

*Supporting Information*

Fluorescent calix[4]triazole for selective fluoride anion  
sensing

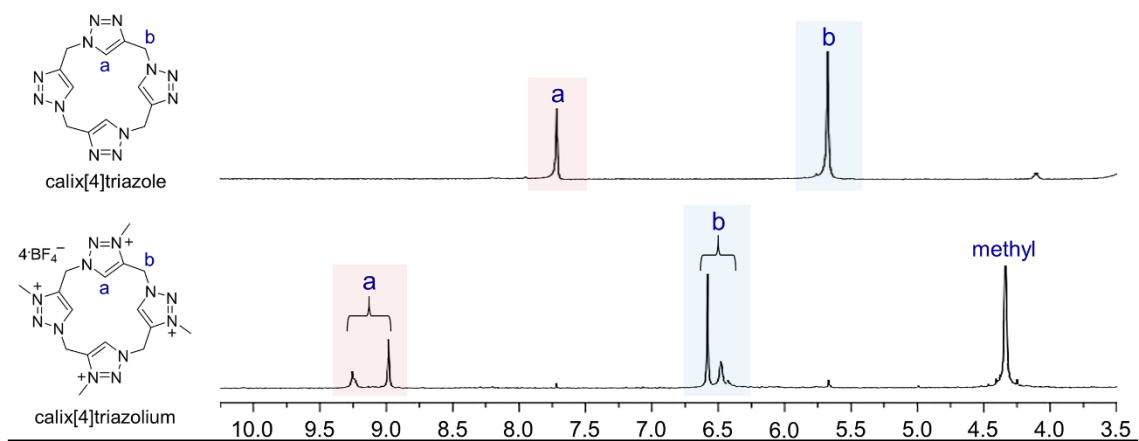
Jihee Cho and Sanghee Kim\*

College of Pharmacy, Seoul National University, Seoul 08826, Republic of Korea

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## 1. $^1\text{H}$ NMR spectrum



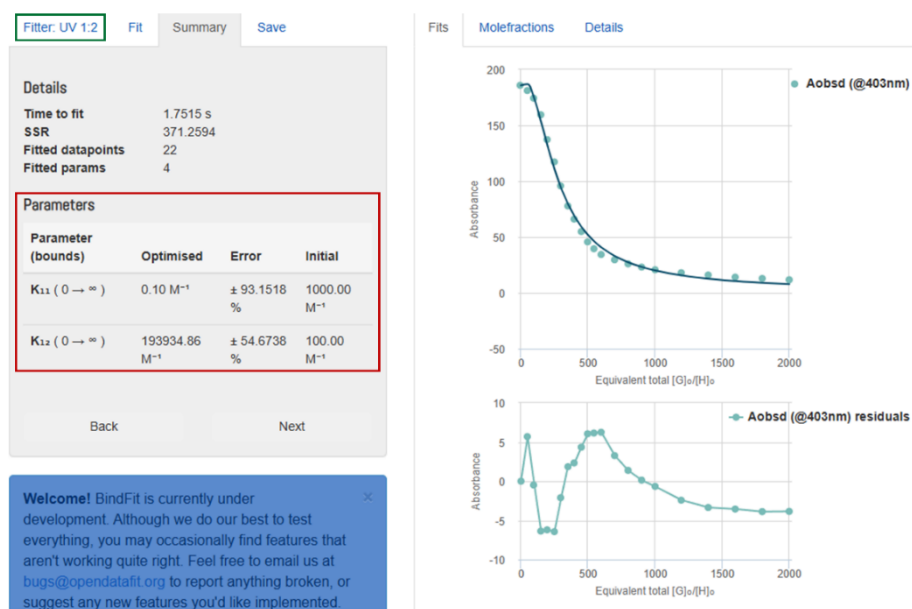
**Fig. S1.**  $^1\text{H}$  NMR spectra of calix[4]triazole and calix[4]triazolium in  $\text{DMSO-}d_6$ .

## 2. Complexation studies between Py-CT4 and F<sup>-</sup>

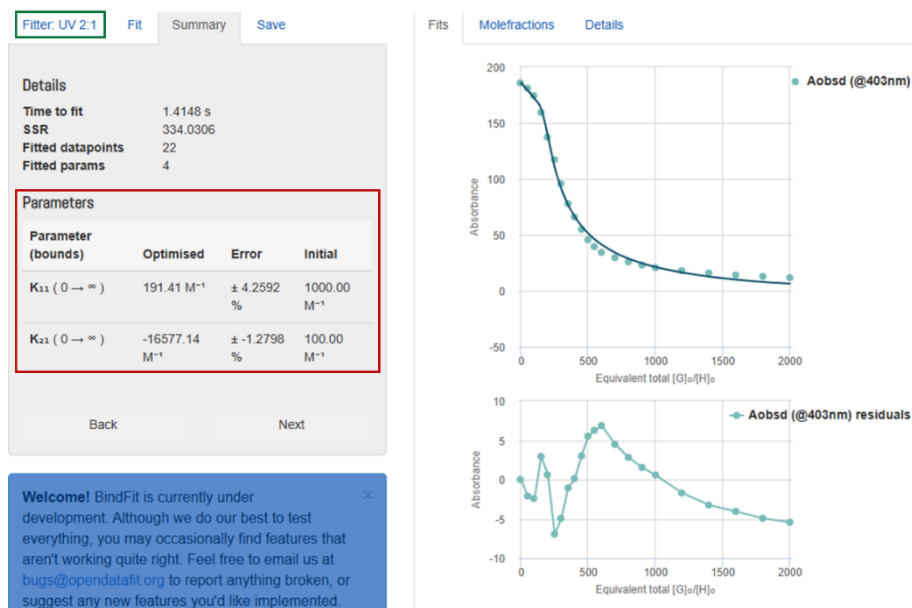
(1) Association constant calculated by bindfit



**Fig. S2.** Screenshot of the summary window of <http://app.supramolecular.org/bindfit/>. This screenshot shows the raw data for the fluorescence titration of **Py-CT4** with F<sup>-</sup> with fluorescence emissions at 403 nm vs. the data fitted to a 1:1 UV binding model and the corresponding residual plot and association constants with the calculated asymptotic standard errors.



**Fig. S3.** Screenshot of the summary window of <http://app.supramolecular.org/bindfit/>. This screenshot shows the raw data for the fluorescence titration of **Py-CT4** with F<sup>-</sup> with fluorescence emissions at 403 nm vs. the data fitted to a 1:2 UV binding model and the corresponding residual plot and association constants with the calculated asymptotic standard errors.



**Fig. S4.** Screenshot of the summary window of <http://app.supramolecular.org/bindfit/>. This screenshot shows the raw data for the fluorescence titration of **Py-CT4** with F<sup>-</sup> with fluorescence emissions at 403 nm vs. the data fitted to a 2:1 UV binding model and the corresponding residual plot and association constants with the calculated asymptotic standard errors.

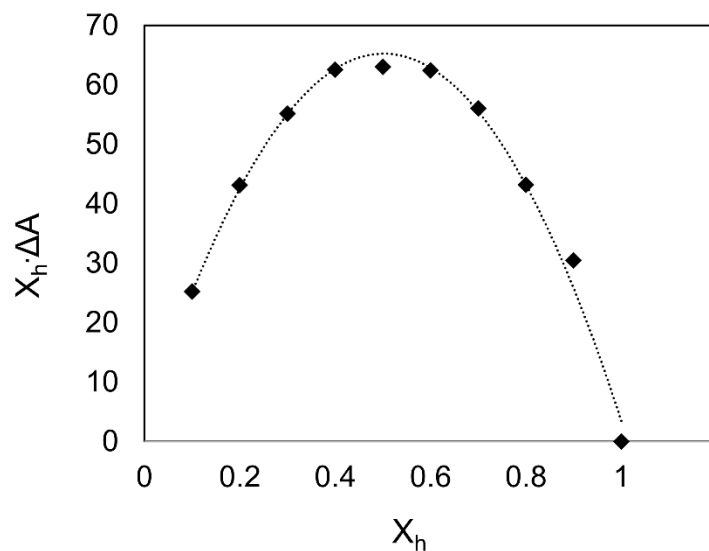
**Table S1.** Summary of association constants between **Py-CT4** and F<sup>-</sup> according to different binding models.<sup>a</sup>

Binding models		
1:1	1:2	2:1
74.30 (±15.50%)	$K_{11}$ 0.10 (±93.15%) $K_{12}$ 193934.86 (±54.67%)	$K_{11}$ 191.41 (±4.26%) $K_{21}$ -16577.14 (±-1.28%)

<sup>a</sup> Bindfit software from *supramolecular.org* was used for data analysis.

(2) Job's plot for determining binding stoichiometric ratio

**Experiment details.** Stock solutions with equal concentrations of **Py-CT4** (10  $\mu\text{M}$ ) and  $\text{F}^-$  (10  $\mu\text{M}$ ) in DMSO were prepared. Ten vials were each filled with a 10 mL solution of **Py-CT4** and  $\text{F}^-$  in the following ratios (**Py-CT4**: $\text{F}^-$ ): 10:0, 9:1, 8:2, 7:3, 6:4, 5:5, 4:6, 3:7, 2:8, 1:9. Job's plot was constructed by plotting the change in the fluorescence at 403 nm of **Py-CT4** against the molar fraction of the host.



**Fig. S5.** Job's plot generated from the fluorescence titration data of **Py-CT4** with  $\text{F}^-$  in a DMSO solution.