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Electronic Supplementary Information

Recognition of naphthoflavone by Calix[4]pyrrole[2]phenanthrene

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1. Materials and Methods.

All reactions were carried out with oven-dried glassware. Commercial reagents were used without further purification. Flash column chromatography was performed on 100-200 mesh silica gel. ¹H NMR, ¹³C NMR spectra were recorded on a Bruker DMX400 NMR spectrometer. Melting points were determined using WRR melting point apparatus and were uncorrected. High Resolution atmospheric-pressure chemical ionization mass spectra (APCI-MS) were determined by Bruker Daltonics. Inc, APEX II. FT-ICRMS. Electrospray ionization mass spectra (ESI-MS) were recorded on the Thermo Fisher® Exactive LC-MS spectrometer.

2. Synthesis of New Compounds.



Compound 2: A mixture of 3,6-dibromo-9,10-dimethoxyphenanthrene^{S1} (3.94 g, 10 mmol), Na₂CO₃ (2.96 g, 28 mmol), N-Boc-2-pyrroleboronic acid (4.00 g, 22 mmol) and tetrakis(triphenylphosphine)palladium (0.23 g, 0.2 mmol) in 100 mL DMF in a flask was stirred at 90 °C for 24 h under N₂. After evaporating the solvents, resulting mixture was extracted with dichloromethane (3×50 mL) and then washed with water and brine successively. The organic layer was dried over anhydrous Na₂SO₄ and evaporated. Then, the residue was dissolved in 100 mL THF/MeOH (V/V = 5:1) and sodium methanolate (3.50g, 64.8mmol) was added. The resulting mixture was quenched by the addition of 50 mL water. The organic layer was separated and dried with anhydrous MgSO₄. The solvent was removed in vacuo and the residue was separated by column chromatography on silica gel (eluent: 1:2 DCM/Petroleum ether) to give compound **2**

(1.99 g, yield 54%) as gray solid. M.p.: 172-174 °C. ¹H NMR (400 MHz, Chloroform-*d*) δ 8.67 (s, 2H), 8.64 (s, 2H), 8.18 (d, J = 8.5 Hz, 2H), 7.74 (d, J = 8.5 Hz, 2H), 6.97 (s, 2H), 6.73 (s, 2H), 6.40 (q, J = 2.7 Hz, 2H), 4.08 (s, 6H). ¹³C NMR (101 MHz, CDCl₃) δ 143.6, 132.5, 130.3, 128.8, 127.8, 123.8, 122.9, 119.3, 117.2, 110.4, 106.6, 61.0. HRMS (APCI) m/z: [M+H]⁺ calcd for C₂₄H₂₁N₂O₂⁺, 369.1598; found, 369.1579.



Compound 1: To a mixture of **2** (184 mg, 0.5 mmol) and acetone (180 mg, 0.52 mmol) in dichloromethane (10 mL) was added trifluoroacetic acid (28.5 mg, 0.25 mmol). The mixture was stirred at room temperature for 20 h under N₂. Then the reaction was quenched by the addition of 10 mL water. The organic layer was separated and dried with anhydrous MgSO4. The solvent was removed in vacuo and the residue was separated by column chromatography on silica gel (eluent: 2:1 DCM/Petroleum ether) to give **1** (73.5 mg, 36%) as a gray solid. M.p.: 218-220 °C. ¹H NMR (400 MHz, Chloroform-*d*) δ 8.52 (s, 2H), 8.42 (s, 2H), 8.13 (d, *J* = 8.5 Hz, 2H), 7.70 (d, *J* = 8.6 Hz, 2H), 6.81 (s, 2H), 6.50 (s, 2H), 4.06 (s, 6H), 1.93 (s, 6H). ¹³C NMR (101 MHz, CDCl₃) δ 143.49, 140.30, 131.49, 129.58, 128.58, 127.58, 122.80, 122.69, 116.17, 106.18, 105.78, 60.99, 36.01, 28.91. HRMS (APCI) m/z: [M+H]⁺ calcd for C₅₄H₄₉N₄O₄⁺, 817.3748; found, 817.3753.



Figure S2.¹³C NMR spectrum (101 MHz, CDCl₃, 298K) of 2



Figure S4.¹³C NMR spectrum (101 MHz, CDCl₃, 298K) of 1

4. ¹H NMR studies of Complexation of the Host and Guest



Figure S5. Partial ¹H NMR spectra (400 MHz, CDCl₃, 298 K) of (I) free **1**, (II) **1** and 1.0 equiv. of **G1**, and (III) free **G1**. $[1]_0 = 4.0$ mM.



Figure S6. Partial ¹H NMR spectra (400 MHz, CDCl₃, 298 K) of (a) free **1**, (b) **1** and 1.0 equiv. of **G2**, and (c) free **G2**. $[1]_0 = 4.0$ mM.



Figure S7. Partial ¹H NMR spectra (400 MHz, CDCl₃, 298 K) of (a) free **1**, (b) **1** and 1.0 equiv. of **G3**, and (c) free **G3**. $[1]_0 = 4.0$ mM.



Figure S8. Partial ¹H NMR spectra (400 MHz, CDCl₃, 298 K) of (I) free **1**, (II) **1** and 1.0 equiv. of **G5**, and (III) free **G5**. $[1]_0 = 4.0$ mM.

5. Determination of the Association Constants of the Complexes

In the ¹H NMR titrations, CDCl₃ was chosen to dissolve the host and the guests. Chemical shifts were reported in parts per million (*ppm*). By a mole ratio plot, each stoichiometry was determined. Titration curve-fitting and association constant values were calculated by employing the BindFit program developed by Prof. Pall Thordarson of UNSW. 1:1 Binding stoichiometry was chosen in the BindFit program. This program employs a nonlinear least-squares regression analysis and is available free of cost online through the following link: http://supramolecular.org.



Figure S9. ¹H NMR spectra (400 MHz, CDCl₃, 298 K) of **1** at a concentration of 1.0 mM with different concentrations of **G4**: (1) 0.00 mM; (2) 0.2 mM; (3) 0.3 mM; (4) 0.4 mM; (5) 0.5 mM; (6) 0.6 mM; (7) 0.7 mM; (8) 0.8 mM; (9) 1.0 mM; (10) 1.2 mM; (11) 1.4 mM; (12) 1.6 mM; (13) 1.8 mM; (14) 2.0 mM; (15) 2.2 mM; (16) 2.4 Mm; (17) 3.0 mM; (18) 10.0 mM.



Figure S10. Mole ratio plot of the complexation of 1 and G4 in CDCl₃ at 298 K.



Figure S11. Plot of chemical shift (ppm) for the H_4 of 1 and G4 in CDCl₃ at 298 K.



Figure S12. Nonlinear least-square analysis of the ¹H NMR binding data corresponding to the formation of [**1**•G4] complex. The data were fitted to a 1:1 binding model to give $K_a = 105.8 \pm 22.2 \text{ M}^{-1}$. The residual distribution is shown below the binding isotherm. All solid lines were obtained from non-linear curve-fitting to a 1:1 binding model using the <u>www.supramolecular.org</u> web applet.



Figure S13. ¹H NMR spectra (400 MHz, CDCl₃, 298 K) of **1** at a concentration of 1.0 mM with different concentrations of **G5**: (1) 0.00 mM; (2) 0.2 mM; (3) 0.3 mM; (4) 0.4 mM; (5) 0.5 mM; (6) 0.6 mM; (7) 0.7 mM; (8) 0.8 mM; (9) 1.0 mM; (10) 1.2 mM;

(11) 1.4 mM; (12) 1.6 mM; (13) 1.8 mM; (14) 2.0 mM; (15) 2.2 mM; (16) 2.4 Mm; (17) 3.0 mM; (18) 10.0 mM.



Figure S14. Mole ratio plot of the complexation of 1 and G5 in CDCl₃ at 298 K.



Figure S15. Plot of chemical shift (ppm) for the H_4 of 1 and G5 in CDCl₃ at 298 K.



Figure S16. Nonlinear least-square analysis of the ¹H NMR binding data corresponding to the formation of [**1**•G5] complex. The data were fitted to a 1:1 binding model to give $K_a = 139.2 \pm 13.2 \text{ M}^{-1}$. The residual distribution is shown below the binding isotherm. All solid lines were obtained from non-linear curve-fitting to a 1:1 binding model using the <u>www.supramolecular.org</u> web applet.

6. ESI MS Studies of the new compounds and complexes



Figure S17. ESI Spectrum of 2



Figure S18. ESI Spectrum of host 1



Figure S19. ESI Spectrum of complex 1@G4



Figure S20. ESI Spectrum of complex 1@G5

7. ¹H-¹H ROESY spectral of the complexes



Figure S21. ¹H-¹H ROESY spectrum (400 MHz, CDCl₃, 298 K) of 1@G4



8. References

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