrum (400 MHz, DMSO-*d*₆, 298 K) of compound G5.



Figure S8. ¹³C NMR spectrum (101 MHz, DMSO- d_6 , 298 K) of compound **G5**. **G6**: white solid, yield: 30%. ¹H NMR (400 MHz, DMSO- d_6 , 298 K): δ [ppm] = 10.06 (s, 2H), 8.80 – 8.78 (m, 2H), 8.59 (d, J = 6.8 Hz, 2H), 8.47 (d, J = 8.3 Hz, 2H), 8.35 (d, J = 8.4 Hz, 2H), 8.27 – 8.25 (m, 2H), 8.09 – 8.07 (m, 2H), 4.69 (t, J = 7.4 Hz, 4H), 2.02

 $-1.99 \text{ (m, 4H)}, 1.33 - 1.20 \text{ (m, 20H)}. {}^{13}\text{C NMR} (101 \text{ MHz, DMSO-}d_6, 298 \text{ K}): \delta \text{ [ppm]} = 149.93, 136.92, 136.87, 134.90, 131.22, 130.35, 127.28, 127.24, 125.87, 60.82, 30.46, 29.04, 28.97, 28.82, 28.47, 25.53. ESI-TOF-HRMS: m/z calcd for [M-PF_6]⁺ C₂₃H₄₂F₆N₂P⁺, 599.2984; found 599.2980 (error = -0.7 ppm).$



Figure S9. ¹H NMR spectrum (400 MHz, DMSO- d_6 , 298 K) of compound G6.



Figure S10. ¹³C NMR spectrum (101 MHz, DMSO-*d*₆, 298 K) of compound G6.

2. Characterization data of pseudo[2]rotaxanes

2.1 ¹H NMR spectra of H1 and G1-G2

Determination of the association constants of the 1:1 complexes

The stoichiometries and association constants between **H1** and **G1-G2** were determined by ¹H NMR titrations. For a typical ¹H NMR titration, a 0.5 mM solution of **H1** was prepared in CD_2Cl_2/CD_3CN (1:1, v/v). This solution (0.5 mL) was placed in an NMR tube and then titrated with a solution of guest. Nonlinear curve-fitting method was then used to obtain the binding constants through the following equation:³

$$\delta = \delta_0 + \Delta \delta (0.5/[H]_0) ([G] + [H]_0 + 1/K_a - (([G] + [H]_0 + 1/K_a)^2 - 4[H]_0[G])^0 - 0.5)$$
(Eq. S1)

where δ is the chemical shift of proton 2 on H1 at [G], δ_0 is the chemical shift of proton 2 on H1 without guest, $\Delta\delta$ is the chemical shift change of proton 2 when the H1 is completely complexed, [H]₀ is the fixed initial concentration of the H1 and [G] is the varying concentration of guest.



Figure S11. Partial ¹H NMR spectra (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of H1 (0.5 mM) with different concentrations of G1 (0~4.17 mM).



Figure S12. The chemical shift of H_2 on H1 upon addition of G1 in the 1:1 mixture of CD_2Cl_2 and CD_3CN at 298 K. The red solid line was obtained from the non-linear curve-fitting using Eq. S1. The titration was repeated twice.



Figure S13. Plots of mole fractions of H1 (red) and G1@H1 (black) when titrating G1 into a solution of H1. The concentration of [G1@H1] was calculated by [HG] = $0.5([H] + [G]_0 + 1/K_a) - 0.5(([H]_0 + [G]_0 + 1/K_a)^2 - 4[H]_0[G]_0)^0 - 0.5)$.³ The concentration of [H1] was calculated by [H] = [H]_0-[HG].



Figure S14. Job plot obtained by plotting the chemical shift change ($\Delta\delta$) of the host's proton 2 in ¹H NMR spectra by varying the ratio of host and guest against the mole fraction of host H1. This experiment supports the 1:1 binding stoichiometry between H1 and G1 in the 1:1 mixture of CD₂Cl₂ and CD₃CN.



Figure S15. ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 1.0 mM, 298 K) of (a) **G2**, (c) **H1**, and (b) their equimolar mixture. In the host-guest mixture, the protons of the guest undergo significant shifts, accompanied with peak broadening or disappearing into the baseline, suggesting a binding event.



Figure S16. Partial ¹H NMR spectra (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of H1 (0.5 mM) with different concentrations of G2 (0~4.17 mM).



Figure S17. The chemical shift of H_2 on H1 upon addition of G2 in the 1:1 mixture of CD_2Cl_2 and CD_3CN at 298 K. The red solid line was obtained from the non-linear curve-fitting using Eq. S1. The titration was repeated twice.



Figure S18. Plots of mole fractions of H1 (red) and G2@H1 (black) when titrating G2 into a solution of H1. The concentration of [G2@H1] was calculated by $[HG] = 0.5([H] _{0}+[G]_{0}+1/K_{a})-0.5(([H]_{0}+[G]_{0}+1/K_{a})^{2}-4[H]_{0}[G]_{0})^{0}-0.5)$.³ The concentration of [H1] was calculated by $[H] = [H]_{0}-[HG]$.



Figure S19. Job plot obtained by plotting the chemical shift change ($\Delta\delta$) of the host's proton 2 in ¹H NMR spectra by varying the ratio of host and guest against the mole fraction of host H1. This experiment supports the 1:1 binding stoichiometry between H1 and G2 in the 1:1 mixture of CD₂Cl₂ and CD₃CN.

2.2 ESI-MS of the 1:1 complexes



Figure S21. ESI-MS of G2@H1.

3. Characterization data of pseudo[3]rotaxanes

Determination of the association constants of the 2:1 complexes: The stoichiometries and association constants between H1 and G3-G6 were determined by ¹H NMR titrations. For a typical ¹H NMR titration, a 0.5 mM solution of guest was prepared in CD₂Cl₂/CD₃CN (1:1, v/v). This solution (0.5 mL) was placed in an NMR tube and then titrated with a solution of H1. Nonlinear curve-fitting method was then used to obtain the binding constants through the following equation:³

 $\delta = \delta_0 + (\Delta \delta_{\text{HG}} K_{a1}[\text{H}] + \Delta \delta_{\text{H2G}} K_{a1} K_{a2}[\text{H}]^2) / (1 + K_{a1}[\text{H}] + K_{a1} K_{a2}[\text{H}]^2)$ (Eq. S2) where δ is the chemical shift of proton on guest (proton b on **G3**, **G5**; proton e on **G4**, **G6**) at [H], δ_0 is the chemical shift of proton on guest without **H1**, $\Delta \delta_{\text{HG}}$ is the chemical shift change of proton on guest when guest is completely complexed by the first H1, $\Delta \delta_{\text{H2G}}$ is the chemical shift change of proton on guest when guest is completely complexed by the second H1. [H] is the varying concentration of H1.

The concentration of [G] can be calculated by $[G] = [G]_0 / (1+K_{a1}[H] + K_{a1}K_{a2}[H]^2).^3$ The concentration of [HG] and [H₂G] can be calculated by [HG] = $K_{a1}[H][G]$ and [H₂G] = $K_{a2}[HG][G]$, respectively. [G]₀ is the fixed initial concentration of the guest.

3.1 ¹H and 2D NMR spectra of H1 and G3-G4



Figure S22. The chemical shift of H_b on G3 upon addition of H1 in the 1:1 mixture of CD_2Cl_2 and CD_3CN at 298 K. The red solid line was obtained from the non-linear curve-fitting using Eq. S2.



Figure S23. Job plot obtained by plotting the chemical shift change ($\Delta\delta$) of the guest's proton b in ¹H NMR spectra by varying the ratio of host and guest against the mole

fraction of guest G3. This experiment supports the 2:1 binding stoichiometry between H1 and G3 in the 1:1 mixture of CD_2Cl_2 and CD_3CN .



Figure S24. Partial ¹H,¹H-ROESY NMR spectrum (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of **G3@H1**₂ ([**H1**] = 16 mM). The peak attributed to protons H_a of the pyridinium rings on **G3** disappears into the baseline, and we can't discriminate it clearly in the NMR spectrum.



Figure S25. ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 1.0 mM, 298 K) of (a) **G4**, (c) **H1**, and (b) their 2:1 (H:G) mixture. In the host-guest mixture, the protons of

the guest undergo significant shifts, accompanied with peak broadening or disappearing into the baseline, suggesting a binding event.



Figure S26. Partial ¹H NMR spectra (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of G4 (0.5 mM) with different concentrations of H1 (0~3.45 mM).



Figure S27. The chemical shift of H_e on G4 upon addition of H1 in the 1:1 mixture of CD_2Cl_2 and CD_3CN at 298 K. The red solid line was obtained from the non-linear curve-fitting using Eq. S2.



Figure S28. Plots of mole fractions of G4 (black), G4@H1 (red) and G4@H1₂ (blue) when titrating H1 into a solution of G4. The concentration of [G4] was calculated by $[G] = [G]_0 / (1+K_{a1}[H]+K_{a1}K_{a2}[H]^2)$.³ The concentration of [G4@H1] and [G4@H1₂] were calculated by [HG] = $K_{a1}[H][G]$ and $[H_2G] = K_{a2}[HG][G]$, respectively. [G]₀ is the fixed initial concentration of the G4.



Figure S29. Job plot obtained by plotting the chemical shift change ($\Delta\delta$) of the guest's proton e in ¹H NMR spectra by varying the ratio of host and guest against the mole fraction of guest G4. This experiment supports the 2:1 binding stoichiometry between H1 and G4 in the 1:1 mixture of CD₂Cl₂ and CD₃CN.



Figure S30. Partial ¹H,¹H-ROESY NMR spectrum (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of G4@H1₂ ([H1] = 16 mM). The peak attributed to H_e on G4 was assigned according the ¹H NMR titrations spectra. Other peaks attributed to protons on G4 exhibit significant broadening effects or disappear into the baseline, and we can't discriminate them clearly in the NMR spectrum.

3.2 ESI-MS of the 2:1 complexes



Figure S31. ESI-MS of G3@H12.



Figure S32. ESI-MS of G4@H12.





Figure S33. ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 1.0 mM, 298 K) of (a) **G5**, (c) **H1**, and (b) their 2:1 (H:G) mixture. In the host-guest mixture, the protons of the guest undergo significant shifts, accompanied with peak broadening or disappearing into the baseline, suggesting a binding event.



Figure S34. Partial ¹H NMR spectra (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of G5 (0.5 mM) with different concentrations of H1 (0~3.45 mM).



Figure S35. The chemical shift of H_b on G5 upon addition of H1 in the 1:1 mixture of CD_2Cl_2 and CD_3CN at 298 K. The red solid line was obtained from the non-linear curve-fitting using Eq. S2.



Figure S36. Plots of mole fractions of G5 (black), G5@H1 (red) and G5@H1₂ (blue) when titrating H1 into a solution of G5.



Figure S37. Job plot obtained by plotting the chemical shift change ($\Delta\delta$) of the guest's proton b in ¹H NMR spectra by varying the ratio of host and guest against the mole fraction of guest G5. This experiment supports the 2:1 binding stoichiometry between H1 and G5 in the 1:1 mixture of CD₂Cl₂ and CD₃CN.



Figure S38. ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 1.0 mM, 298 K) of (a) **G6**, (c) **H1**, and (b) their 2:1 (H:G) mixture. In the host-guest mixture, the protons of the guest undergo significant shifts, accompanied with peak broadening or disappearing into the baseline, suggesting a binding event.



Figure S39. Partial ¹H NMR spectra (500 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of G6 (0.5 mM) with different concentrations of H1 (0~3.45 mM).



Figure S40. The chemical shift of H_e on G6 upon addition of H1 in the 1:1 mixture of CD_2Cl_2 and CD_3CN at 298 K. The red solid line was obtained from the non-linear curve-fitting using Eq. S2.



Figure S41. Plots of mole fractions of G6 (black), G6@H1 (red) and G6@H1₂ (blue) when titrating H1 into a solution of G6.



Figure S42. Job plot obtained by plotting the chemical shift change ($\Delta\delta$) of the guest's proton e in ¹H NMR spectra by varying the ratio of host and guest against the mole fraction of guest G6. This experiment supports the 2:1 binding stoichiometry between H1 and G6 in the 1:1 mixture of CD₂Cl₂ and CD₃CN.

3.4 ESI-MS of the 2:1 complexes



Figure S43. ESI-MS of G5@H12.



Figure S44. ESI-MS of G6@H12.

4. X-ray single crystallography

Suitable single crystals of $G4@H1_2 \cdot 3CH_3CN$ were successfully obtained by slow evaporation of their saturated solutions in 1, 2-dichloroethane and acetonitrile (1:1, v/v). Single crystal X-ray data were collected on a Bruker D8 VENTURE with Ga K α radiation ($\lambda = 1.34139$ Å) at 100 K. The structure was solved by intrinsic phasing methods (SHELXT) and refined by full-matrix least-squares F^2 using SHELXL⁴ in the OLEX2 program package.⁵ All non-hydrogen atoms were refined with anisotropic thermal parameters and the hydrogen atoms were fixed at calculated positions and refined by a riding mode.

entry	G4@H12•3CH3CN				
Moiety formula	2(C ₆₆ H ₇₀ N ₂ O ₈), C ₂₈ H ₃₄ N ₂ , 3(C ₂ H ₃ N)				
	$2(F_6P),$				
Empirical formula	$C_{166}H_{183}F_{12}N_9O_{16}P_2$				
Formula weight	2850.14				
Temperature/K	100				
Crystal system	triclinic				
Space group	P-1				
a/Å	15.6160(7)				
b/Å	22.8099(10)				
c/Å	24.2217(10)				
α/°	65.842(2)				
β/°	74.382(2)				
$\gamma^{ m o}$	77.047(2)				
Volume/Å ³	7517.4(6)				
Z	2				
$\rho_{calc}g/cm^3$	1.259				
μ/mm^{-1}	0.601				
F(000)	3016				
Reflections collected	194297				
Independent reflections	$26565 \ [R_{int} = 0.0708, R_{sigma} = 0.0446]$				
Data/restraints/parameters	26565/2814/2289				
Goodness-of-fit on F ²	1.077				
Final R indexes [I>= 2σ (I)]	$R_1 = 0.0918, wR_2 = 0.2653$				
Final R indexes [all data]	$R_1 = 0.1070, wR_2 = 0.2831$				
CCDC number	2370291				

Crystal data and structure refinement for G4@H12•3CH3CN



Crystal structure of G4@H1₂•3CH₃CN.



Optical microscopic pictures of G4@H1₂ single crystals.

5. UV-Vis spectra of the complexes



Figure S45. Optical images show the solution color change because of the charge transfer between H1 (2.0 mM or 4.0 mM) and G1-G6 (2.0 mM) in 1:1 (v/v) CHCl₃/CH₃CN solution.



Figure S46. UV-vis spectra in 1:1 (v/v) CHCl₃/CH₃CN solution at 298 K of (a) H1 (2.0 mM) upon complexation with 1.0 equiv. of G1-G2 and (b) H1 (4.0 mM) upon complexation with G3-G6 (2.0 mM).

6. Acid/base stimuli-responsive complexation



Figure S47. UV-vis spectra of G1@H1 ([G1]=[H1]=2.00 mM, 298 K) in CHCl₃/CH₃CN (1:1, v/v) in the acid/base stimuli-responsive cycles.



Figure S48. UV-vis spectra of G2@H1 ([G2]=[H1]=2.00 mM, 298 K) in CHCl₃/CH₃CN (1:1, v/v) in the acid/base stimuli-responsive cycles. Inset: acid/base stimuli-responsive color changes of G2@H1.



Figure S49. UV-vis spectra of **G3@H1**₂ ([**G3**]=2.00 mM, [**H1**]=4.00 mM, 298 K) in CHCl₃/CH₃CN (1:1, v/v) in the acid/base stimuli-responsive cycles.



Figure S50. UV-vis spectra of **G5@H1**₂ ([**G5**]=2.00 mM, [**H1**]=4.00 mM, 298 K) in CHCl₃/CH₃CN (1:1, v/v) in the acid/base stimuli-responsive cycles.



Figure S51. UV-vis spectra of **G6@H1**₂ ([**G6**]=2.00 mM, [**H1**]=4.00 mM, 298 K) in CHCl₃/CH₃CN (1:1, v/v) in the acid/base stimuli-responsive cycles. Inset: acid/base stimuli-responsive color changes of **G6@H1**₂.



Figure S52. Partial ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 298 K) of a) 1.0 mM free guest **G1**, b) after addition of 1 equiv. of **H1** (1.00 mM) to a), c) after addition of 3 equiv. of TFA to b), d) after addition of 3 equiv. of Et₃N to c).



Figure S53. Partial ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 298 K) of a) 1.0 mM free guest **G2**, b) after addition of 1 equiv. of **H1** (1.00 mM) to a), c) after addition of 3 equiv. of TFA to b), d) after addition of 3 equiv. of Et₃N to c).



Figure S54. Partial ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 298 K) of a) 1.0 mM free guest **G3**, b) after addition of 2 equiv. of **H1** (2.00 mM) to a), c) after addition of 6 equiv. of TFA to b), d) after addition of 6 equiv. of Et₃N to c).



Figure S55. Partial ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 298 K) of a) 1.0 mM free guest **G4**, b) after addition of 2 equiv. of **H1** (2.00 mM) to a), c) after addition of 6 equiv. of TFA to b), d) after addition of 6 equiv. of Et₃N to c).



Figure S56. Partial ¹H NMR spectra (400 MHz, CD₂Cl₂:CD₃CN=1:1, 298 K) of a) 1.0 mM free guest **G5**, b) after addition of 2 equiv. of **H1** (2.00 mM) to a), c) after addition of 6 equiv. of TFA to b), d) after addition of 6 equiv. of Et₃N to c).



Figure S57. Partial ¹H NMR spectra (400 MHz, $CD_2Cl_2:CD_3CN=1:1$, 298 K) of a) 1.0 mM free guest **G6**, b) after addition of 2 equiv. of **H1** (2.00 mM) to a), c) after addition of 6 equiv. of TFA to b), d) after addition of 6 equiv. of Et₃N to c).

7. Computational data

Computational Methods. Geometry optimizations were performed with Gaussian 16 software package,⁶ using the (B3LYP-D3(BJ)) method⁷ with 6-31G* basis set⁸ in vacuum without applying any geometry constraints (C1 symmetry). Frequency calculations were then conducted at the same computational level to confirm the nature of all located stationary points. Single point energy was calculated with Gaussian 16 software package using the (B3LYP-D3(BJ)) method with 6-311+G** basis set in vacuum. Non-covalent interaction (Independent gradient model based on Hirshfeld partition, IGMH) analysis⁹ was carried out with Multiwfn 3.8 (dev) program¹⁰ and visualized by the VMD 1.9.3 program.¹¹ (set isovalue = 0.005)

Energy-minimized structures of G1@H1:				Ene	ergy-minimized	l structures of (G2@H1:	
Car	rtesian coordina	ates of G1@H1	l:	Cai	Cartesian coordinates of G2@H1:			
С	2.9805992	-1.6613262	-3.5946312	С	-3.0979952	-3.7318013	-1.8694391	
С	0.1967830	-1.2460181	-3.7236753	С	-0.5176780	-4.1222883	-0.8159501	
С	2.4683802	-0.4093861	-3.9983113	С	-2.9331242	-4.2374853	-0.5608380	
С	2.1497851	-2.7179322	-3.2752012	С	-2.0189681	-3.4545572	-2.6889702	
С	0.7359790	-2.5423812	-3.4102152	С	-0.7016821	-3.6846183	-2.1732252	
С	1.1126441	-0.1988760	-4.0286463	С	-1.6759071	-4.4075113	-0.0408840	
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				Η	0.8797201	-1.1472901	0.6896590
				Η	0.5854550	0.5845930	0.8916921

Н	-0.8762531	-1.1729901	-1.1572121
Н	-1.1542741	0.5317890	-0.8900581
Н	-1.7512731	-1.7284641	1.0705081
Н	-1.8272551	-0.0311450	1.5222281
Н	-3.3741072	-1.2143341	-0.8432311
Н	-3.5128513	0.4191700	-0.2104990
Н	-4.3642623	-2.1639932	1.1965631
Η	-4.3290963	-0.5925620	2.0046361

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