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Supporting Information for

Ratiometric Fluorescence Detection of a Tag Fused Protein Using the Dual-Emission Artificial Molecular Probe

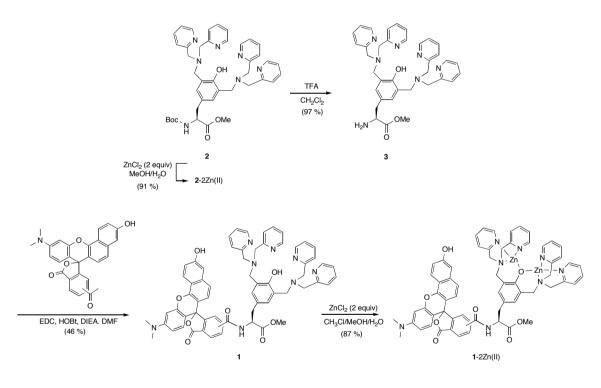
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General methods

Unless otherwise noted, all chemical reagents were purchased from commercial suppliers and used without further purification. ¹H-NMR spectra were recorded using a JNM-EX400 (JEOL, 400 MHz) spectrometer and the chemical shifts (§ppm) are referenced to the respective solvent. FAB mass spectra were recorded using a QP5050A (Shimadzu). Fluorescent spectra and Absorption spectra were recorded on a Perkin-Elmer LS55 spectrometer and U-2550 (Shimadzu), respectively. Reverse phase HPLC was conducted with a Lachrom (Hitachi) with C18 columns.

Synthesis and Compound Characterizations



Scheme S1 Synthesis of 1-2Zn(II).

The details of syntesis of **2**, **2**-2Zn(II) and **3** were described previously.⁸

SNARF-DpaTyr (1) A mixture of **3** (20 mg, 0.033 mmol), carboxy SNARF (mixture of 5- and 6- carboxylate mixture) (15 mg, 0.033 mmol), HOBt \cdot H₂O (7.5 mg, 0.050 mmol), EDC \cdot HCl (9.4 mg, 0.050 mmol) and *N*,*N*'-diisopropylethylamine (30 mL, 0.17 mmol) in dry DMF (2 mL) was stirred at room temperature for 6 h. The mixture

was poured into water and extracted with ethyl acetate. The organic layer was washed with brine and dried over Na_2SO_4 . After removal of the solvent in vacuo, the residue was dissolved in CH₃Cl (containing a small amount of MeOH) and the solution was diluted with Hexane to form precipitate. The precipitate was filtered and washed with Hexane to give 1 (16 mg, 46%) as a purple powder. 1 was used in the next reaction without further purification. Due to assignment difficulty, ¹H NMR of 1 was measured after isolation of each isomer by reverse phase HPLC.

¹H NMR (400 MHz; CD₃OD; solvent)

isomer-1 : $\delta_{\rm H}$ 3.00 – 3.06 (2H, m, CH₂), 3.43 (6H, s, NMe), 3.76 (3H, s, Me), 4.15 – 4.28 (12H, m, CH₂), 6.91 – 6.96 (1H, m, Ar-H), 7.24 – 7.29 (5H, m, Ar-H), 7.37 – 7.44 (12H, m, Ar-H), 7.56 – 7.61 (2H, m, Ar-H), 7.82 (4H, td, J = 8.0, 1.6 Hz, Ar-H), 8.07 - 8.12 (1H, m, Ar-H), 8.57 - 8.58 (5H, m, Ar-H), 8.83 (1H, d, J = 9.2 Hz, Ar-H).

isomer-2 : $\delta_{\rm H}$ 2.82 – 2.97 (2H, m, CH₂), 3.42 (6H, s, NMe), 3.70 (3H, d, J = 4.0, Me), 4.00 – 4.20 (12H, m, CH₂), 6.58 (1H, dd, J = 7.2, 1.2 Hz, Ar-H), 6.87 – 7.00 (3H, m, Ar-H), 7.07 – 7.16 (5H, m, Ar-H), 7.22 – 7.31 (3H, m, Ar-H), 7.36 – 7.44 (10H, m, Ar-H), 7.55 (1H, t, J = 8.8 Hz, Ar-H), 7.66 (1H, dd, J = 14.2, 1.2 Hz, Ar-H), 7.82 (4H, qd, J = 7.8, 1.6 Hz, Ar-H), 7.98 – 8.05 (2H, m, Ar-H), 8.25 - 8.31 (1H, m, Ar-H), 8.53 -8.63 (4H, m, Ar-H), 8.81 (1H, d, J = 8.8 Hz, Ar-H).

FAB-HRMS m/e calcd for $[M + H]^+$ 1053.4299, found 1053.4312.

SNARF-DpaTyr(Zn) (1-2Zn(II)) To a solution of **1** (mixture of the isomers 7.7 mg, 7.3 μ mol) in distilled MeOH / CHCl₃ (1.5 mL / 0.5 mL) was added aqueous solution of ZnCl₂ (100 mM; 136 mL, 0.014 mmol), and the solution was stirred at room temperature for 1 h. The solution was concentrated by evaporation, and then lyophilized under vacuum. The obtained solid was suspended in ethyl acetate, filtered, and dried in vacuo to give 1-2Zn(II) (7.9 mg, 87%) as a purple powder.

FAB-HRMS m/e calcd for [SNARF-DpaTyr + 2Zn + 2Cl⁻]⁺ 1249.2103, found 1249.2106.

Preparation of D4 peptide (Boc-DDDD-NH₂)

The Boc-DDDD-NH₂ was synthesized by the standard Boc chemistry in solution phase from Boc-Asp(Obzl)-OH and H-ASP(Obzl)-CONH₂. Detailed synthetic procedure and compound characterization was described previously.⁸

Preparations of the tag-fused RNases

The tagged RNases were constructed by the self-assembly of a S-protein (purchased from Aldrich) with a S-peptide^{S1} tethering a D4-tag (DDDD) or His₆-tag (HHHHHH) at its N-terminus, which was synthesized by the automated peptide synthesizer and purified by reverse-phase HPLC. Detailed synthetic procedures and compound characterizations were described previously.⁸

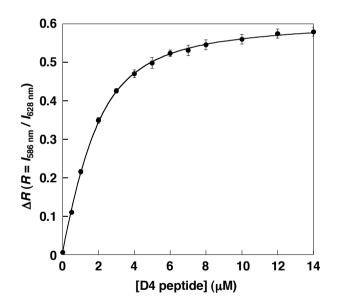


Fig. S1 Fluorescence titration curve of the emission intensity ratio R ($R = I_{586 \text{ nm}} / I_{628}$ nm) of 1-2Zn(II) (2 μ M) upon addition of D4 peptide in 50 mM HEPES buffer, pH 7.2, at 25 °C.

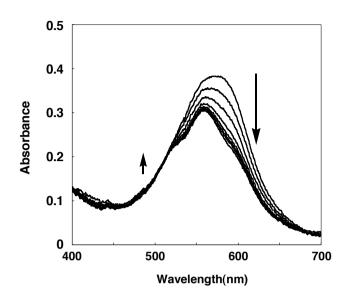


Fig. S2 Absorption spectral change of 1-2Zn(II) (2 μ M) upon the addition of D4 peptide : [D4 peptide] = 0, 0.5, 1, 2, 3, 4, 5, 6, 7, 8, 10, 12 and 14 μ M in 50 mM HEPES buffer, pH 7.2, at 25 °C.

References

S1. (a) P. R. Connelly, R. Varadarajan, J. M. Sturtevant and F. M. Richards, *Biochemistry*, 1990, 29, 6108-6114. (b) I. Hamachi, R. Eboshi, J. Watanabe and S. Shinkai, *J. Am. Chem. Soc.*, 2000, 122, 4530-4531.