

Electronic Supplementary Information

**Supramolecular optimization of the visual contrast in a colorimetric chemosensing assay
that releases resorufin dye**

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Materials and Instrumentation

All the solvents and chemicals were purchased from Sigma-Aldrich, Alfa-Aesar, or VWR international and used without further purification unless otherwise stated. ^1H and ^{13}C spectra were recorded on Bruker AVANCE III HD 400, 500 MHz spectrometers. Mass spectrometry (MS) was performed using a Bruker microTOF II spectrometer. Synthesized compounds were purified using Biotage flash purification system with SNAP Ultra flash chromatography cartridges.

N-acetyl- β -D-glucosaminidase (NAG) Stock Solution

A 0.9 mg/mL solution of NAG enzyme (Sigma-Aldrich) from bovine kidney was prepared in 1 M PBS Buffer + 0.1 M BSA and the vendor's guarantee of enzymatic activity was confirmed using the standard chromogenic substrate p-nitrophenyl-N-acetyl- β -D-glucosaminide.

HOCl/ OCl⁻ Stock Solution

Hypochlorite stock solution was prepared using a previously reported method.¹ To a 1500 μL solution of 154 mM NaCl, 250 μL of 10-14% w/w NaOCl was added followed by dropwise addition of 6M HCl to obtain a pH range of 3.92. The concentration of active total chlorine species in solution expressed as $[\text{HOCl}]_{\text{T}}$ (where $[\text{HOCl}]_{\text{T}} = [\text{HOCl}] + [\text{Cl}_2] + [\text{Cl}_3^-] + [\text{OCl}^-]$) in HPLC Grade water was determined by converting all the active chlorine species to OCl^- with 0.1 M NaOH and measuring the concentration of OCl^- . The concentration of OCl^- was determined spectrophotometrically at 292 nm ($\epsilon = 362 \text{ M}^{-1} \text{ cm}^{-1}$) with a UV-visible spectrophotometer. Calculation: $A = \epsilon cl$; where $l = 1 \text{ cm}$, $A = 0.6359$, $\epsilon = 362 \text{ M}^{-1} \text{ cm}^{-1}$. Thus, $c = 1.76 \text{ mM HOCl/ OCl}^-$

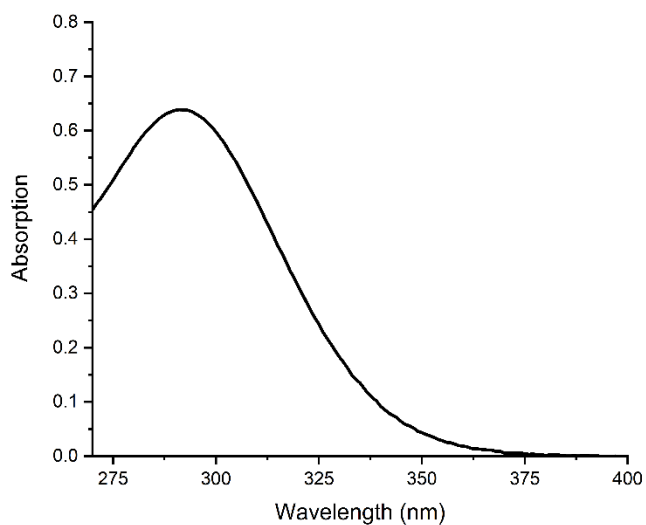


Figure S1: Absorption spectrum of HOCl/ OCl⁻ for stock solution concentration determination.

Synthesis

The chemosensor **RT-1** and enzyme substrate **NHPO** were synthesized as previously described,^{2,3} and the following ¹H NMR and HR-MS data demonstrate high purity. Tetralactam macrocycles **M1** and **M2** were synthesized as part of previous studies^{4,5} and the purity was confirmed by ¹H NMR.

RF-TBA: Resorufin sodium salt (50 mmol, 10.6 mg) and 40 % wt tetrabutylammonium hydroxide solution (50 mmol, 33 μ L) were dissolved in 50 mL of PBS. The resulting mixture was extracted with chloroform (3 \times 50 mL). The combined chloroform layers were dried under vacuum to obtain pure **RF.TBA** as a dark pink solid.

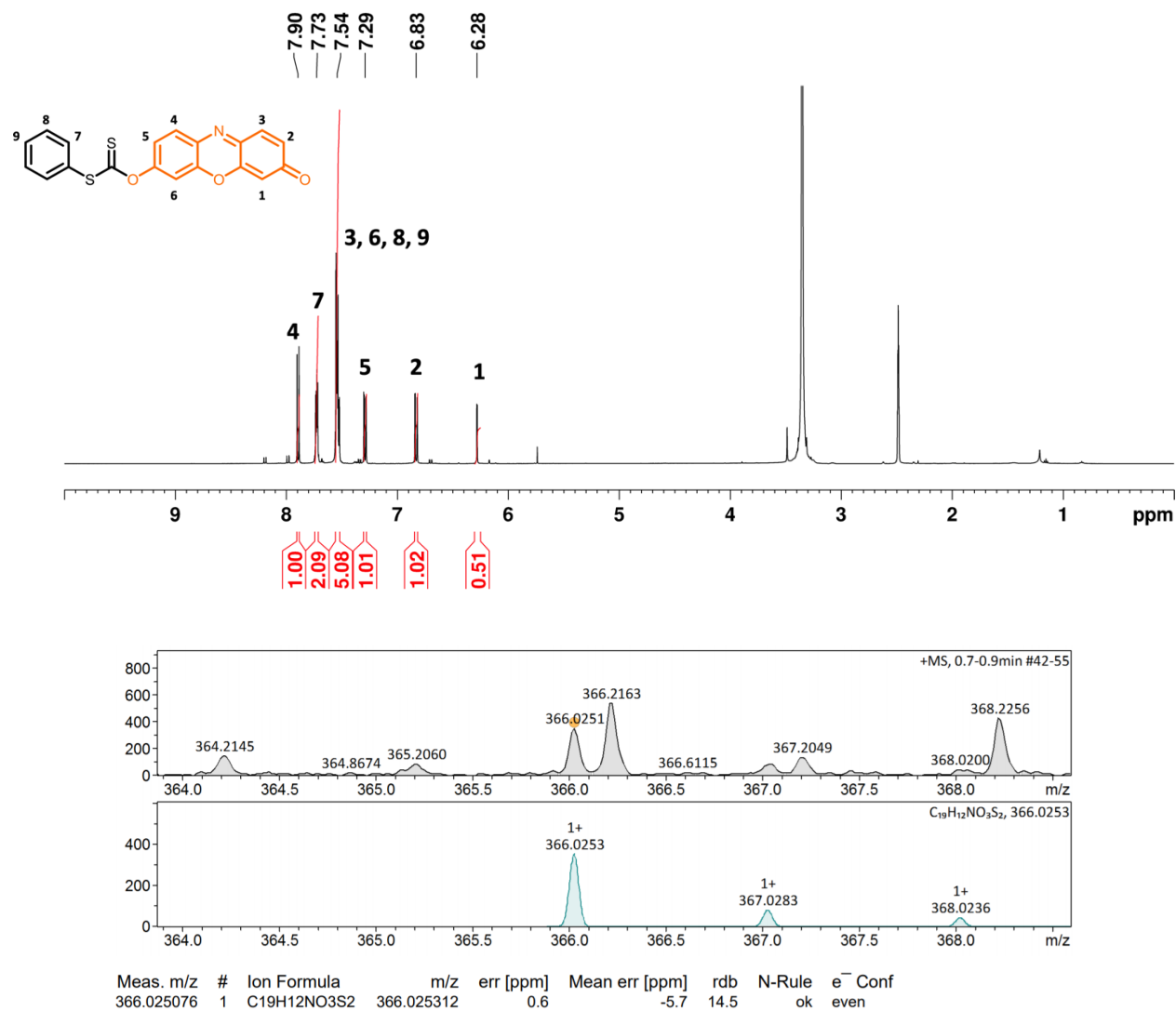


Figure S2. ¹H NMR (500 MHz; DMSO-d₆; Me₄Si) and HR-ESI mass spectrum of **RT-1**.

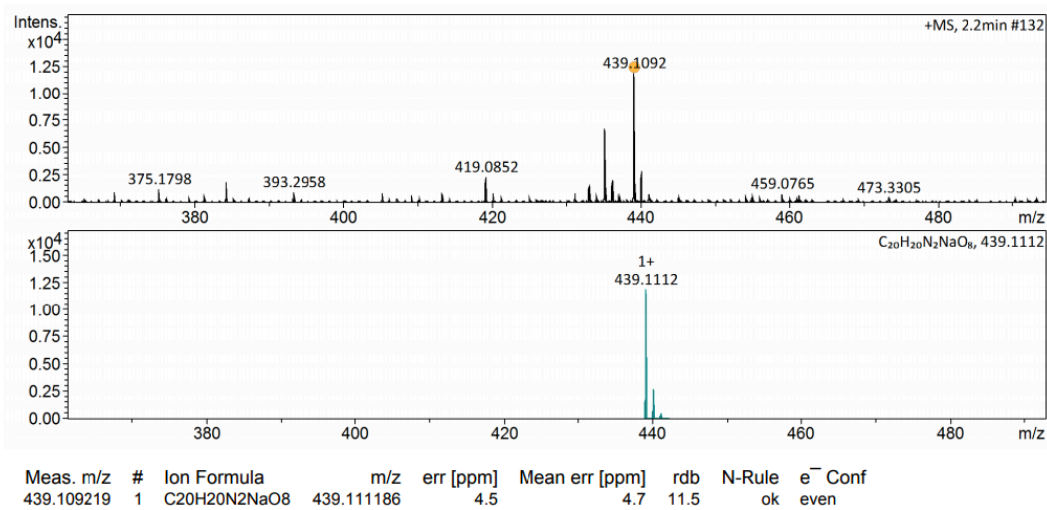
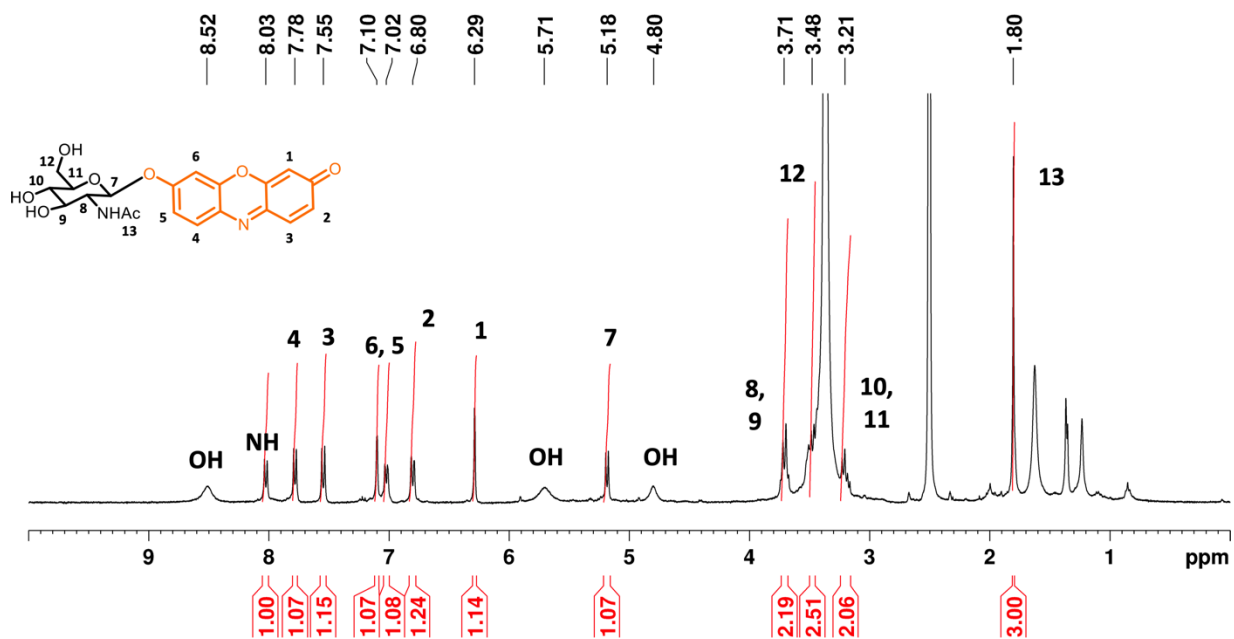


Figure S3. ¹H NMR (400 MHz; DMSO-d₆; Me₄Si) and HR-ESI mass spectrum of NHPO. The broad OH peaks in the ¹H NMR spectrum are due to exchange promoted by adventitious water in the DMSO-d₆.

¹H NMR Titration Data

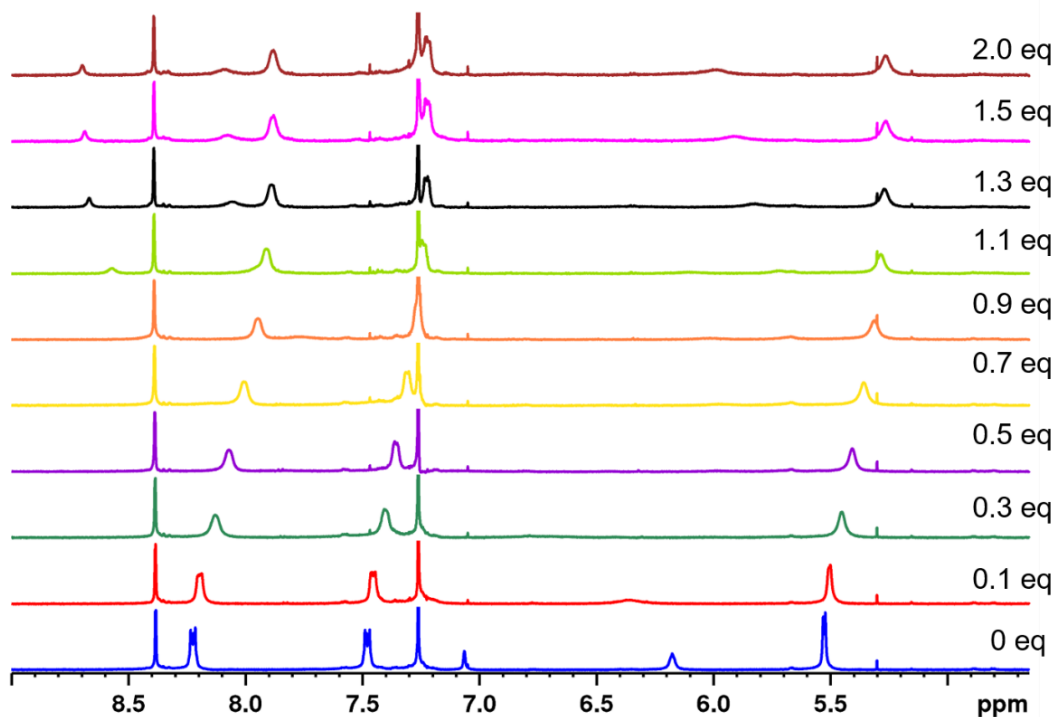


Figure S4. ¹H NMR titration (500 MHz, CDCl₃, 25° C) that added aliquots from a stock solution containing 10 mM **RF** (tetrabutylammonium salt)/ 0.5 mM **M1** to a solution of **M1** (0.5 mM).

K_a Determination by Fluorescence Titration

Previously described titration method was employed.⁶ Stock solutions of the guest, **RF** (1 mM) and host, **M2** (1 mM) were made in pure water. A solution of the guest was placed in a cuvette (10 μM) and aliquots of the host (**M2**) were added fluorescence spectra were acquired (ex: 540 nm, em: 585 nm). The data was plotted and association constant for **RF** binding to **M2** was determined by non-linear squares fitting of the titration points to a model for 1:1 binding within the Origin software.⁷

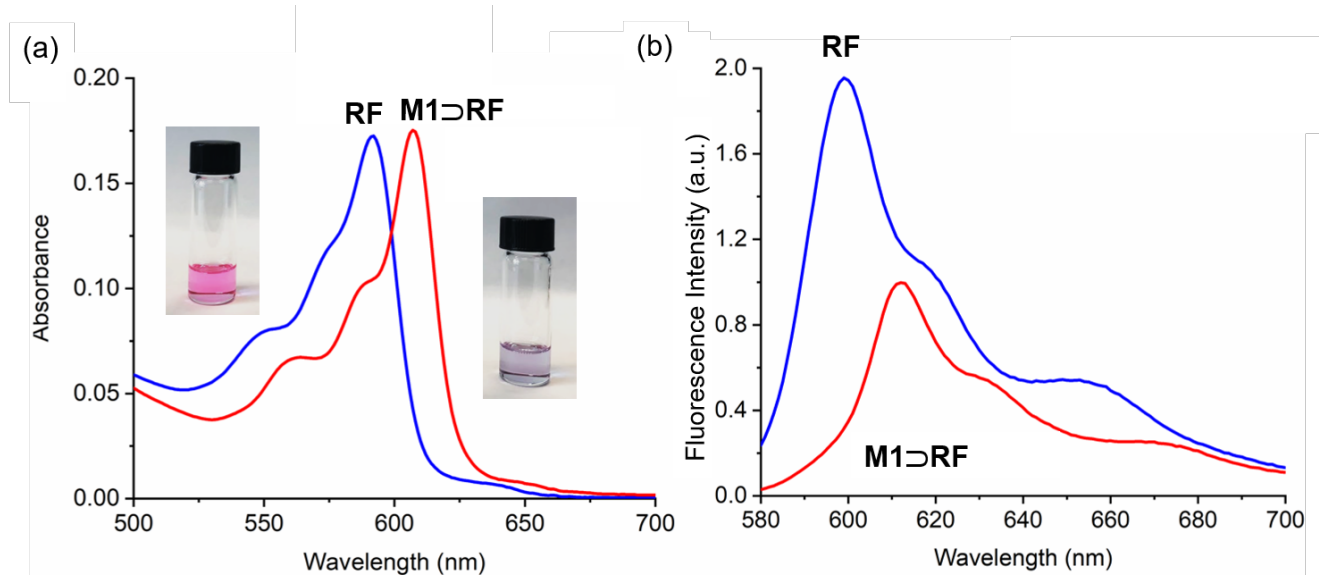


Figure S5. (a) Absorption (b) fluorescence emission of 10 μM RF (tetrabutylammonium salt) and M1D RF in chloroform at 25°C.

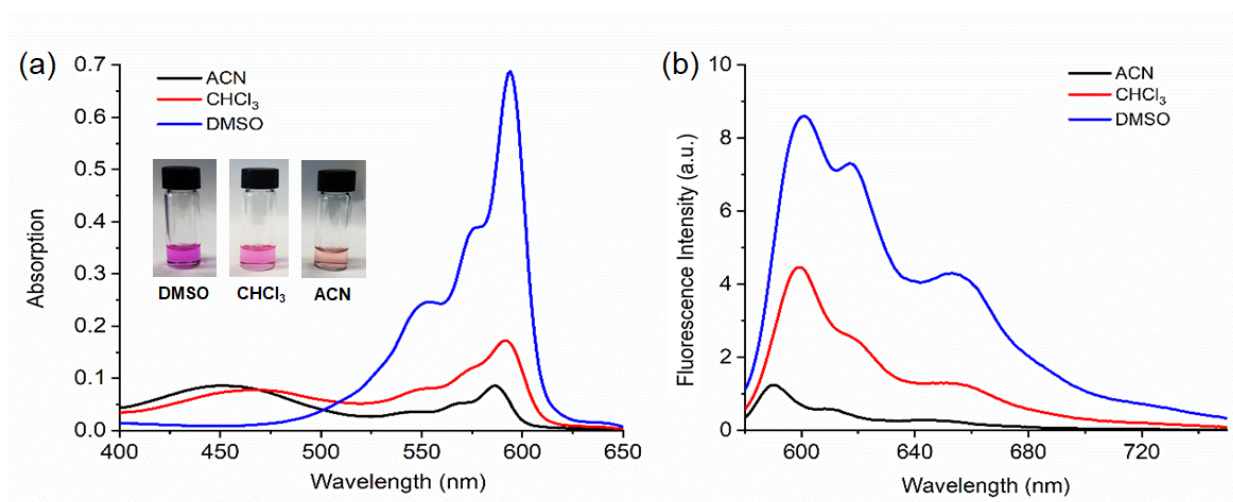


Figure S6. Absorption and emission spectra of 10 μM RF (tetrabutylammonium salt) in different organic solvents at 25°C, along with photographs of the solutions.

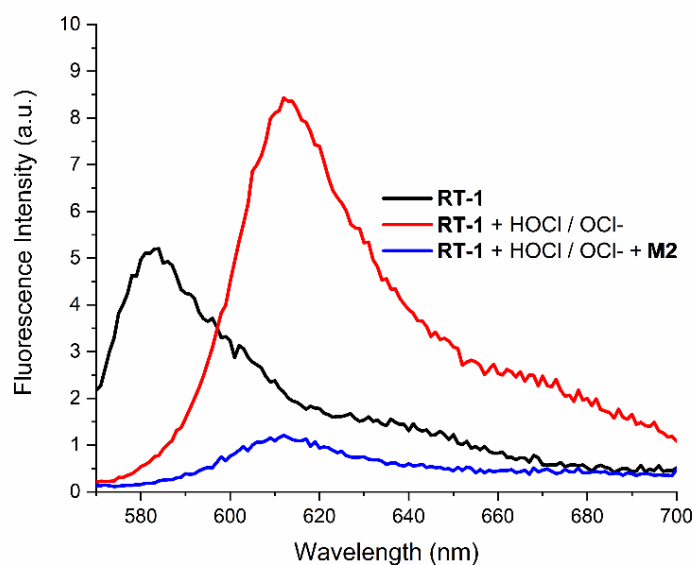


Figure S7. Fluorescence spectra ($\lambda_{\text{ex}} = 550 \text{ nm}$) of a sample, initially containing **RT-1** ($50 \mu\text{M}$, black line), and 3 minutes after addition of HOCl/OCl⁻ ($5 \mu\text{M}$, red line), or 3 minutes after a two-step addition sequence of HOCl/OCl⁻ ($5 \mu\text{M}$) and then **M2** ($500 \mu\text{M}$) (blue line). In 200 mM PBS, pH 7.4 at 25°C.

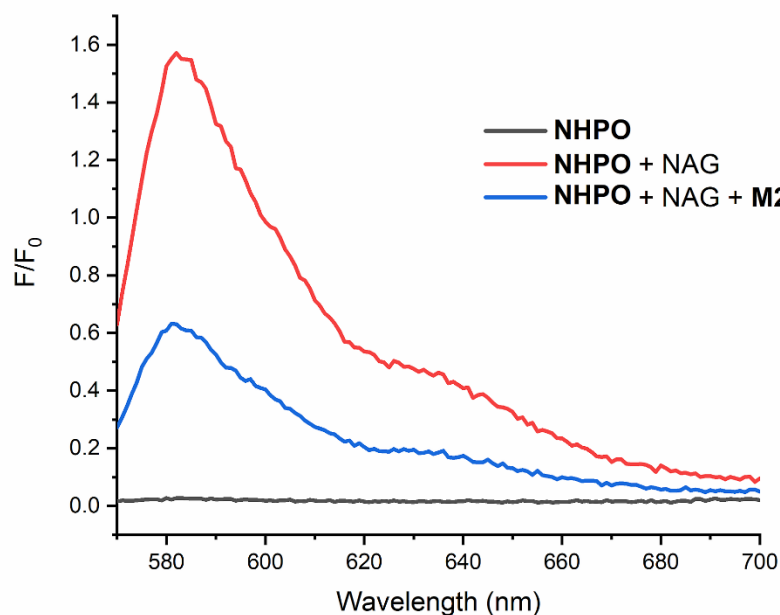


Figure S8. Fluorescence spectra ($\lambda_{\text{ex}} = 550 \text{ nm}$) of a sample initially containing **NHPO** ($50 \mu\text{M}$, black line), 30 minutes after addition of $0.9 \mu\text{g/mL}$ NAG (red line), or 45 minutes after a two-step addition sequence of $0.9 \mu\text{g/mL}$ NAG and then **M2** ($500 \mu\text{M}$) (blue line). In 100 mM PBS + 100 μM BSA, pH 7.4 at 25°C.

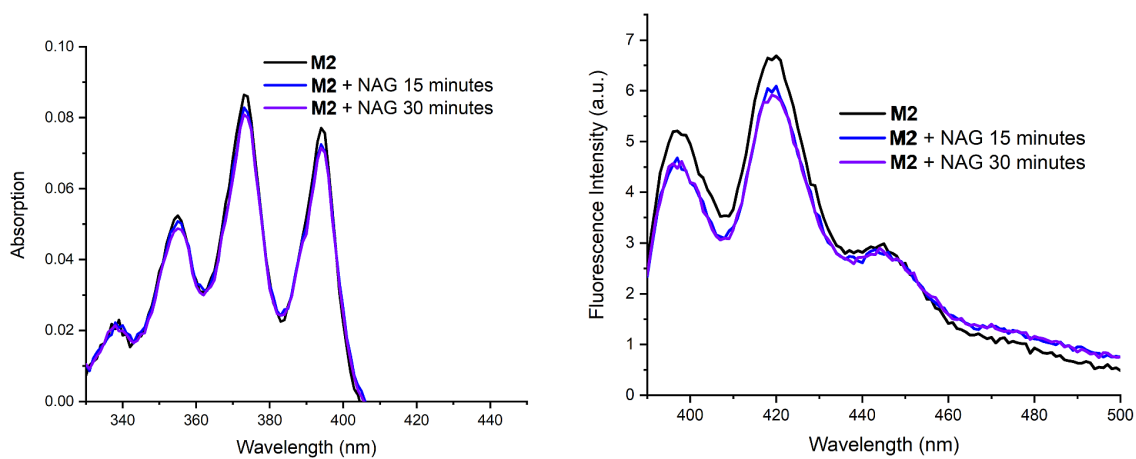


Figure S9. Absorption and fluorescence emission (ex: 370 nm, em: 390 nm) of a solution containing 15 μM **M2** in the presence and absence of (0.1 $\mu\text{g}/\text{mL}$ NAG enzyme plus ~ 10 μM BSA), in water and 25°C. The very small intensity decrease upon protein addition is due to sample dilution, and it appears there is negligible interaction of NAG or BSA with **M2**.

Molecular Modeling

The semiempirical PM7 method was employed within the MOPAC program. (J. J. P. Stewart, MOPAC; Stewart Computational Chemistry: Colorado Springs, CO, 2008.) The dielectric constant of the solvent was set at 78.4 for water and 25 °C. Solubilizing groups are shortened to hydrogens.

Cartesian Coordinates at the PM7 Level

TOTAL ENERGY = -10938.84941 EV

FINAL GEOMETRY OBTAINED

EPS=78.4 PM7 CHARGE=-1 EF xyz GNORM=0.100 SHIFT=80

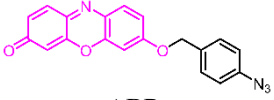
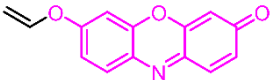
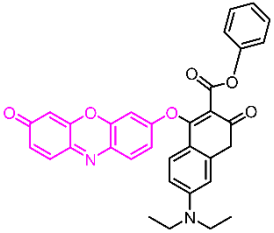
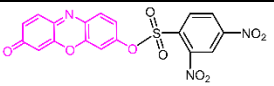
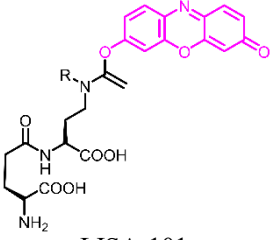
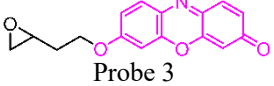
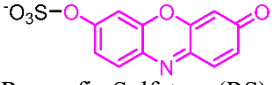
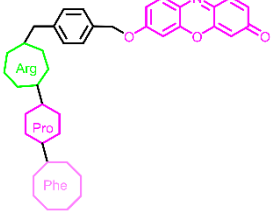
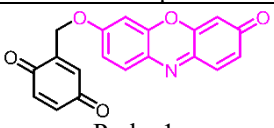
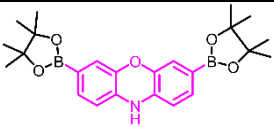
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C	4.52278289	+1	-0.98838362	+1	3.30059312	+1
C	5.10257652	+1	0.82824274	+1	4.78463703	+1
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C	11.11361398	+1	2.10070945	+1	0.95557584	+1
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C	11.63203900	+1	-0.25491023	+1	0.83943026	+1
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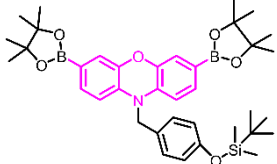
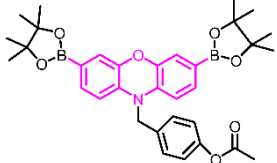
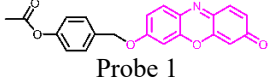
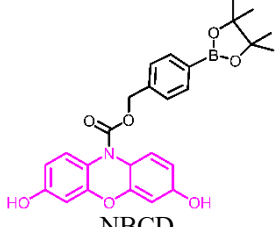
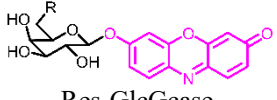
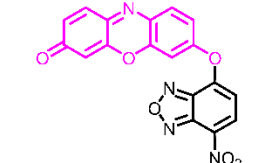
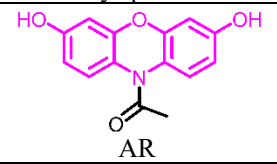
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C	8.32636327	+1	1.31144390	+1	3.19984307	+1
C	5.51597253	+1	0.58717751	+1	-2.47873205	+1
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O	4.34109097	+1	3.75401182	+1	-3.61738014	+1
O	8.82536173	+1	1.65456233	+1	4.27007678	+1
H	4.46014140	+1	2.78371788	+1	4.23936159	+1
H	4.55184649	+1	-2.04663149	+1	3.03732730	+1
H	5.61015438	+1	1.14825144	+1	5.69459216	+1
H	5.64279222	+1	-1.24860737	+1	5.10493309	+1
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H	7.81422108	+1	5.04548630	+1	-4.59408984	+1
H	8.97856143	+1	5.24510619	+1	-2.42709583	+1
H	2.23997220	+1	-1.01948292	+1	-1.23589298	+1
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H	1.16035048	+1	3.03050105	+1	-2.15043929	+1
H	2.04594398	+1	3.79118971	+1	0.02934829	+1
H	2.90765132	+1	3.72837743	+1	3.49225535	+1
H	2.00585196	+1	4.06961559	+1	2.01658031	+1

H	4.88030841	+1	3.94571460	+1	1.48305548	+1
H	6.62622777	+1	5.09489682	+1	1.45217784	+1
H	7.85352421	+1	8.87211527	+1	-0.23473706	+1
H	5.59220871	+1	9.80464951	+1	0.20547888	+1
H	3.81447609	+1	8.35585677	+1	1.16485293	+1
H	8.22454506	+1	4.36372020	+1	0.32136289	+1
H	10.96703491	+1	4.54264697	+1	0.48862585	+1
H	10.66878882	+1	5.28384485	+1	-1.08673499	+1
H	10.83345957	+1	-1.08709250	+1	-3.01392265	+1
H	9.52843975	+1	-0.53460490	+1	-4.06752197	+1
H	8.27262727	+1	-1.10292388	+1	-1.59182849	+1
H	6.38000558	+1	-2.24100260	+1	-1.54902425	+1
H	5.64420075	+1	-6.15715447	+1	0.11590198	+1
H	7.80038316	+1	-6.96446016	+1	-0.82338215	+1
H	9.26662705	+1	-5.40591135	+1	-2.09040512	+1
H	2.34140368	+1	-2.23234920	+1	0.23727619	+1
H	2.98204237	+1	-2.57440552	+1	1.84340073	+1
H	5.19314717	+1	-1.72297880	+1	0.21319645	+1
H	7.62270163	+1	3.29617288	+1	2.65361514	+1
H	7.88608749	+1	-1.54385129	+1	1.28306589	+1
H	8.88914855	+1	-0.77683924	+1	3.44729399	+1
H	5.56946817	+1	-0.46567424	+1	-2.77011967	+1
H	4.51120938	+1	1.26814086	+1	-4.23614250	+1
H	5.39113145	+1	4.35269899	+1	-1.31209333	+1

Table S1: Abridged collection of enzyme substrates and chemosensors that release resorufin (RF).

Probe	Analyte	Method of Detection	Reference Number
<p>Resorufin Thionocarbonate (RT)</p>	Mercury Hg^{2+}	Chromogenic	8
<p>Resorufin β-D- glucuronide (REG)</p>	E. coli	Chromogenic	9
<p>Novel Probe 1</p>	Alkaline Phosphatase (ALP)	Fluorescence	10
<p>Resorufin turn on Probe (RTP-1)</p>	Hydrazine (N_2H_4)	Fluorescence	11
<p>Resorufin-β-D- Galactopyranoside</p>	Biotinylated DNA	Fluorescence	12
<p>F-Chemosimeter 1</p>	Fluorine (F^-)	Chromogenic/ Fluorescence	13
<p>Sulfite Selective Probe</p>	Sulfite (SO_3^{2-})	Chromogenic/ Fluorescence	14
<p>R1</p>	Perborate (BO_3^-) /Hydrazine (N_2H_4)	Chromogenic/ Fluorescence	15,16
<p>Re-SS</p>	Polysulfides	Fluorescence	17
<p>Ozone Probe 1</p>	Ozone (O_3)	Chromogenic/ Fluorescence	18
<p>PC1</p>	Hydrogen Peroxide (H_2O_2)	Fluorescence	19

 <p>ABR</p>	Hydrogen Sulfide (H ₂ S)	Fluorescence	20
 <p>Mercury Probe 1</p>	Mercury (Hg ²⁺)	Chromogenic/Fluorescence	21
 <p>RC</p>	Hydrogen sulfide (H ₂ S), Cysteine (Cys), Homocysteine (Hcy), Glutathione (GSH)	Fluorescence	22
 <p>Probe 1</p>	Acetylcholinesterase	Fluorescence	23
 <p>LISA-101</p>	γ -glutamyl cyclotransferase	Fluorescence	24
 <p>Probe 3</p>	Epoxy-hydrolase	Fluorescence	25
 <p>Resorufin Sulfatase (RS)</p>	Sulfatase	Fluorescence	26
 <p>Pro-fluoroprobe</p>	Thrombin protease	Fluorescence	27
 <p>Probe 1</p>	Glucose	Fluorescence	28
 <p>PR1</p>	Reactive Oxygen, Nitrogen (ONOO ⁻)	Fluorescence and Colorimetric	29

 <p>Pinkment-OTBS</p>	(ONOO ⁻ and F ⁻)	Fluorescence and Colorimetric	29
 <p>Pinkment-OAC</p>	Esterase and H ₂ O ₂	Fluorescence and Colorimetric	29
 <p>Probe 1</p>	Carboxylesterase	Fluorescence	30
 <p>NBCD</p>	H ₂ O ₂	Fluorescence	31
 <p>Res-GlcGcase</p>	β-Glucocerebrosidase	Fluorescence	32
 <p>Cys probe</p>	Cysteine	Fluorescence	33
 <p>AR</p>	Phosphite and Nickel oxide	Fluorescence	34

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