

Naphthalimide-tyrosine-based dicationic amphiphile for intracellular 'turn-on' simultaneous detection of ATP and CTP

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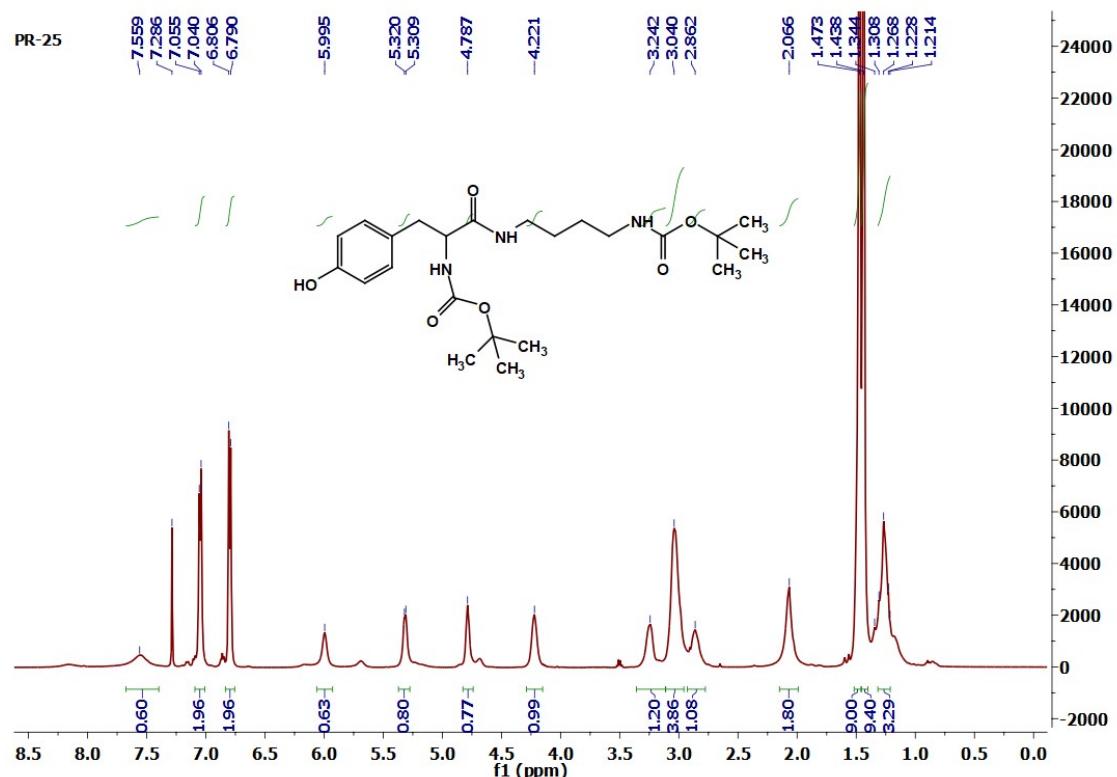


Figure 1a: ¹H NMR Spectrum of compound 2.

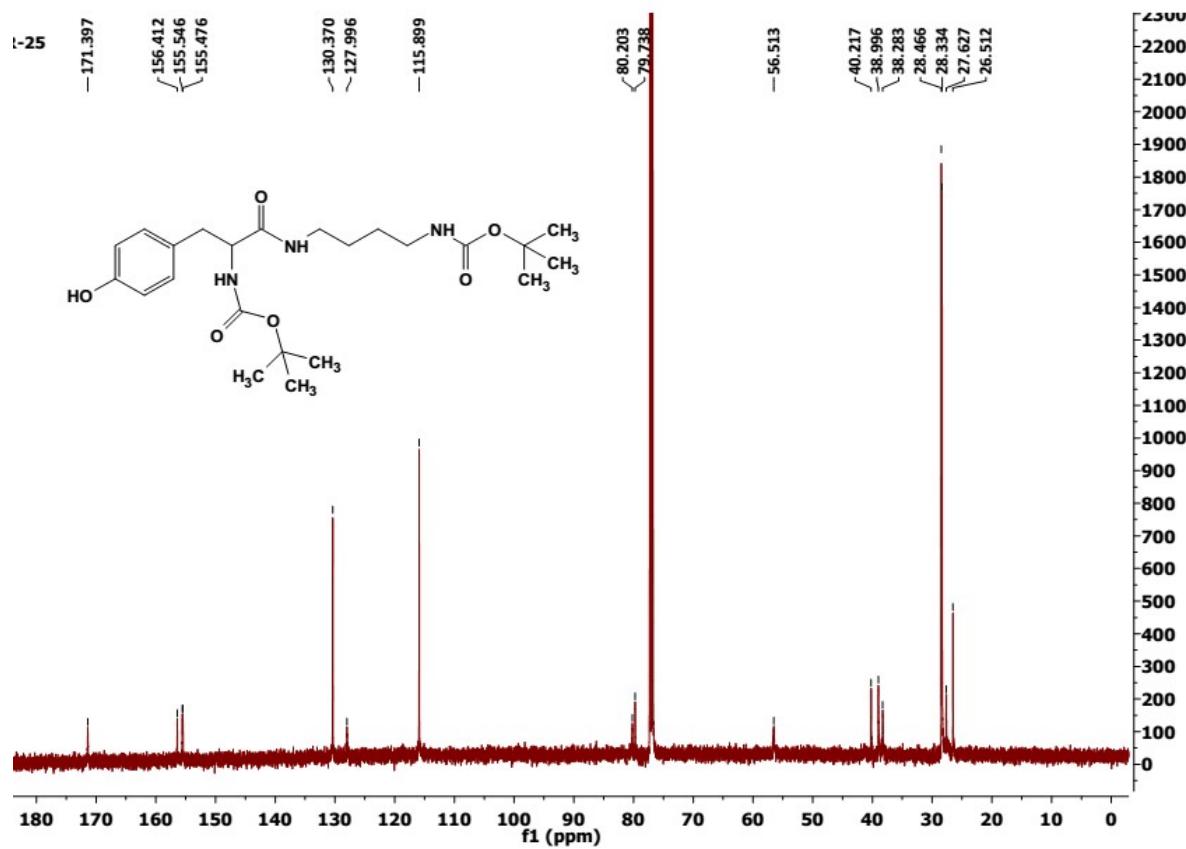


Figure 1b: ^{13}C NMR Spectrum of compound 2.

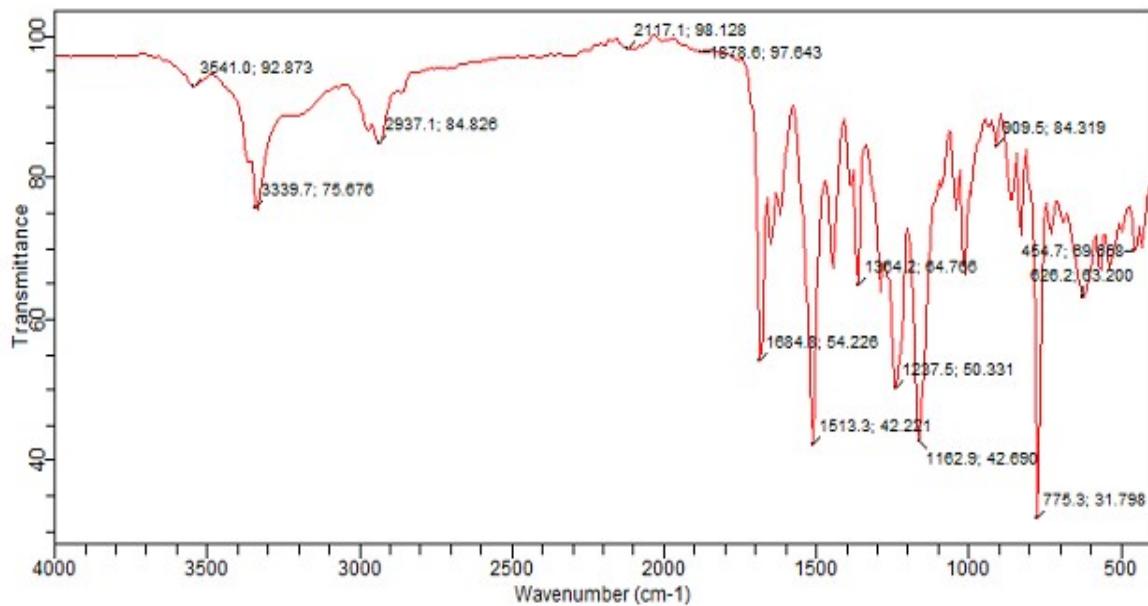


Figure 1c: FTIR Spectrum of compound 2.

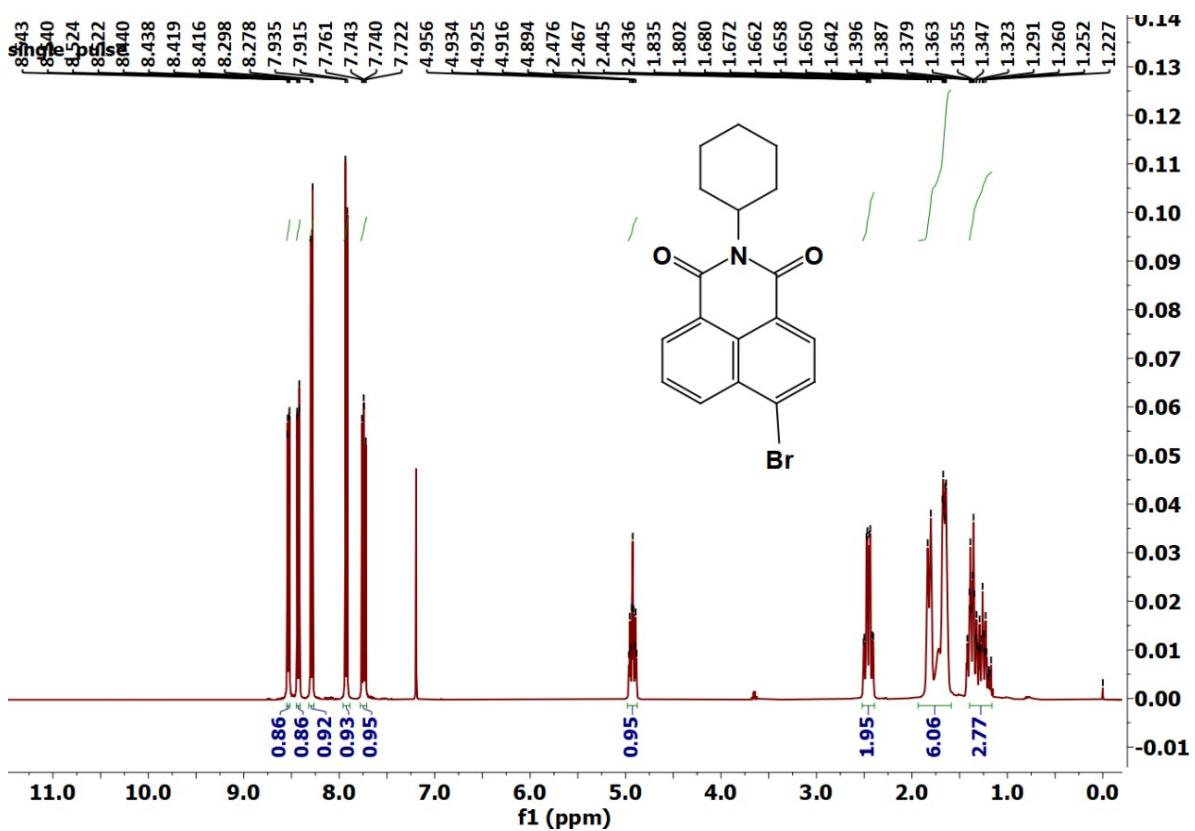


Figure 2a: ¹H NMR Spectrum of compound 3.

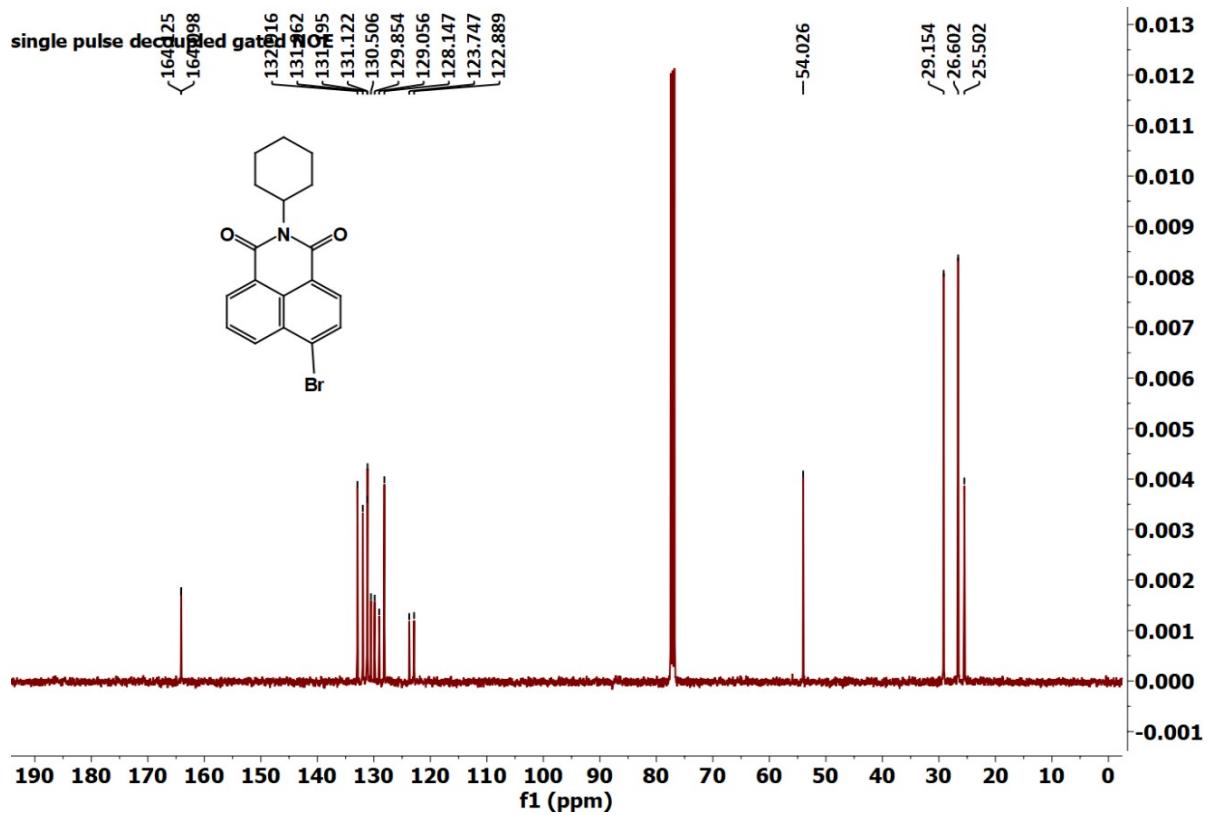


Figure 2b: ¹³C NMR Spectrum of compound 3.

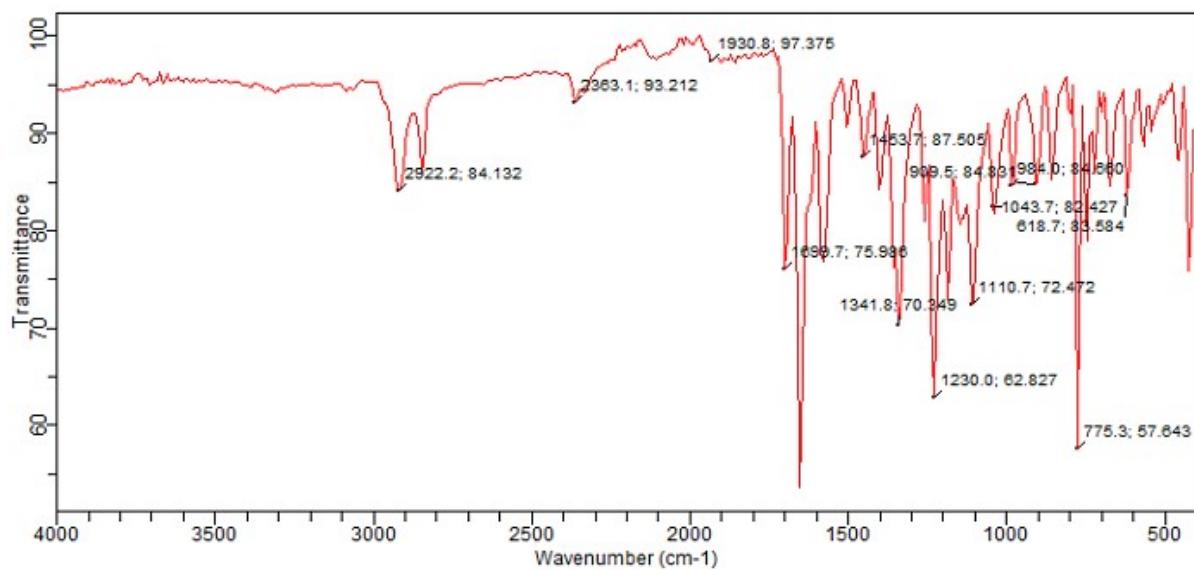


Figure 2c: FTIR Spectrum of compound 3.

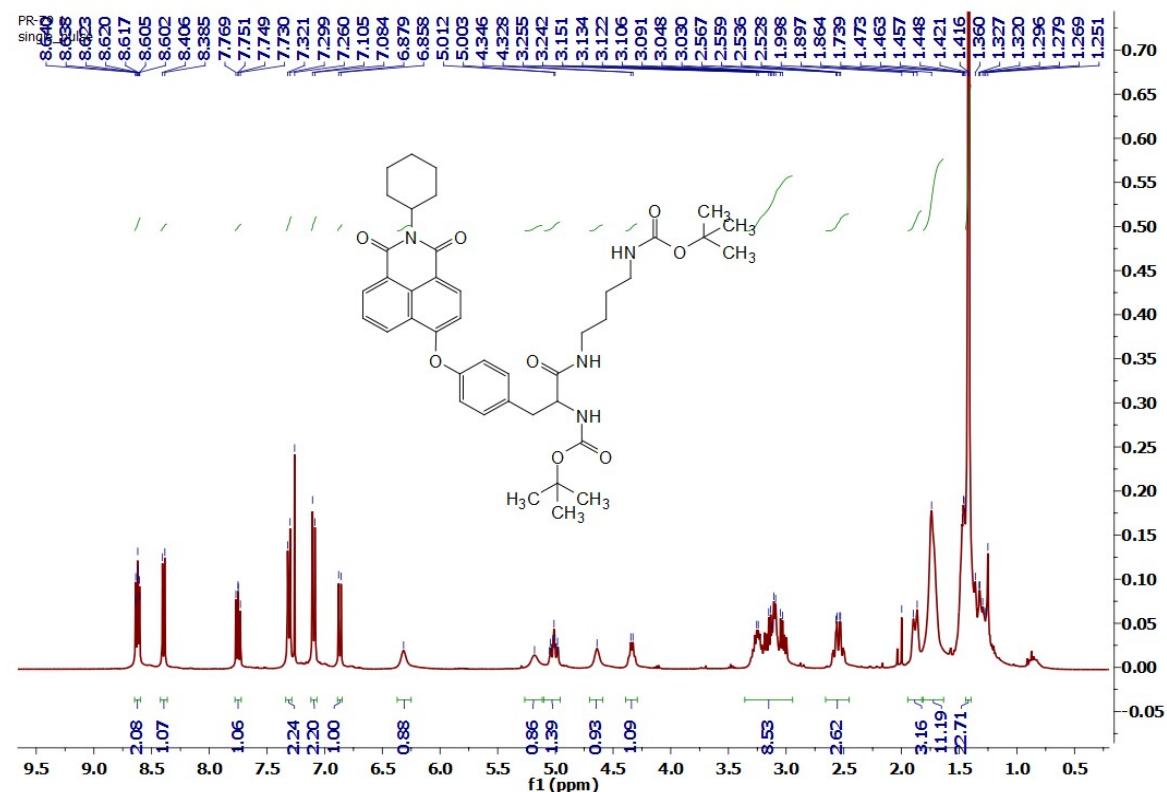


Figure 3a: ^1H NMR Spectrum of compound 4.

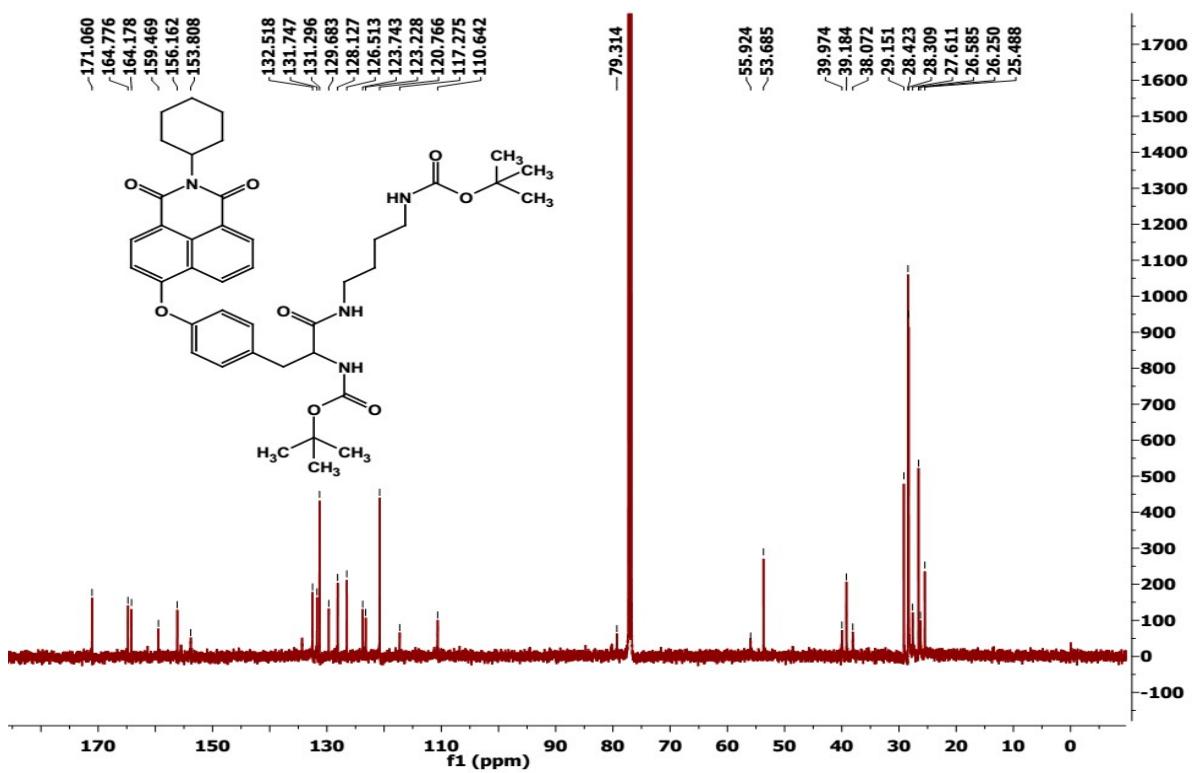


Figure 3b: ^{13}C NMR Spectrum of compound 4.

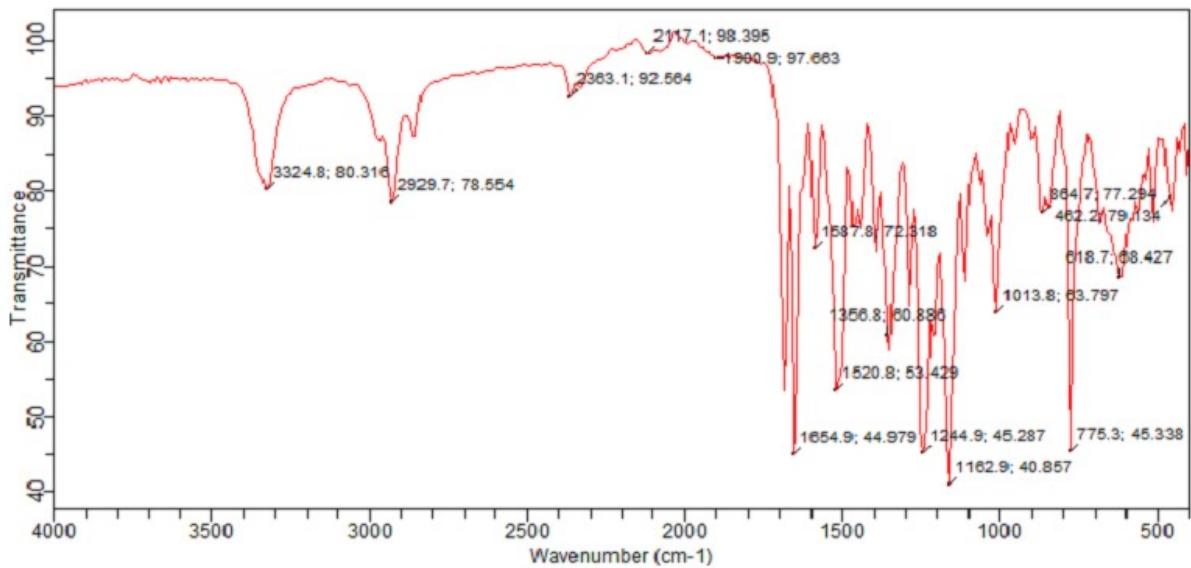


Figure 3c: FTIR Spectrum of compound 4.

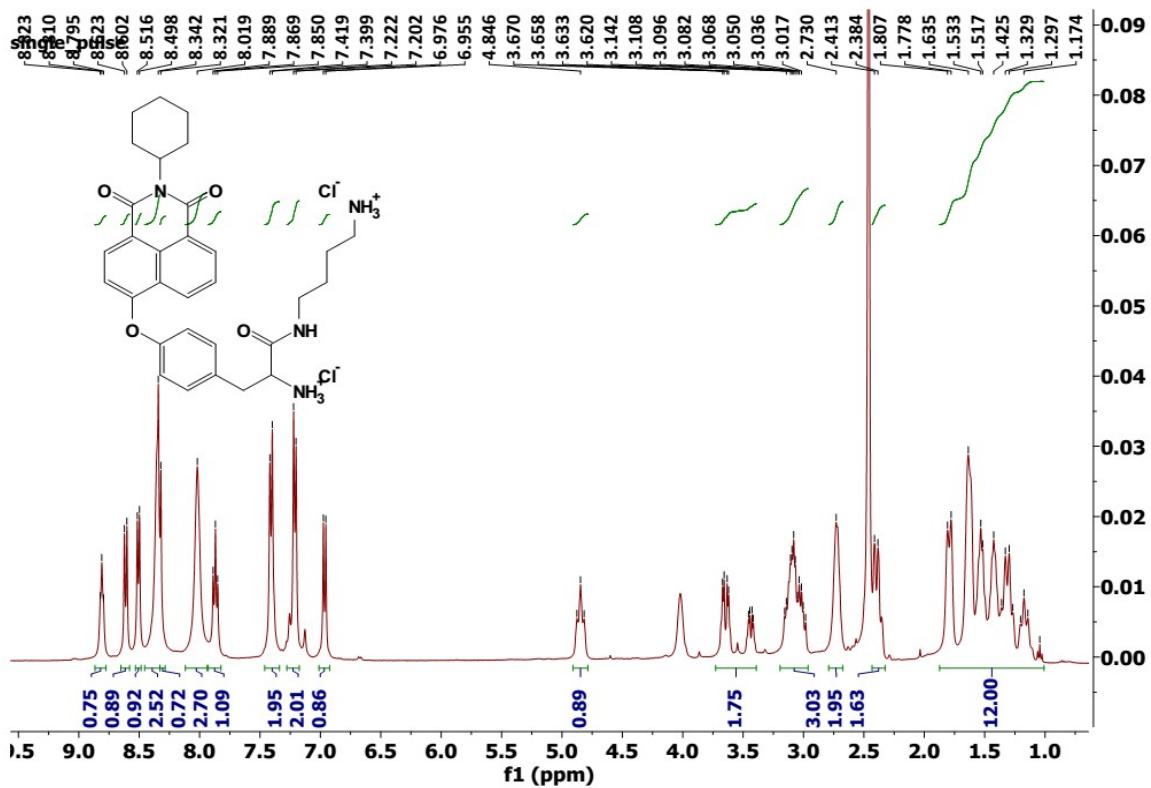


Figure 4a: ^1H NMR Spectrum of compound YN-1.

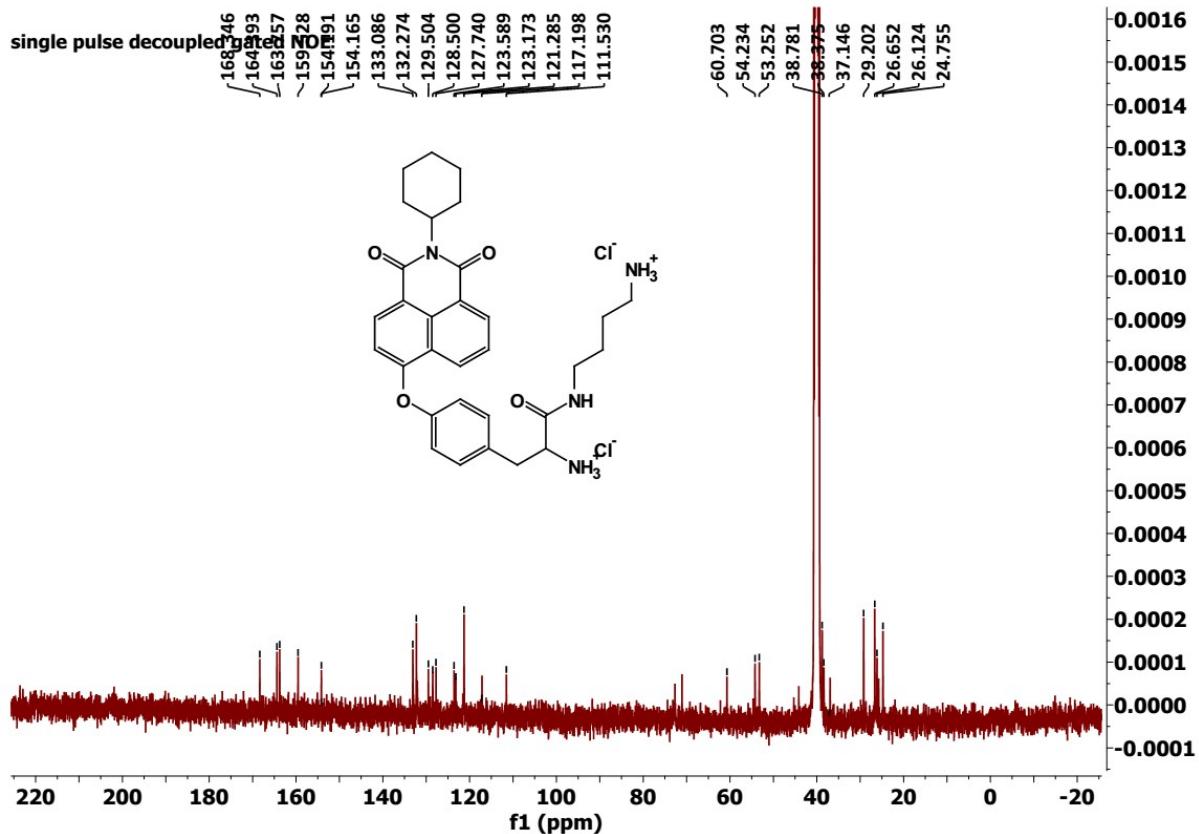


Figure 4b: ^{13}C NMR Spectrum of compound YN-1.

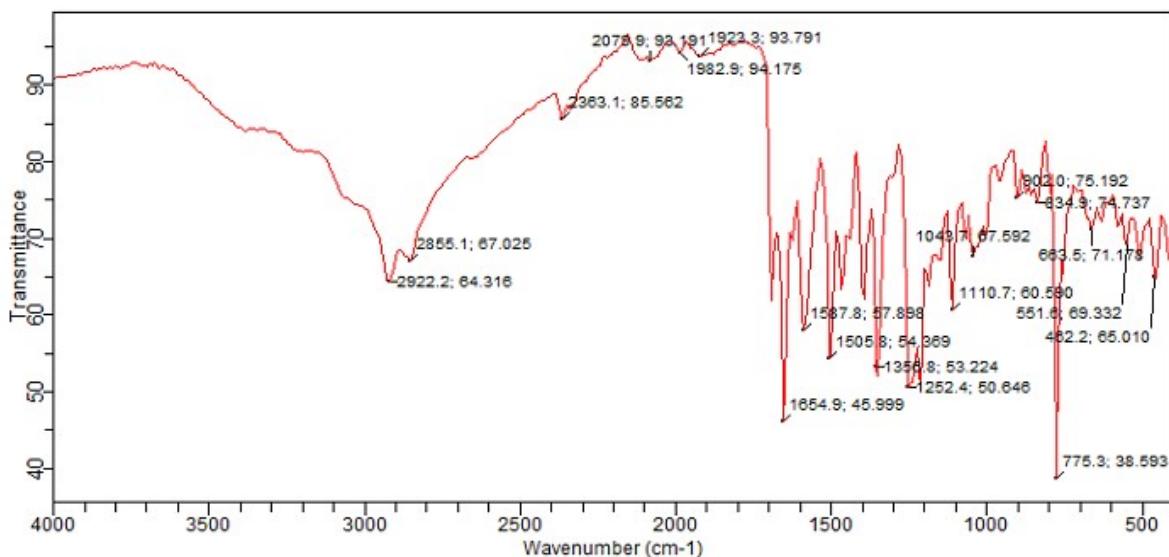


Figure 4c: FTIR Spectrum of compound YN-1.

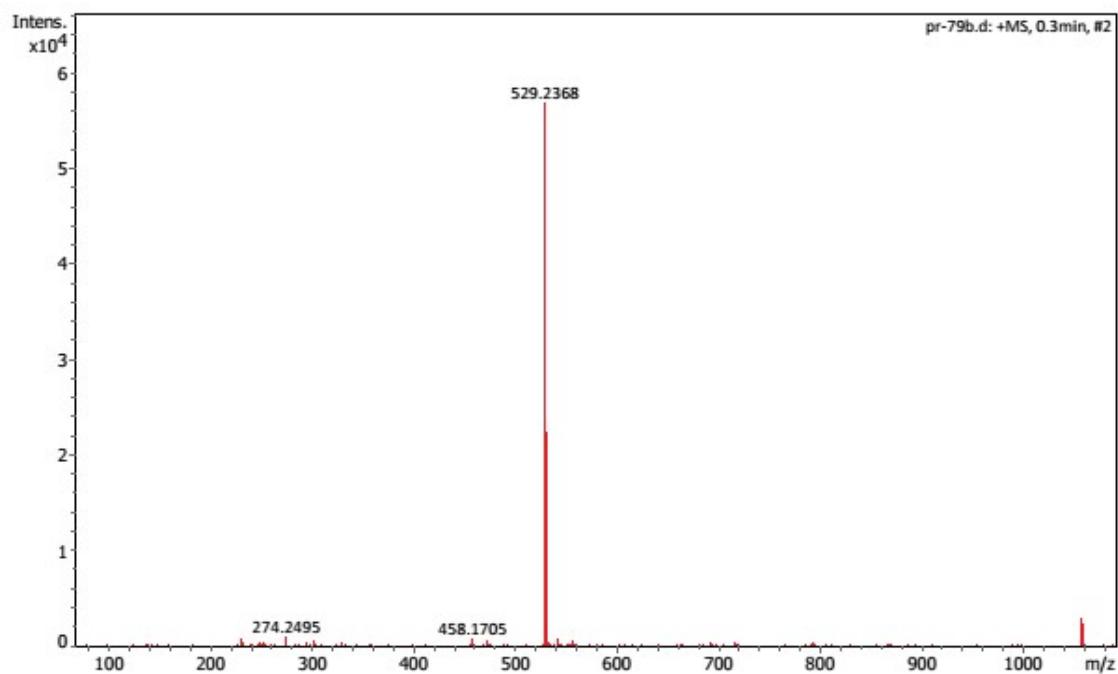


Figure 4d: Mass Spectrum of YN-1.

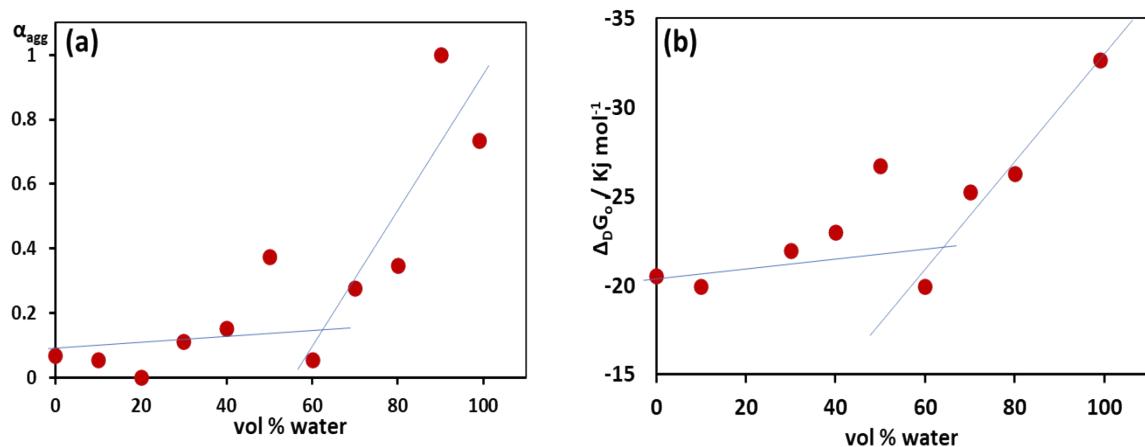


Figure S5: determination of degree of aggregation α_{agg} and Gibbs free energy (ΔG) from absorbance data recorded in 0–100 % water fraction in DMSO.

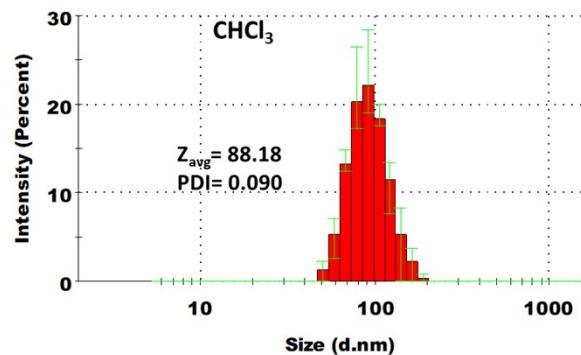


Figure S6a: Dynamic light scattering (DLS) data of YN-1 (5 μM) in CHCl_3 .

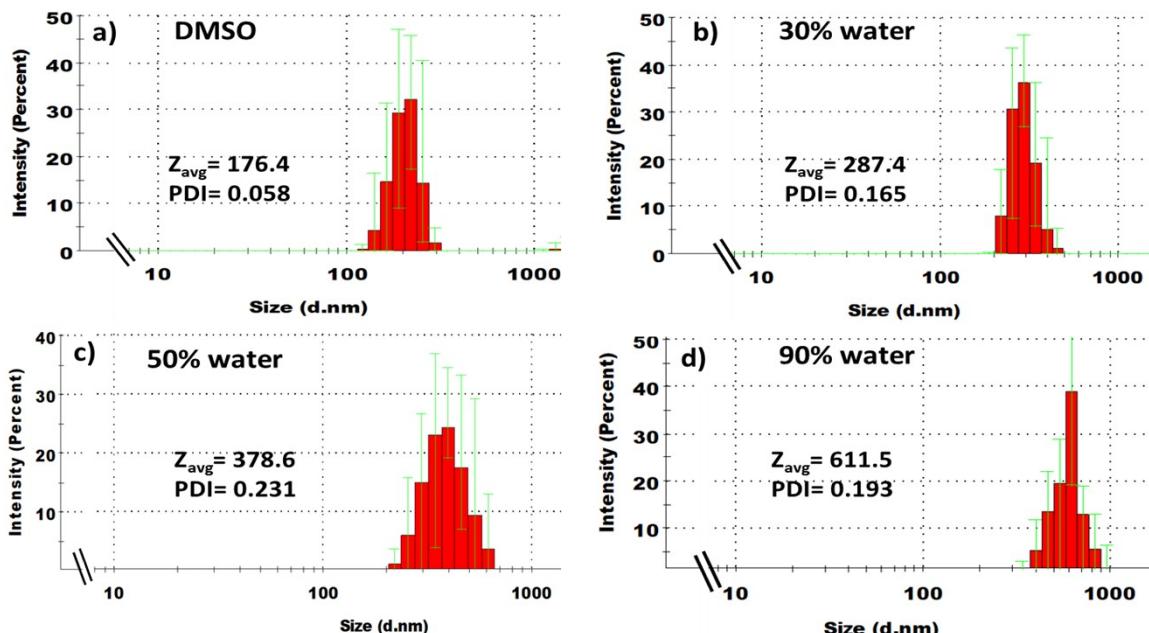


Figure S6b: Dynamic light scattering (DLS) data of YN-1 (5 μM) in (a) DMSO and in different water fractions such as (b) 30% water; c) 50% water and; d) 90% water fractions in DMSO for measuring the size of aggregates.

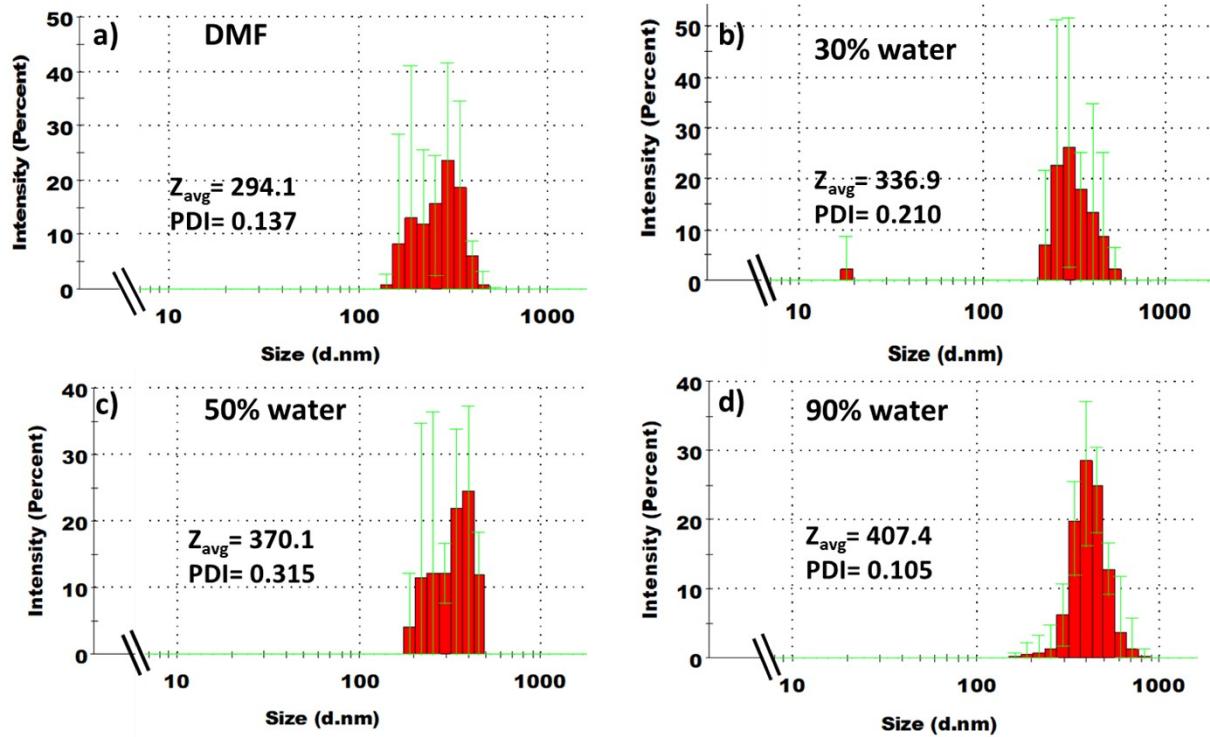


Figure S6c: Dynamic light scattering (DLS) data of YN-1 (5 μM) in (a) DMF; and different water fractions such as (b) 30%; (c) 50% and (d) 90% in DMF for measuring the size of the aggregates.

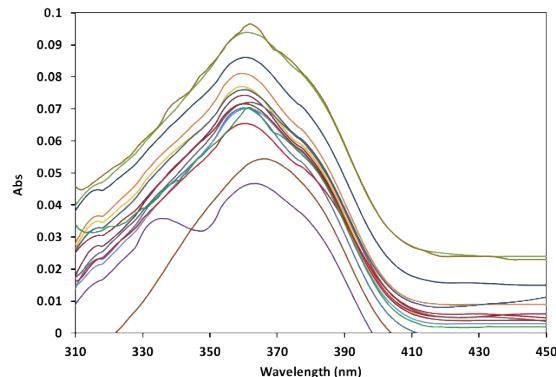


Figure S7: Absorbance spectrum of YN-1 (10 μM) upon addition of NPPs in HEPES buffer: DMSO (2:8, v/v, pH 7.2) solution.

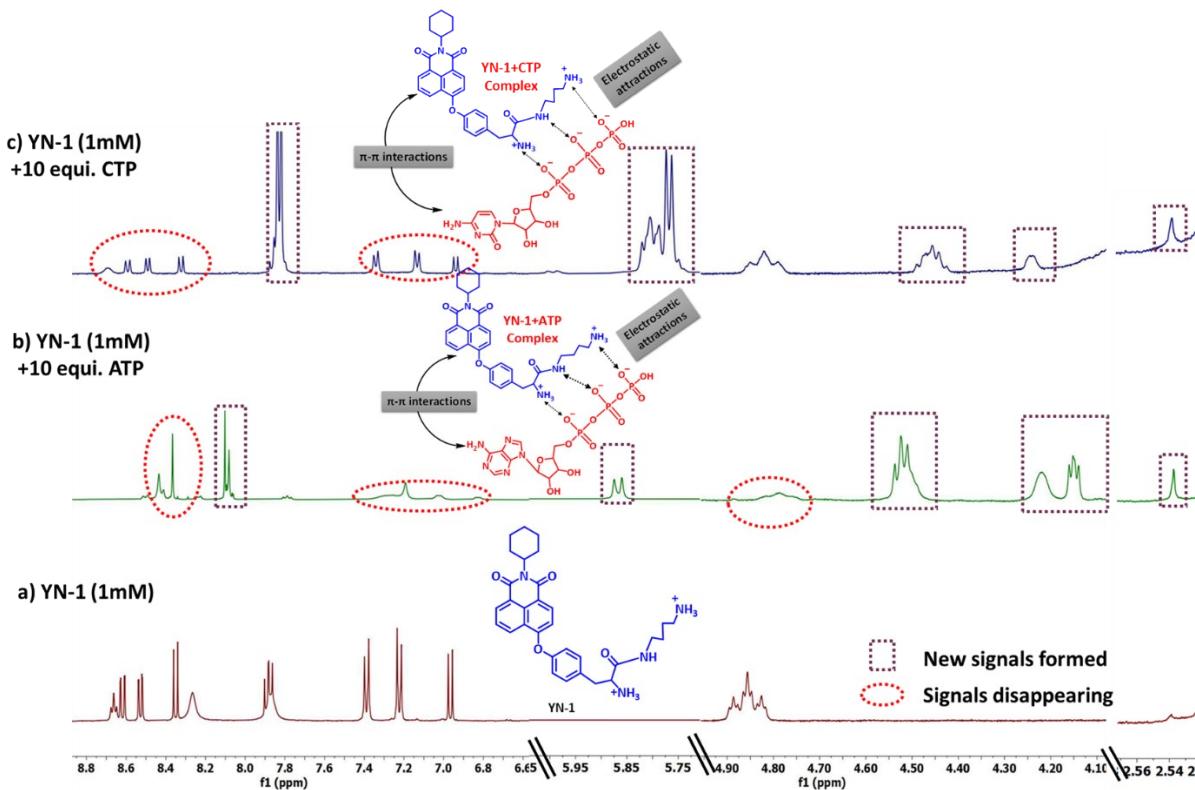


Figure S8: Partial ^1H NMR spectra of a) YN-1 (1 mM) and upon addition of 10 equivalents of b) YN-1+ATP and; c) YN-1+CTP in $\text{DMSO}-d_6$ solvent showing proposed mechanism and comparison of concentration-based discrimination between ATP and CTP (10 equivalents).

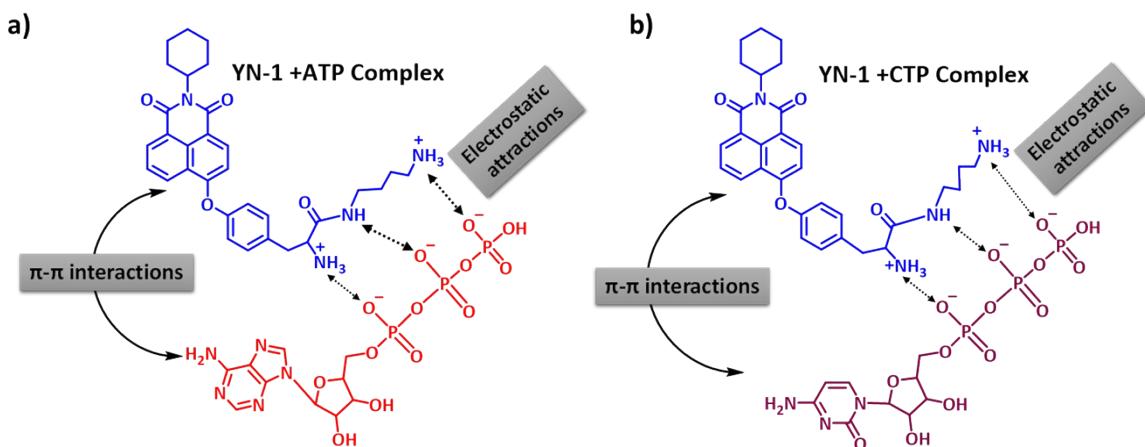


Figure S9: Proposed mechanism of a) YN-1 (1 mM) +ATP; and b) YN-1+CTP.

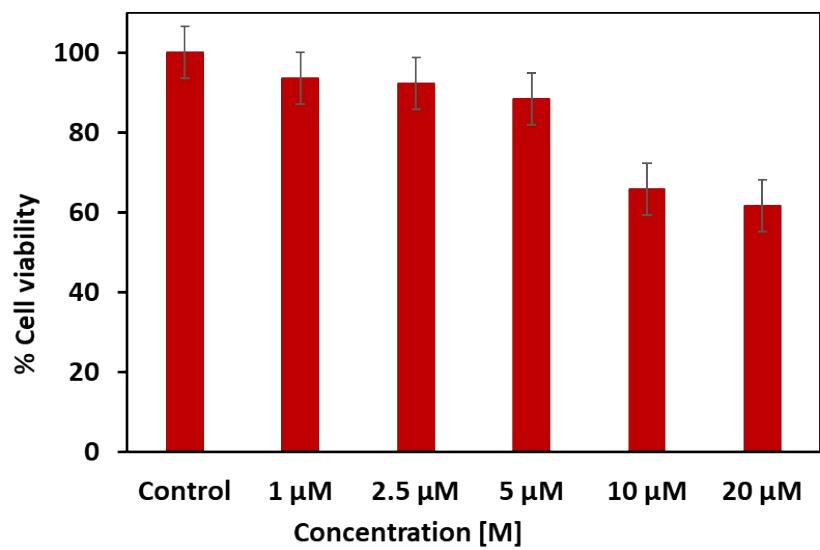
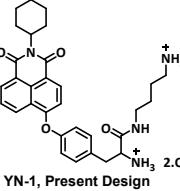
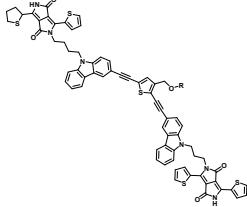
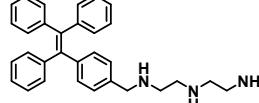
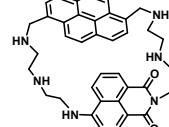
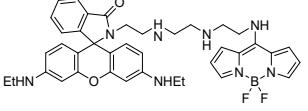
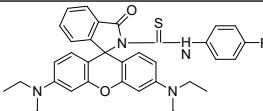
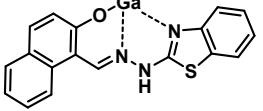
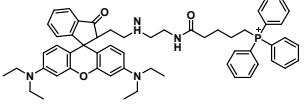
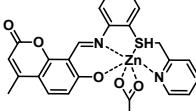
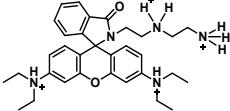


Figure S10. The MTT assay of YN-1.

Table S1. Summary of the reported fluorescent probes for ATP and CTP. HEPES: 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid; PBS: phosphate-buffered saline; Tris: tris(hydroxymethyl) aminomethane; DMSO: dimethyl sulfoxide; DMF: dimethylformamide; THF: tetrahydrofuran; λ_{ex} = excitation wavelength; λ_{em} = emission wavelength; Sensitivity: lowest detection limit.

Structure	Type	Selectivity	Media	Sensitivity	Applications	Ref.
 YN-1, Present Design	Turn-on ($\lambda_{\text{ex}}=360$ nm, $\lambda_{\text{em}}=440$ nm)	ATP and CTP	HEPES buffer: DMSO (2:8, v/v, pH 7.2)	1.72 nM (ATP); 7.1 nM (CTP)	Cell imaging	Present Manuscript
	Turn-on ($\lambda_{\text{ex}}=323$ nm, $\lambda_{\text{em}}=550$ nm)	ATP	THF: H ₂ O (1:1, v/v)	19.6 μM	-	1.
	Turn-on ($\lambda_{\text{ex}}=315$ nm, $\lambda_{\text{em}}=564$ nm)	ATP	PBS buffer (pH = 7.4)	5 μM	Cell imaging	2.
	Turn-on ($\lambda_{\text{ex}}=350$ nm, $\lambda_{\text{em}}=380$ nm and 550 nm)	ATP CTP	DMSO: TRIS buffer+100 mM NaCl (0.2:9.8, v/v, pH = 7.4)	0.08 mM	-	3.

	Turn-on ($\lambda_{\text{ex}} = 405 \text{ nm}$, $\lambda_{\text{em}} = 560 \text{ nm}$)	ATP	H_2O	0.43 μM	Cell imaging	4.
	Turn-on ($\lambda_{\text{ex}} = 425 \text{ nm}$, $\lambda_{\text{em}} = 602 \text{ nm}$)		DMF: PBS buffer (0.1:9.9, v/v, pH = 7.4)	67.4 nM	Cell imaging	5.
	Turn-on ($\lambda_{\text{ex}} = 500 \text{ nm}$, $\lambda_{\text{em}} = 560 \text{ nm}$)	ATP	HEPES (pH 7.4+10% $\text{C}_2\text{H}_5\text{OH}$)	3 mM	Cell imaging	6.
	Turn-on ($\lambda_{\text{ex}} = 500 \text{ nm}$, $\lambda_{\text{em}} = 560 \text{ nm}$)	Zn ²⁺ and ATP	$\text{CH}_3\text{CN}: 0.01 \text{ M HEPES (9:1, v/v; pH 7.4)}$	6.7 μM (for 1) and 1.7 μM (for 2)	-	7.
	Turn-on ($\lambda_{\text{ex}} = 480 \text{ nm}$, $\lambda_{\text{em}} = 634 \text{ nm}$)	ATP	$\text{C}_2\text{H}_5\text{OH}:\text{H}_2\text{O}: \text{PBS (0.1:9.9, v/v)}$	0.05 μM	Cell imaging	8.
	Turn-on ($\lambda_{\text{ex}} = 345 \text{ nm}$, $\lambda_{\text{em}} = 471 \text{ nm}$)	ATP	H_2O	1.5 μM	Cell imaging	9.

	Turn-on ($\lambda_{\text{ex}}=403$ nm, $\lambda_{\text{em}}=557$ nm)	ATP	HEPES buffer (pH 5.5)	-	-	10.
	Turn-on ($\lambda_{\text{ex}}=510$ nm, $\lambda_{\text{em}}=591$ nm)	ATP ADP CTP	DMSO: PBS (4:6, v/v)	-	Cell imaging	11.
	Turn-off ($\lambda_{\text{ex}}=440$ nm, $\lambda_{\text{em}}=518$ nm)	ATP ADP	DMSO:water (9:1, v/v)	-	Cell imaging and In vivo images of zebrafish.	12.
	Turn-on ($\lambda_{\text{ex}}=520$ nm, $\lambda_{\text{em}}=583$ nm)	ATP	PBS buffer (pH 7.4)	0.033 mM	Cell imaging	13.
	Turn-off ($\lambda_{\text{exc}}=315$ nm, $\lambda_{\text{emi}}=512$ nm)	ATP	CH ₃ OH:water (3:1, v/v)	6.6 μM	Cell imaging	14.
	Turn-on ($\lambda_{\text{ex}}=510$ nm, $\lambda_{\text{em}}=583$ nm)	ATP	HEPES buffer (pH 7.2)	-	Cell imaging	15.

	Turn-off ($\lambda_{\text{ex}} = 457 \text{ nm}$, $\lambda_{\text{em}} = 560 \text{ nm}$)	ATP and CTP	TRIS HCl buffer (pH 7.2)	3.9 μM (ATP) and 8.3 μM (CTP)	-	16.
	Turn-on ($\lambda_{\text{ex}} = 420 \text{ nm}$, $\lambda_{\text{em}} = 580 \text{ nm}$)	ATP	HEPES (pH = 7.2)	0.1 μM	Cell imaging	17.
Terbium(III)-organic framework	Turn-off ($\lambda_{\text{ex}} = 330 \text{ nm}$, $\lambda_{\text{em}} = 493 \text{ nm}, 546 \text{ nm}, 586 \text{ nm}$ and 623 nm)	CTP	DMF:water (1 : 1, v/v)	-	-	18.
	Turn-on ($\lambda_{\text{ex}} = 370 \text{ nm}$, $\lambda_{\text{em}} = 538 \text{ nm}$)	ATP ADP	HEPES buffer (pH 7.4, 1% MeCN)	1.0 μM	Cell imaging	19.
	Turn-on ($\lambda_{\text{ex}} = 342 \text{ nm}$, $\lambda_{\text{em}} = 482 \text{ nm}$)	ATP	Polycation buffer (pH 10.2, 0.5 mM Na2CO3, 0.5 mM NaHCO3)	0.1 μM	In vitro	20.

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