

Supporting Information

A Near-infrared Fluorescent Probe Based on Hemicyanine Dye with Oxazolidine Switch for Mitochondrial pH Detection

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1 NMR Spectra for Probe AH+

1.1 ¹H NMR Spectra for probe AH+

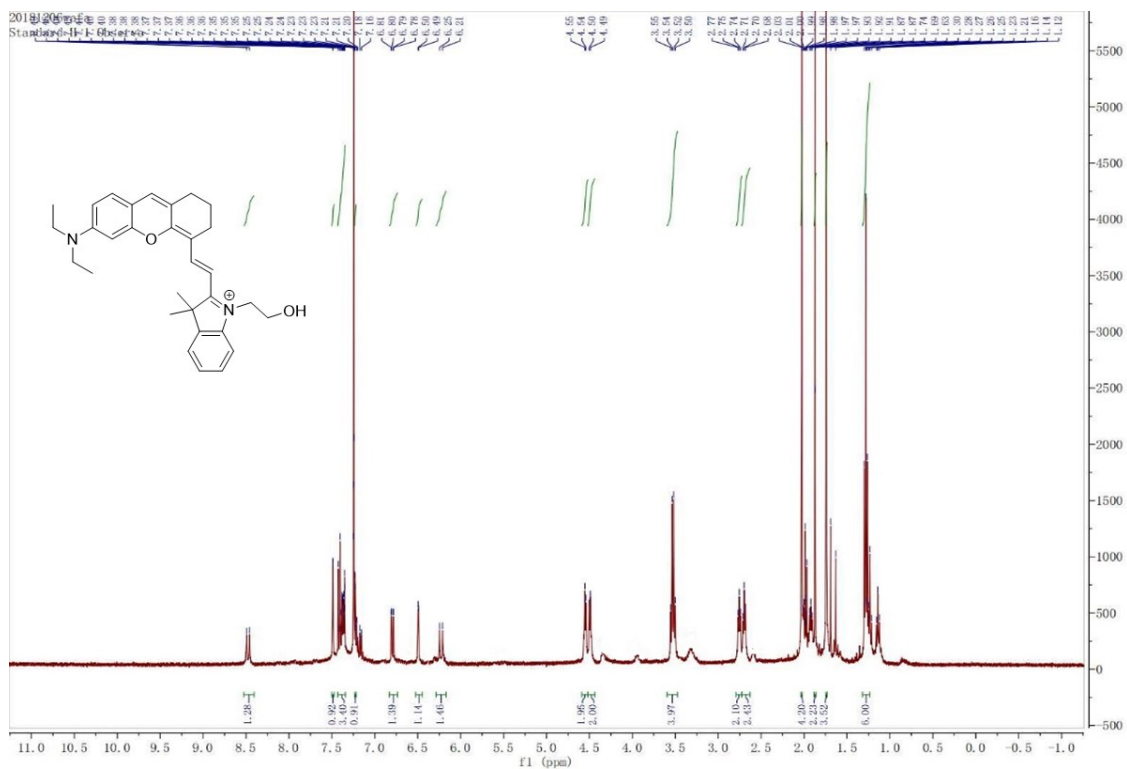


Figure S1: ¹H NMR spectrum of probe AH+ based on rhodamine dye in chloroform-d solution.

1.2 ^{13}C NMR Spectra for probe AH+

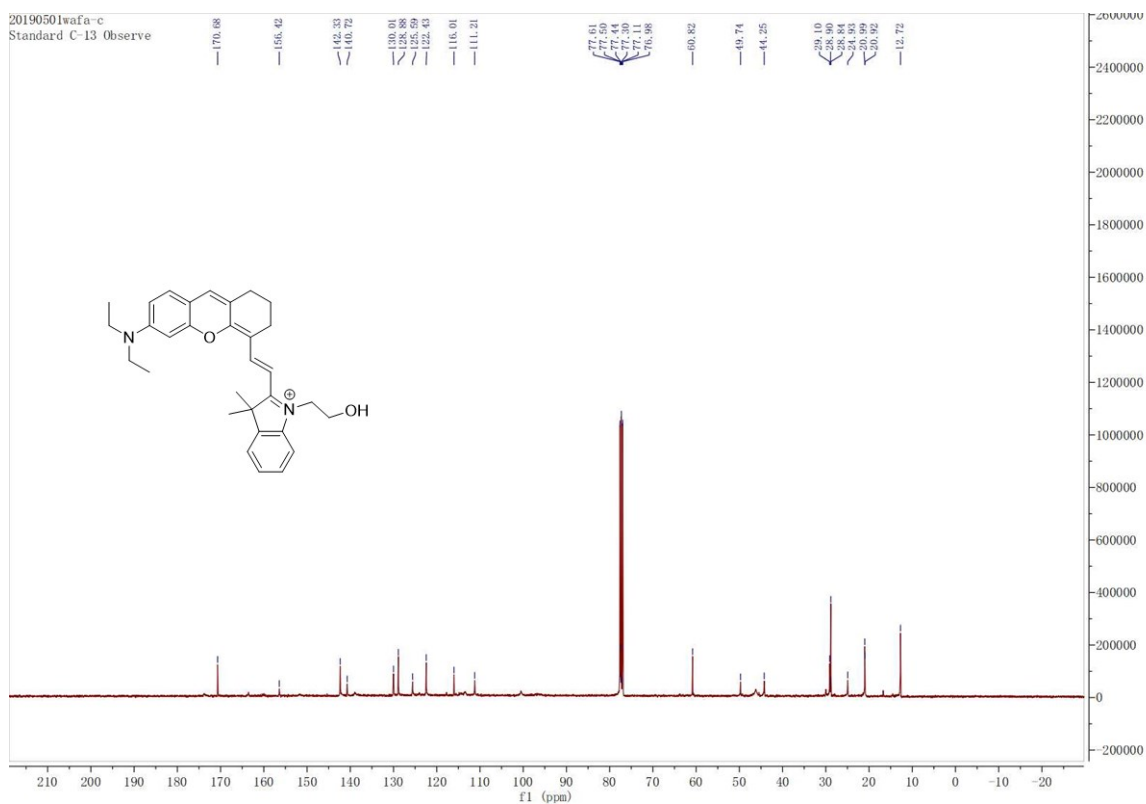


Figure S2: ^{13}C NMR spectrum of probe AH+ based on rhodamine dye in chloroform-d solution.

1.3 ESI-MS spectra for probe AH+

LCQ Instrument Control

S#: 7437 IT: 5.139 ST: 1.56

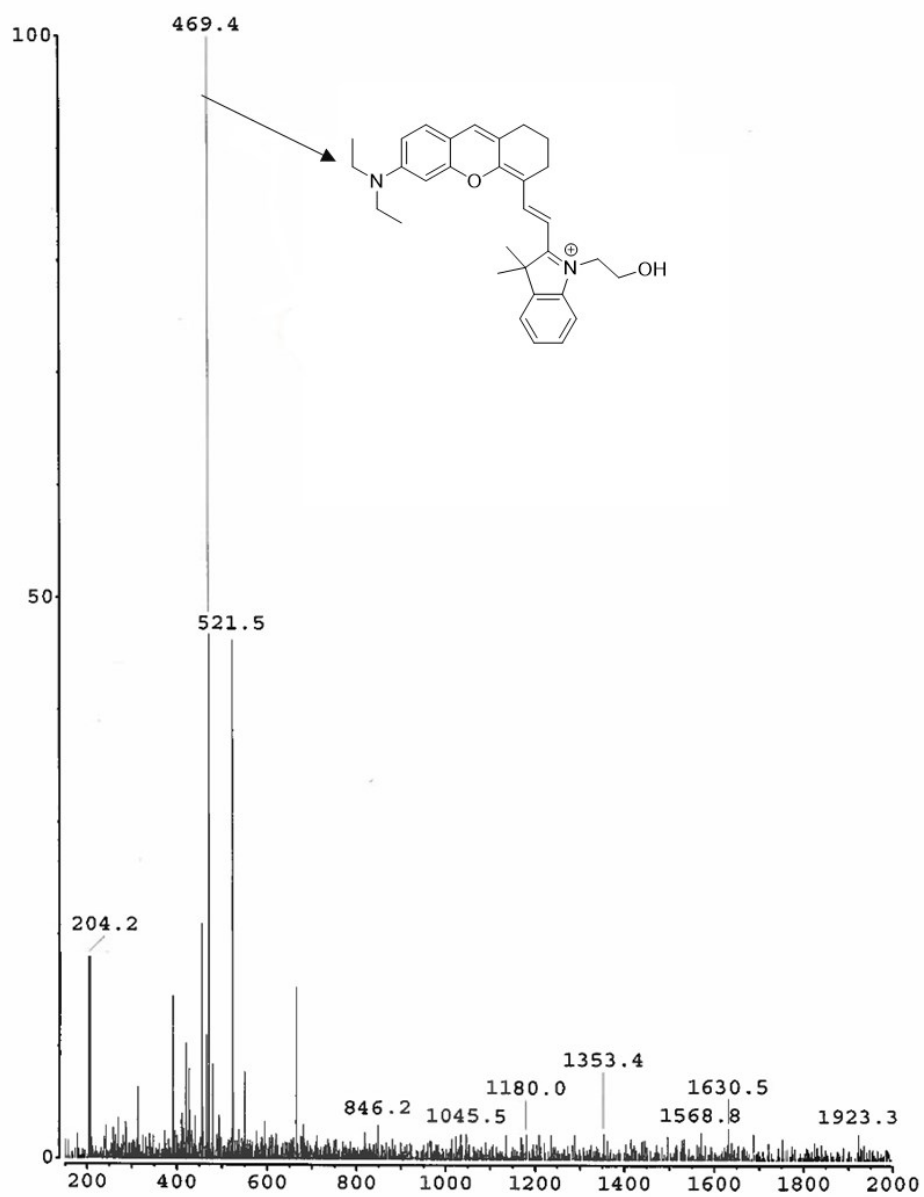


Figure S3. Electrospray Ionization mass spectrum of probe AH+.

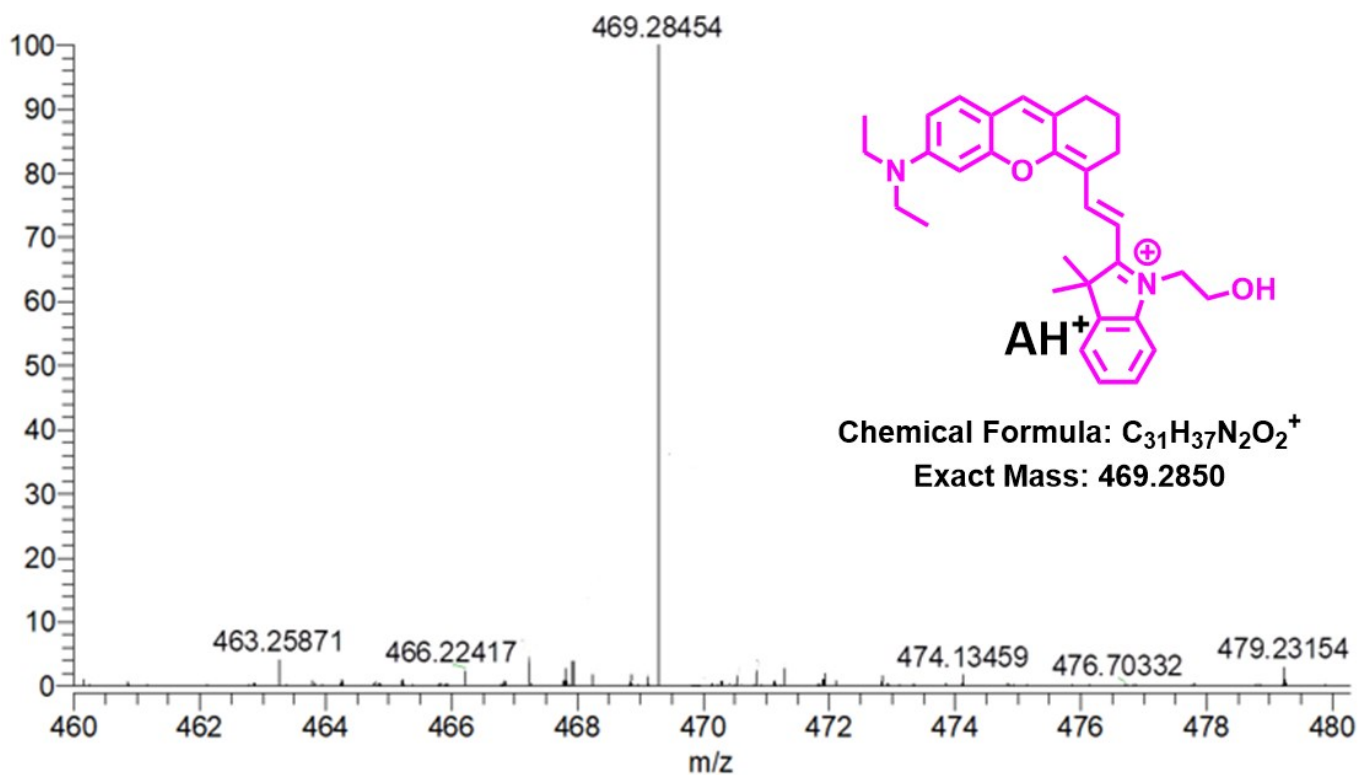


Figure S4. High-resolution electrospray ionization mass spectrum of probe **AH⁺**.

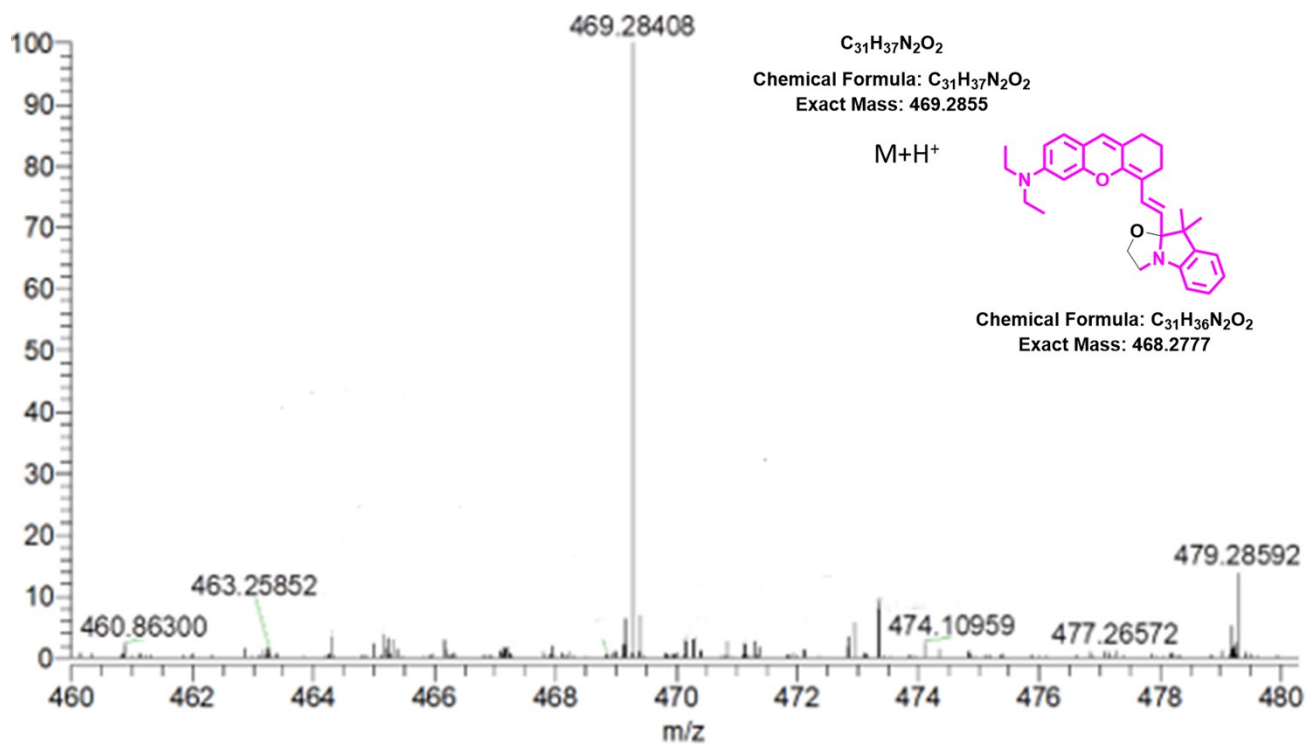


Figure S5. High-resolution electrospray ionization mass spectrum of probe **A** under a basic condition.

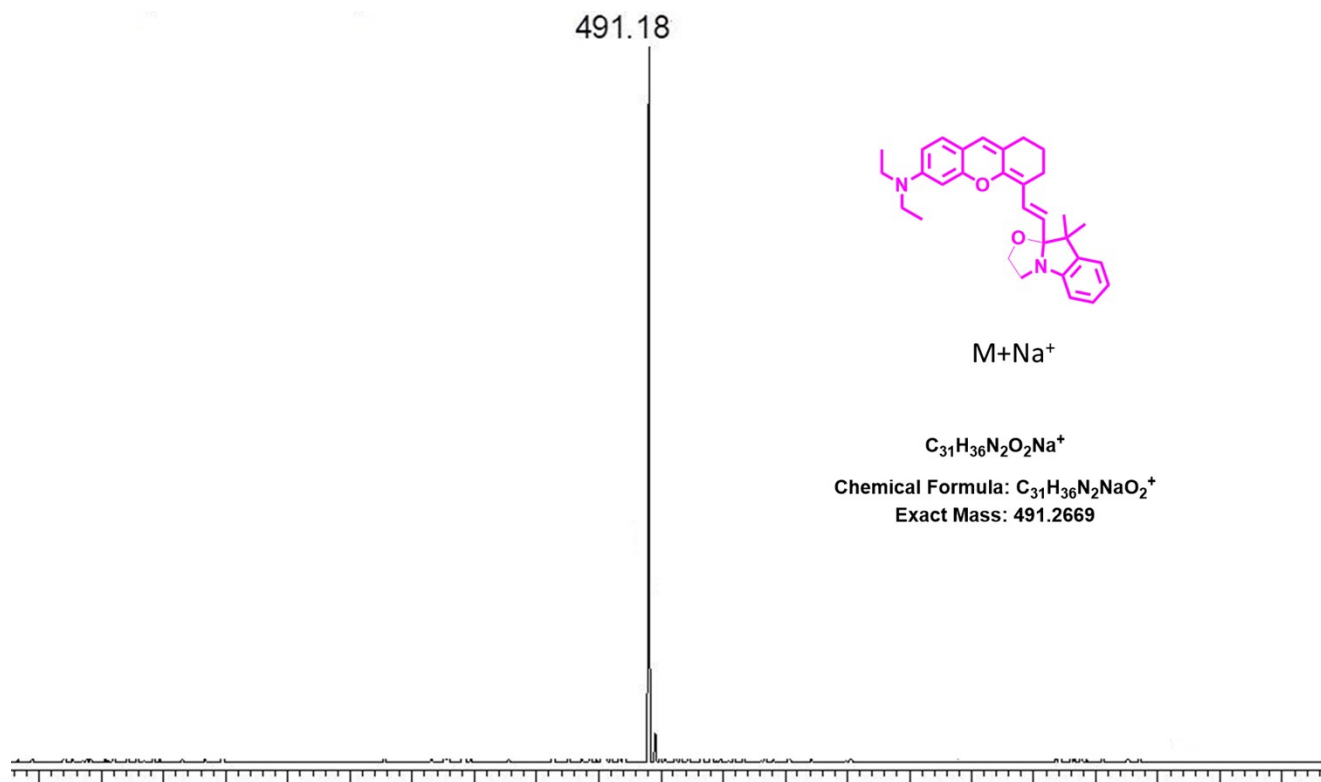
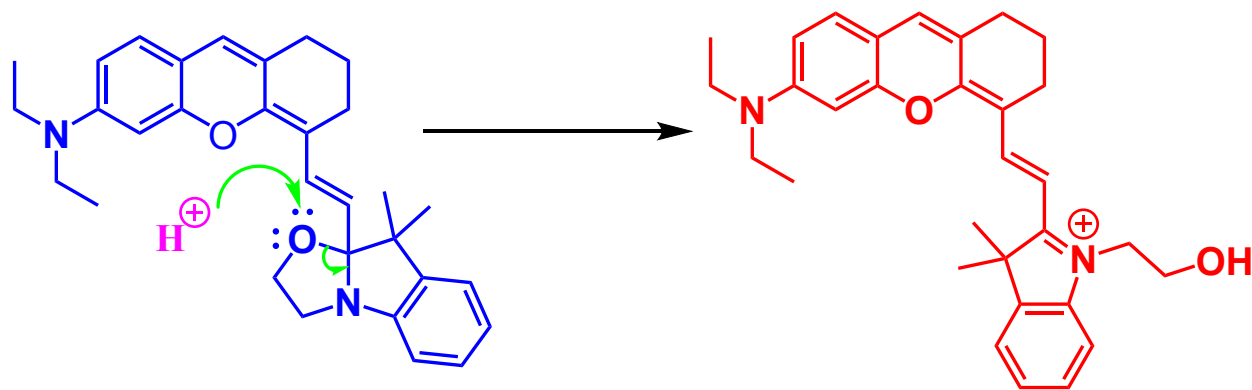


Figure S6. Matrix-assisted Laser Desorption Ionization-time of flight mass spectrum of probe **A** under a basic condition.

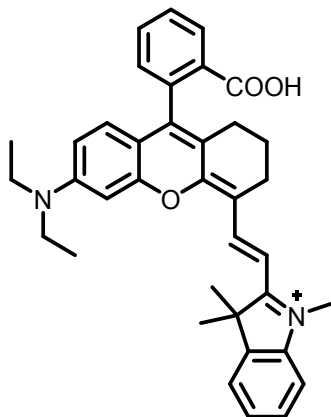


Scheme S1. Ring opening of an oxazolidine switch of the probe is triggered by acidic pH condition.

2 Calculation

2.1 Calculation of Quantum Yield

We chose the near-infrared hemicyanine dye shown in Figure S3 as fluorescent standard dye¹ to calculate fluorescence quantum yields of probe **AH**⁺.



Hemicyanine dye ($\Phi = 0.41$ in EtOH)

The structure of near- infrared hemicyanine¹

The UV-Vis absorption spectrum of probe **AH**⁺ was collected in the range from 300 to 800 nm with increments of 1 nm. The UV-Vis absorption spectra measured in freshly prepared buffer. Citrate-phosphate buffer was used for acidic pH 5.0, and phosphate buffer for pH 10.7. The fluorescence spectra of probe **AH**⁺ was collected under the excitation wavelength of 670 nm. The excitation and emission slit widths were set to 5 nm. NIR hemicyanine was chosen as reference standards to calculate the fluorescence quantum yield of probe **AH**⁺, in ethanol and buffer solutions. The absorbance and fluorescence spectra of the standard dye was measured in pH 7.4 PBS buffer with 5% ethanol. The absorbance and fluorescence spectra of the probe **A** was measured in pH 5.0 citrate-phosphate buffer and in pH 7.4 PBS buffer containing 5% EtOH. The absorbance was kept between 0.05 and 0.1 in order to obtain optimized data. The probe samples and reference were freshly prepared under identical conditions. The fluorescence quantum yields were calculated according to literature⁴ using the equation 1 below¹ :

$$\Phi_{F(X)} = \Phi_{F(S)} \left(\frac{A_S F_X}{A_X F_S} \right) \left(\frac{n_X}{n_S} \right)^2 \quad (1)$$

Where ΦF is the fluorescence quantum yield, A is the absorbance at the excitation wavelength, F is the area under the corrected emission curve, and n is the refractive index of the solvents used. Subscripts s and x refer to the standard and to the unknown, respectively.

2.2 Calculation of probe pK_a value by fluorometric titration

The fluorometric titration as a function of pH was obtained fluorescence spectra. The equation (2) below was used to calculate the pK_a value of probe AH^+ .

$$F = \frac{F_{\min} [H^+]^n + F_{\max} K_a}{K_a + [H^+]^n} \quad (2)$$

The expression of the steady-state fluorescence intensity F as a function of the proton concentration has been extended for the case of n : complex between H^+ and a fluorescent dye. Where F_{\min} and F_{\max} are the fluorescence intensities at maximal and minimal H^+ concentrations, respectively.

n is apparent stoichiometry of H^+ binding to the probe A . Nonlinear fitting of equation expressed above to the fluorescence titration data was plotted as a function of H^+ concentration²⁻³.

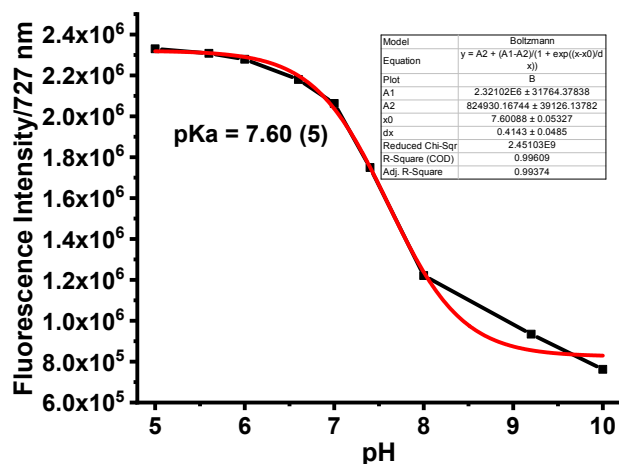


Figure S7. Plot curve of fluorescence intensity of 5 μ M probe AH^+ versus pH

3 Solvent Effect

The effect of ethanol percentage was investigated in water-ethanol mixed solution on dye fluorescence intensity (Figure S5). Increase of the percentages of ethanol from 10% to 90% resulted in enhancement of fluorescence intensity of the dyes because water increase percentages can effectively prevent fluorescence quenching due to dye aggregation in aqueous solutions.

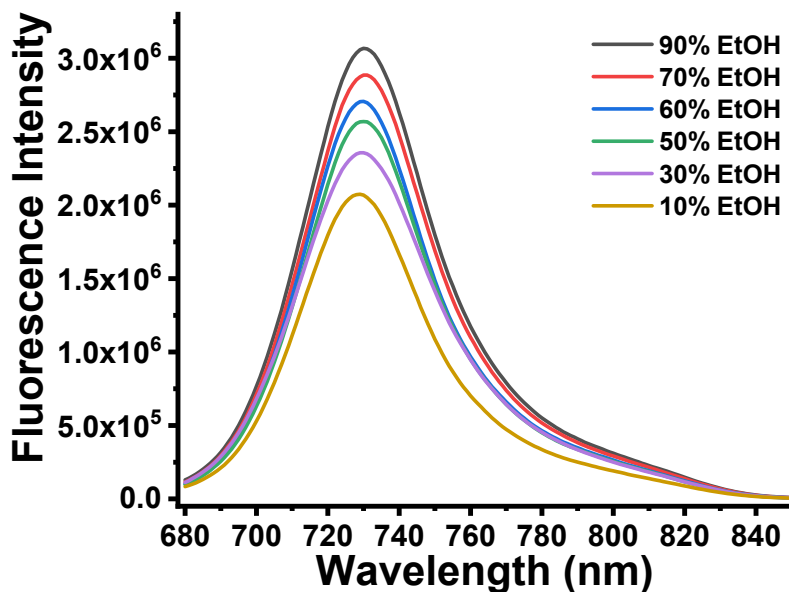
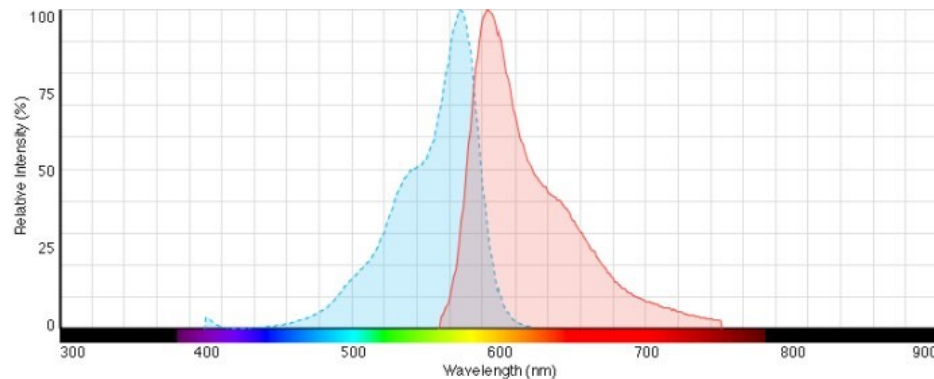


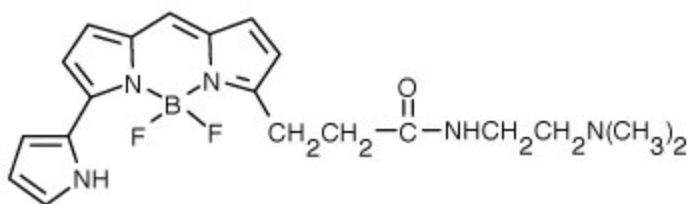
Figure S8. Fluorescence spectra of 5 μM probe AH⁺ in pH 7.0 buffers with different percentages of ethanol under excitation at 670 nm.

4. Commercial LysoTracker Red DND-99 emission spectra from Thermo Fisher Scientific

<https://www.thermofisher.com/order/catalog/product/L7528#/L7528>



Absorption and fluorescence spectra of LysoTracker Red DND-99 which was obtained from the link above.

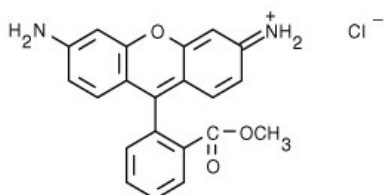


Chemical structure of Lysotracker Red DND-99.

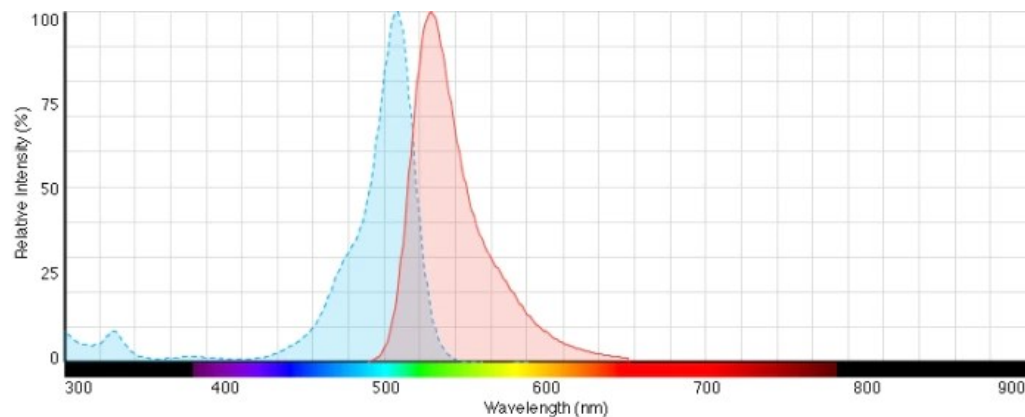
In the cellular fluorescence imaging experiments, we excited Lysotracker Red DND-99 at 559 nm and collected fluorescence from 600 nm to 650 nm while we excited the probe at 635 nm and collected fluorescence from 725 nm to 775 nm. The absorption of Lysotracker Red DND-99 at 635 nm is extremely small according to its absorption peak above which was obtained from the website link above. The collected fluorescence of Lysotracker Red DND-99 from 600 nm to 650 nm is not overlapped with that of the probe from 725 nm to 775 nm (Figure 7).

5. Mitochondria-specific Rhodamine 123.

Rhodamine dye 123 has been commercialized as a mitochondria-targeting probe by ThermoFisher Scientific. The detailed information is available from the link below.



<https://www.thermofisher.com/order/catalog/product/R302?SID=srch-srp-R302#/R302?SID=srch-srp-R302>



Absorption and fluorescence spectra of Rhodamine 123 which was obtained from the link above.

In cellular fluorescence imaging experiments, we excited Rhodamine 123 at 488 nm and collected fluorescence from 500 nm to 550 nm while we excited the probe at 635 nm and collected fluorescence from 725 nm to 775 nm. The absorption of Rhodamine 123 at 635 nm is almost zero according to its absorption peak above which was obtained from the website link above. The collected fluorescence of Rhodamine 123 from 500 nm to 550 nm is not overlapped with that of the probe from 725 nm to 775 nm (Figure 7).

6. Golgi-specific fluorescent probe (Golgi-GFP).

A Golgi-specific fluorescent probe (Golgi-GFP, BacMam) from ThermoFisher Scientific was used for colocalization experiment with the probe.

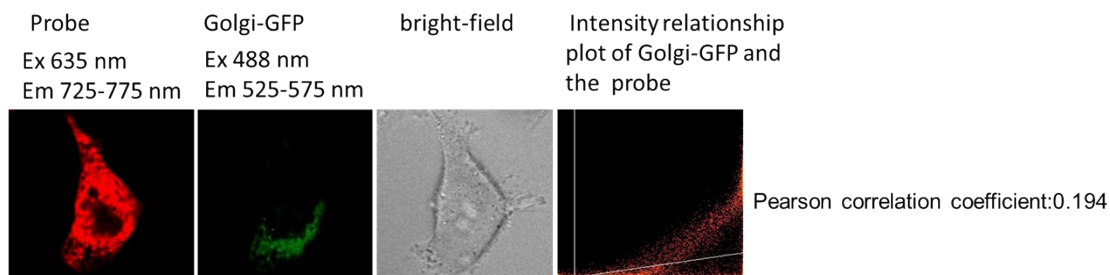


Figure S9. Confocal microscopic fluorescence images and merged images of the probe colocalized with Golgi-GFP, BacMam in HeLa cells. Colocalization scatterplot of the probe with Golgi-GFP, BacMam. Pearson correlation coefficient: 0.194

7. ER-specific fluorescent probe (ER-Tracker™ Green, glibenclamide BODIPY® FL).

ER-Tracker™ Green dye from TheromoFisher Scientific™ was employed for colocalization experiment with the probe.

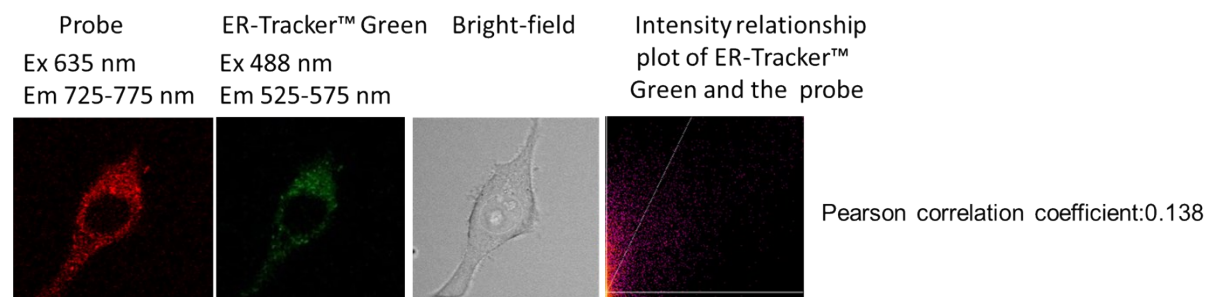


Figure S10. Confocal microscopic fluorescence images and merged images of the probe colocalized with Golgi-GFP, BacMam in HeLa cells. Colocalization scatterplot of the probe with ER-Tracker™ Green (glibenclamide BODIPY® FL). Pearson correlation coefficient: 0.138

8. Computationally derived structures for probe A and AH⁺.

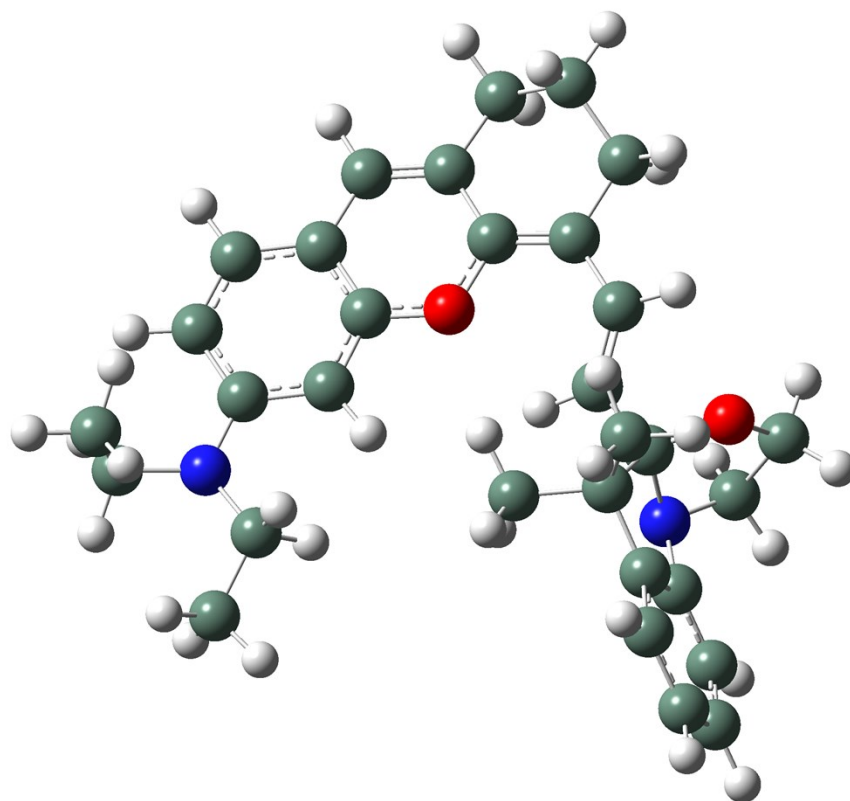


Figure S11. Drawing of probe A with atoms represented as spheres of arbitrary size (H-white, C-grey, N-blue and O-red) using the GaussView program.

Table S1. Atomic coordinates for probe A.

| Row | Symbol | X | Y | Z |
|-----|--------|----------|----------|----------|
| 1 | C | 5.065562 | -0.30195 | 0.231392 |
| 2 | C | 3.918238 | -1.1465 | 0.093227 |
| 3 | C | 2.655842 | -0.51606 | 0.026288 |
| 4 | C | 2.570472 | 0.857508 | 0.108113 |
| 5 | C | 3.691188 | 1.695034 | 0.248157 |
| 6 | C | 4.948799 | 1.060704 | 0.30816 |
| 7 | O | 1.328055 | 1.402961 | 0.032912 |
| 8 | C | 1.107847 | 2.734944 | 0.080026 |
| 9 | C | 2.208399 | 3.618407 | 0.21323 |
| 10 | C | 3.47284 | 3.086059 | 0.292677 |
| 11 | C | -0.22955 | 3.161402 | -0.01656 |
| 12 | C | -0.4802 | 4.657831 | -0.00213 |
| 13 | C | 0.703405 | 5.460642 | -0.52496 |
| 14 | C | 1.956349 | 5.098755 | 0.261899 |
| 15 | C | -1.3436 | 2.329105 | -0.09771 |
| 16 | C | -1.44635 | 0.936341 | -0.05956 |
| 17 | N | 4.038025 | -2.50149 | 0.029835 |
| 18 | C | 5.348935 | -3.1135 | -0.15602 |
| 19 | C | 2.849696 | -3.2905 | -0.31294 |
| 20 | C | 2.999117 | -4.79088 | -0.13598 |
| 21 | C | 5.839974 | -3.04814 | -1.59754 |
| 22 | C | -2.64245 | 0.23876 | -0.12328 |
| 23 | C | -4.06793 | 0.769153 | -0.23452 |
| 24 | C | -4.86679 | -0.51027 | -0.22335 |
| 25 | C | -3.99988 | -1.59284 | -0.11677 |
| 26 | N | -2.68327 | -1.11415 | -0.07904 |
| 27 | C | -6.23192 | -0.71838 | -0.28953 |
| 28 | C | -6.71414 | -2.02998 | -0.24242 |
| 29 | C | -5.83195 | -3.10373 | -0.12632 |
| 30 | C | -4.45217 | -2.9037 | -0.05959 |
| 31 | C | -4.45013 | 1.626452 | 0.982585 |
| 32 | C | -4.29159 | 1.521784 | -1.55506 |
| 33 | C | -1.5289 | -1.9787 | 0.048135 |
| 34 | C | -1.10646 | -2.12267 | 1.502644 |
| 35 | O | 0.054533 | -2.92997 | 1.504442 |
| 36 | H | 6.057687 | -0.73129 | 0.266281 |
| 37 | H | 1.741027 | -1.07871 | -0.09343 |
| 38 | H | 5.840598 | 1.672691 | 0.408444 |
| 39 | H | 4.328261 | 3.750475 | 0.389404 |

| Row | Symbol | X | Y | Z |
|-----|--------|----------|----------|----------|
| 40 | H | -1.37601 | 4.87539 | -0.59212 |
| 41 | H | -0.70675 | 4.979214 | 1.024217 |
| 42 | H | 0.499696 | 6.532647 | -0.44503 |
| 43 | H | 0.860348 | 5.240429 | -1.58812 |
| 44 | H | 2.835042 | 5.629837 | -0.11595 |
| 45 | H | 1.832084 | 5.40953 | 1.308209 |
| 46 | H | -2.27694 | 2.876612 | -0.18611 |
| 47 | H | -0.5422 | 0.357509 | 0.04278 |
| 48 | H | 6.059746 | -2.64612 | 0.526935 |
| 49 | H | 5.282479 | -4.14878 | 0.175449 |
| 50 | H | 2.030745 | -2.96412 | 0.328549 |
| 51 | H | 2.552468 | -3.06533 | -1.34825 |
| 52 | H | 2.028251 | -5.25478 | -0.32872 |
| 53 | H | 3.714325 | -5.24028 | -0.82917 |
| 54 | H | 3.290911 | -5.05344 | 0.884998 |
| 55 | H | 6.820545 | -3.52354 | -1.69084 |
| 56 | H | 5.149922 | -3.56559 | -2.27037 |
| 57 | H | 5.930477 | -2.01398 | -1.94166 |
| 58 | H | -6.92092 | 0.117394 | -0.37411 |
| 59 | H | -7.78302 | -2.21377 | -0.2923 |
| 60 | H | -6.22068 | -4.11683 | -0.08422 |
| 61 | H | -3.77821 | -3.74774 | 0.040142 |
| 62 | H | -3.88934 | 2.561422 | 1.020171 |
| 63 | H | -5.51325 | 1.874639 | 0.932159 |
| 64 | H | -4.27199 | 1.083253 | 1.913689 |
| 65 | H | -3.71512 | 2.446922 | -1.60169 |
| 66 | H | -4.0121 | 0.901822 | -2.41011 |
| 67 | H | -5.34914 | 1.779607 | -1.65286 |
| 68 | H | -1.7752 | -2.95251 | -0.3764 |
| 69 | H | -0.71222 | -1.57685 | -0.55485 |
| 70 | H | -0.90405 | -1.13474 | 1.935272 |
| 71 | H | -1.91692 | -2.5848 | 2.080605 |
| 72 | H | 0.349497 | -3.06125 | 2.410781 |

Table S2. Excitation energies and oscillator strengths for probe A.

| Excited State | Nature | E (eV) | λ (nm) | f | Orbital transitions | Normalized coefficient |
|---------------|-----------|--------|----------------|--------|---------------------------------------------------------------|--------------------------------------------------------|
| 1 | Singlet-A | 2.9705 | 417.39 | 0.5427 | 126 ->127 | 0.70254 |
| 2 | Singlet-A | 3.8205 | 324.52 | 0.0945 | 124 ->127 125 ->127 126 ->128 126 ->129 | 0.12029 -0.16699 0.49384 -0.44206 |
| 3 | Singlet-A | 3.9032 | 317.65 | 0.0432 | 124 ->127 125 ->127 126 ->128 126 ->130 | -0.27488 0.45655 0.10080 -0.42340 |
| 4 | Singlet-A | 4.0144 | 308.85 | 0.0071 | 124 ->127 125 ->127 | 0.55750 0.41595 |
| 5 | Singlet-A | 4.1347 | 299.86 | 0.0063 | 126 ->128 126 ->129 | 0.48025 0.51226 |
| 6 | Singlet-A | 4.2653 | 290.68 | 0.3107 | 124 ->127 125 ->127 126 ->129 126 ->130 126 ->132 | -0.28800 0.27224 -0.11983 0.50848 -0.10521 |

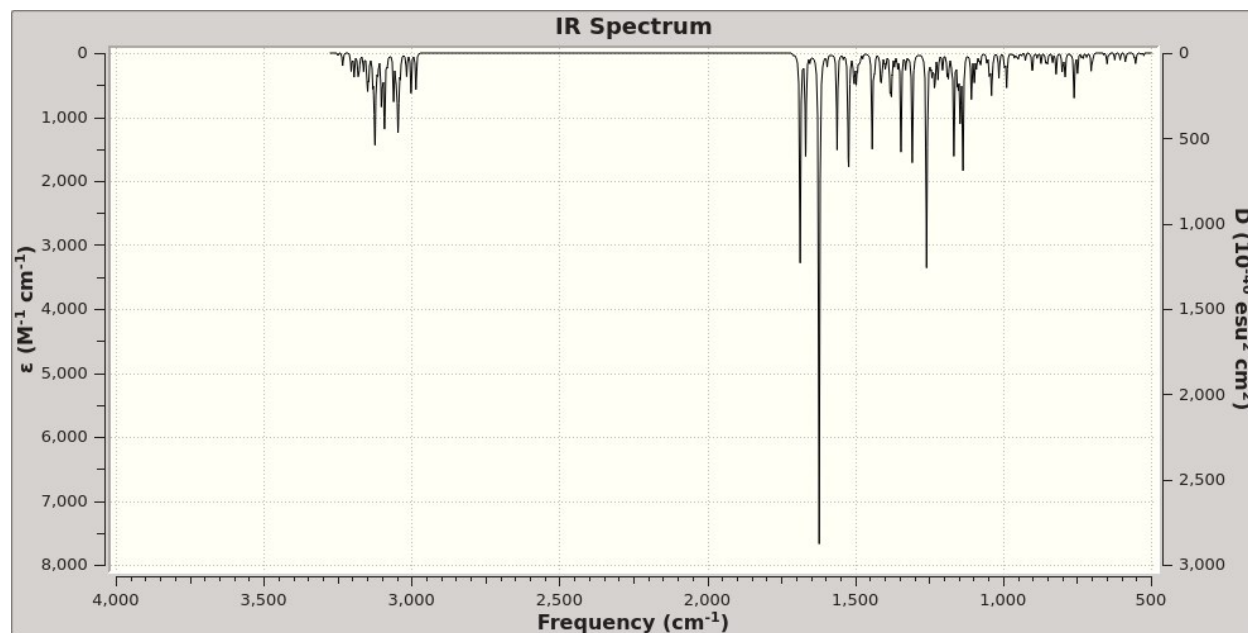


Figure S12. Calculated (top) FTIR spectrum of probe A.

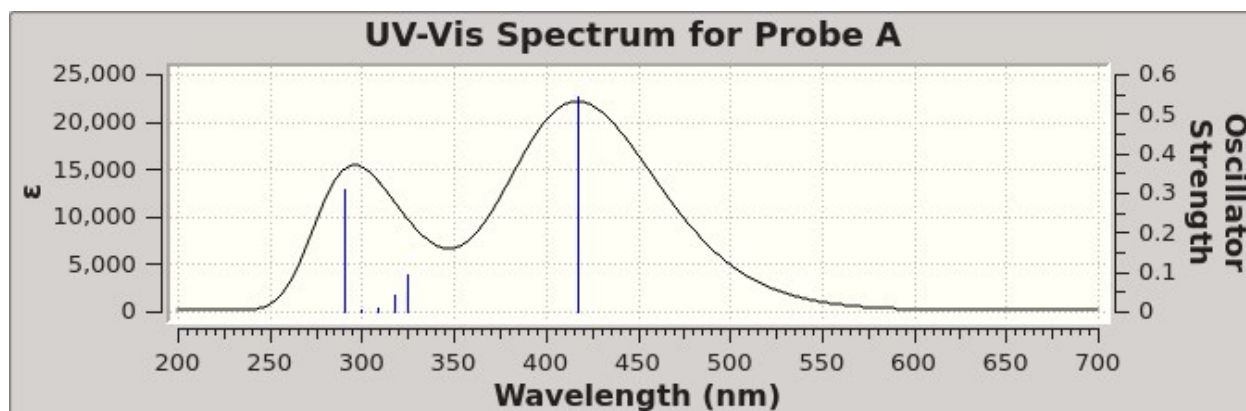


Figure S13. Calculated UV-Vis spectrum for probe A in water.

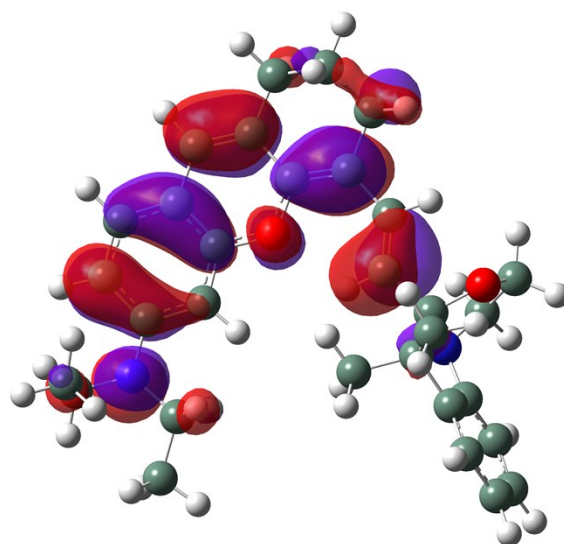


Figure S14. LCAO for level 126 for probe A involved with the transition noted as Excited State 1 in Table S2.

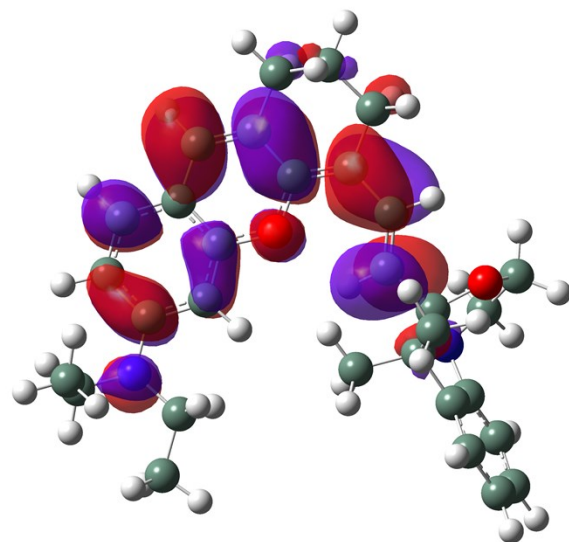


Figure S15. LCAO for level 127 for probe **A** involved with the transition noted as Excited State 1 in Table S2.

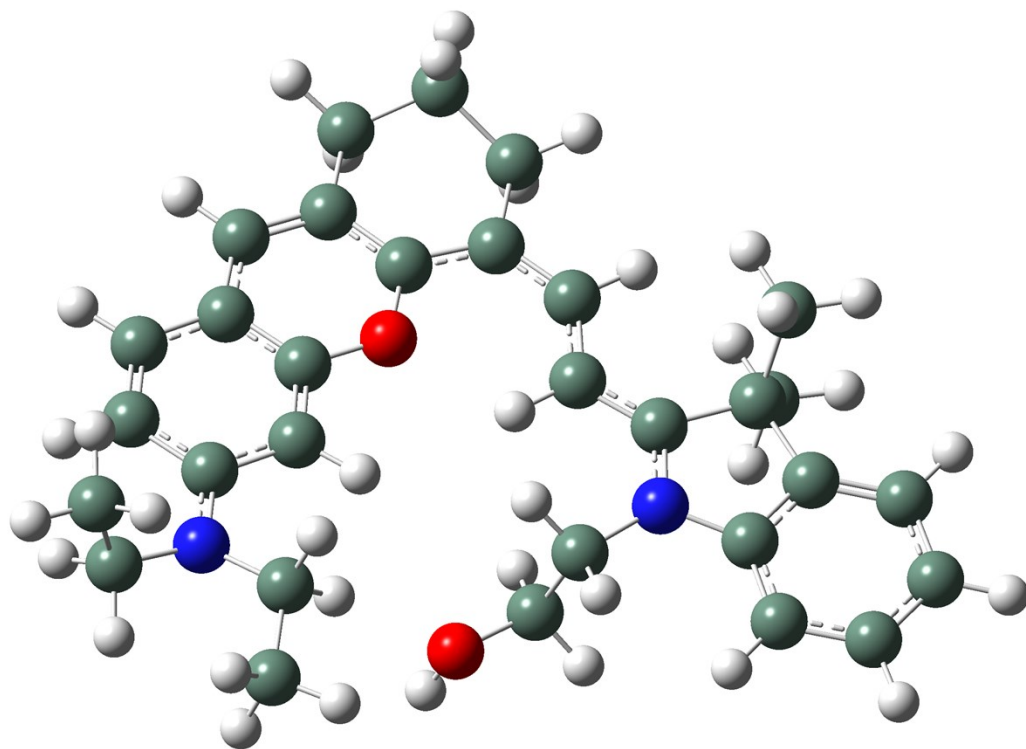


Figure S16. Drawing of probe AH^+ with atoms represented as spheres of arbitrary size (H-white, C-grey, N-blue and O-red) using the GaussView program.

Table S3. Atomic coordinates for probe **AH⁺**.

| Row | Symbol | X | Y | Z |
|-----|--------|----------|----------|----------|
| 1 | C | 5.065562 | -0.30195 | 0.231392 |
| 2 | C | 3.918238 | -1.1465 | 0.093227 |
| 3 | C | 2.655842 | -0.51606 | 0.026288 |
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| 7 | O | 1.328055 | 1.402961 | 0.032912 |
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| 9 | C | 2.208399 | 3.618407 | 0.21323 |
| 10 | C | 3.47284 | 3.086059 | 0.292677 |
| 11 | C | -0.22955 | 3.161402 | -0.01656 |
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| 13 | C | 0.703405 | 5.460642 | -0.52496 |
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| 16 | C | -1.44635 | 0.936341 | -0.05956 |
| 17 | N | 4.038025 | -2.50149 | 0.029835 |
| 18 | C | 5.348935 | -3.1135 | -0.15602 |
| 19 | C | 2.849696 | -3.2905 | -0.31294 |
| 20 | C | 2.999117 | -4.79088 | -0.13598 |
| 21 | C | 5.839974 | -3.04814 | -1.59754 |
| 22 | C | -2.64245 | 0.23876 | -0.12328 |
| 23 | C | -4.06793 | 0.769153 | -0.23452 |
| 24 | C | -4.86679 | -0.51027 | -0.22335 |
| 25 | C | -3.99988 | -1.59284 | -0.11677 |
| 26 | N | -2.68327 | -1.11415 | -0.07904 |
| 27 | C | -6.23192 | -0.71838 | -0.28953 |
| 28 | C | -6.71414 | -2.02998 | -0.24242 |
| 29 | C | -5.83195 | -3.10373 | -0.12632 |
| 30 | C | -4.45217 | -2.9037 | -0.05959 |
| 31 | C | -4.45013 | 1.626452 | 0.982585 |
| 32 | C | -4.29159 | 1.521784 | -1.55506 |
| 33 | C | -1.5289 | -1.9787 | 0.048135 |
| 34 | C | -1.10646 | -2.12267 | 1.502644 |
| 35 | O | 0.054533 | -2.92997 | 1.504442 |
| 36 | H | 6.057687 | -0.73129 | 0.266281 |
| 37 | H | 1.741027 | -1.07871 | -0.09343 |
| 38 | H | 5.840598 | 1.672691 | 0.408444 |
| 39 | H | 4.328261 | 3.750475 | 0.389404 |

| Row | Symbol | X | Y | Z |
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| 40 | H | -1.37601 | 4.87539 | -0.59212 |
| 41 | H | -0.70675 | 4.979214 | 1.024217 |
| 42 | H | 0.499696 | 6.532647 | -0.44503 |
| 43 | H | 0.860348 | 5.240429 | -1.58812 |
| 44 | H | 2.835042 | 5.629837 | -0.11595 |
| 45 | H | 1.832084 | 5.40953 | 1.308209 |
| 46 | H | -2.27694 | 2.876612 | -0.18611 |
| 47 | H | -0.5422 | 0.357509 | 0.04278 |
| 48 | H | 6.059746 | -2.64612 | 0.526935 |
| 49 | H | 5.282479 | -4.14878 | 0.175449 |
| 50 | H | 2.030745 | -2.96412 | 0.328549 |
| 51 | H | 2.552468 | -3.06533 | -1.34825 |
| 52 | H | 2.028251 | -5.25478 | -0.32872 |
| 53 | H | 3.714325 | -5.24028 | -0.82917 |
| 54 | H | 3.290911 | -5.05344 | 0.884998 |
| 55 | H | 6.820545 | -3.52354 | -1.69084 |
| 56 | H | 5.149922 | -3.56559 | -2.27037 |
| 57 | H | 5.930477 | -2.01398 | -1.94166 |
| 58 | H | -6.92092 | 0.117394 | -0.37411 |
| 59 | H | -7.78302 | -2.21377 | -0.2923 |
| 60 | H | -6.22068 | -4.11683 | -0.08422 |
| 61 | H | -3.77821 | -3.74774 | 0.040142 |
| 62 | H | -3.88934 | 2.561422 | 1.020171 |
| 63 | H | -5.51325 | 1.874639 | 0.932159 |
| 64 | H | -4.27199 | 1.083253 | 1.913689 |
| 65 | H | -3.71512 | 2.446922 | -1.60169 |
| 66 | H | -4.0121 | 0.901822 | -2.41011 |
| 67 | H | -5.34914 | 1.779607 | -1.65286 |
| 68 | H | -1.7752 | -2.95251 | -0.3764 |
| 69 | H | -0.71222 | -1.57685 | -0.55485 |
| 70 | H | -0.90405 | -1.13474 | 1.935272 |
| 71 | H | -1.91692 | -2.5848 | 2.080605 |
| 72 | H | 0.349497 | -3.06125 | 2.410781 |

Table S4. Excitation energies and oscillator strengths for probe AH^+ .

| Excited State | Nature | E (eV) | λ (nm) | f | Orbital transitions | Normalized coefficient |
|---------------|-----------|--------|----------------|--------|---------------------------------------------------------------|-------------------------------------------------------|
| 1 | Singlet-A | 2.0899 | 593.26 | 0.7993 | 126 ->127 | 0.70631 |
| 2 | Singlet-A | 2.8527 | 434.62 | 0.2023 | 125 ->127 126 ->128 | 0.68787 0.12101 |
| 3 | Singlet-A | 3.6147 | 343.00 | 0.1318 | 122 ->127 124 ->127 126 ->128 | -0.24296 -0.42811 0.49106 |
| 4 | Singlet-A | 3.7149 | 333.75 | 0.3055 | 124 ->127 125 ->127 126 ->128 | 0.53865 -0.10258 0.41981 |
| 5 | Singlet-A | 3.9046 | 317.54 | 0.0383 | 123 ->127 126 ->129 | 0.66762 -0.16028 |
| 6 | Singlet-A | 3.9614 | 312.98 | 0.1464 | 122 ->127 123 ->127 125 ->128 126 ->128 126 ->130 | 0.58588 0.11095 -0.22459 0.19025 -0.14800 |

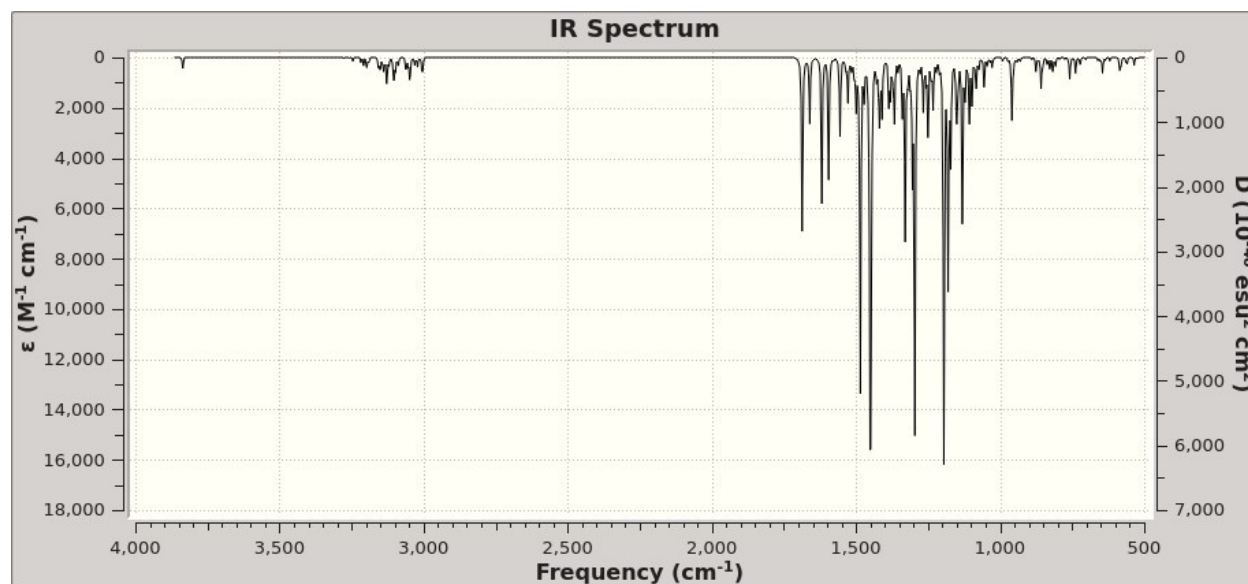


Figure S17. Calculated (top) FTIR spectrum of probe AH^+ .

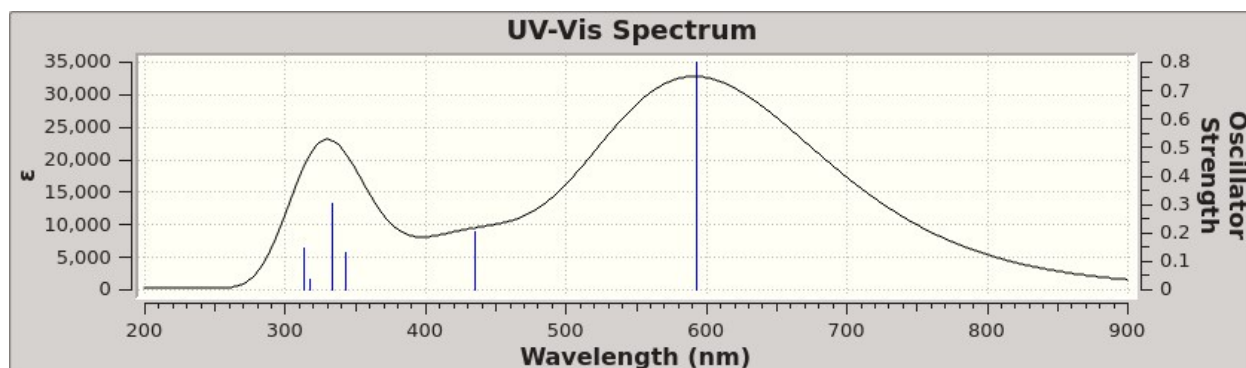


Figure S18. Calculated UV-Vis spectrum for probe AH^+ in water.

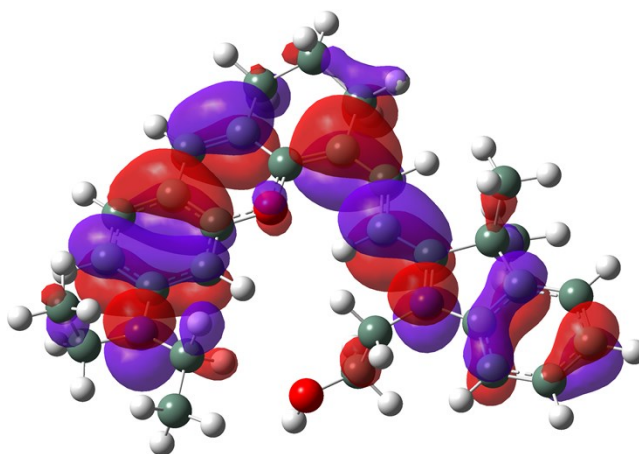


Figure S19. LCAO for level 126 for probe AH^+ involved with the transition noted as Excited State 1 in Table S4.

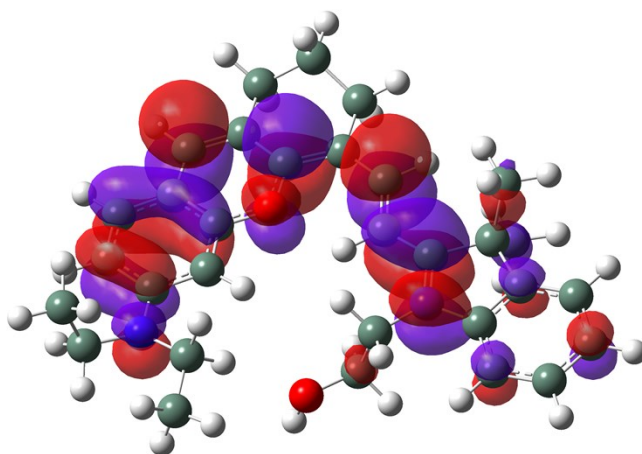


Figure S20. LCAO for level 127 for probe AH^+ involved with the transition noted as Excited State 1 in Table S4.

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