Supporting Information

Fluorescence Turn-On Probe for Homocysteine and Cysteine in Water

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Experimental

General Procedures. ¹H and ¹³C NMR spectra were recorded using an Advance 300MHz Bruker spectrometer in chloroform- d_3 , D₂O. ¹H NMR chemical shifts are reported as parts per million (δ) and referenced to residual solvent peak (δ 4.80 for D₂O and 7.28 for CDCl₃). and ¹³C NMR chemical shifts in CDCl₃ were reported relative to CHCl₃ (77.23 ppm). UV visible spectra were recorded on a Beckman DU 800 spectrophotometer. Fluorescence spectra were recorded on a Jasco FP-6500 spectrophotometer. Analytical thin-layer chromatography was performed using Kieselgel 60_F-254 plates from Merck. Column chromatography was carried out on Merck silica gel 60 (70 - 230 mesh). All solvents and reagents were commercially available and used without further purification unless otherwise noted.

Fluorescence titration experiment

Probe 1 was dissolved in DMSO to afford a concentration of 10 mM stock solution, which was diluted with distilled water up to 10 μ M. Guests were dissolved in 10 μ M host (1) solution and used for the fluorescence titration experiment. Fluorescence titration was performed with Jasco FP-6500 spectrofluorometer at rt.

Human blood serum treatment

Venous blood specimens were collected in EDTA-treated tubes after a 12-hour fast. The tubes were immediately covered with aluminum foil and placed on ice until they arrived at the laboratory room (within 1-3 hours) for separating plasma. The tubes centrifuged at for 15minutes at 2-8°C at 1,000×g. The separated plasmas were immediately stored at -70°C until analysis.

Fig S1. ¹H NMR spectra of **1** after addition of Cys. (a) **1** only (b) **1** and 2 equiv. L-cysteine in D_2O with HEPES buffer (pH=7.4). Peaks below 4.79 ppm are from those of HEPES.



Fig S2. ¹H NMR spectra of **1** after addition of Cys. (a) **1** only (b) **1** and 2 equiv. D/L-homocysteine in D₂O with HEPES buffer (pH=7.4). Peaks below 4.79 ppm are from those of HEPES.



Fig S3. UV-vis absorbance and fluorescence spectra of probe 1.



Abs & Flu (Ex.365nm) of Host 1

Fig S4. Fluorescence emission change of probe 1 vs D/L-homocysteine equiv at 500 nm.



Ъ

-50

0

[G]/[H]

1000

1500

500

Fig S5. Fluorescence titration spectra of the probe **1** upon addition of L-cysteine. ([**1**] = 10 μ M, $\lambda_{ex} = 365$ nm, $\lambda_{em} = 450$ nm)



Fig S6. Fluorescence emission change of probe 1 vs L-cysteine equiv at 450 nm.



cysteine Kd = 1.203 x 10 ⁻³ M Fig S7. Homocysteine and cysteine selectivity of 1 in the presence of other amino acids and glutathione. $[1] = 10 \mu M$, Hcy, Cys = 500 equiv in the presence of each amino acids and GSH (500 equiv). Light blue bar upon the addition of each analyte to 1 and brown bar upon the addition of Hcy to a solution of 1 and each analyte.



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Fig S8. (a) Proposed 1-Hcy binding mode, (b) its computational model structure.



⁽a)



(b)

Fig S9. The HOMO and LUMO orbitals of probe **1** and its homocysteine adduct. DFT calculation results based on B3LYP $6-31G^*$ in Spartan 2004, Wavefunction Inc.



LUMO (-2.33 eV)



HOMO (-6.45 eV)



LUMO (-1.5" eV)



НОМО (-5.89 eV) 1∙Нсу

1