

SUPPORTING INFORMATION

Overriding Phthalate Decomposition when Exploring Mycophenolic acid

Intermediates as Selenium-based ROS Biological Probes

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Standard protocol for preparation of Various ROS and RNS:

Generation of OCl⁻: The source of NaOCl was commercial bleach. The concentration of the OCl⁻ stock solution was determined by measuring the absorbance at 209 nm with a molar extinction coefficient of 350 M⁻¹ cm⁻¹.

Generation of H₂O₂: H₂O₂ solution was added directly. The concentration of H₂O₂ was determined by measuring the absorbance at 240 nm with a molar extinction coefficient of 43.6 M⁻¹cm⁻¹.

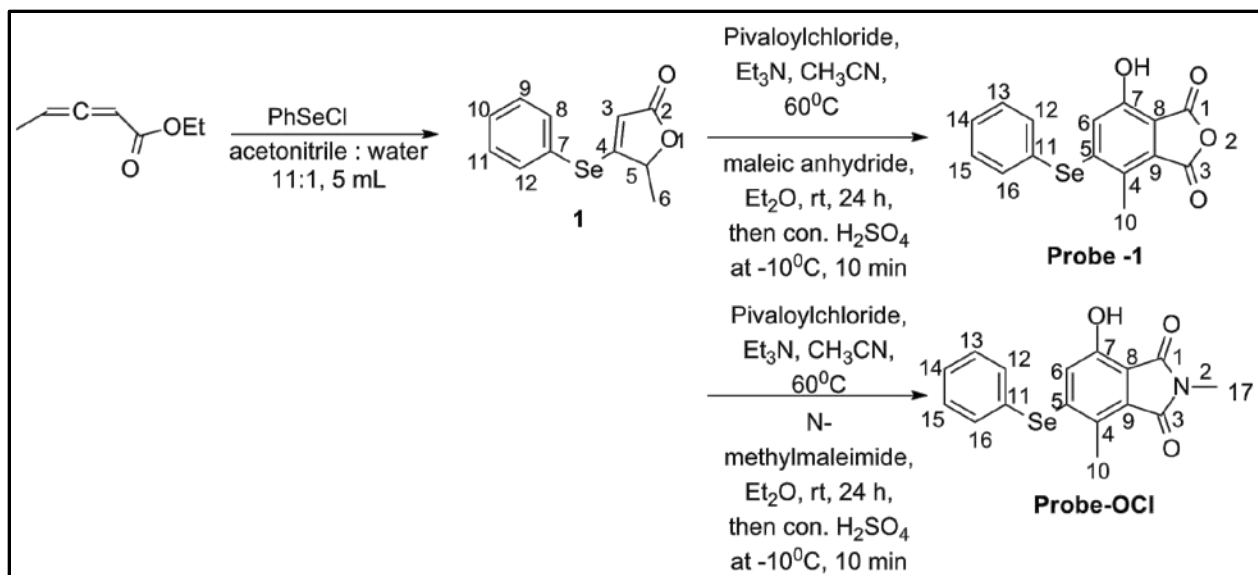
Generation of ^tBuOOH: The commercial available *tert*-Butyl hydroperoxide solution was diluted with deionized water.

Generation of O₂^{•-}: Solid potassium superoxide was used as superoxide radical anion source.

Generation of •OH: Hydroxyl radical (•OH) was generated by the Fenton reaction. To generate •OH, ferrous chloride was added in the presence of 10 equiv of H₂O₂. The concentration of •OH was equal to the Fe (II) concentration.

Generation of peroxynitrite (ONOO⁻): A mixture of sodium nitrite (0.6 M) and hydrogen peroxide (0.7 M) was acidified with hydrochloric acid (0.6 M), and sodium hydroxide (1.5 M) was added within 1~2 s to make the solution alkaline. The resulting solution was stored at lower than -18 °C. The solution was thawed immediately before use. The concentration of the stock solution was determined in 0.1 M NaOH by measuring the absorbance at 302 nm with a molar extinction coefficient of 1670 M⁻¹cm⁻¹.

Generation of NO•: Nitric oxide was generated from SNP (Sodium nitroferricyanide (III) dehydrate). The experiments were performed under anaerobic conditions. The deionized water was degassed with Ar for 30 min and then SNP was added into it under Ar atmosphere and stirred for 30 min at room temperature. The probe solution was also degassed before the reaction with NO•.



Scheme 1. Synthesis of **Probe-1** and **Probe-OCI**.

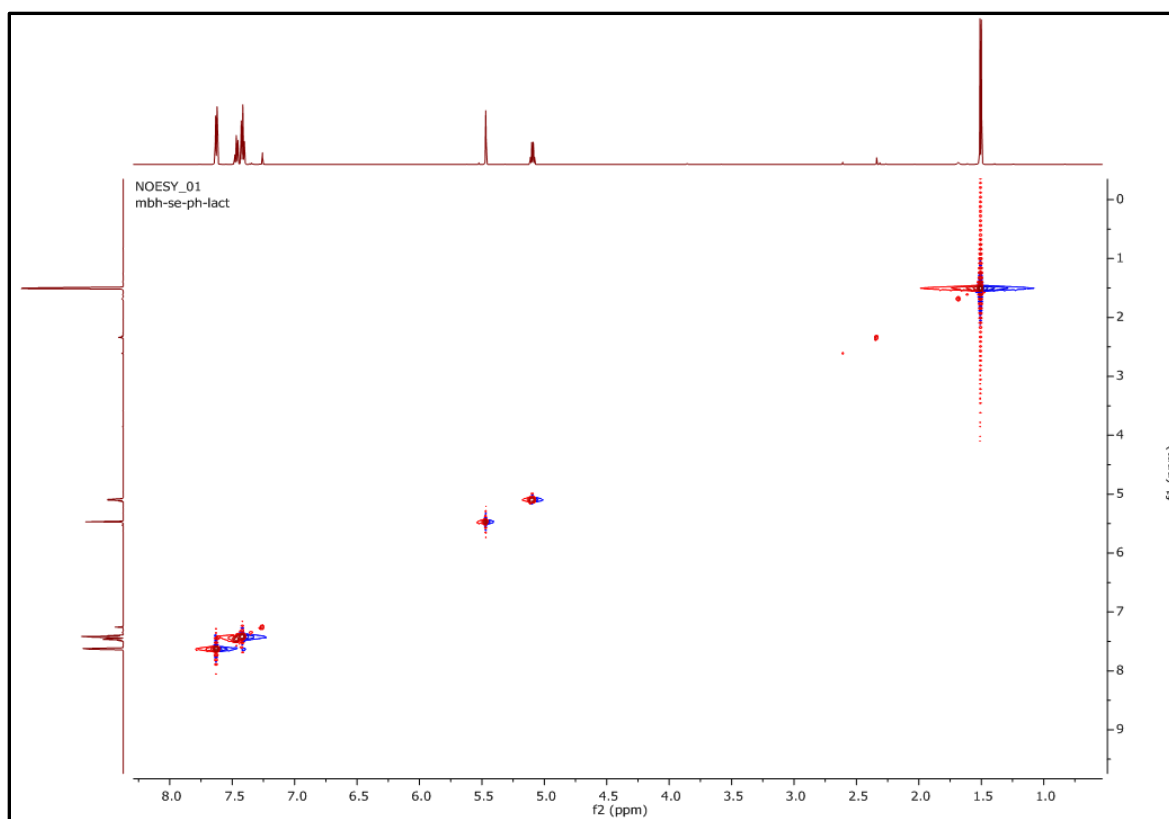
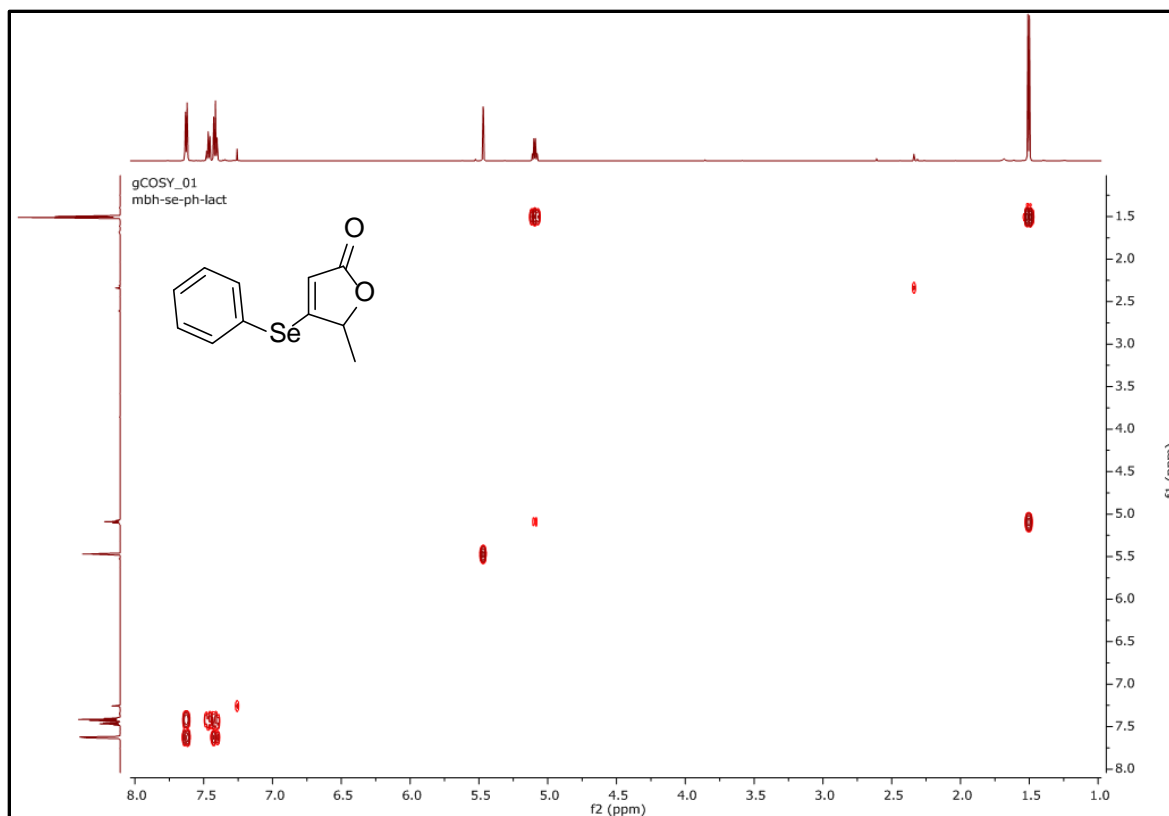


Figure S2. (top) COSY and (bottom) ^1H NOESY NMR spectrum of **1**.

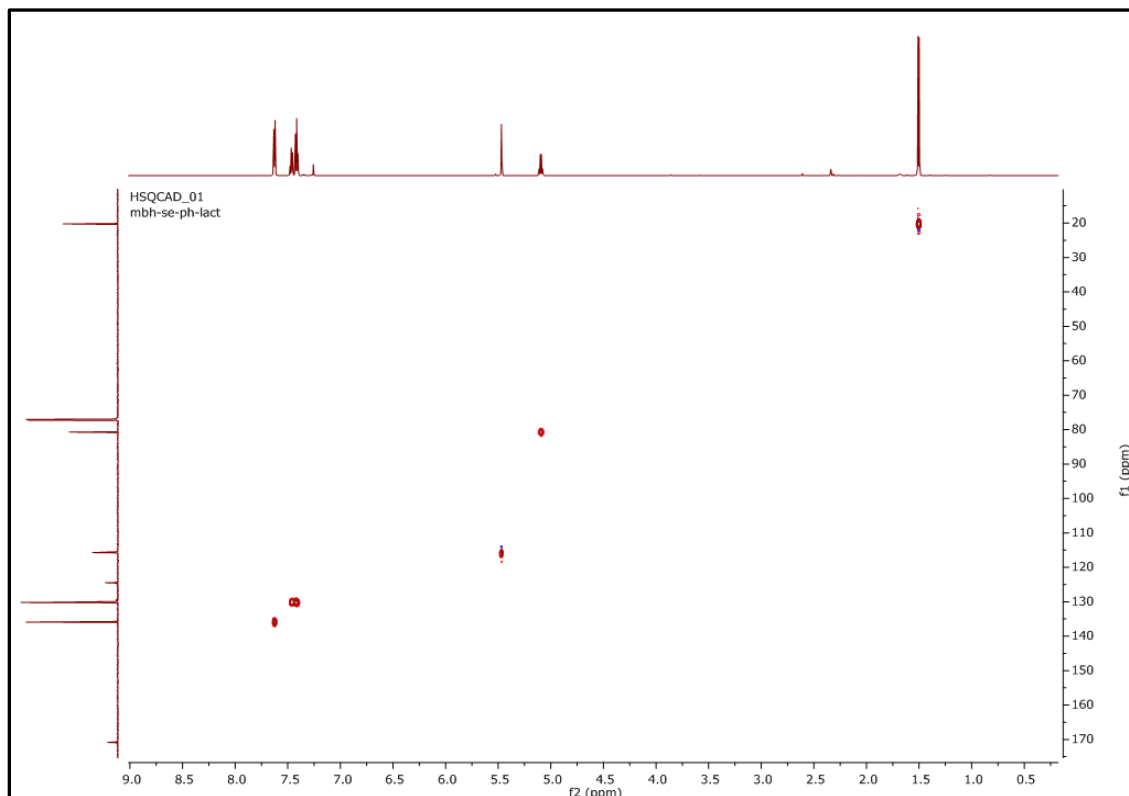
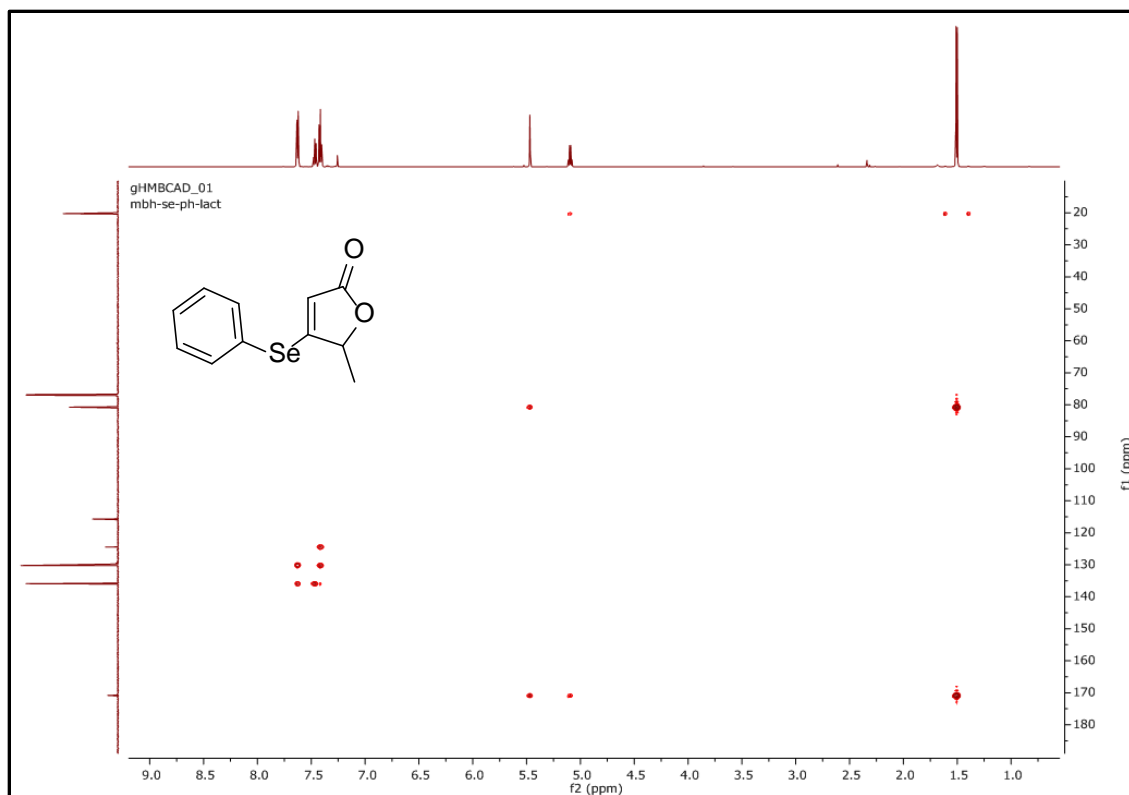


Figure S3. (top) $^1\text{H} - ^{13}\text{C}$ HMBC NMR spectrum of **1** and (bottom) $^1\text{H} - ^{13}\text{C}$ HSQC NMR spectrum of **1**.

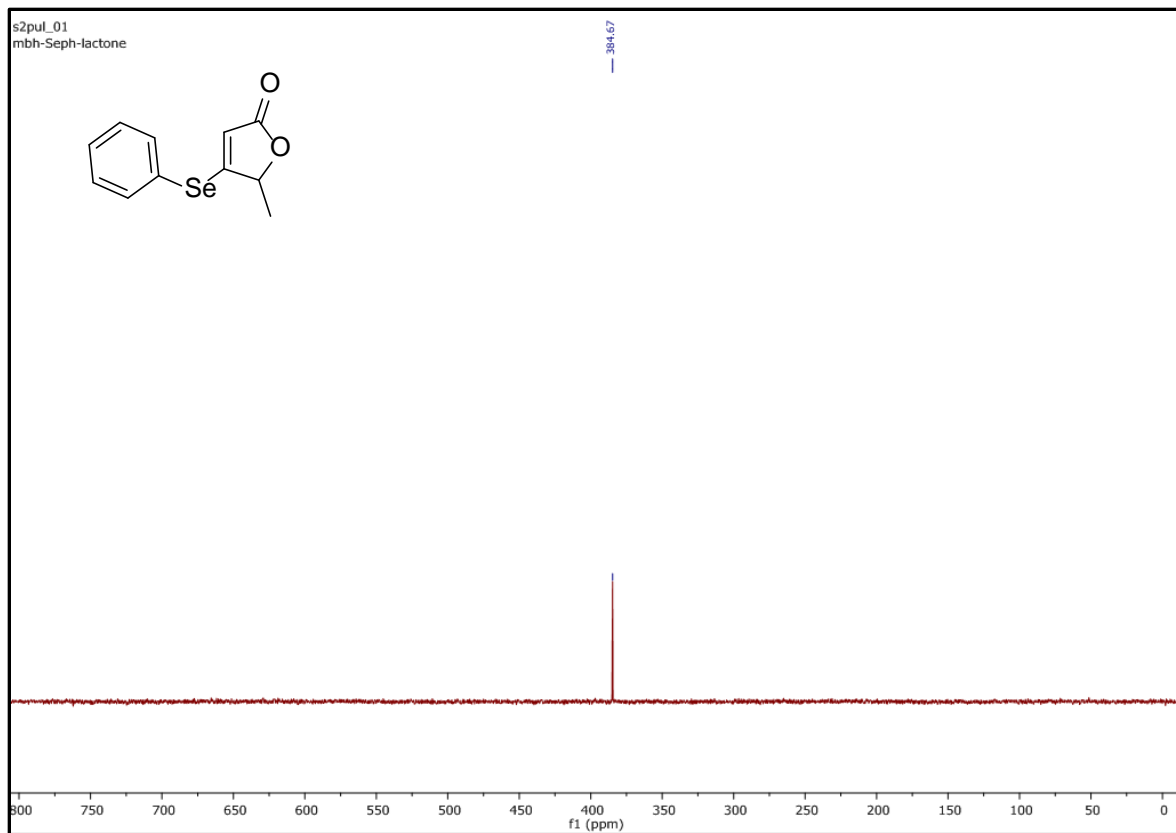


Figure S4. ^{77}Se NMR spectrum of **1**.

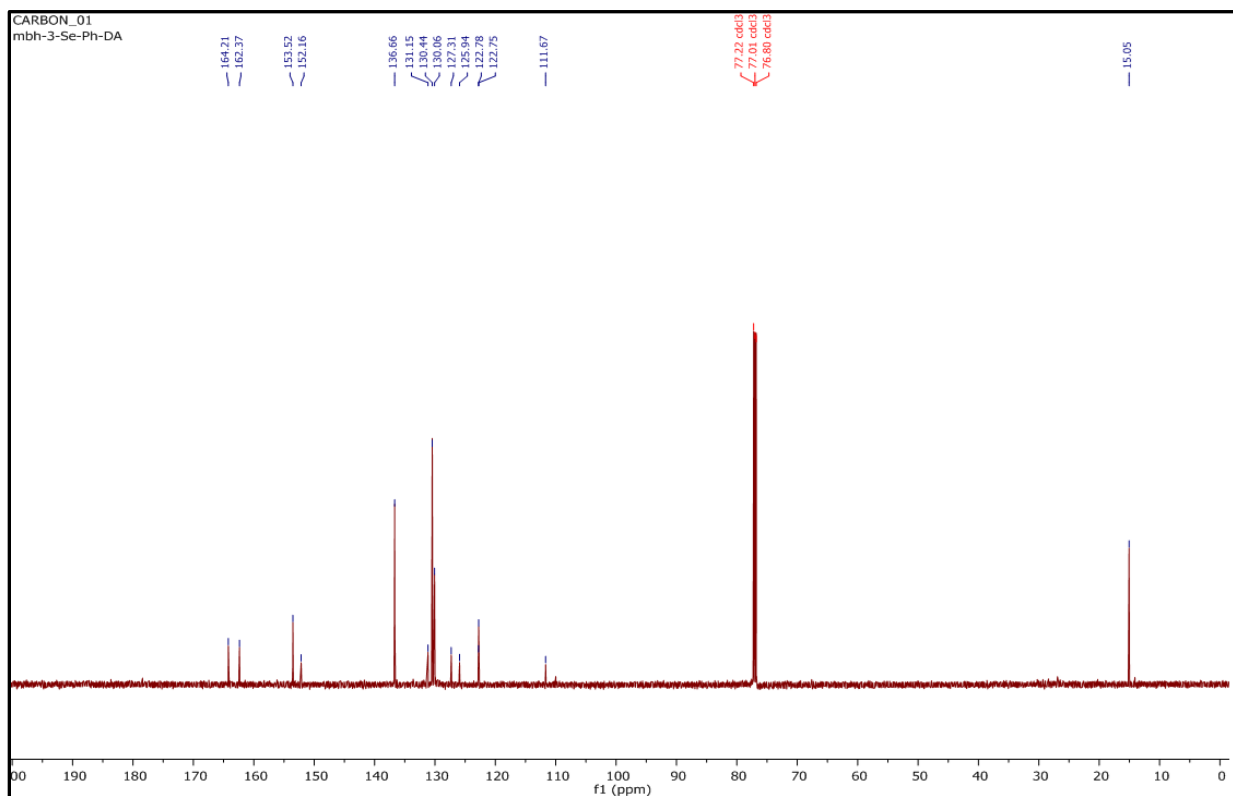
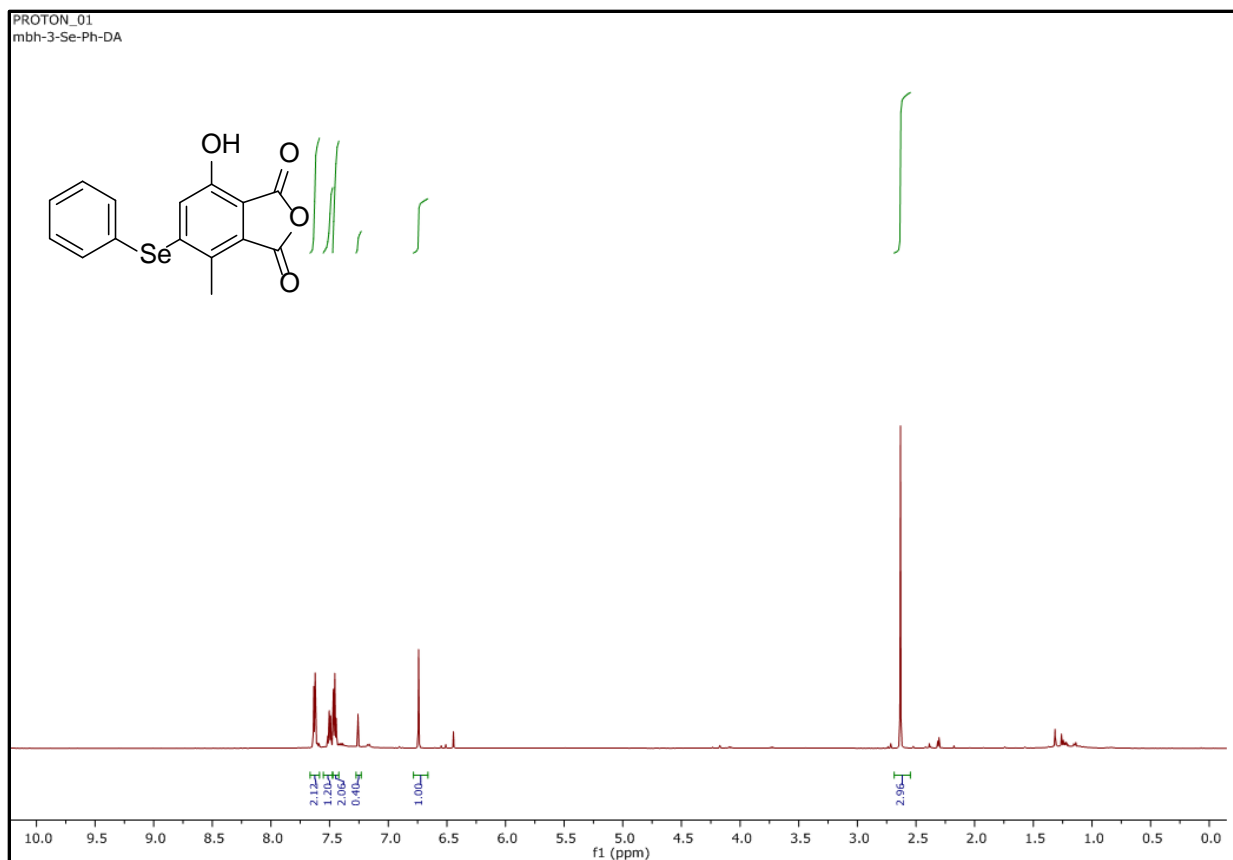


Figure S5. (top) ¹H and (bottom) ¹³C NMR spectrum of **Probe-1**.

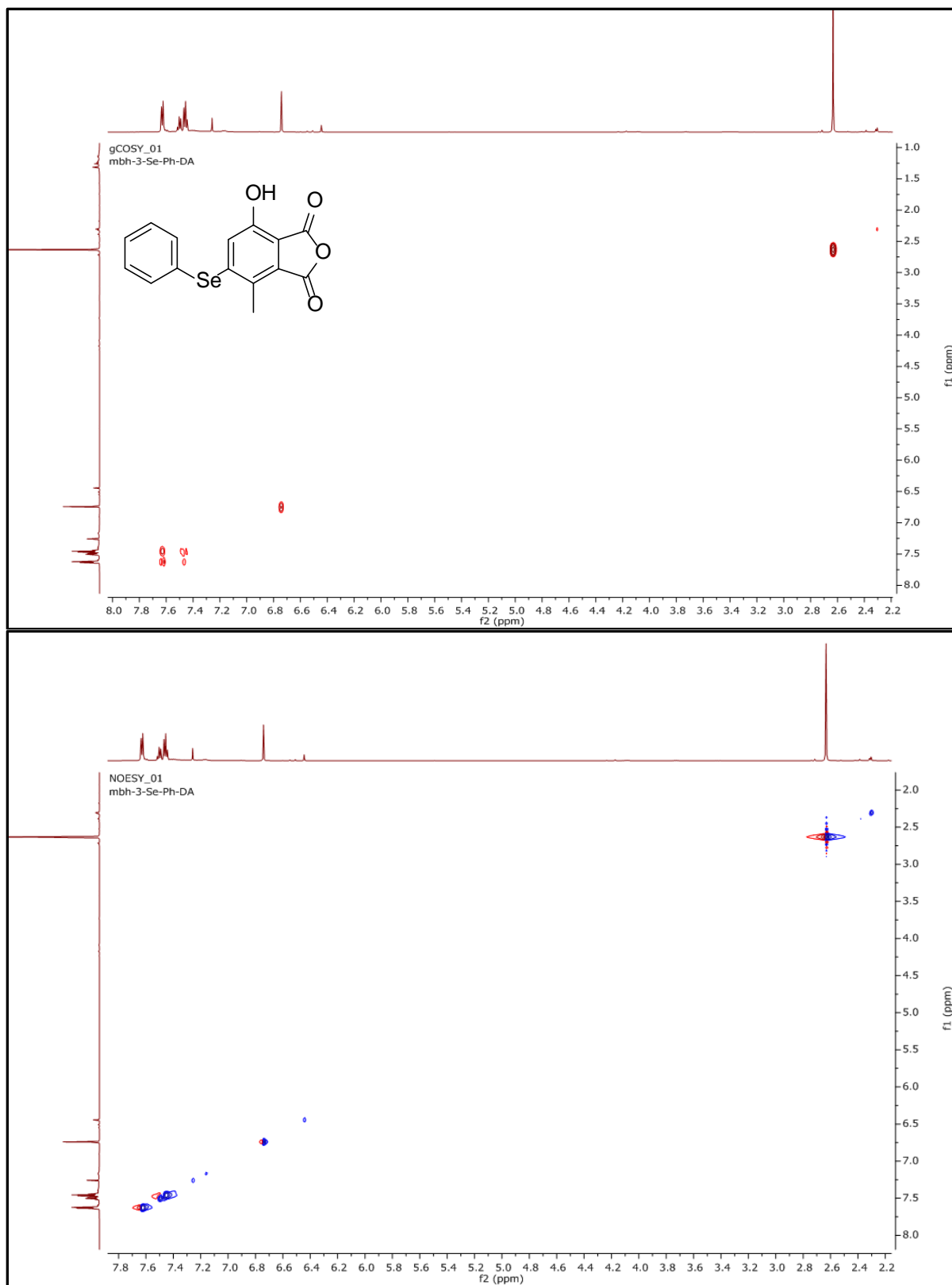


Figure S6. (top) ^1H COSY NMR spectrum of **Probe-1** and (bottom) ^1H NOESY NMR spectrum of **Probe-1**.

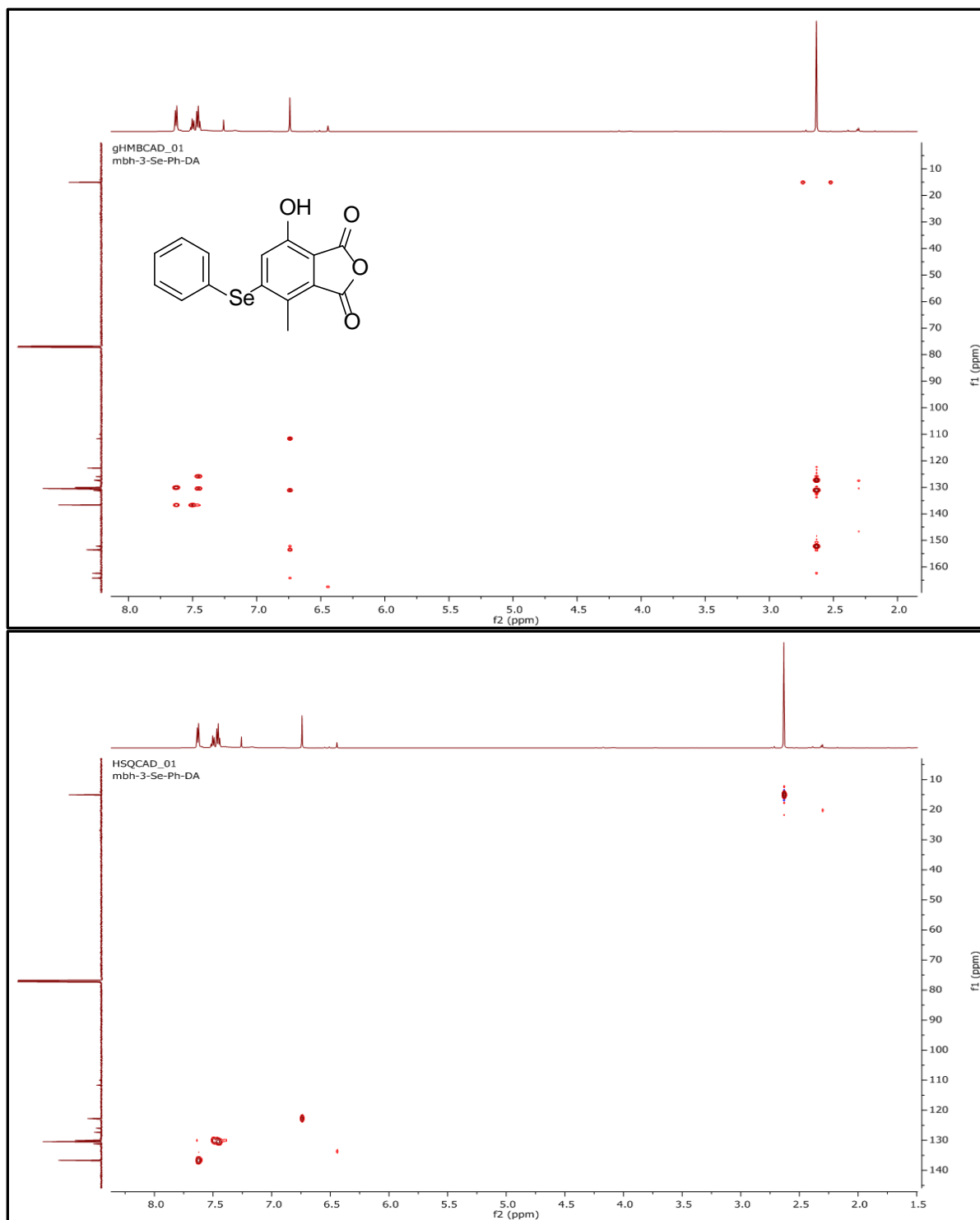


Figure S7. (top) ^1H – ^{13}C HMBC NMR spectrum of **2** and (bottom) ^1H – ^{13}C HSQC NMR spectrum of **Probe-1**.

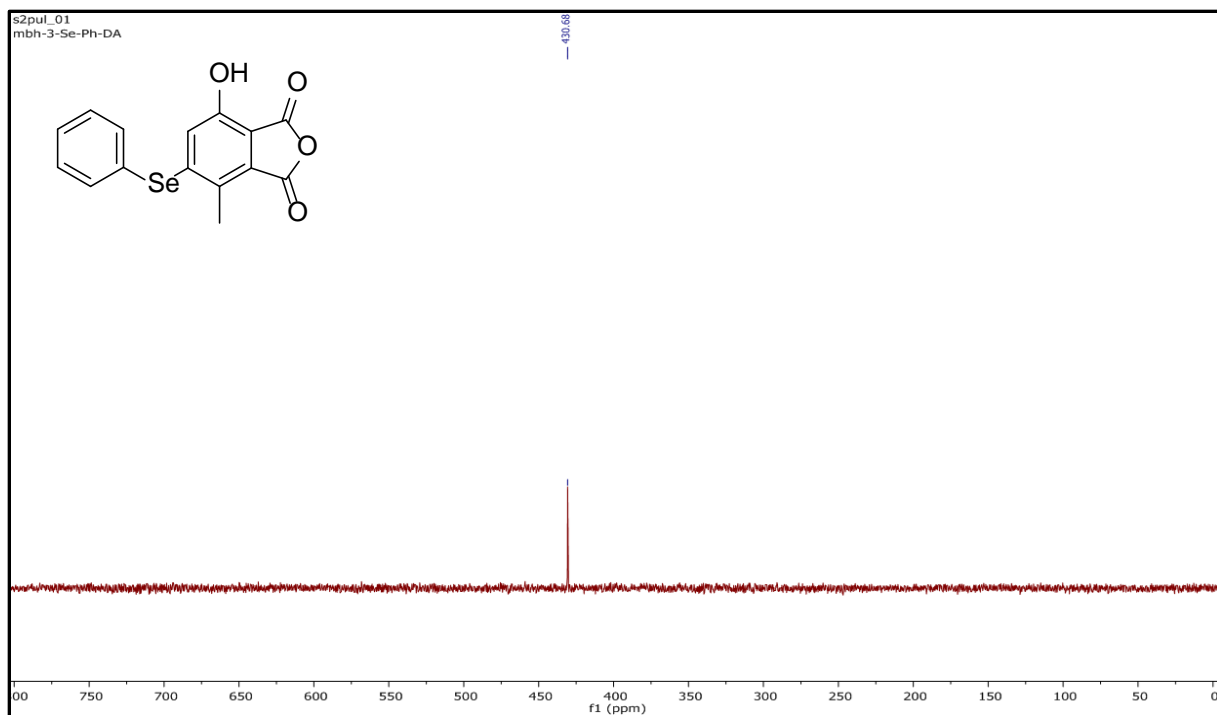


Figure S8. ^{77}Se NMR spectrum of **Probe-1**.

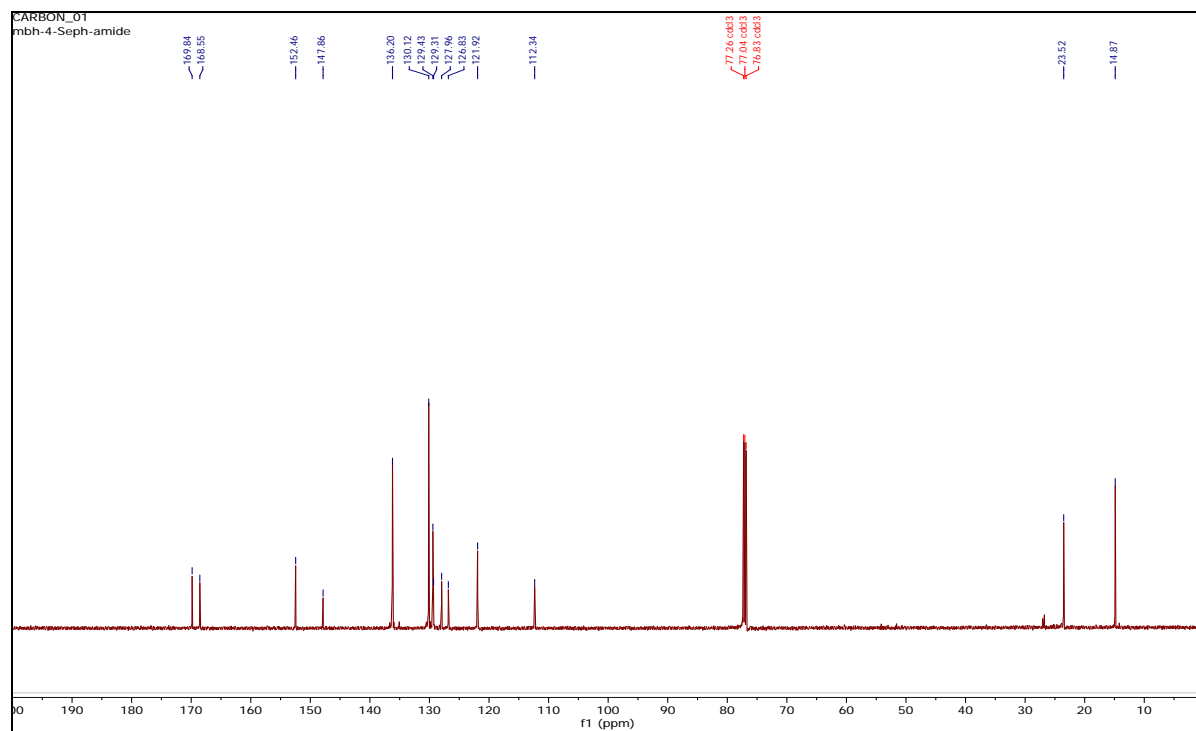
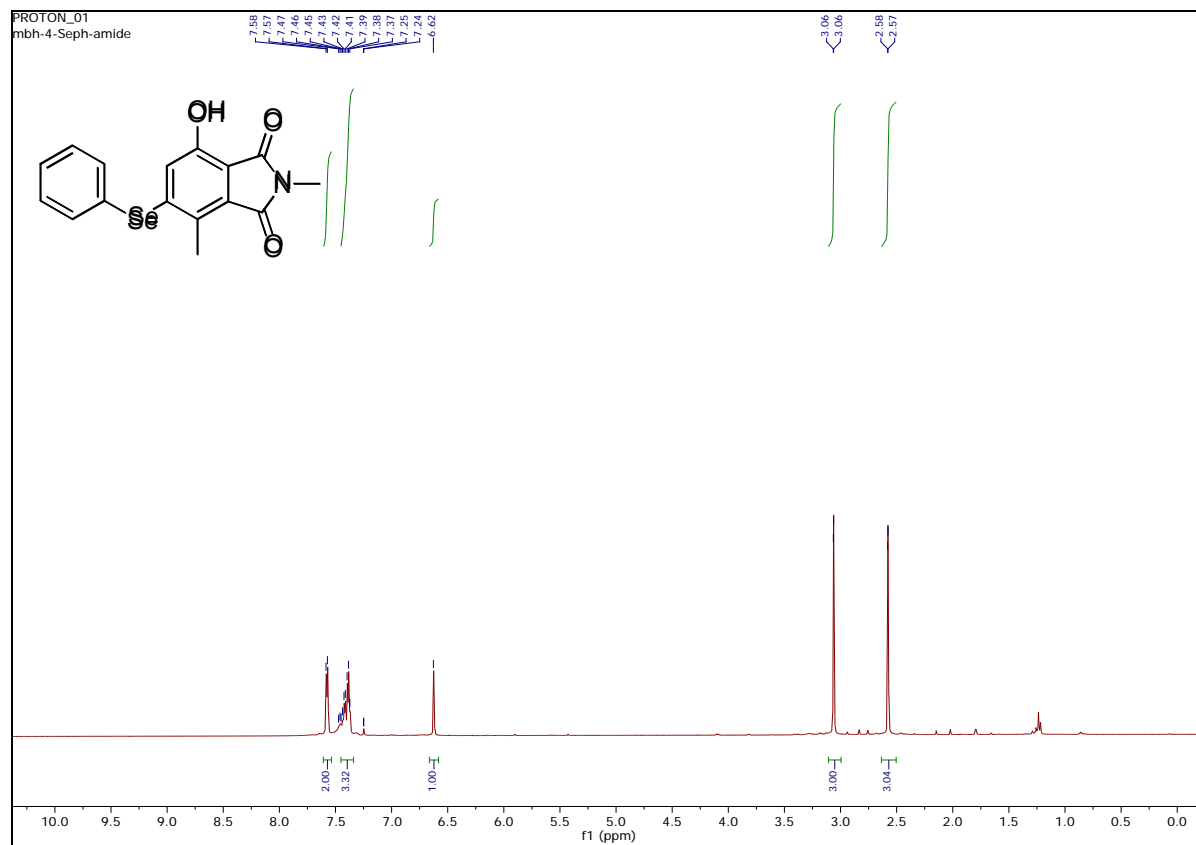


Figure S9. (top) ^1H and (bottom) ^{13}C NMR spectrum of **Probe-OCI**.

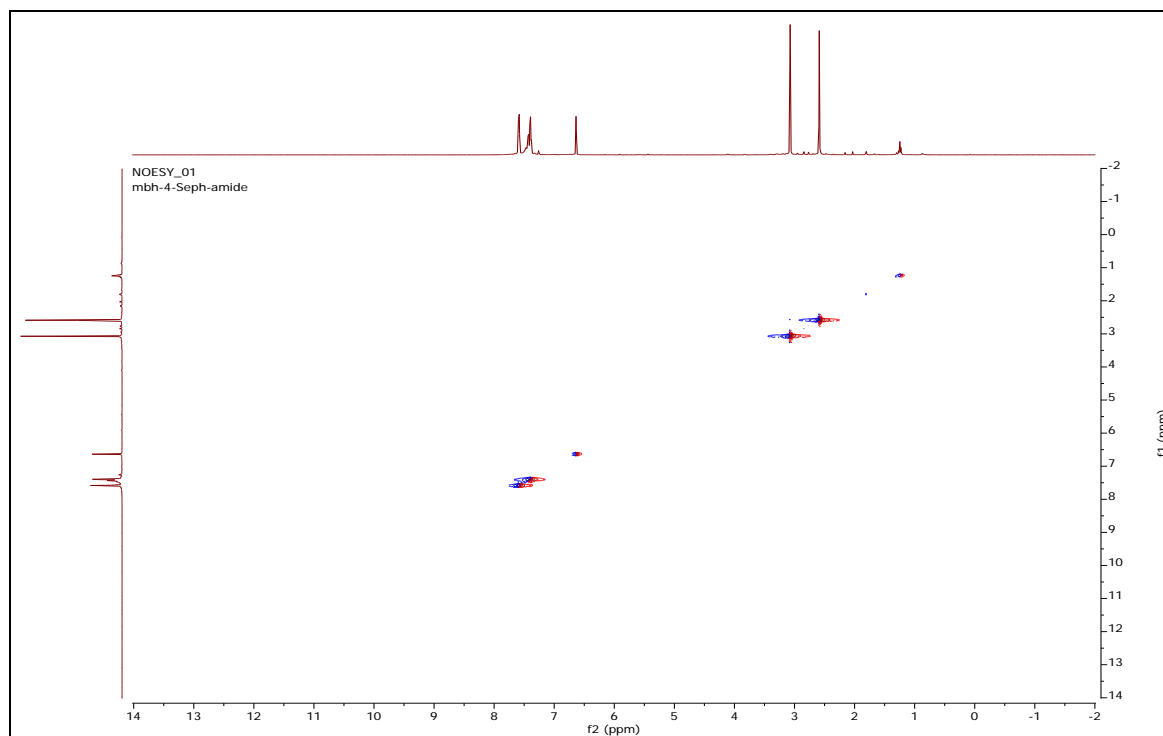
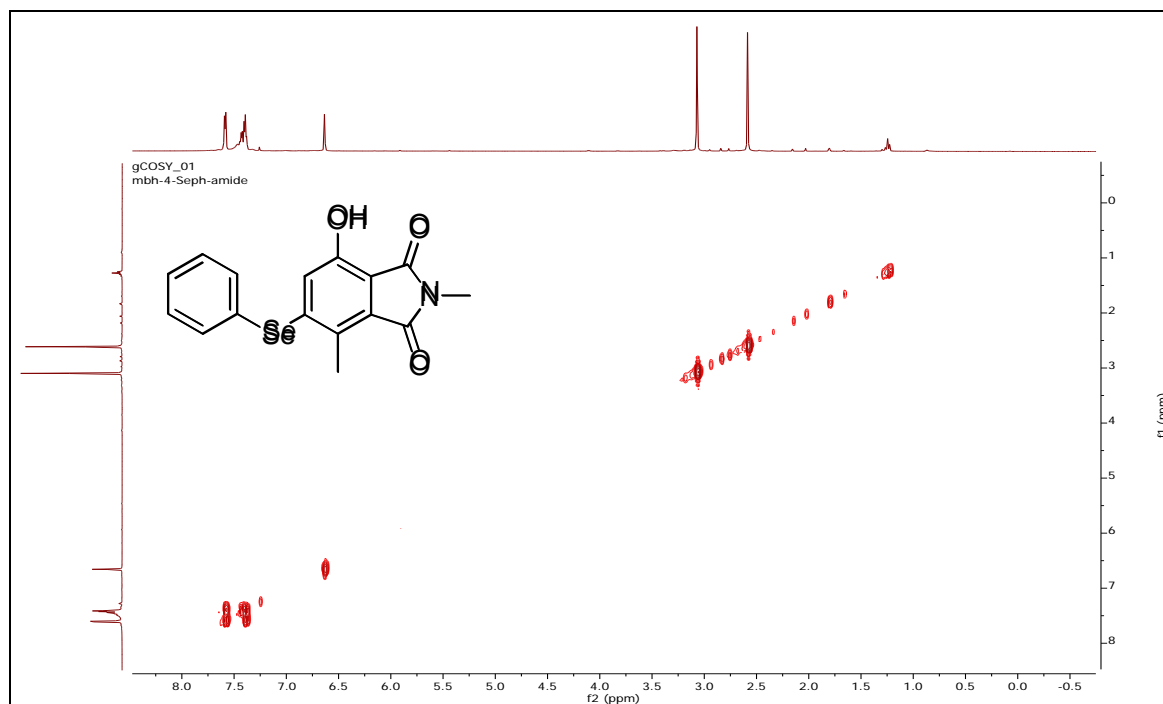


Figure S10. (top) ^1H COSY NMR spectrum of **Probe-OCI** and (bottom) ^1H NOESY NMR spectrum of **Probe-OCI**

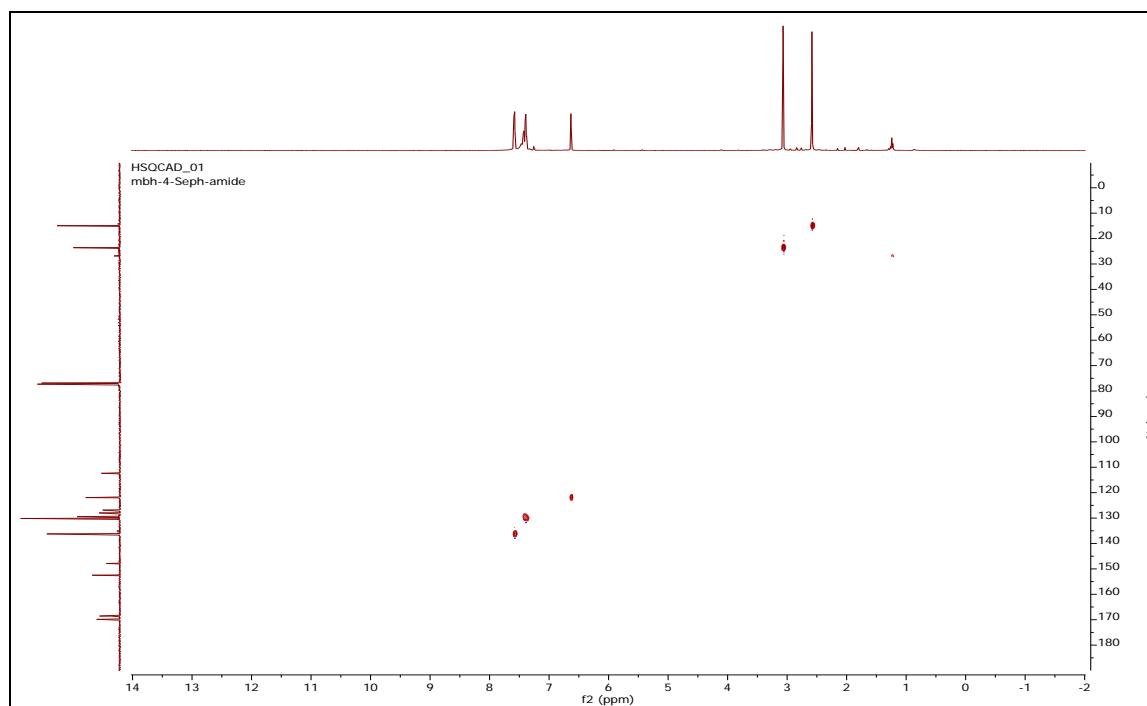
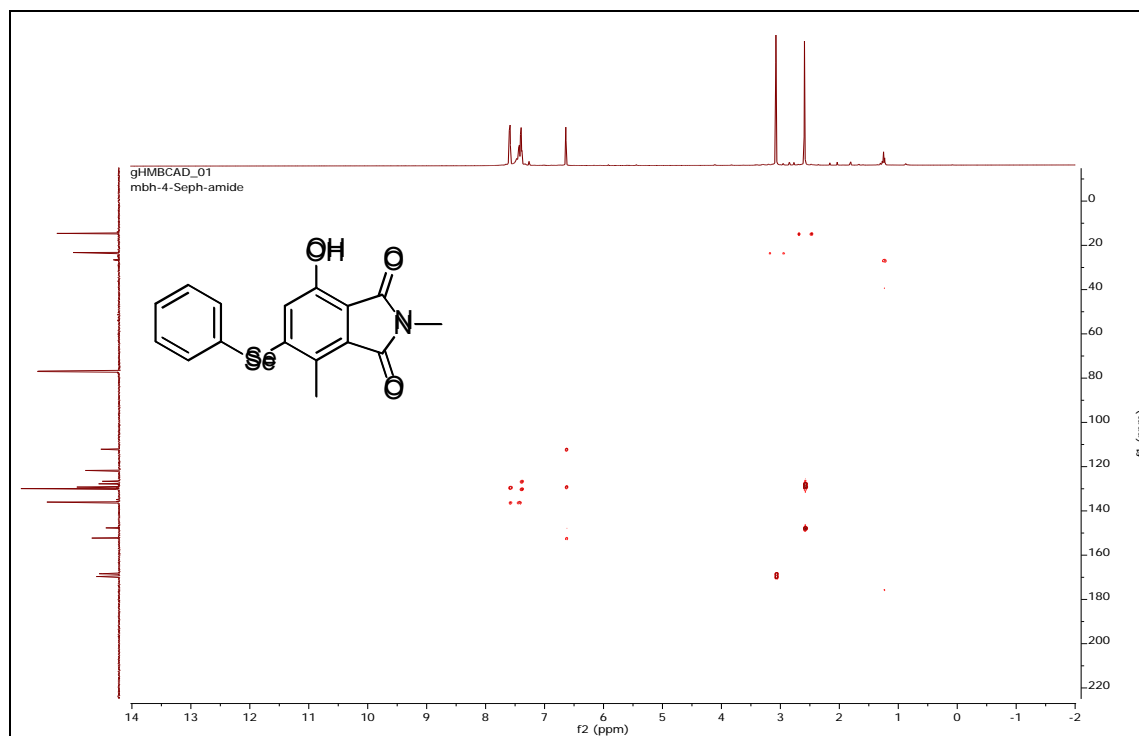


Figure S11. (top) ^1H – ^{13}C HMBC NMR spectrum of **Probe-OCI** and (bottom) ^1H – ^{13}C HSQC NMR spectrum of **Probe-OCI**.

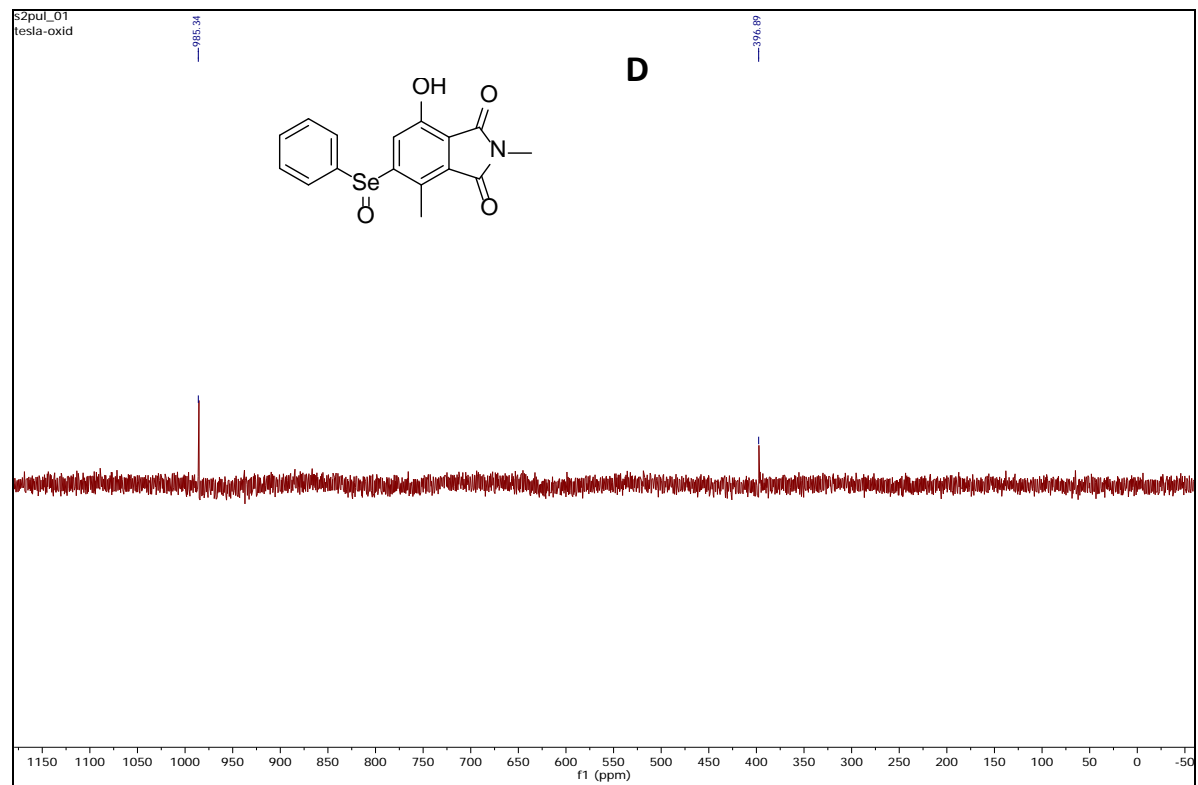
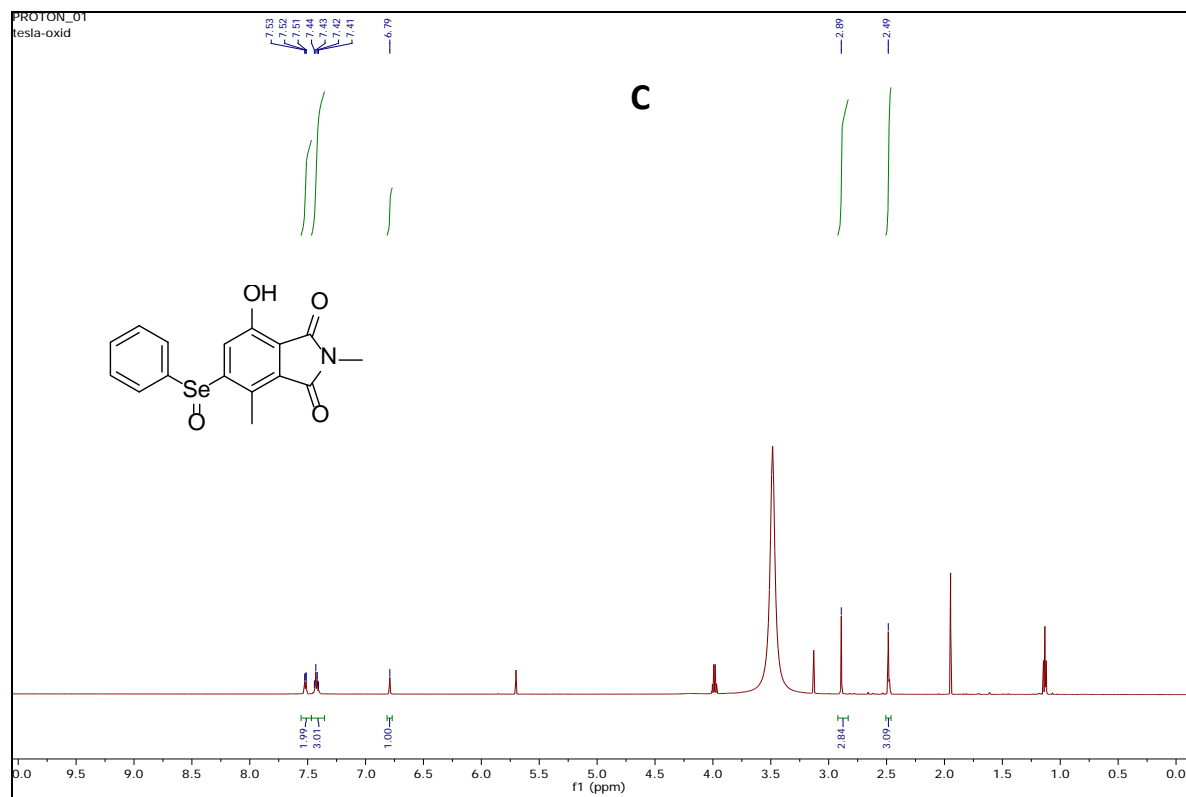


Figure S12. (A) ^1H NMR spectrum of **Probe-OCI** and (B) ^{77}Se NMR spectrum of **Probe-OCI** (C) ^1H NMR spectrum of **Probe-OCI[O]** (D) ^{77}Se NMR spectrum of **Probe-OCI[O]**

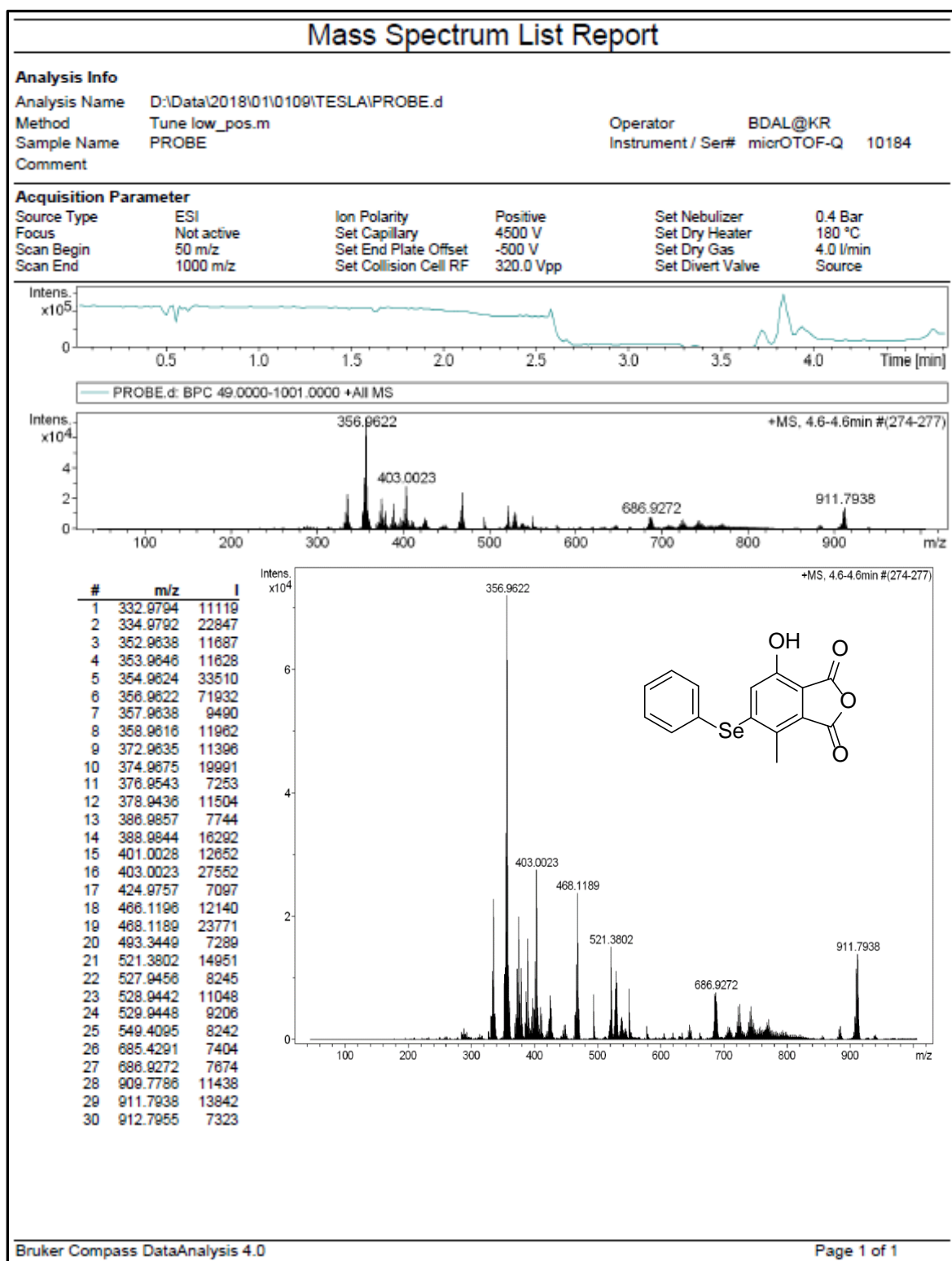


Figure S13. HR-MS (ESI) spectrum of **Probe-1**.

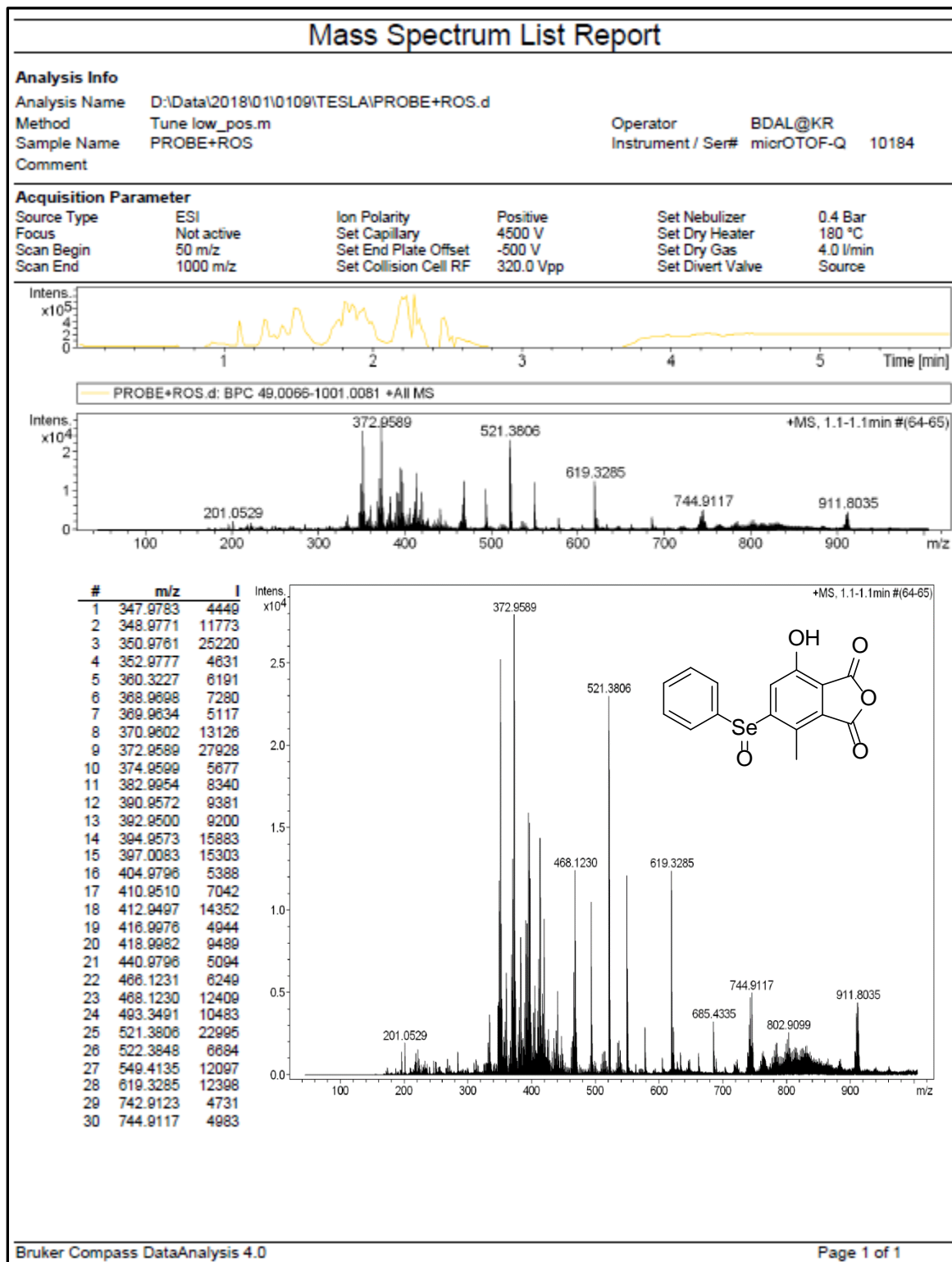


Figure S14. HR-MS (ESI) spectrum of **Probe-1+OCl₂**.

Mass Spectrum List Report

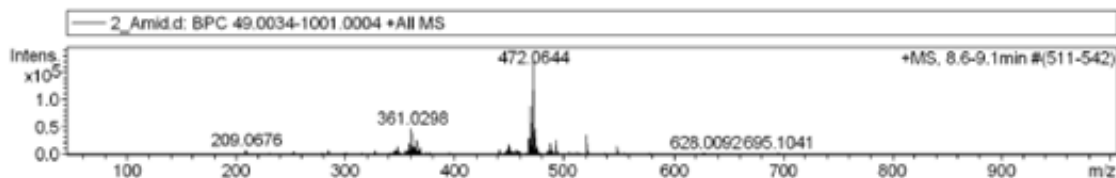
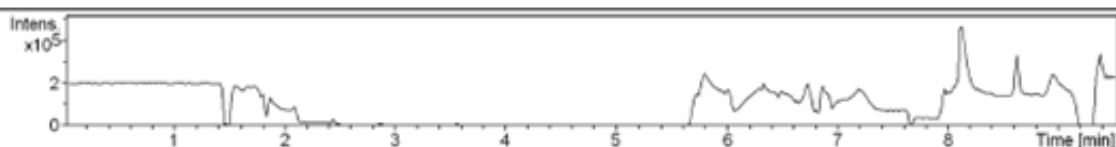
Analysis Info

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 Method Tune_low_pos.m
 Sample Name 2_Amid
 Comment

Operator BDAL@KR
 Instrument / Ser# micrOTOF-Q 10184

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	0.4 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	50 m/z	Set End Plate Offset	-500 V	Set Dry Gas	4.0 l/min
Scan End	1000 m/z	Set Collision Cell RF	320.0 Vpp	Set Divert Valve	Source



#	m/z	I
1	209.0676	7185
2	345.0426	6734
3	348.0119	13756
4	357.0318	4750
5	359.0303	20811
6	361.0298	46663
7	362.0277	11278
8	363.0293	11798
9	364.0223	13775
10	366.0216	25995
11	368.4221	12727
12	441.2938	9284
13	448.0803	7676
14	450.0794	17383
15	456.0848	8167
16	466.0639	27021
17	469.0651	29697
18	470.0645	87817
19	471.0700	16588
20	472.0644	179783
21	473.0664	40361
22	474.0713	46028
23	475.0731	8095
24	476.0612	14437
25	486.0388	9023
26	488.0396	20340
27	493.3474	26174
28	521.3785	35777
29	522.3817	9007
30	549.4093	13257

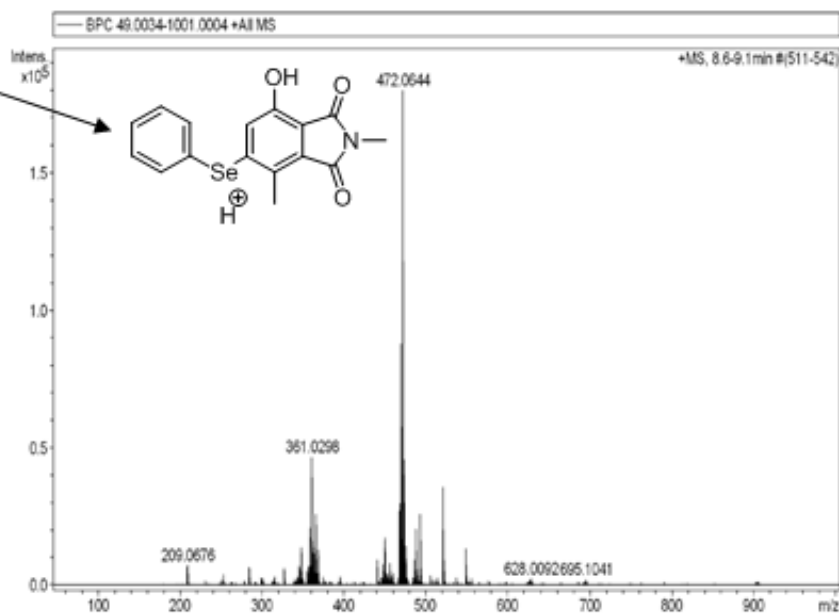


Figure S15. HR-MS (ESI) spectrum of Probe-OCI.

Mass Spectrum List Report

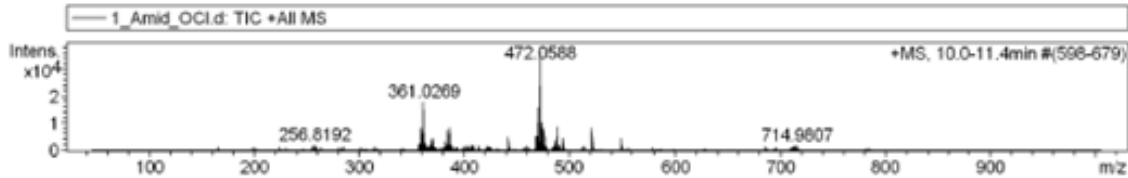
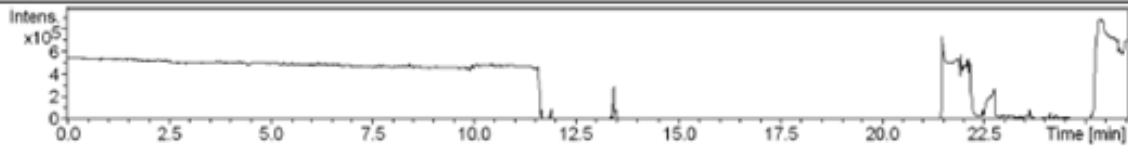
Analysis Info

Analysis Name D:\Data\2018\04\0412\1_Amid_OCI.d
 Method Tune_low_pos.m
 Sample Name 1_Amid_OCI
 Comment

Operator BDAL@KR
 Instrument / Ser# micrOTOF-Q 10184

Acquisition Parameter

Source Type	ESI	Ion Polarity	Positive	Set Nebulizer	0.4 Bar
Focus	Not active	Set Capillary	4500 V	Set Dry Heater	180 °C
Scan Begin	50 m/z	Set End Plate Offset	-500 V	Set Dry Gas	4.0 l/min
Scan End	1000 m/z	Set Collision Cell RF	320.0 Vpp	Set Divert Valve	Source



#	m/z	I
1	256.8192	2052
2	357.0288	2659
3	358.0282	2598
4	359.0273	8163
5	361.0269	17757
6	362.0154	3018
7	363.0253	2876
8	367.9912	2279
9	368.4203	3465
10	369.9903	4599
11	381.9776	3065
12	383.0780	7483
13	385.9843	8268
14	405.9563	1973
15	441.2916	4697
16	468.0593	5080
17	469.0613	5381
18	470.0592	15945
19	471.0617	3115
20	472.0588	36474
21	473.0604	8376
22	474.0575	10217
23	476.0526	8026
24	486.0541	3770
25	488.0528	8859
26	490.0558	2012
27	493.3444	4264
28	521.3745	8297
29	522.3782	2144
30	549.4056	4359

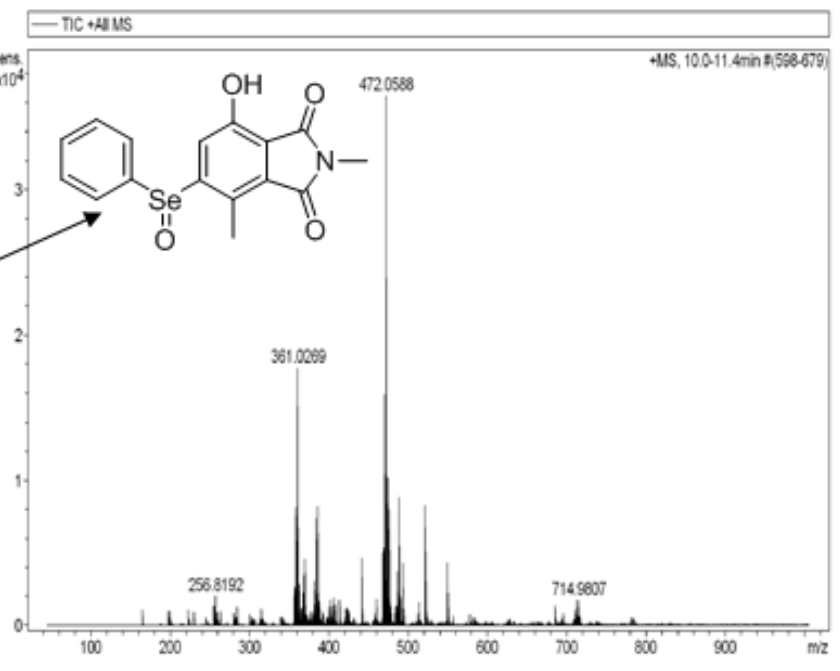


Figure S16. HR-MS (ESI) spectrum of Probe-OCI+ OCl⁻

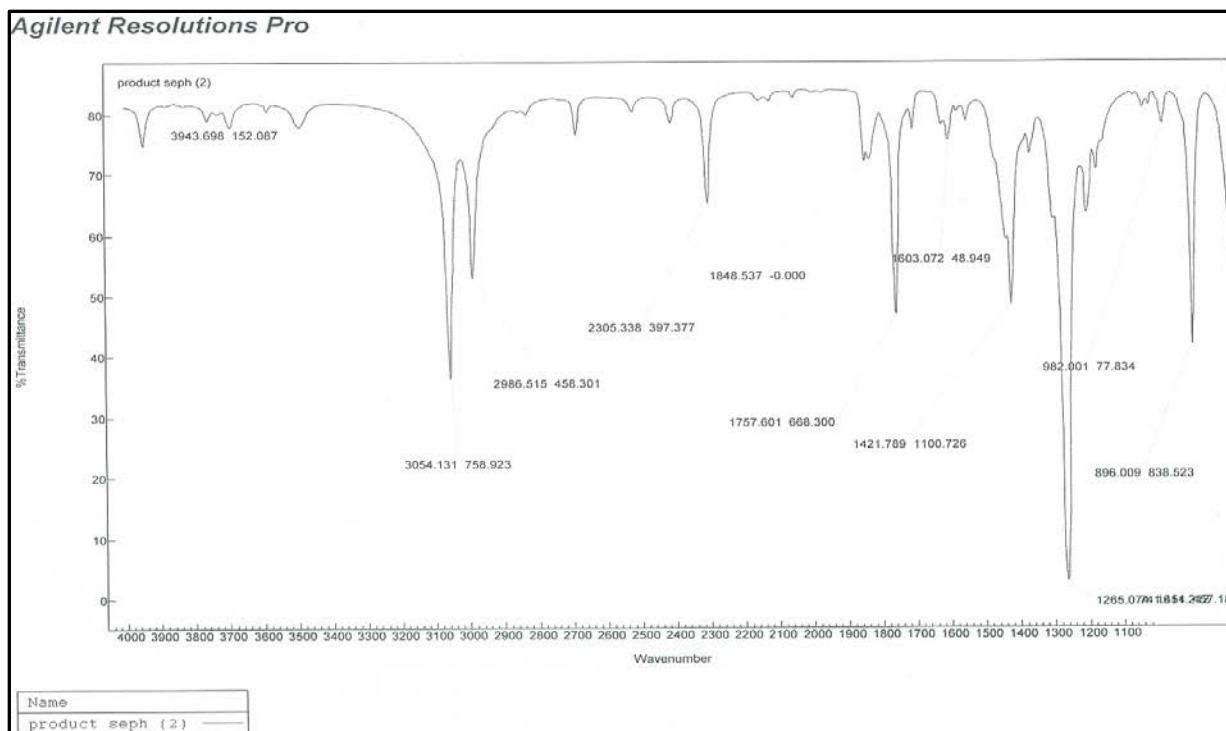
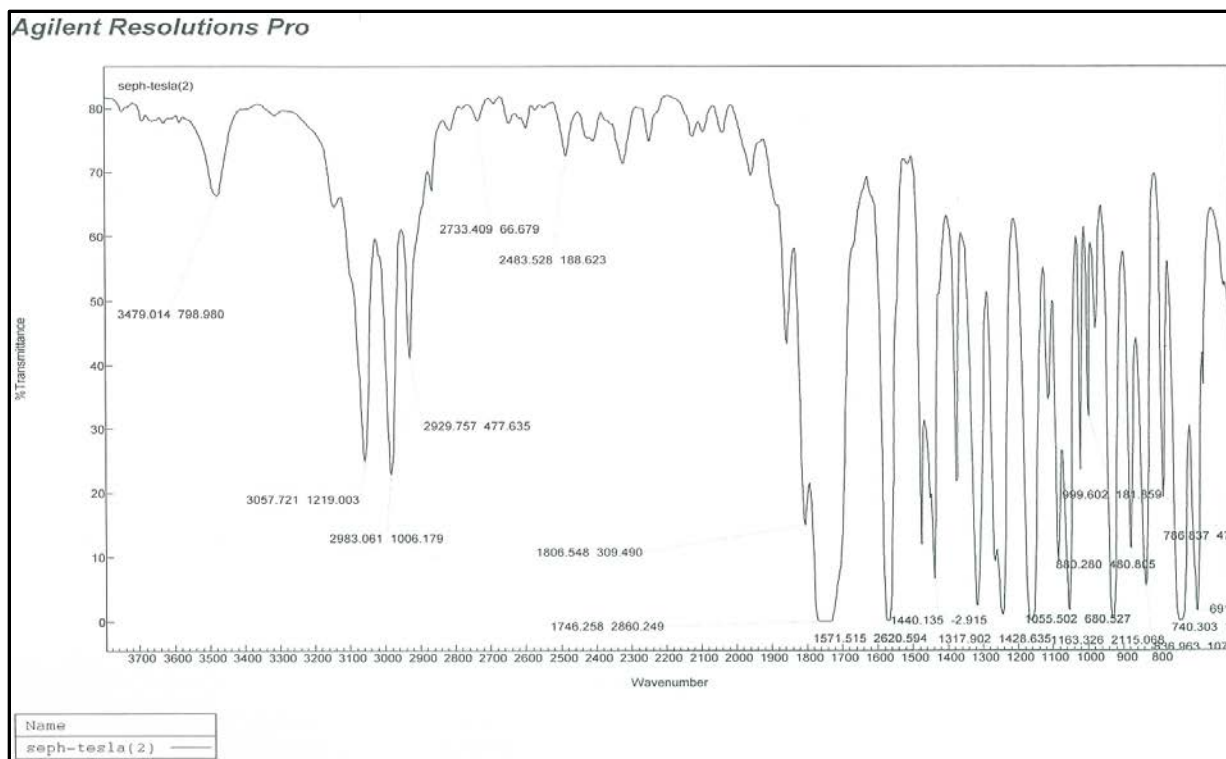


Figure S17. (top) IR spectrum of **1** and (bottom) **Probe-1**.

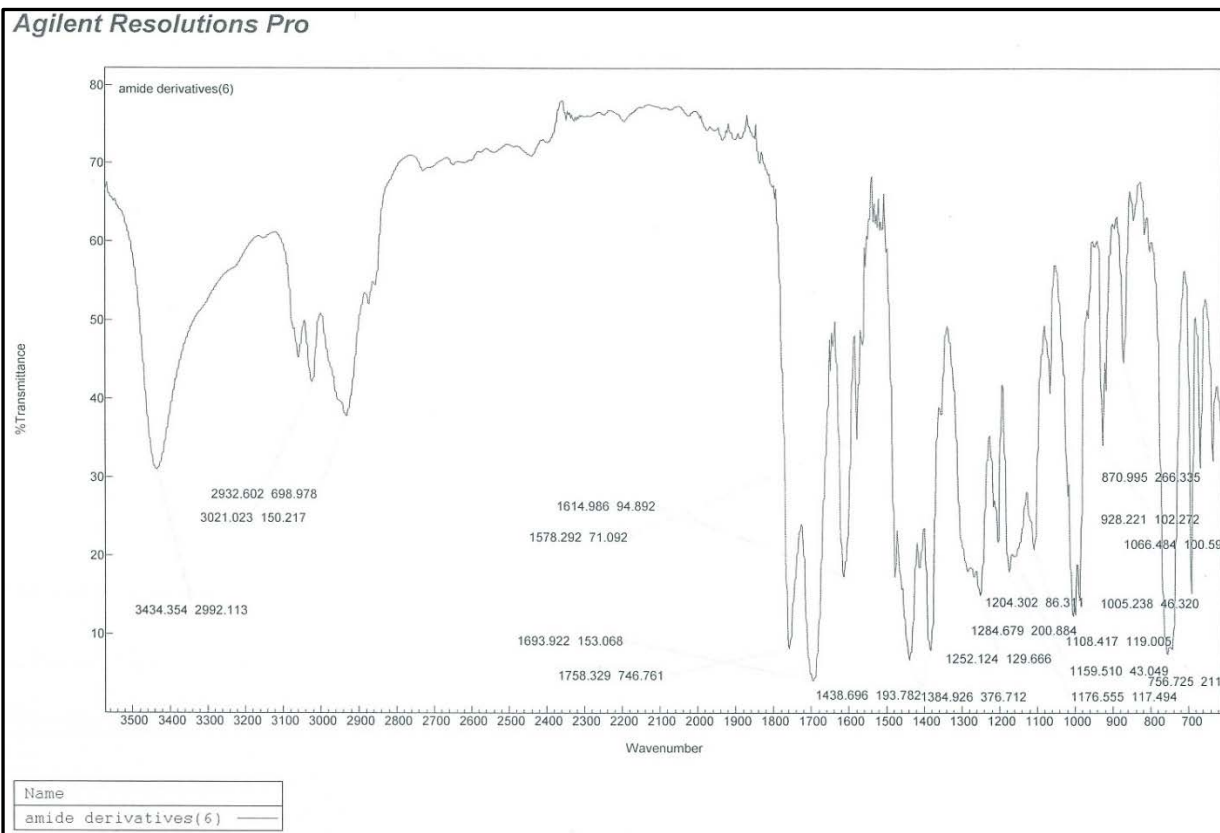


Figure S18. IR spectrum of **Probe-OCI**.

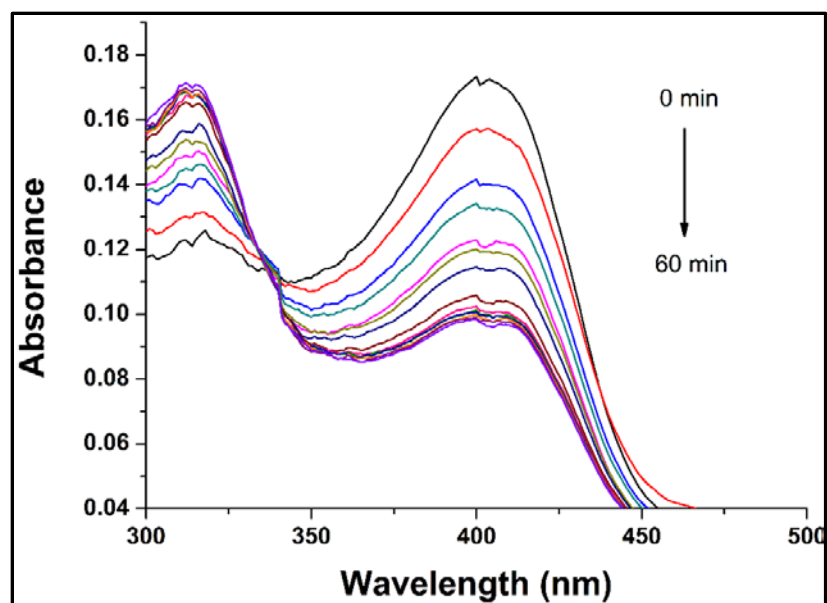


Figure S19. Absorption spectra of the **Probe-1** (15.0 μM) with 1.0 equiv OCl⁻ in time from 0 to 60 minutes in the solution (10 mM PBS pH 7.4) incubated for 1.0 min at r.t.

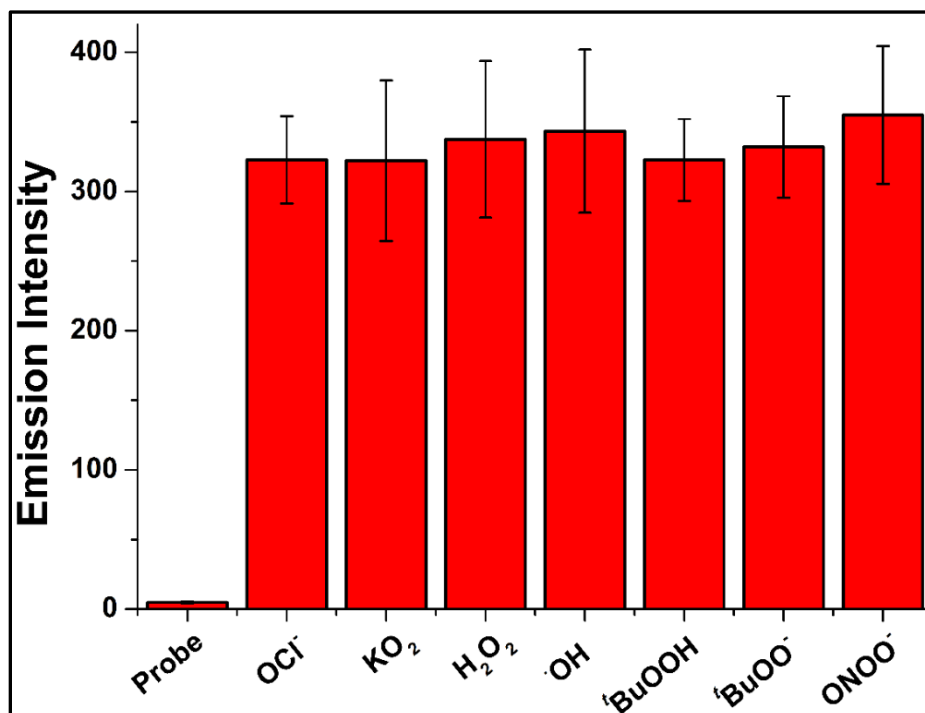


Figure S20. Fluorescence intensity of **Probe-1** (15.0 μM) with ROS/RNS in the solution (10 mM PBS pH 7.4) incubated for 2.0 min. λ_{ex} : 404 nm, λ_{em} : 502 nm, slit width 3.0 nm/ 3.0 nm.

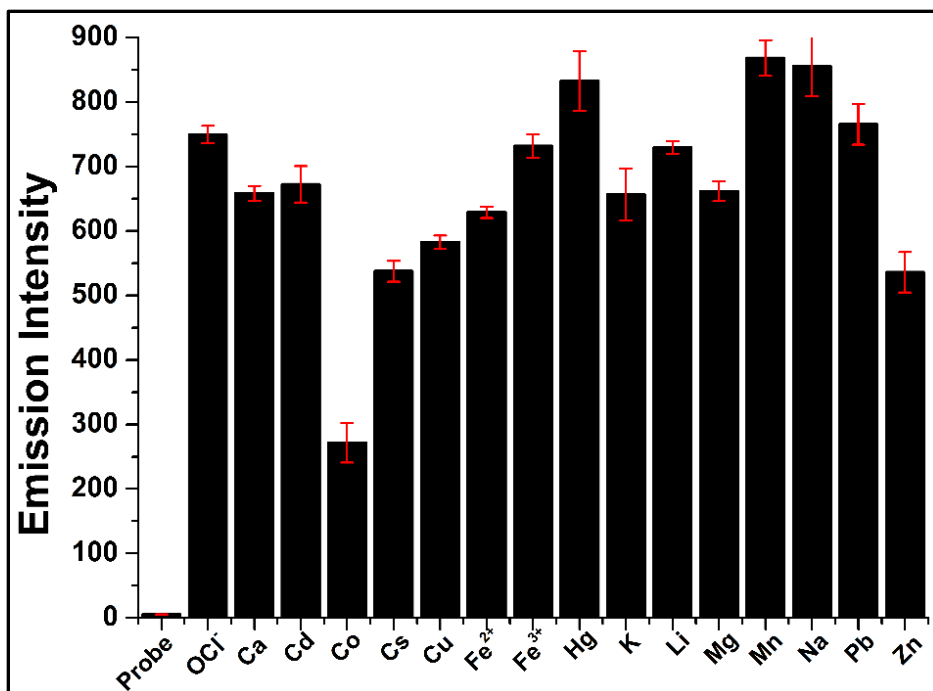


Figure S21. Fluorescence intensity of **Probe-1** (15.0 μM) with various metals in solution (10 mM PBS pH 7.4) incubated for 1.0 min. λ_{ex} : 404 nm, λ_{em} : 512 nm, slit width 3.0 nm/3.0nm.

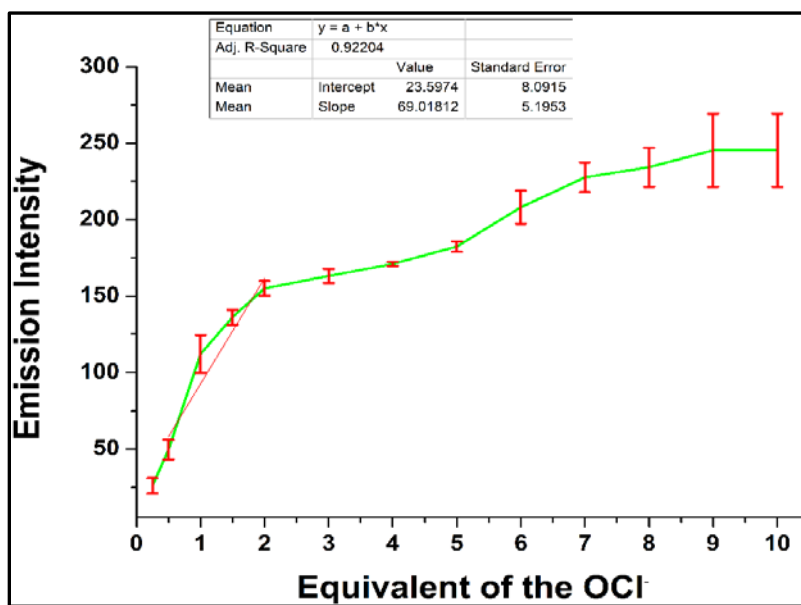


Figure S22. Plot for the calculation of limit of detection from the emission of **Probe-1** (15.0 μM) in the solution (10 mM PBS pH 7.4, 3:7 v/v) with increasing concentration of OCl^- (0.0 to 10.0 μM in water) incubated for 1.0 min at r.t, λ_{ex} . 404 nm and λ_{em} . 502 nm slith width 3.0 nm/ 3.0 nm (average of three experiments).

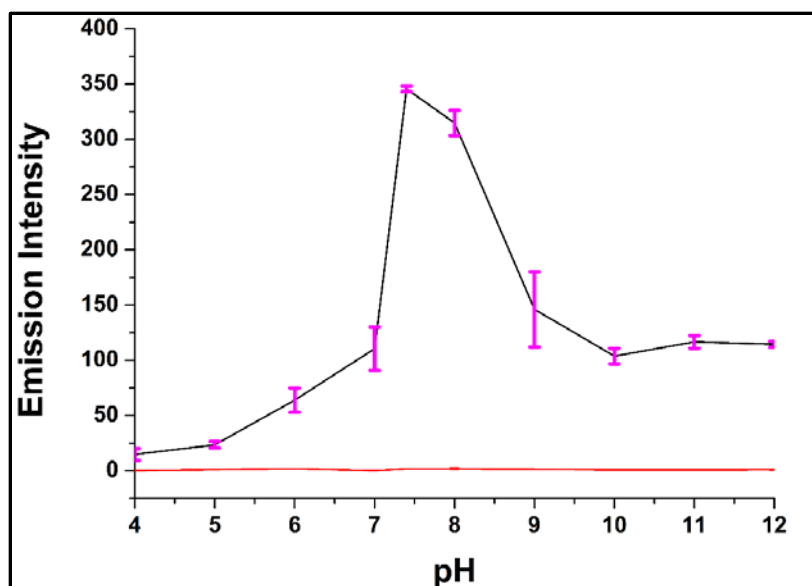


Figure S23. pH-dependent spectral changes of **Probe-1** (15.0 μM) with 1.0 equiv of OCl^- in various pH values in solution (10 mM PBS, pH 7.4); λ_{ex} : 502 nm, λ_{em} : 512 nm. Slit width 3.0 nm/3.0 nm.

Table S1. Results of experimental determination of log P value by the “shake flask” method for the **Probe-1**

C_1	V	A	\bar{A}	C_2	P	Log P
1.3×10^{-4}	35	0.3289 0.28425 0.24004	0.2844	7.49×10^{-5}	29.7	1.47
1.6×10^{-4}	48	0.25045 0.25801 0.20986	0.23944	6.21×10^{-5}	59.1	1.77
2.0×10^{-4}	60	0.25146 0.26715 0.27875	0.2658	7.44×10^{-5}	63.3	1.80

※ C_1 = Concentration (mol L^{-1}) of the stock solution in *n*-octanol before partition; V = volume (μL) of stock solution; A = absorbance in buffer solution after the partition ($\lambda = 500$ nm); \bar{A} = arbitrary absorbance in buffer solution after partitioning ($\lambda = 500$ nm); C_2 = concentration (mol L^{-1}) in buffer solution after partitioning; P = partition coefficient; log P = logarithm of the partition coefficient.

Table S2. Information of the **Probe-1** calculated through the 'molinspiration property engine v2011.04' at the website, <http://www.molinspiration.com>.

Properties (Probe-1) (NOTE: abbreviation same as website)	Value
milogP	3.41
TPSA	67.51
Natom	20
MW	333.20
nON	4
nOHNH	1
nviolations	0
nrotb	2
volume	239.63

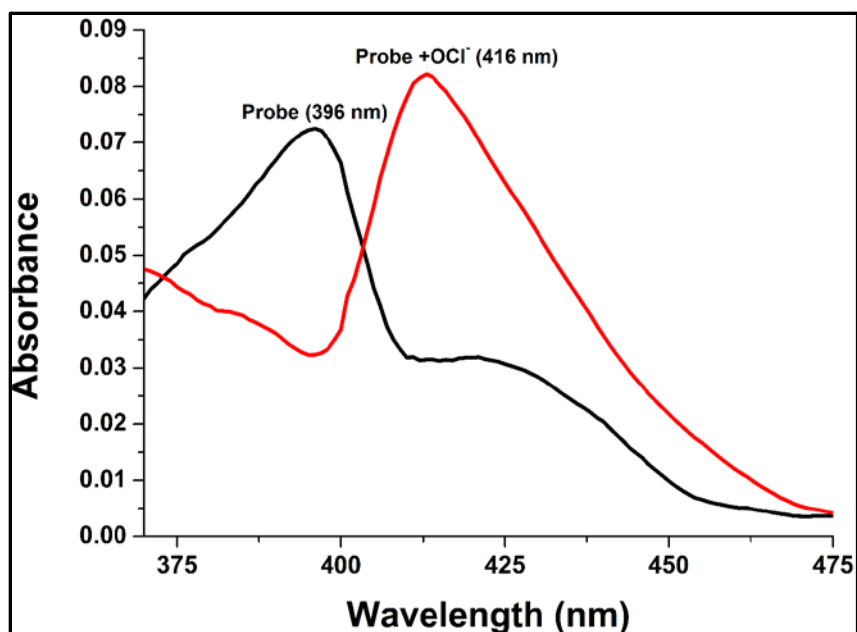


Figure S24. Absorption spectra of the **Probe-OCI** (15.0 μM) and **Probe-OCI** + OCI^- (15.0 μM with 1.0 equiv OCI^-) in the solution (10 mM PBS pH 7.4) incubated for 1.0 min at r.t.

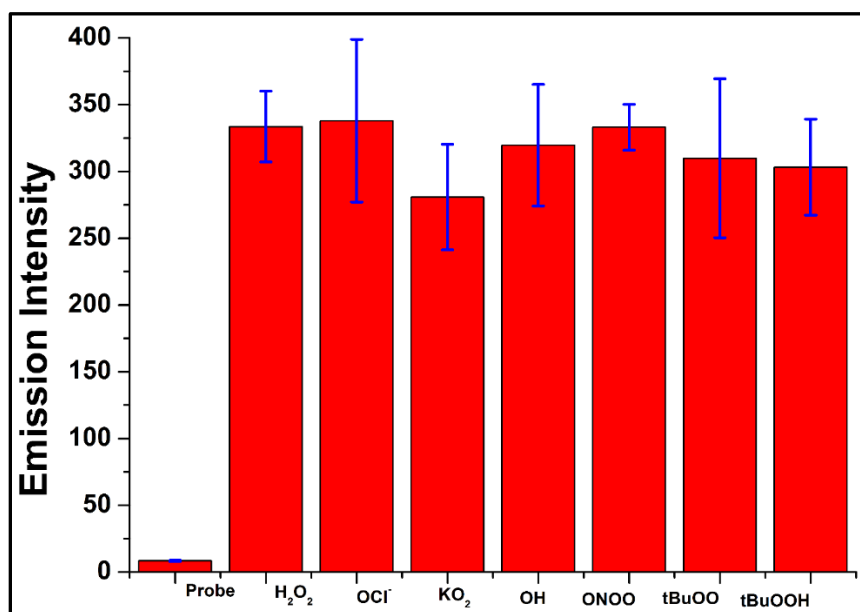


Figure S25. Fluorescence intensity of **Probe-OCI** (15.0 μM) with addition of OCI^- followed by addition of other ROS/RNS in solution (10 mM PBS pH 7.4) incubated for 1.0 min. λ_{ex} : 416 nm, λ_{em} : 523 nm, slit width 3.0 nm/ 3.0 nm.

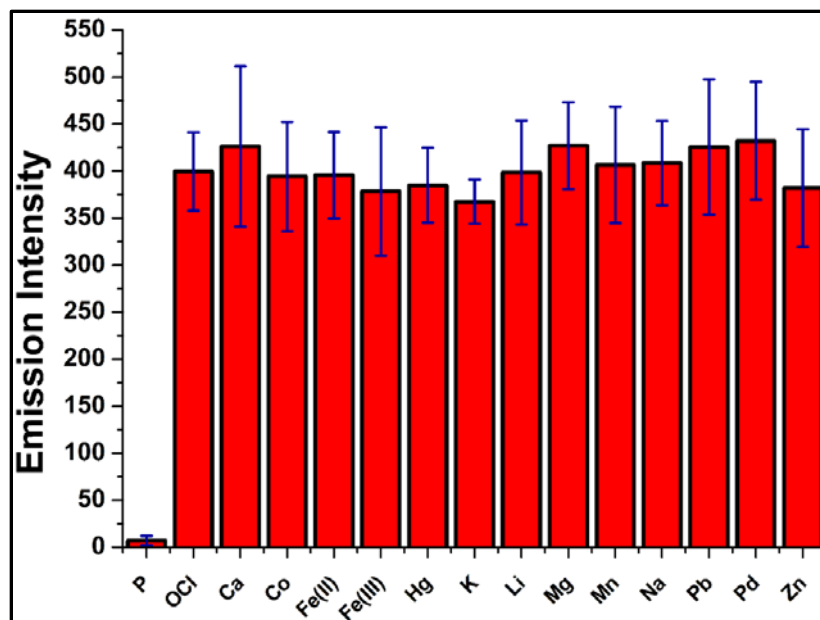


Figure S26. Fluorescence intensity of **Probe-OCl** (15.0 μM) with addition of OCl⁻ followed by the addition of various metals in solution (10 mM PBS pH 7.4) incubated for 1.0 min. λ_{ex} : 416 nm, λ_{em} : 523 nm, slit width 3.0 nm/3.0 nm.

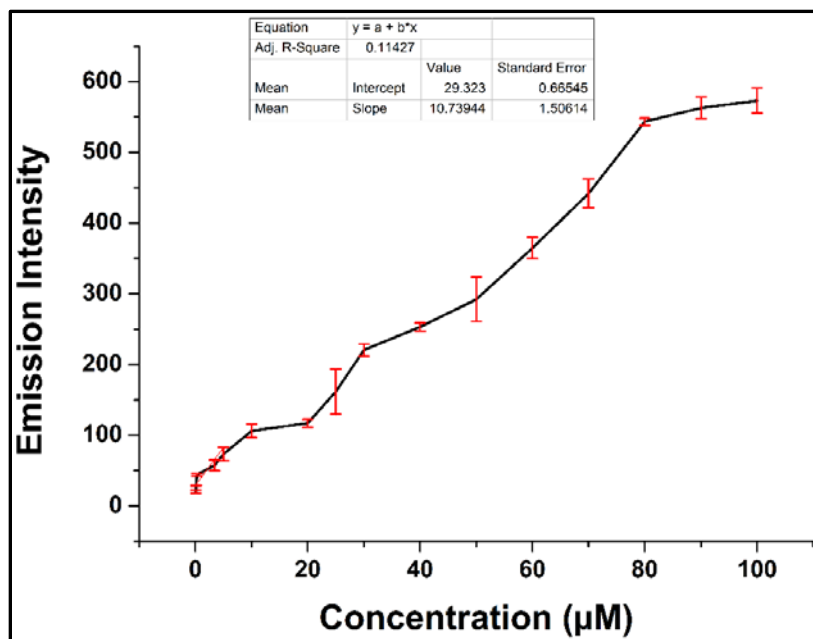


Figure S27. Plot for the calculation of limit of detection from the emission of **Probe-OCl** (15.0 μM) in the solution (10 mM PBS pH 7.4, 3:7 v/v) with increasing concentration of OCl⁻ (0.0 to 100.0 μM in water) incubated for 1.0 min at r.t, λ_{ex} . 416 nm and λ_{em} . 523 nm slit width 3.0 nm/3.0 nm (average of three experiments).

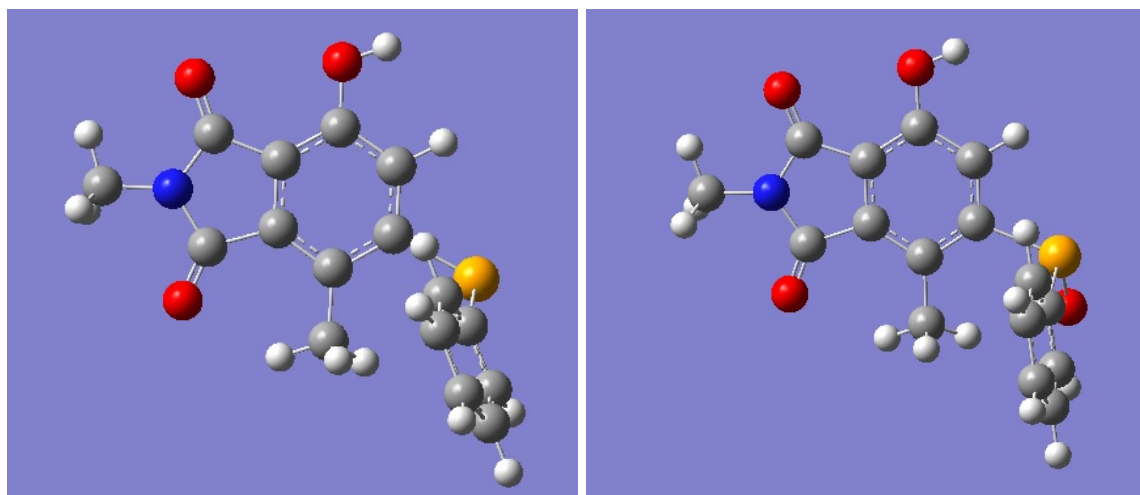


Figure S28. DFT-optimized geometries of (left) probe, and (right) oxidized probe (B3LYP/6-31g* basis set, G09).

	Probe	Oxidized probe
LUMO+2		
LUMO+1		

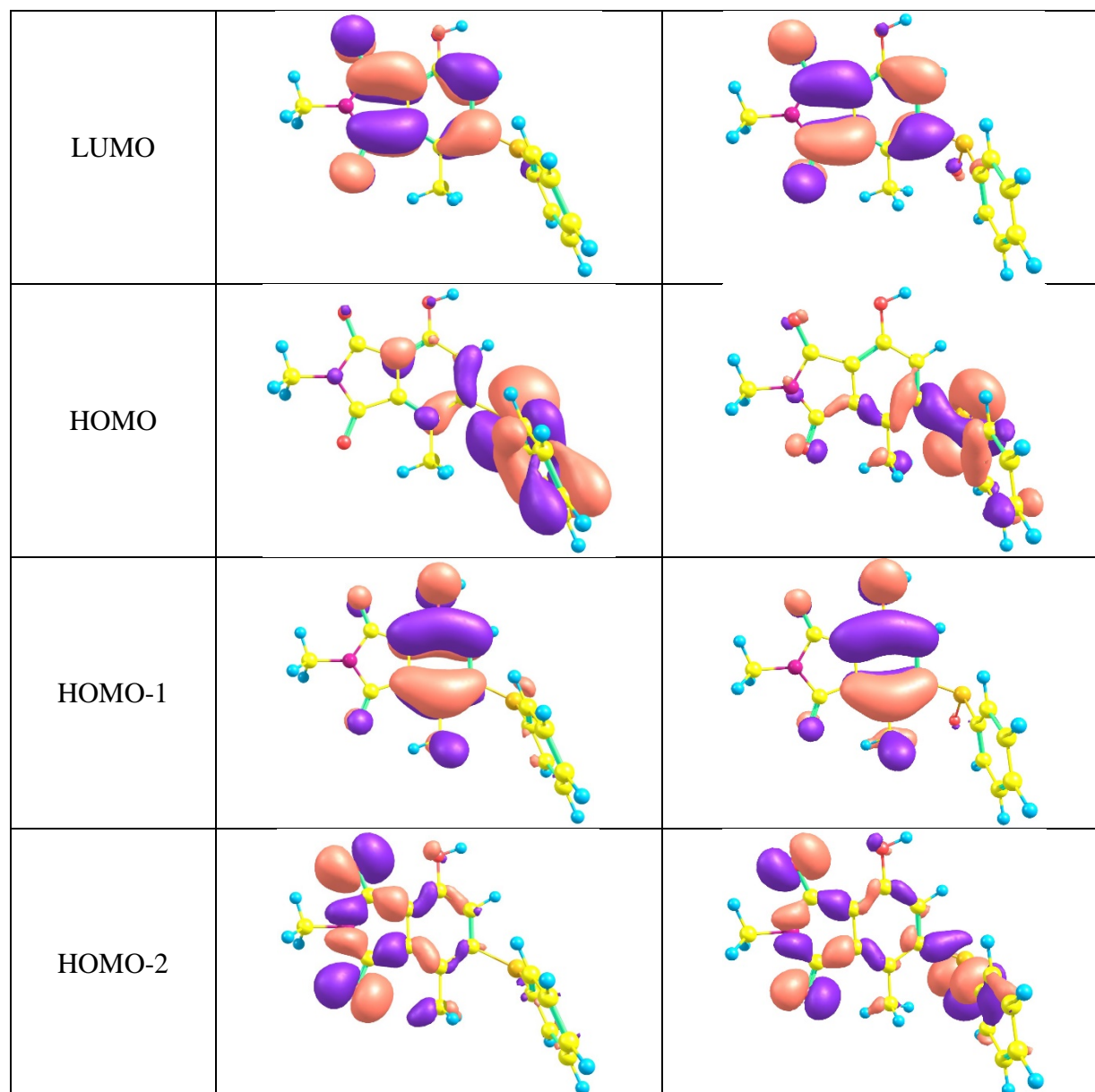


Figure S30 HOMO-LUMO of DFT-optimized geometries of probe and oxidized probe (B3LYP/6-31g* basis set and 6-311g* only for Se, G09).

	<i>f</i>	Composition	CI(%)
Probe	0.0358	HOMO → LUMO	98.4
	0.1158	HOMO-2 → LUMO	2.37
		HOMO-1 → LUMO	86.7
		HOMO → LUMO+1	5.29
Oxidized probe	0.0020	HOMO-2 → LUMO	7.47
		HOMO → LUMO	88.1
	0.1438	HOMO-1 → LUMO	90.7

Table S3. Absorption energies with largest oscillator strength for Probe and oxidized Probe (B3LYP/6-31g* basis set, G09)

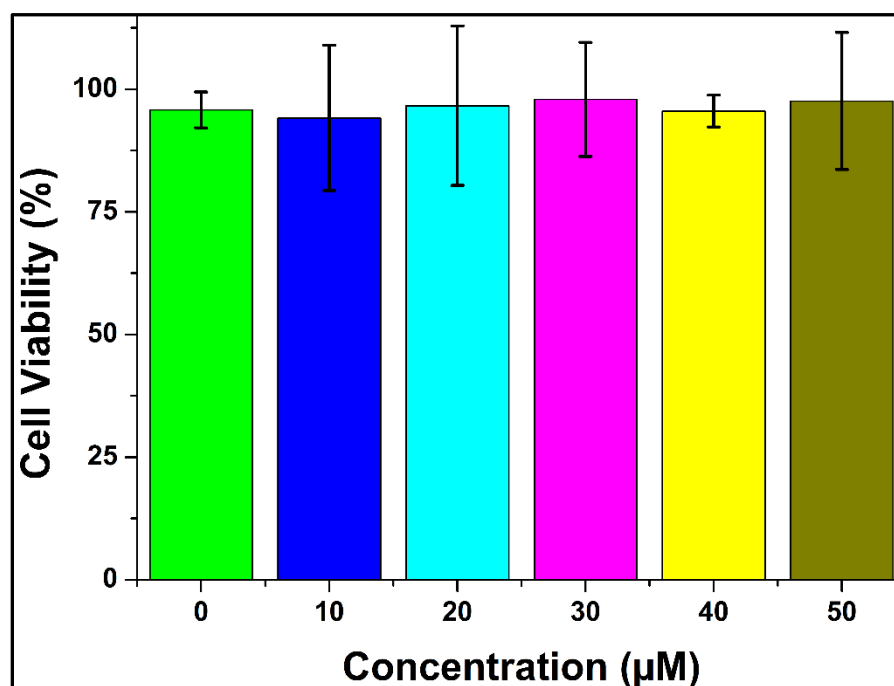


Figure S31. Cell viability was measured by MTT assay for concentration of **Probe-OCI** at 0 µM, 10 µM, 20 µM, 30 µM, 40 µM, 50 µM in 1 hour incubation in U-2 OS cells. Absorbance was determined at 540 nm. Data expressed as a mean ± SD for four experiments.

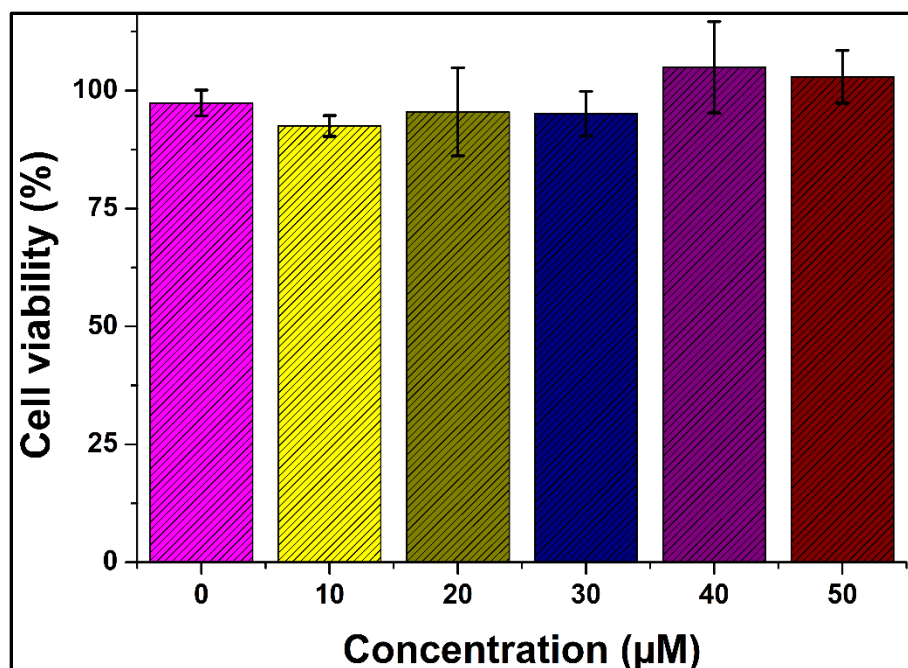


Figure S32. Cell viability was measured by MTT assay for concentration of **Probe-OCI** at 0 µM, 10 µM, 20 µM, 30 µM, 40 µM, 50 µM after 6 hours of incubation with U-2 OS cells. Absorbance was determined at 540 nm. Data is expressed as mean \pm SD for four experiments.

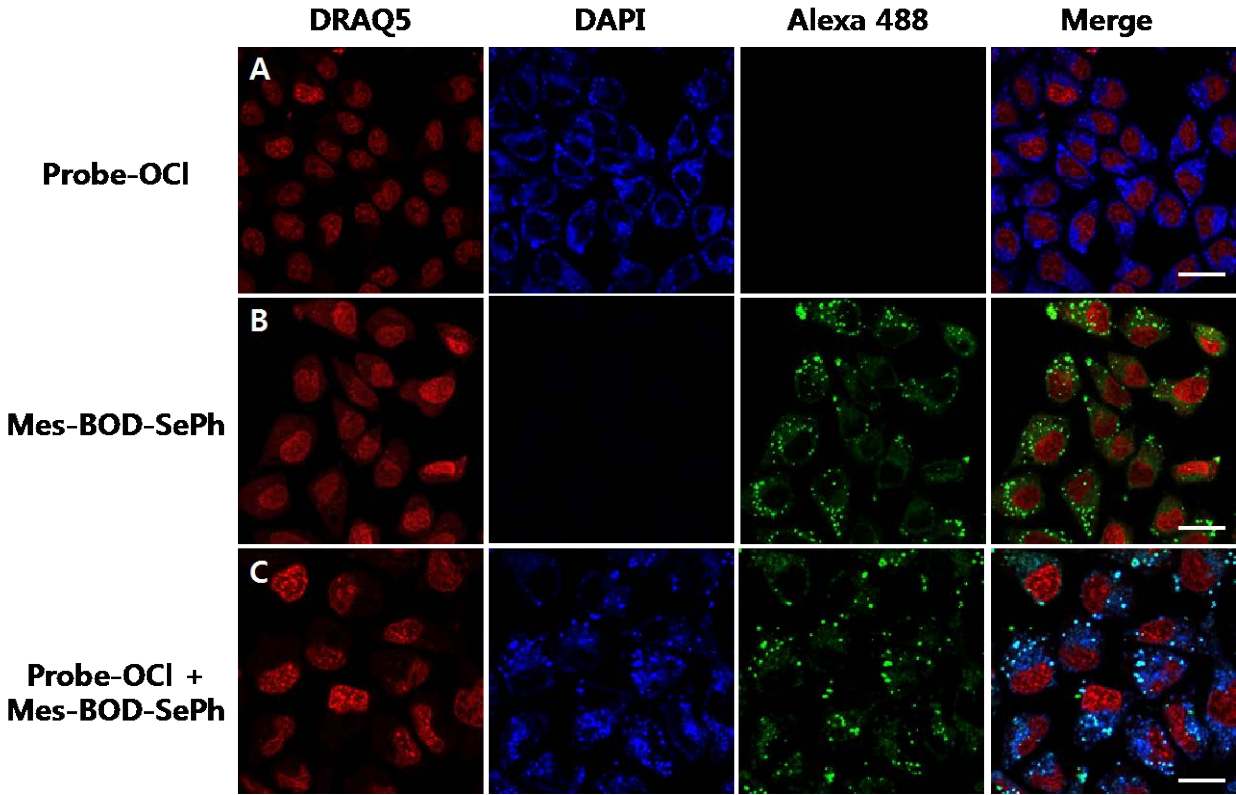


Figure S32. Live cell imaging in HeLa cells with **Probe-OCI** and **Mes-BOD-SePh**. [A] HeLa cells were incubated for 10 min with 25 μ M of **Probe-OCI** (blue) in PBS [B] 25 μ M **Mes-BOD-SePh** (green) [C] and counterstained for 10 min. **Probe-OCI** signals were consistent with lipid droplet staining dye **MES-BOD-SePh**. Scale bar, 10 μ m

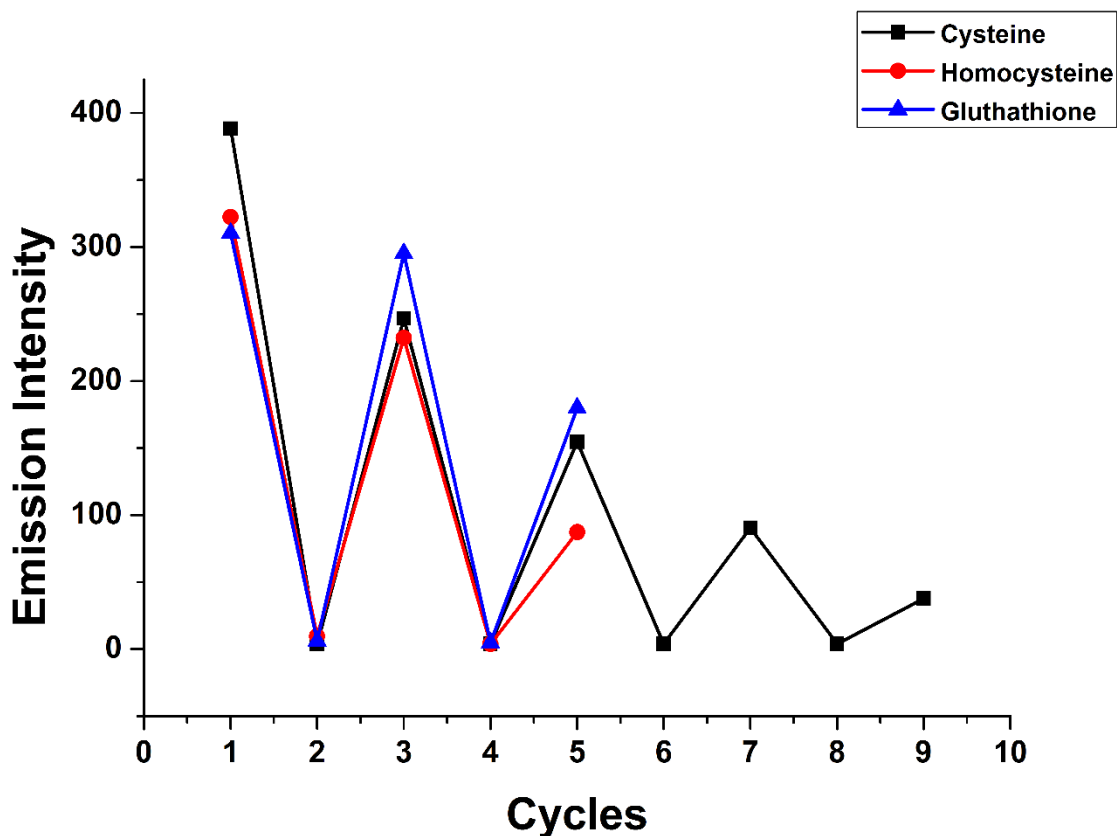


Figure S33. Reversibility of the **Probe-OCI** (15 μ M) with addition of 10 equiv. of L-cysteine (5 minutes incubation time), homocysteine, and glutathione (5 hours incubation time) in solution (10 mM PBS pH 7.4) incubated for 5 hours. λ_{ex} : 416 nm, λ_{em} : 523 nm. Width 3.0 nm/3.0 nm.

Reversibility of **Probe-OCI** was studied by added OCI^- (5 equiv) in three vials with 5 minutes of incubation time. The solution become green fluorescent (measured with 365 nm UV lamp) following the addition of cysteine, homocysteine, and glutathione with an incubation time of 5 minutes for cysteine and 5 hours for homocysteine and glutathione. We observed a significant decrease in fluorescence intensity after the addition of cysteine into the vial after 4 cycles.

Table S4. Results of experimental determination of log P value by the shake flask method for the **Probe-OCI**

C₁	V	A	\bar{A}	C₂	P	Log P
1.3×10^{-5}	3.5	0.0399 0.0491 0.0454	0.0415	1.24×10^{-5}	1.89	0.276
1.6×10^{-5}	4.8	0.0475 0.0429 0.0461	0.0455	1.36×10^{-5}	6.96	0.842
2.0×10^{-5}	6.0	0.0439 0.0409 0.0440	0.0428	1.28×10^{-5}	22.20	1.34

※ C₁ = Concentration (mol L⁻¹) of the stock solution in *n*-octanol before partition; V = volume (μL) of stock solution; A = absorbance in buffer solution after the partition (λ = 500 nm); \bar{A} = arbitrary absorbance in buffer solution after partitioning (λ = 500 nm); C₂ = concentration (mol L⁻¹) in buffer solution after partitioning; P = partition coefficient; log P = logarithm of the partition coefficient.

Properties (Probe-1) (NOTE: abbreviation same as website)	Value
milogP	3.37
TPSA	59.30
Natom	21
MW	346.24
nON	4
nOHNH	1
nviolations	0
nrotb	2
volume	259.99

Table S5. Information of the **Probe-OCI** calculated through 'molinspiration property engine v2011.04' at the website, <http://www.molinspiration.com>.

References:

1. C. A. Lipinski, F. Lombardo, B. W. Dominy, P. J. Feeney, *Adv. Drug Del. Rev.*, **1997**, 23, 3-25.