Supporting Information for:

Rapid Macrocycle Threading by a Fluorescent Dye-Polymer Conjugate in Water with Nanomolar Affinity

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A. Synthesis and Characterization

Macrocycles **M1**^{S1} and **M2**,^{S2} alkyne-substituted aminothiophenes **1**^{S3} and **2**,^{S4} triethyleneglycol azide,^{S5} and mPEG2000-azide^{S5} were synthesized using previously reported methods.



Compound **S1**^{S6}: Aminothiophene alkyne **1** (401 mg, 1.9 mmol) and squaric acid (110 mg, 0.96 mmol) were dissolved in a mixture of anhydrous butanol (20 mL) and anhydrous benzene (60 mL). The reaction mixture was refluxed for 8 hours under argon, equipped with a Dean-Stark apparatus. The solvent was removed under reduced pressure and the residue was purified *via* column chromatography using 0-30% acetone in chloroform as eluent to yield **S1** (367 mg, 0.76 mmol, 77%) as a dark blue-green solid. ¹H NMR (500 MHz, CDCl₃, 25 °C): δ 7.97 (d, *J* = 4.6 Hz, 2H), 6.28 (d, *J* = 4.4 Hz, 2H), 4.18 (d, *J* = 2.8 Hz, 4H), 3.81 (t, *J* = 5.4 Hz, 4H), 3.69 (t, *J* = 5.1 Hz, 4H), 3.62 (q, *J* = 7.2 Hz, 4H), 2.45 (t, *J* = 2.4 Hz, 2H), 1.31 (t, *J* = 7.2 Hz, 6H). ¹³C NMR (150 MHz, CDCl₃, 25 °C): δ 170.8, 169.5, 138.2, 115.4, 108.8, 79.0, 75.1, 66.9, 58.5, 53.4, 50.1, 12.0. HRMS (ESI-TOF): calculated for C₂₆H₂₈N₂O₄S₂ [M+H]⁺ 496.1485, found 496.1462.



Compound **S2**: Squaraine dye **S1** (10 mg, 0.02 mmol) and triethyleneglycol azide (9 mg, 0.05 mmol) were dissolved in chloroform (2 mL) and placed in a small vial. Copper catalyst (2.6 mg, 20 mol%), and *N*,*N*-diisopropylethylamine (10 μ L, 0.08 mmol) were added to the vial and the reaction mixture was stirred at room temperature for 6 hours. The solvent was removed under reduced pressure and the residue was purified *via* column chromatography using 10% methanol in chloroform as eluent to yield pure **S2** (12 mg, 0.014 mmol, 80%), as a dark blue, sticky solid. ¹H NMR (500 MHz, Acetonitrile-d₃, 25°C): δ 7.80 (d, *J* = 4.7 Hz, 2H), 7.79 (s, 2H), 6.40 (d, *J* = 5.0 Hz, 2H), 4.58 (s, 4H), 4.47 (t, *J* = 5.1 Hz, 4H), 3.80 (t, *J* = 5.1 Hz, 4H), 3.77 (t, *J* = 5.3 Hz, 4H), 3.68 (t, *J* = 5.3 Hz, 4H), 3.47-3.59 (m, 16H), 3.41-3.44 (m, 4H), 2.93-3.01 (br s, 2H), 1.23 (t, *J* = 7.2 Hz, 6H). ¹³C NMR: Small quantity of material prevented ¹³C analysis. HRMS (ESI-TOF):

calculated for $C_{38}H_{55}N_8O_{10}S_2$ [M+H]⁺ 846.3399, found 846.3399. LC-MS (ESI-TOF): calculated for $C_{38}H_{54}N_8NaO_{10}S_2$ [M+Na]⁺ 869.3297, found 846.3306.



Compound **S3**: Squaraine dye **S1** (5 mg, 0.01 mmol) and mPEG2000-azide (50 mg, 0.03 mmol) were dissolved in chloroform (1.5 mL) and placed in a small vial. Copper catalyst (1.3 mg, 20mol%), and *N*,*N*-diisopropylethylamine (5 μ L, 0.04 mmol) were added to the vial and the reaction mixture was stirred at room temperature for 24 hours. The solvent was removed under reduced pressure and the residue was purified *via* column chromatography using 10-20% methanol in chloroform as eluent to yield **S3** (20 mg, 4.4 μ mol, 45%), as a dark blue solid. ¹H NMR (600 MHz, CDCl₃, 25°C): δ 7.94 (d, *J* = 4.7 Hz, 2H), 7.71 (s, 2H), 6.26 (d, *J* = 4.4 Hz, 2H), 4.64 (s, 4H), 4.54 (t, *J* = 5.0 Hz, 4H), 3.86 (t, *J* = 5.0 Hz, 4H), 3.45-3.85 (m, ~304H), 3.38 (s, 6H), 1.28 (t, *J* = 3.4 Hz, 6H). ¹³C NMR: Small quantity of material prevented ¹³C analysis. MS (MALDI-TOF): The compound is a distribution of molecular weights, due to polydispersion of the PEG lengths, centered around 4261.36.





Compound **3**: Aminothiophene bisalkyne **2** (125 mg, 0.47 mmol) and squaric acid (28 mg, 0.24 mmol) were dissolved in a mixture of anhydrous butanol (10 mL) and anhydrous benzene (30 mL). The reaction mixture was refluxed for 8 hours under argon, equipped with a Dean-Stark apparatus. The solvent was removed under reduced pressure and the residue was purified *via* column chromatography using 0-30% acetone in chloroform as eluent to yield the product **2** (108 mg, 0.18 mmol, 73%) as a dark blue-green solid. ¹H NMR (500 MHz, CDCl₃, 25°C): δ 7.95 (d, *J* = 4.6 Hz, 2H), 6.33 (d, *J* = 4.6 Hz, 2H), 4.15 (d, *J* = 1.8 Hz, 8H), 3.75-3.85 (m, 16H), 2.44 (t, *J* = 2.4 Hz, 4H). ¹³C NMR (150 MHz, CDCl₃, 25°C): δ 171.4, 169.9, 138.4, 115.7, 109.3, 79.0, 75.1, 66.8, 58.5, 54.8. HRMS (ESI-TOF): calculated for C₃₂H₃₂N₂O₆S₂ [M+H]⁺ 604.1696, found 604.1738.



Compound **S4**: Squaraine dye **3** (30 mg, 0.051 mmol) and triethyleneglycol azide (48 mg, 0.27 mmol) were dissolved in chloroform (5 mL) and placed in a small vial. Copper catalyst (6.4 mg, 20 mol%), and *N*,*N*-diisopropylethylamine (43 μ L, 0.26 mmol) were added to the vial and the reaction mixture was stirred at room temperature for 16 hours. The solvent was removed under reduced pressure and the residue was purified *via* column chromatography using 90:10:0 CHCl₃:MeOH:NH₄OH to 80:19:1 CHCl₃:MeOH:NH₄OH as eluent to yield **S4** (53 mg, 0.041 mmol, 83%), as a dark blue, sticky solid. ¹H NMR (500 MHz, CDCl₃, 25°C): δ 7.93 (d, *J* = 4.7 Hz, 2H), 7.77 (s, 4H), 6.30 (d, *J* = 5.0 Hz, 2H), 4.62 (s, 8H), 4.53 (t, *J* = 5.0 Hz, 8H), 3.85 (t, *J* =

5.1 Hz, 8H), 3.78 (t, J = 4.9 Hz, 8H), 3.70-3.76 (m, 16H), 3.58-3.63 (m, 16H), 3.56 (t, J = 4.7 Hz, 8H), 1.80-2.14 (br s, 4H). ¹³C NMR: Small quantity of material prevented ¹³C analysis. HRMS (ESI-TOF): calculated for C₅₆H₈₅N₁₄O₁₈S₂ [M+H]⁺ 1305.5602, found 1305.5625. LC-MS (ESI-TOF): calculated for C₅₆H₈₄N₁₄NaO₁₈S₂ [M+Na]⁺ 1327.5422, found 1327.5397, two observed peaks are *cis/trans* rotamers (identical mass).





3.83–3.75 (m, 4H), 3.69–3.61 (m, 16H), 3.58 (t, J = 5.3 Hz, 2H), 3.55 (td, J = 5.4, 4.2 Hz, 4H). ¹³C NMR (126 MHz, CDCl₃, 25°C): δ 165.2, 133.1, 72.8, 70.5, 70.5, 69.7, 61.7, 39.7. HRMS (ESI-TOF): calculated for C₁₆H₃₀N₂NaO₈ [M+Na]⁺ 401.1894, found 401.1927.



1,5-BisPEGylated Naphthalene (**N1**): 1,5-Bishydroxynaphthalene (36.8 mg, 0.23 mmol) was added to a degassed mixture of mPEG_{2k}-OTs (1.0 g, 0.46 mmol), K₂CO₃, (0.127 g, 129.2 mmol), and a catalytic amount of LiBr in dry MeCN (20 mL). The reaction mixture was heated under reflux for 3 days and filtered after cooling. The residue was dissolved in water (40 mL), and the resulting solution was extracted with CH₂Cl₂ (2 x 10 mL). The organic layers were combined and evaporated. The resulting brown solid was dissolved in CH₂Cl₂ (15 mL), and the solution was washed carefully with a mixture of brine and 10% aq. NaOH (3:1,3 x 20mL). The organic layer was dried over MgSO₄, and the solvent removed by rotary evaporation. The resulted solid was dissolved in H₂O and dialyzed (MWCO: 3500-5000) for 3 days. The water was removed by lyophilization to afford 0.44 g (47% yield) of **N1** as a brown powder. ¹H NMR (500 MHz, CDCl₃) δ 7.87 – 7.82 (m, 2H), 7.34 (dd, J = 8.5, 7.6 Hz, 2H), 6.84 (dd, J = 7.7, 0.9 Hz, 2H), 4.32 – 4.17 (m, 4H), 3.99 (dd, J = 5.5, 4.4 Hz, 4H), 3.82 – 3.46 (m, 458H), 3.38 (s, 6H). ¹³C NMR: Small quantity of material prevented ¹³C analysis. MS (MALDI-TOF): The compound is a distribution of molecular weights, due to polydispersion of the PEG lengths, centered around 3778.20 [M+Na⁺].



Isomerism of Encapsulated Squaraine Dye

To elucidate the source of the isomerism of the encapsulated squaraine dye (see section C below), the rotaxane structure **SR1** was prepared. The structure has a dye with symmetric disubstituted amino groups on the thiophene units. The NMR spectrum of **SR1** shows two sets of peaks for the encapsulated dye demonstrating that the isomerism is due to *cis* or *trans* orientation of the thiophene units and not due to hindered C-N bond rotation. It is worth noting that the *cis* or *trans* dye conformers are only observed as separate NMR signals when the dye is encapsulated by the macrocycle, presumably due to sterically restricted C-C bond rotation.



Compound SR1: Squaraine dye 3 (31 mg, 0.052 mmol) was dissolved in anhydrous CHCl₃ (100 mL). Solutions of 5-tert-butylisophthalic acid chloride (67 mg, 0.26 mmol) and a mixture of 9,10anthracenemethylenediamine (59 mg, 0.25 mmol) and N,N-diisopropylethylamine (70 µL, 0.40 mmol) dissolved in anhydrous CHCl₃ (40 mL for each solution) were drawn into separate syringes. The components were added dropwise over a period of 6 hours by syringe pump to the solution containing squaraine dye 3. At this time, the solvent was removed under reduced pressure and the residue was purified via column chromatography using 2-5% acetone in chloroform as eluent to yield **SR1** (32 mg, 0.022 mmol, 43%) as a dark green solid. ¹H NMR (600 MHz, CDCl₃, 25°C); δ 9.12-9.38 (m, 2H), 8.50-8.55 (m, 4H), 7.97-8.13 (m, 4H), 7.66-7.90 (m, 8H), 6.79-7.00 (m, 8H), 6.37-6.59 (m, 2H), 5.68-5.75 (m, 2H), 5.07-5.36 (m, 8H), 4.20-4.30 (m, 8H), 3.45-3.70 (m, 16H), 2.55-2.60 (m, 4H), 1.48-1.59 (m, 18H). ¹³C NMR (150 MHz, CDCl₃, 25°C): δ 179.8, 178.8, 178.2, 169.8, 169.7, 166.8, 166.3, 166.0, 163.6, 163.5, 152.9, 152.5, 152.4, 139.3, 138.1, 133.0, 132.7, 132.6, 130.5, 130.4, 130.3, 130.2, 129.2, 129.1, 129.0, 128.7, 126.6, 126.1, 125.7, 125.4, 124.9, 124.4, 124.4, 124.0, 123.8, 123.2, 123.2, 122.0, 113.3, 112.5, 109.2, 109.1, 79.0, 79.0, 75.5, 75.5, 69.5, 66.8, 66.7, 58.7, 58.7, 54.6, 53.7, 37.7, 37.3, 35.3, 35.3, 31.5, 31.4, 29.3 [60 out of a theoretical 176 signals (two rotamers)]. HRMS (ESI-TOF): calculated for $C_{88}H_{85}N_6O_{10}S_2 [M+H]^+$ 1449.5763, found 1449.5766.

B. Photophysical Properties

All measurements were made at 25 °C, unless otherwise noted. Quantum yield measurements were performed in triplicate. An absorbance of 0.10 (3.0 μ M of dye) and bis[4-(dimethylamino)phenyl]squaraine ($\Phi_f = 0.70$ in chloroform) and methylene blue ($\Phi_f = 0.02$ in H₂O) were used as standards for quantum yield measurements.^{S7}



Figure S1: Absorption and emission maxima of a) squaraine dye **S1**; b) complex **M1**₂**S1**; c) squaraine dye **S2**; d) complex **M1**₂**S2**; e) squaraine dye **S3**; f) complex **M1**₂**S3**; g) squaraine rotaxane **SR1**; and h) squaraine dye **S4** in CHCl₃ (3.0 μ M).

Dye	λ _{abs} (nm)	log ε	λ _{em} (nm)	Φ_{f}
S1	660	5.51	678	0.15
M1>S1	678	5.14	715	0.34
S2	662	5.42	678	0.11
M1>S2	678	5.15	719	0.34
S3	662	5.36	679	0.09
M1>S3	677	5.13	718	0.26
S4	659	5.46	714	0.12
SR1	680	5.11	720	0.29

Table S1: Photophysical data for squaraine dyes in CHCl₃ (3.0 μ M).



Figure S2: Absorption and emission maxima of a) squaraine dye **S2**; b) complex **M2** $_{2}$ **S2**; c) squaraine dye **S3**; and d) complex **M2** $_{2}$ **S3** in H₂O (3.0 μ M).

Compound	λ _{abs} (nm)	λ_{em} (nm)	Φ_{f}
S2	660	678	0.04
M2>S2	676	715	0.26
S3	663	679	0.03
M25S3	678	714	0.27

Table S2: Photophysical data for squaraine dyes in H_2O (3.0 μ M).

C. Dye Isomerism and Computational Structure of Threaded Complex

The squaraine dyes exist as conformational isomers, based on the relative orientation of the thiophene units (*cis* or *trans*). When free in solution, the squaraine isomerism occurs rapidly on the NMR timescale at room temperature. Upon complexation within the macrocycle, the barrier to C-C bond rotation that exchanges the squaraine conformations is slowed and both conformations are observed in the ¹H NMR spectrum, with the *trans* isomer preferred.



Figure S3: Cis/trans conformational isomers for thiophene-containing squaraine dyes.

A computational structure of the **M1**2**S1** analogue complex (the macrocycle lacks *tert*-Butyl groups) was optimized using the BLYP-gCP-D3 method in combination with def2-TZVP basis set and def2-TZVP/J auxiliary basis sets.^{S12} The calculation was executed using the ORCA3.0.3 program. The structure is very similar to analogous aniline-containing squaraine rotaxane crystal structures, and shows bifurcated hydrogen bonds between the macrocycle amide NH residues and the squaraine oxygens. Furthermore, the macrocycle adopts a macrocyclic chair conformation with the two anthracene units stacked against the encapsulated squaraine dye.



Figure S4: Computational structure (BLYP-gCP-D3/def2-TZVP) of **M151** analogue (most hydrogens are hidden) highlighting the hydrogen bonding (*left*) and aromatic stacking (*right*).

D. Evidence for Host-Guest Complexation

¹H NMR Studies of Host-Guest Complexation

For complex formation studies, vials containing guest and macrocycle were dissolved in the solvent indicated. The samples were allowed to equilibrate for 10 minutes at room temperature, then ¹H NMR spectra of the complexes were acquired.



Figure *S5*: Partial ¹H NMR (600 MHz, CDCl₃, 25 °C) of a) **S1** (2.0 mM); b) **M1** (2.5 mM); and c) **M1** S1 (2.0 mM). Red and blue lines indicate major changes in chemical shift upon complexation. *indicates signals corresponding to the minor **M1** S1 complex with **S1** encapsulated as *cis* conformer (*trans:cis* ratio for encapsulated **S1** ~1.5:1). *indicates signals corresponding to a very slight excess of **M1**.



Figure S6: Partial ¹H-NMR (600 MHz, CD₃OD, 25 °C) of a) **S1** (2.0 mM); b) **M2** (2.0 mM); and c) **M2**₂**S1** (2.0 mM). Red and blue lines indicate major changes in chemical shift upon complexation. *indicates signals corresponding to the minor **M2**₂**S1** complex with **S1** encapsulated as *cis* conformer (*trans:cis* ratio for encapsulated **S1** ~3:1).



Figure S7: Partial ¹H NMR (500 MHz, D₂O, 25°C) of a) **F1** (2.0 mM); b) **M2**5**F1** (2.0 mM); and c) **M2** (2.0 mM). Red and blue lines indicate major changes in chemical shifts upon complexation.



Figure *S8*: Partial ¹H NMR (500 MHz, D₂O, 25°C) of a) **N1** (2.0 mM); b) **M2**₂**N1** (2.0 mM); and c) **M2** (2.0 mM). Red and blue lines indicate major changes in chemical shifts upon complexation.

Fluorescence Studies of Host-Guest Complexation and Job Plot

A solution of squaraine dye **S3** (3.0 μ M) in water was placed in a cuvette and allowed to equilibrate for 20 minutes. A 1.0 mM stock solution of **M2** was prepared and aliquots of this stock solution were titrated into the cuvette containing **S3**; the sample was excited at 690 nm and the emission spectra was collected after each addition.



Figure S9: Fluorescence emission of **S3** (3.0 μ M) upon addition of **M2** (up to 2 mol. equiv.) showing dramatic turn-on effect (25 °C, H₂O, Excitation = 690 nm).

Solutions containing **S3** + **M2** in H₂O at 25 °C were prepared from 250 μ M stock solutions and placed in a 3.0 mL cuvette. In all cases, the total concentration [**S3**+**M2**] was kept constant at 2.0 μ M, while the mole fraction of dye was increased in increments of 0.1 from 0.0 to 1.0. The solutions were excited at 710 nm and the emission at 725 nm was recorded. A plot of the fluorescence intensity at 725 nm versus the mole fraction yielded the Job Plot, confirming 1:1 binding between **S3** and **M2**.



Figure *S10*: Job Plot indicating 1:1 binding between **S3** and **M2** ([**S3+M2**] = 2.0 μ M, H₂O, 25 °C, Excitation = 710 nm, Emission = 725 nm).

Squaraine Stability Studies

Vials containing **S3** (5.0 μ M) in 100% H₂O showed <5.0% decrease in absorption of maxima band after 24 hours. But absorption for a solution of **S3** (5.0 μ M) in the presence of strong nucleophile Na₂S (10 mM) decreased by 55% after 20 minutes (curve B). In contrast, there was essentially no change in absorbance when Na₂S was added to a sample of **S3** that had been premixed with **M2** (forming **M2**₂**S3**) (curve A).



Figure *S11*: Change in absorbance of maxima band for solutions of A) complex **M2**₂**S3** (5.0 μ M **S3**, 7.5 μ M **M2**) or B) squaraine dye **S3**, in the presence of external nucleophile Na₂S (10 mM) in water at 25°C.

TLC Studies of Squaraine/Macrocycle Complexation

Separate solutions of S3, S3+M2, S4, and S4+M2 (100 μ M each) in water were spotted on a silica gel plate and allowed to dry. The plate was eluted with methanol:chloroform (1:9) as the mobile phase. The plate was removed, allowed to dry, and observed under laboratory light and also under UV light. The presence of M2 completely prevented elution of S3 up the plate, indicating formation of a very polar M2>S3 complex. In contrast, the presence of M2 had no effect on the retention factor for control dye S4, indicating that a threaded complex was not formed.



Figure S12: TLC plates after elution with methanol:chloroform (1:9) as the mobile phase.

E. Determination of Association Constants (K_a)

Algorithms for determining association constants^{S8-S11}

For 1:1 Host-Guest System:

 $H + G \xleftarrow{K_a} HG$

The desired association constant is expressed in equation (1.1)

$$K_a = \frac{[HG]}{[H][G]}$$
 (1.1)

Assigning the total concentration of host and guest as $[H]_0$ and $[G]_0$, respectively, gives mass balance equations (1.2) and (1.3)

$$[H]_0 = [H] + [HG](1.2)$$

 $[G]_0 = [G] + [HG] (1.3)$

Equation (1.2) is rearranged to define [H] and equation (1.3) is rearranged to define [G]. The newly defined [H] and [G] are used to replace [H] and [G] in equation (1.1) and the resulting equation is rearranged to yield equation (1.4)

$$[HG]^{2} - ([H]_{0} + [G]_{0} + 1/K_{a})[HG] + [H]_{0}[G]_{0} = 0 (1.4)$$

The real root of equation (1.4) is expressed in equation (1.5), which defines [HG] based on K_a and experimentally determined values ([H]₀ and [G]₀)

$$[HG] = 0.5 \left\{ \left([H]_0 + [G]_0 + 1/K_a \right) - \sqrt{\left([H]_0 + [G]_0 + 1/K_a \right)^2 - 4[H]_0[G]_0} \right\} (1.5)$$

UV/Vis titration method

All absorbance measurements were made in triplicate on a Perkin-Elmer Lambda 25 spectrometer.

.According to Beer's law:

$$\begin{split} &A = \varepsilon_{_{H}} b[H] + \varepsilon_{_{G}} b[G] + \varepsilon_{_{HG}} b[HG](2.1) \\ &\text{If at the selected wavelength, } \varepsilon_{\text{H}} = 0, \text{ then equation (2.1) can be simplified as equation (2.2)} \\ &A = \varepsilon_{_{G}} b[G] + \varepsilon_{_{HG}} b[HG](2.2) \\ &\text{According to equation (1.3), [G] can be defined by [G]_{0} and [HG], \text{ thus (2.2) gives equation (2.3)} \\ &A = \varepsilon_{_{G}} b[G]_{_{0}} + (\varepsilon_{_{HG}} - \varepsilon_{_{G}}) b[HG](2.3) \\ &\text{At the beginning of the titration,} \\ &A_{_{0}} = \varepsilon_{_{G}} b[G]_{_{0}} (2.4) \\ &\text{Then} \\ &\Delta A = A - A_{_{0}} = (\varepsilon_{_{HG}} - \varepsilon_{_{G}}) b[HG] = \varepsilon b[HG](2.5) \end{split}$$

Here $\varepsilon = \varepsilon_{HG} - \varepsilon_G$ From Equation (1.5) and (2.5), we get $\Delta A = f([H]_0, [G]_0, \varepsilon, K_a)$ (2.6) Here we established the relationship between absorption data ΔA and the concentration [H]₀, [G]₀, the absorption coefficients ε_{HG} and ε_{G} , and the association constant K_a . In a typical titration experiment, [G]₀ is kept constant and [H]₀ incrementally increased. OriginLabTM (version 8.6) software was used for nonlinear fitting. In our analysis, we treated either parameter G₀ or H₀ as slightly adjustable to reach the best fitting, since real molar concentration is inherently dependent on a variety of experimental conditions (polydispersity of polymer modified dye, balance error, presence of water as impurity both in our macrocycle and TEG or PEG modified molecule, photobleaching of squaraine dye). Varying this parameter, and related values in later analyses, greatly improves the accuracy of the fitting result. We define host guest ratio:

$$\frac{[H]_0}{[G]_0} = r \ (2.7)$$

We further introduce concentration correction coefficient m into equation (2.7)

$$\frac{[H]_0}{m[G]_0} = r \ (2.8)$$

From equation (2.6) and (2.8) we get

$$\Delta A = f([G]_0, r, m, \varepsilon, K_a) (2.9)$$

Equation (2.9) was used for the nonlinear fitting. To better initialize the parameters, we bounded the fitted parameter within reasonable scope (0.5 < m < 1.5, $0 < K_a < 1000/[G]_0$). The resulting m value was used to correct the real concentration ratio.

Fluorescence titration method

All fluorescence measurements were made in triplicate on a Horiba-Yovan Fluoromax 4 spectrometer with a slit width of 2.0 nm.

 $[G]_t$ was kept constant and $[H]_t$ incrementally increased. The fluorescence signal was related to the concentration of [G] and [HG], assuming that the host has negligible fluorescence at the wavelength studied. The fluorescence intensity can be described by equation (3.1)

$$F = k_1[G] + k_2[HG]$$
 (3.1)

Combining equations (1.3) and (3.1) gives equation (3.2)

$$F = (k_2 - k_1)[HG] + k_1[G]_0 (3.2)$$

When [H]_t=0,

 $F_0 = k_1[G]_0$ (3.3)

$$F/F_0 = 1 + \frac{(k_2 - k_1)[HG]}{k_1[G]_0} = 1 + \frac{k[HG]}{[G]_0}$$
(3.4)

Here $k=(k_2-k_1)/k_1$

Combining equations (3.4), (2.8) (1.5), we relate the fluorescence signal to concentration by $F/F_0 = f([G]_0, r, m, k, K_a)$ (3.5)

Equation (3.5) is used in OriginLabTM (version 8.6) software for nonlinear fitting to obtain the association constant K_a .

Determination of Squaraine Association Constants in Chloroform

Solutions of dye (3.0 μ M) with varying equivalents of macrocycle were equilibrated in chloroform in the dark for 24 hours at 20 °C, at which time the samples were placed in cuvettes and the absorbance at 660 nm was recorded. A plot of the change in absorbance versus the concentration of macrocycle was fitted to the 1:1 binding model expressed above:



Figure S13: Determination of K_a for the association of squaraine S1 with macrocycle M1 in CHCl₃.



Figure S14: Determination of K_a for the association of squaraine S3 with macrocycle M1 in CHCl₃.

Determination of Squaraine Association Constants in Methanol

Solutions of dye (3.0 μ M) with varying equivalents of macrocycle were equilibrated in methanol in the dark for 2 hours at 20 °C, at which time the samples were placed in cuvettes and the absorbance at 660 nm was recorded. A plot of the change in absorbance versus the H/G ratio was fitted to the 1:1 binding model expressed above:



Figure S15: Determination of K_a for the association of squaraine S1 with macrocycle M2 in MeOH.



Figure S16: Determination of K_a for the association of squaraine S3 with macrocycle M2 in MeOH.

Determination of Squaraine Association Constants in Water

Solutions of dye (30 nM) with varying equivalents of macrocycle were equilibrated in H_2O for 5 minutes at 20 °C, at which time the samples were placed in cuvettes and the dye fluorescence intensity at 720 nm was recorded (Excitation = 710 nm, slit width 2 nm). A plot of the fluorescence (F/F_o) versus the H/G ratio was fitted to the 1:1 binding model expressed above:



Figure S17: Determination of K_a for the association of squaraine **S2** with macrocycle **M2** in H₂O.



Figure S18: Determination of K_a for the association of squaraine S3 with macrocycle M2 in H₂O.

Fluorescence Studies of Fumaride (F1) Complexation

Macrocycle fluorescence is quenched upon association with F1.





Determination of fumaride association constant by fluorescence titration.

In the fluorescence quenching titration experiment, $[H]_0$ was kept constant and $[G]_0$ incrementally increased. The macrocycle and complex were treated as fluorescent while the fumaride guest was assumed to be transparent at the selected excitation wavelength. The following equation was used for the nonlinear fitting.

$$[HG] = 0.5 \left\{ \left([H]_0 + [G]_0 + 1/K_a \right) - \sqrt{\left([H]_0 + [G]_0 + 1/K_a \right)^2 - 4[H]_0[G]_0} \right\}$$

$F/F_{0} = 1 - k \times [HG]/[H]_{0}$

In the above equation, $[H]_0$ is the total concentration of macrocycle used for the titration, $[G]_0$ is the total concentration of fumaride guest used for the titration, [HG] is the concentration of complex in the titration, K_a is the association constant and k is the emission coefficient. The OriginLabTM 8.6 software was used for the nonlinear fitting.



Figure S20: Determination of K_a for the association of fumaride F1 with macrocycle M2 in H₂O at 20 °C.





Figure S21: Partial ¹H NMR (600 MHz, D_2O , 25°C) of (*top*) M25F1 (2.0 mM); (*middle*) M25F1 (2.0 mM) treated with S3 (2.0 mM) which displaced F1; (*bottom*) fumaride F1 (2.0 mM). Red line indicates change in chemical shift for M2 peak B upon displacement of F1 by S3.

Determination of Squaraine Association Constant by Competitive Titration method

In the competitive titration experiment, S2 or S3 was titrated into a mixture of F1 + M2 (F1:M2 molar ratio was 100:1). The squaraine/macrocycle complex was treated as fluorescent and the macrocycle and squaraine dye as transparent at the selected excitation wavelength. The association constant for M2₂F1 was predetermined by the fluorescence quenching titration above to be 1.6×10^4 M⁻¹. Newton's method was used to iterate the concentration of macrocycle ([H]) during the displacement titration.^{S9}

The following equation was used for the nonlinear fitting:

 $F = k \times K_a \times [H] \times [G]_0 / (1 + K_a * [H])$

In the above equation, $[G]_0$ is the total concentration of **S2** or **S3** used for the titration, [H] is the concentration of macrocycle in the titration, K_a is the association constant between **S2** or **S3** and **M2** and k is the emission coefficient. The nonlinear fitting was conducted using OriginLabTM (version 8.6) software.



Figure S22: Determination of K_a by the competitive displacement of F1 from M2₂F1 by addition of squaraine S2 to a mixture of M2+ F1 (F1:M2 molar ratio was 100:1) in H₂O at 20 °C.



Figure S23: Determination of K_a by the competitive displacement of F1 from M2₂F1 by addition of squaraine S3 to a mixture of M2+ F1 (F1:M2 molar ratio was 100:1) in H₂O at 20 °C.

F. Thermodynamic Measurements

ITC Titration of M2 and F1

An aqueous solution of **M2** (300 μ L, 0.1-0.5 mM) was injected into the sample cell of a Malvern Microcal iTC200 Microcalorimeter. An aqueous solution of **F1** (30-35 μ L, 2-10 mM) was loaded into the titration syringe and 20-25 injections (1 μ L each) were added stepwise into the sample cell at 27 °C. The heat generated was monitored over time. To adjust for the dilution effect, the **F1** solution was titrated into the pure H₂O and heat generated by this process was also monitored over time. The net heat was obtained by subtracting this dilution value from the heat of complexation. The resulting data was analyzed using the OriginTM program to fit a 1:1 binding model and values for K_a , Δ H, and Δ S were determined directly.



Figure S24: ITC titration of F1 into an aqueous solution containing M2 at 27 °C.

ITC Titration of M2 and S3

The association constant for **M2** and **S3** was too high for accurate measurement using our microcalorimeter (C>1000) but the step-like isotherm did confirm that N=1. A single injection method was used to measure the enthalpy of complexation. A single aliquot of **S3** (5 μ L of 2.0 mM stock solution) was injected into the cell containing **M2** (50 μ M, 200 μ L) and the generated heat was monitored over time. To adjust for the dilution effect, an identical aliquot of **S3** solution was added to pure H₂O and heat generated was monitored over time. The net heat effect was obtained by appropriate subtraction.



Figure S25: ITC single injection of S3 into H_2O or an aqueous solution containing M2 (50 μ M, 27 °C) enabled determination of the enthalpy of complexation.

The association constant at 27 °C was determined by fluorescence titration. Separate aliquots of **S3** (300 nM) and varying molar equivalents of **M2** were equilibrated in H₂O for 5 minutes and the fluorescence intensity at 720 nm was recorded (ex: 710 nm) in temperature controlled cuvettes. The isotherm was fitted to the 1:1 binding model expressed above:



Figure S26: Determination of K_a for the association of S3 with M2 in H₂O at 27 °C.

A competitive ITC titration experiment that added **S3** to a solution of **M2** $_{2}$ **F1** produced very little heat and the titration isotherm could not be fitted reliably to a binding model. The small change in heat for this displacement equilibrium is consistent with the similar values of Δ H for each guest (**S3** or **F1**) binding to **M2**.

ITC Titration of M2 and N1

An aqueous solution of **M2** (300 µL, 0.5 mM) was injected into the sample cell of a Malvern Microcal iTC200 Microcalorimeter. An aqueous solution of **N1** (30-35 µL, 2-10 mM) was loaded into the titration syringe and 20-25 injections (1 µL each) were added stepwise into the sample cell at 27 °C. The heat generated was monitored over time. To adjust for the dilution effect, the **N1** solution was titrated into the pure H₂O and heat generated by this process was also monitored over time. The net heat was obtained by subtracting this dilution value from the heat of complexation. The resulting data was analyzed using the OriginTM program to fit a 1:1 binding model and values for K_a , Δ H, and Δ S were determined directly.



Figure S27: ITC titration of N1 into an aqueous solution containing M2 in H₂O at 27 °C.

G. Kinetic Studies

Determination of Association Rate Constant (kon) by Absorption Method

Molar absorptivities at 690 nm were determined from absorbance of the dye before association $(\mathcal{E}_{\lambda \text{Dye}})$ and after association with macrocycle $(\mathcal{E}_{\lambda \text{Complex}})$.

 $\begin{array}{l} \epsilon_{\lambda \text{S1}} \ (690 \ nm, \ CHCl_3): \ 6.02 \times 10^4 \\ \epsilon_{\lambda \text{M1} \text{JS1}} \ (690 \ nm, \ CHCl_3): \ 1.41 \times 10^5 \\ \epsilon_{\lambda \text{S3}} \ (690 \ nm, \ CHCl_3): \ 1.66 \times 10^4 \\ \epsilon_{\lambda \text{M2} \text{JS3}} \ (690 \ nm, \ CHCl_3): \ 1.17 \times 10^5 \end{array}$

When [D] = 5.0 μ M; [M] = 15.0 μ M (CHCl₃ study)

Because the equilibrium constant is high ($K_a > 10^5 \text{ M}^{-1}$) the formation process is treated as an irreversible second-order reaction that obeys the following relationship:

$$\ln\left(\frac{[\mathbf{M}] \cdot [\mathbf{D}]_0}{[\mathbf{D}] \cdot [\mathbf{M}]_0}\right) = ([\mathbf{M}]_0 - [\mathbf{D}]_0) \cdot k\mathbf{t}$$

Rearrangement leads to:

$$\frac{\ln\left(\frac{[M]\cdot[D]_0}{[D]\cdot[M]_0}\right)}{([M]_0 - [D]_0)} = kt$$

The [M] is given by:

$$[M] = 3[D]_0 - [R]$$

Since b = 1 cm, the [R] at a specific wavelength λ is given by the modified Beer's Law:

$$[R] = \frac{A_{\lambda} - \mathcal{E}_{\lambda D}[D]_{0}}{\mathcal{E}_{\lambda R} - \mathcal{E}_{\lambda D}}$$

Substitution into the equation gives:

$$\frac{\ln\left(\frac{(3[D]_0 - [R]) \cdot [D]_0}{([D]_0 - [R]) \cdot [M]_0}\right)}{([M]_0 - [D]_0)} = kt$$

The plot of the ratio versus time yields a linear relationship whose slope = k_{on} .



Figure S28: Second-order kinetic plot for the rate of association between S1 and M1 in $CHCI_3$ at 20 °C. N = 3.



Figure S29: Second-order kinetic plot for the rate of association between S3 and M1 in $CHCI_3$ at 20 °C. N = 3.

Determination of Association Rate Constant (kon) by Stopped-flow Method

All stopped-flow experiments were run in triplicate at 20 °C and averaged to obtain kinetic curves which were fitted according to the following method:

Algorithms for the determination of k_{on}

In a second order reaction, we have

$$H + G \xleftarrow{k_{on}}{k_{off}} HG$$

The reaction rate can be written as

$$Rate = -\frac{d[G]}{[G]} = k_{on}[H][G] - k_{off}[HG] = k_{on}[H][G] - \frac{k_{on}}{K_{a}}[HG](4.1)$$

For strong association system where $K_a > 100 \text{ M}^{-1}$, the contribution of reverse reaction on reaction rate is negligible. So equation (4.1) can be simplified as (4.2)

$$Rate = -\frac{d[G]}{[G]} = k_{on}[H][G](4.2)$$

In the case $[H]_0=[G]_0$, since H and G react with a 1:1 stoichiometry, $[H]=[H]_0-x$ and $[G]=[G]_0-x$. At any time t, [H]=[G] and equation (4.2) can be further simplified as (4.3)

$$Rate = -\frac{d[G]}{[G]} = k_{on}[G]^2 (4.3)$$

Solving the differential rate equation (4.3) gives the integrated rate equation (4.4)

$$\frac{1}{[G]_{t}} - \frac{1}{[G]_{0}} = k_{on} t (4.4)$$

Measuring *k*on by fluorescence

Equimolar solutions of the host and guest were mixed, and the fluorescence signal was related to the concentration of [G] and [HG], assuming that the host has negligible fluorescence at the wavelength studied. The fluorescence intensity can be described by equation (5.1)

$$F = k_1[G] + k_2[HG]$$
 (5.1)

When t=0,

$$F_0 = k_1[G]_0$$
 (5.2)

Combining equations (1.3), (5.1) and (5.2) gives equation (5.3)

$$F/F_0 = 1 + \frac{(k_2 - k_1)[HG]}{k_1[G]_0}$$
(5.3)

Combining equation (1.3) and equation (4.4) gives equation (5.4)

$$\frac{1}{[G]_0 - [HG]} - \frac{1}{[G]_0} = k_{on} t (5.4)$$

Replacing [HG] in equation (5.4) by equation (5.3) gives the equation (5.5), which describes the relationship between fluorescence and time

$$F/F_0 = 1 + \frac{k_2 - k_1}{k_1} \left(1 - \frac{1}{k_{on} t[G]_0 + 1} \right) (5.5)$$

Equation (5.5) was used in OriginLabTM 8.6 version for the nonlinear fitting. The association rate k_{on} can be determined by using the stopped-flow data and nonlinear regression analysis.

In the case where the guest molecule is also transparent, especially at nanomolar concentration, the fluorescence intensity can be described by equation (5.6)

$$F = k_2[HG]$$
 (5.6)

Replacing [HG] in equation (5.4) by equation (5.6) gives the equation (5.7)

$$F = k_2 G_0 \left(1 - \frac{1}{k_{on} t [G]_0 + 1} \right) (5.7)$$

Equation (5.7) is used in OriginLabTM (version 8.6) for nonlinear fitting to determine the association rate k_{on} .

Determination of *k*on in Methanol

In methanol, $[D]_{\circ} = [M]_{\circ} = 0.5 \ \mu$ M, and the fluorescence response at 720 nm (Excitation = 710 nm) was monitored as a function of time.



Figure S30: Determination of k_{on} for association of squaraine S1 with macrocycle M2 in MeOH at 20 °C.



Figure *S31*: Determination of k_{on} for the association of squaraine **S3** with macrocycle **M2** in MeOH at 20 °C.

Determination of *k*on in Water

In water, $[D]_o = [M]_o = 30$ nM and the fluorescence response at 720 nm (Excitation = 710 nm) was monitored as a function of time.



Figure S32: Determination of k_{on} for association of squaraine S2 with macrocycle M2 in H₂O at 20 °C.



Figure S33: Determination of k_{on} for association of squaraine S3 with macrocycle M2 in H₂O at 20 °C.

Determination of k_{on} in H₂O/Methanol Mixtures



Table S3: Influence of solvent on k_{on} for M25S3 at 20 °C.

Figure S34: Influence of solvent on k_{on} for M25S3 at 20 °C. Also shown is a fit of the data to an exponential equation that reflects the dependence on mole fraction of water.

H. Cell Studies

Cell Toxicity

Cell toxicity was measured using the common MTT cell viability assay. CHO-K1 (Chinese hamster ovary) cells (ATCC CCL-61) were cultured following ATCC protocols and maintained at 37 °C, in a humidified incubator with 5% CO₂ atmosphere. The cells were seeded into 96-microwell plates, and grown to confluence using F-12K medium (supplemented with 10 % fetal bovine serum; 1 % penicillin streptomycin) in a humidified incubator (37 °C; 5 % CO₂). The Vybrant MTT cell proliferation assay kit (Invitrogen, Eugene, USA) was performed according to the manufacturer's protocol. The cells were incubated with different concentrations of each compound in F-12K media for 24 h at 37 °C, 5% CO₂. The medium was removed and replaced with 110 μ L of F-12K medium containing [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] (MTT, 1.2 mM) and the sample incubated for 4 h at 37 °C and 5 % CO₂. An SDS-HCl detergent solution was added and the samples incubated for an additional 12 h. The absorbance of each well was measured at 570 nm (n = 4), and the readings were normalized relative to untreated cells.



Figure S35: MTT assay of toxicity against CHO-K1 cells.

Fluorescence Microscopy

Cultured CHO-K1 cells were plated out on 8-well microscope chambered slides (Nunc) in F-12K medium and allowed to grow to approximately 70% confluence. The medium was replaced with solutions of **S3** (10 μ M) or **S3** + **M2** (10 μ M) in OptiMEM media (Life Technologies, catalogue number 11058-021) and the cells allowed to incubate for 8 hours at 37 ° C. The cells were then washed twice with PBS and imaged under OptiMEM media. Fluorescence microscopy was conducted on a Nikon Eclipse TE-2000U epifluorescence microscope equipped with an X-cite 120 fluorescence illumination system and either: (a) red filter set (Nikon C53375, Exciter: HQ620/60X, Dichroic: 660LP, Emitter: HQ700/75m), (b) near-infrared filter set (Nikon C81153, Exciter: HQ710/75X, Dichroic: 750LP, Emitter: HQ810/90m), (c) blue excitation, red emission filter set (Exciter: D360/40x, Dichroic: 400DCLP, Emitter: 610LP). Images were collected in NIS-Elements using a Hamamatsu ORCA-Flash4.0 V2 Digital CMOS camera with 2x2 binning and 2 second integration times. Images were processed in ImageJ, and a 100 pnt rolling background was subtracted.

Shown in Figure S36 are representative micrographs of living CHO-K1 cells containing either **S3** or **M2**₂**S3** complex and imaged using the blue excitation, red emission filter set (c) described above. The images show that the **S3** dye cannot be imaged using this filter set (although it can be visualized using using the red excitation/emission filter set (a) described above), whereas the cells containing **M2**₂**S3** are easily observed. The Supporting Information contains a movie of live cells <ja5b03573_si_002.avi> imaged for 2 minutes at 100x speed and 90x magnification. The movie shows time resolved tracking of endosomes containing the **M2**₂**S3** complex which is selectively visualized using the energy transfer imaging filter set.



Figure S36. Epifluorescence micrographs of live CHO cells after incubation for 8 hours with S3 (10 μ M) or M2 + S3 (10 μ M each) in culture media. The micrographs were acquired using a blue excitation, red emission filter set (Exciter: D360/40x, Dichroic: 400DCLP, Emitter: 610LP) and the same acquisition settings; thus, the image intensities are directly comparable. The length bar is 10 μ m.

The same type of energy transfer imaging (blue excitation, red emission) was achieved on the mesoscale using a Multispectral FX Pro *in vivo* imaging system equipped with a CCD camera (filter set: 410/10 excitation; 700/17.5 emission). Shown in Figure S37 is an image (30 second exposure) of a microwell plate with separate solutions (200 μ L) of culture media containing; no additives; **M2** (10 μ M); **S3** (10 μ M); or a mixture of **M2+S3** (10 μ M each). The fluorescence intensity of the last well was 20 times brighter than any other due to formation of the **M2>S3** complex which enables energy transfer from the surrounding macrocycle anthracene units to the encapsulated squaraine dye.



Figure S37. Image of a microwell plate using a CCD camera and an illumination filter set that allowed blue excitation (410/10 nm) and captured red emission (700/17.5 nm). Each microwell (4 mm diameter) contained a solution of additive (10 μ M) in culture media (200 μ L), and the image is an overlay of a photograph and a fluorescence intensity pixel map. The fluorescence intensity bar is in arbitrary units.

I.¹H and ¹³C NMR Spectra



Figure S38: ¹H NMR (500 MHz, CDCl₃, 25 °C) spectrum of squaraine dye **S1**.



Figure S39: ¹³C NMR (150 MHz, CDCl₃, 25 °C) spectrum of squaraine dye **S1**.



Figure S40: ¹H NMR (500 MHz, CD₃CN, 25 °C) spectrum of squaraine dye **S2**.



Figure *S41*: ¹H NMR (600 MHz, CDCl₃, 25 °C) spectrum of squaraine dye **S3**.



Figure S42: ¹H NMR (500 MHz, CDCI₃, 25 °C) spectrum of squaraine dye **3**.



Figure S43: ¹³C NMR (150 MHz, CDCl₃, 25 °C) spectrum of squaraine dye **3**.



Figure *S44*: ¹H NMR (600 MHz, CDCl₃, 25 °C) spectrum of squaraine rotaxane **SR1**.



Figure S45: ¹³C NMR (150 MHz, CDCl₃, 25 °C) spectrum of squaraine dye **SR1**.



Figure S46: ¹H NMR (500 MHz, CDCl₃, 25 $^{\circ}$ C) spectrum of squaraine dye **S4**.



Figure S47: ¹H NMR (500 MHz, CDCl₃, 25 °C) spectrum of fumaride **F1**.



Figure S48: ¹³C NMR (150 MHz, CDCl₃, 25 °C) spectrum of fumaride F1.



Figure S49: ¹H NMR (500 MHz, CDCl₃, 25 °C) spectrum of **N1**.



Figure *S50*: Partial COSY spectrum (600 MHz, D₂O, 25°C) of complex **M2**₂**S3**.



Figure *S51*: Partial HSQC spectrum (600 MHz, D₂O, 25 °C) of complex **M2**₂**S3**.



Figure S52: Partial ROESY spectrum (600 MHz, D₂O, 25 °C) of complex M2₂S3.

J. References

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