## **Supporting Information**

## Monitoring Self-Sorting by Electrospray Ionization Mass Spectrometry: Formation Intermediates and Error-Correction during the Self-Assembly of Multiply Threaded Pseudorotaxanes

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*Figure S1.* Changes of <sup>1</sup>H NMR spectra (700 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 5:1, 1.0 mM) with increasing reaction time after mixing **1**-H·PF<sub>6</sub> and **2**-H·PF<sub>6</sub> with **4** in a 1:1:1 ratio, and <sup>1</sup>H NMR spectra of independently generated **18b**-H·PF<sub>6</sub> and **19**-H·PF<sub>6</sub>. Complexed and uncomplexed species are denoted by "c" and "uc" in parentheses, respectively. Fast processes occurring at a timescale less than 4 min are not observable with these NMR experiments. The signals for **18**-H·PF<sub>6</sub> were not identifiable due to significant signal overlapping and broadening and its low concentration. The complexation of **1**-H·PF<sub>6</sub> to the 24-crown-4 unit of **4** has almost finished within 4 min, while most of **2**-H·PF<sub>6</sub> is still free. It is then gradually consumed to afford **11**-2H·2PF<sub>6</sub> over time.



*Figure S2.* <sup>1</sup>H NMR spectra (500 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 2:1, 10.0 mM) of (a) C7, (c) **3**-2H·2PF<sub>6</sub>, and (b) an equimolar mixture of **3**-2H·2PF<sub>6</sub> and C7. Complexed and uncomplexed species are denoted by "c" and "uc" in the parentheses, respectively. Asterisk = solvent. The NMR results suggest **15**-2H·2PF<sub>6</sub> to be the dominating species in a 1:1 mixture of **3**-2H·2PF<sub>6</sub> and C7 as indicated by the significant complexation-induced shifts of H<sub>c</sub>, H<sub>d</sub>, and H<sub>e</sub> while H<sub>a</sub> and H<sub>b</sub> remain almost unaffected with respect to the free axle. This should be even more so, when the binding constants increase upon changing the solvent to a less polar one (CH<sub>2</sub>Cl<sub>2</sub> : CH<sub>3</sub>CN = 8:1, as used in the MS experiments).



*Figure S3.* ESI-FTICR mass spectrum of 1:1 mixture (295 K,  $CH_2Cl_2:CH_3CN = 8:1, 250 \mu M$ ) of **3-**2H**·**2PF<sub>6</sub> and **C7**. The result supports the conclusion from the NMR experiments.



*Figure S4.* ESI-FTICR mass spectra (295 K,  $CH_2Cl_2:CH_3CN = 8:1$ , 250 µM) of (a) 1:1 mixture of **2**-H·PF<sub>6</sub> and **4**, (b) 1:1 mixture of **1**-H·PF<sub>6</sub> and **4**, (c) 2:1 mixture of **2**-H·PF<sub>6</sub> and **4** ([**4**] = 250 µM), and (d) 1:2:1 mixture of **1**-H·PF<sub>6</sub>, **2**-H·PF<sub>6</sub>, and **4** ([**4**] = 250 µM). These results indicate: (i) **19**-H·PF<sub>6</sub> and **18b**-H·PF<sub>6</sub> are the by far major species in the 1:1 mixtures of **4** with **1**-H·PF<sub>6</sub> and **2**-H·PF<sub>6</sub>, respectively; (ii) **20**-2H·2PF<sub>6</sub> is dominant in the 2:1 mixture of **2**-H·PF<sub>6</sub> and **4**, and **18**-H·PF<sub>6</sub> may exist as minor species even when considering the fragmentation of [**20**-2H]<sup>2+</sup> into [**2**-H]<sup>+</sup> and [**18**-H]<sup>+</sup> during ionization process since the intensity of [**18**-H]<sup>+</sup> is much higher than that of [**2**-H]<sup>+</sup>; (iii) self-sorted pseudorotaxane **11**-2H·2PF<sub>6</sub> is thermodynamically favored over mismatched structure **20**-2H·2PF<sub>6</sub> in a 1:2:1 mixture of **1**-H·PF<sub>6</sub>, **2**-H·PF<sub>6</sub>, and **4**.



*Figure S5.* <sup>1</sup>H NMR spectra (500 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 2:1, 10.0 mM) of (a) **2-H·PF**<sub>6</sub>, (d) **4**, (b) 1:1 mixture of **2-H·PF**<sub>6</sub> and **4**, and (c) 2:1 mixture of **2-H·PF**<sub>6</sub> and **4**. The "uc" in the parentheses denotes the signal from uncomplexed species. Asterisk = solvent. The similar shifting of H<sub>c</sub>, H<sub>d</sub>, H<sub>e</sub>, and H<sub>f</sub> as observed for [**2-H@C7**]·PF<sub>6</sub> (Figure 1) and the absence of free H<sub>d</sub> in (b) indicate that **18b-H·PF**<sub>6</sub> is dominant in this solution. The assignment of **20-**2H·2PF<sub>6</sub> in the 2:1 mixture of **2-**H·PF<sub>6</sub> and **4** is inconclusive due to serious signal overlapping and broadening. But considering the MS result in Figure S4, **20-**2H·2PF<sub>6</sub> is believed to be the major species as indirectly suggested by the low peak intensity of free H<sub>d</sub> in Figure S5d.



*Figure S6.* <sup>1</sup>H-<sup>1</sup>H COSY NMR spectrum (500 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 2:1, 10.0 mM) of 1:1 mixture of **2-H·PF**<sub>6</sub> and **4**. The COSY spectrum supports the assignments for the peaks in Figure S5b.



*Figure S7.* <sup>1</sup>H-<sup>1</sup>H COSY NMR spectrum (500 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 2:1, 10.0 mM) of 2:1 mixture of **2**-H·PF<sub>6</sub> and **4**. Due to significant signal overlapping and broadening, the COSY spectrum still could not contribute to the identification of **20**-2H·2PF<sub>6</sub> in the 2:1 mixture of **2**-H·PF<sub>6</sub> and **4**.



*Figure S8.* <sup>1</sup>H NMR spectra (500 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 2:1, 10.0 mM) of (a) **4**, (c) **1**-H·PF<sub>6</sub>, and (b) 1:1 mixture of **1**-H·PF<sub>6</sub> and **4**. The "c" in the parentheses denotes the signal from complexed species. Asterisk = solvent. The obvious complexation-induced shift of H<sub>1</sub>, H<sub>a</sub>, and H<sub>b</sub> indicate **19**-H·PF<sub>6</sub> to be the predominant complex in this solution, which is in line with the MS results.



*Figure S9.* <sup>1</sup>H NMR spectra (500 MHz, 298 K, CDCl<sub>3</sub>:CD<sub>3</sub>CN = 2:1, 10.0 mM) of (a) 1:1:1 mixture of **1**-H·PF<sub>6</sub>, **2**-H·PF<sub>6</sub>, and **4**, (b) 1:2:1 mixture of **1**-H·PF<sub>6</sub>, **2**-H·PF<sub>6</sub>, and **4**, and (c) 2:1 mixture of **2**-H·PF<sub>6</sub> and **4**. Complexed and uncomplexed species are denoted by "c" and "uc" in the parentheses, respectively. Asterisk = solvent impurity. The significant signal overlapping and broadening hamper the unambiguous assignment of all species in 1:2:1 mixture of **1**-H·PF<sub>6</sub>, **2**-H·PF<sub>6</sub>, and **4**. But with the previous knowledge about the related systems in mind, complexed H<sub>a</sub> and H<sub>b</sub> and uncomplexed H<sub>d</sub> and their peak integrations suggest **11**-2H·2PF<sub>6</sub> to be the primary species. With respect to **20**-2H·2PF<sub>6</sub>, the NMR result is inconclusive but MS results indicate that it coexist with **11**-2H·2PF<sub>6</sub> in this solution as a minor species (Figure S4d).



*Figure S10.* <sup>1</sup>H NMR spectra (500 MHz, 298 K,  $CDCl_3:CD_3CN = 5:1$ , 2.0 mM) of (a) **3-**2H**·**2PF<sub>6</sub>, (c) **6**, and (b) 2:1 mixture of **3-**2H**·**2PF<sub>6</sub> and **6**. The "uc" in the parentheses denotes the signal from uncomplexed species. The NMR results suggest **25-**4H**·**4PF<sub>6</sub> is dominant in this solution since most of H<sub>a</sub> on **3-**2H**·**2PF<sub>6</sub> experience complexation-induced shift after mixing with **6** in 2:1 ratio.



*Figure S11.* ESI-FTICR mass spectrum of 2:1 mixture of **3**-2H·2PF<sub>6</sub> and **6** (295 K, CH<sub>2</sub>Cl<sub>2</sub>:CH<sub>3</sub>CN = 8:1, 250  $\mu$ M) under identical ionization condition as used for the kinetics study of **13**-4H·4PF<sub>6</sub>. The mass spectrum shows a low intensity of **25**-4H·4PF<sub>6</sub> with a charge distribution from dication to tetracation. Some fragments (*m*/*z* 413, 559, and 872) are also observed. This is presumably derived from the instability (or metastability) of multiply charged ions of **25**-4H·4PF<sub>6</sub> in the gas phase, which fragment quickly after their generation in the ion source.