Supplementary information

Targeted Isolation of Sulfur-containing Metabolites from *Lsr2*-deletion Mutant Strain of *Streptomyces roseosporus*

Lina Deng,^a Rui Wang,^a Guowei Wang,^{*a} Mingxu Liu,^a Guojian Liao,^a Zhihua Liao,^b and Min Chen^{*a}

^a College of Pharmaceutical Sciences, Key Laboratory of Luminescent and Real-Time Analytical Chemistry (Southwest University), Ministry of Education, Southwest University, Chongqing 400715, P.R. China

^b School of Life Sciences, Southwest University, Chongqing 400715, P.R. China

List of supplementary information

Supplementary Experimental section

- Table S1. Sulfur-containing metabolites detected and characterized in $\Delta Srlsr2$ strain by UPLC-QTOF-MS and HR-MS/MS analysis.
- Table S2. Cytotoxic activities of 1-13 isolated from the $\Delta Srlsr2$ strain of S roseosporus
- Figure S1. HR-MS/MS spectra and proposed fragmentation pathway of peaks 17 and 18 (1 and 2).
- Figure S2. HPLC guided isolation of sulfur-containing metabolites.
- Figure S3. X-ray crystal structure of 10.
- Figure S4. ¹H NMR (400 MHz, DMSO) spectrum of compound 1.
- Figure S5. ¹³C NMR (100 MHz, DMSO) spectrum of compound 1.
- Figure S6. DEPT 135 (100 MHz, DMSO) spectrum of compound 1.
- Figure S7. HSQC (400 MHz, DMSO) spectrum of compound 1.
- Figure S8. HMBC (400 MHz, DMSO) spectrum of compound 1.
- Figure S9. HRESIMS spectrum of compound 1.
- Figure S10. ¹H NMR (400 MHz, DMSO) spectrum of compound **2**.
- Figure S11. ¹³C NMR (100 MHz, DMSO) spectrum of compound **2**.
- Figure S12. HSQC (400 MHz, DMSO) spectrum of compound 2.
- Figure S13. HMBC (400 MHz, DMSO) spectrum of compound 2.
- Figure S14. HRESIMS spectrum of compound 2.
- Figure S15. ¹H NMR (400 MHz, DMSO) spectrum of compound **3**.
- Figure S16. ¹³C NMR (100 MHz, DMSO) spectrum of compound **3**.
- Figure S17. DEPT 135 (100 MHz, DMSO) spectrum of compound 3.
- Figure S18. HSQC(400 MHz, DMSO) spectrum of compound 3.
- Figure S19. HMBC (400 MHz, DMSO) spectrum of compound 3.
- Figure S20. HRESIMS spectrum of compound 3.
- Figure S21. ¹H NMR (400 MHz, DMSO) spectrum of compound 4.
- Figure S22. ¹³C NMR (100 MHz, DMSO) spectrum of compound 4.
- Figure S23. DEPT 135 (100 MHz, DMSO) spectrum of compound 4.
- Figure S24. HSQC (400 MHz, DMSO) spectrum of compound 4.

Figure S25. HMBC (400 MHz, DMSO) spectrum of compound 4.

Figure S26. HRESIMS spectrum of compound 4.

Figure S27. ¹H NMR (400 MHz, DMSO) spectrum of compound 5.

Figure S28. ¹³C NMR (100 MHz, DMSO) spectrum of compound 5.

Figure S29. HSQC (400 MHz, DMSO) spectrum of compound 5.

Figure S30. HMBC (400 MHz, DMSO) spectrum of compound 5.

Figure S31. HRESIMS spectrum of compound 5.

Figure S32. ¹H NMR (400 MHz, DMSO) spectrum of compound 6.

Figure S33. ¹³C NMR (100 MHz, DMSO) spectrum of compound 6.

Figure S34. DEPT 135 (100 MHz, DMSO) spectrum of compound 6.

Figure S35. HSQC (400 MHz, DMSO) spectrum of compound 6.

Figure S36. HMBC (400 MHz, DMSO) spectrum of compound 6.

Figure S37. HRESIMS spectrum of compound 6.

Supplementary Experimental section

Cytotoxicity was evaluated against SGC7901, MDA-MB-231, A549 and HepG 2 cell lines by the MTT colorimetric method.¹ Cell lines were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% (v/v) fetal bovine serum, 2mM L-glutamine, 100 units/mL penicillin, and 100 ug/mL streptomycin at 37 °C in humidified 5% CO₂. For cytotoxicity assays, cells were seeded in 96-well plates at optimal cell denisity (10000 cells per well) to ensure exponential growth for the duration of the assay. After a 24h preincubation growth, the medium was replaced with experimental medium containing the appropriate drug concentration or vehicle controls (0.1% v/v DMSO). After 48 h incubation, cell viability was measured using Alamar Blue reagent according to the manufacturer's instructions. Absorbance was measured at 490 nm. Results were expressed as the mean \pm standard error for six replicates as a percentage of vehicle control (take as 100%). The percent growth was standardized to controls (0.1% DMSO as negative control) using Microsoft Excel 2013. A statistical analysis including IC₅₀ determination and graphical output was perform in GraphPad Prism 5 using nonlinear regression variable slope curve fitting.

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Peaks	t_R (min)	[M+H] ⁺ or [M+Na] ⁺	MS/MS	Molecular Formula	ppm	Tentatively Identification[Ref]	Molecular formula is previous reported in <i>Streptomyces</i> spp.
1	8.67	398.1508	266.1075,248.0944,231.0710,182.0502,160.1115,134.0465	$C_{24}H_{19}N_3O_3$	-0.2	unknown	no
2	9.57	215.1195	187.1225,170.0967,160.1130,143.0841,132.0812,103.0569	$C_{13}H_{14}N_2O$	5.0	unknown	no
3	10.25	261.1249	233.1293,215.1190,170.0687,120.0818,114.0560	$C_{14}H_{16}N_2O_3$	3.8	cyclo(L-6-Hyp-L-Phe)[1]	Streptomyces sp. DA18
4	10.64	280.0531#	244.0552,228.0589,214.0445,189.0488,188.0386, 182.0722,157.0761,130.0654	$C_{13}H_{11}N_3OS$	3.5	SF2738D [2-3]	Streptomyces sp.
5	11.04	285.0668#	249.0688,233.0752,219.0598,203.0646,189.0480, 188.0385,157.0765,130.0657	$C_{13}H_{14}N_2O_2S$	-2.1	unknown	Streptomyces sp. TPU1236A and other Streptomycesspecies
6	11.31	285.0669#	247.0915,233.0745,219.0595,203.0647,189.0481, 188.0389,157.0760,130.0655	$C_{13}H_{14}N_2O_2S$	-1.7	SF2738C [2-3]	Streptomyces sp. TPU1236A and other Streptomyces species
7	12.69	292.0751	274.0642,260.0487,244.0541,230.0391,214.0431,189.0479 ,188.0385,182.0712,157.0771,130.0658	$C_{13}H_{13}N_3O_3S$	1.7	Pyrisulfoxin A [3-4]	Streptomyces californicus
8	13.23	271.0598	243.0619,215.0697,197.0591,169.0630,153.0174, 149.0234,91.0547,65.0384	$C_{15}H_{10}O_5$	3.0	unknown	Streptomyces sp. AC35 and other Streptomyces species
9	13.75	291.0794	273.0695,259.0539,243.0593,229.0446,204.0485, 188.0537,181.0759,156.0802,130.0660	$C_{14}H_{14}N_2O_3S$	-3.1	unknown	no
10	14.39	219.1751	203.1433,188.1218,161.1317,147.1147,133.1024,119.0842	$C_{15}H_{22}O$	0.9	unknown	no
11	15.01	304.1129	290.0972,262.1018,260.0850,247.0900,233.0741,232.0915 ,203.0647,189.0490,188.0385,157.0770,130.0655	$C_{15}H_{17}N_3O_2S$	3.3	SF2738E[2-3]	Streptomyces sp.
12	16.41	252.0763#	212.0831,198.0672,182.0717,157.0766,130.0654	$C_{12}H_{11}N_3O_2$	4.7	Caerulomycin A [3,5]	Streptomyces caeruleus
13	17.23	261.1613	233.1660,170.1052,130.1229,120.0811	$C_{15}H_{20}N_2O_2$	3.8	Cyclo-(L-phe-L-Leu) [6]	Streptomyces sp. H7372 and other Streptomyces species
14	18.14	274.0644	244.0543,230.0342,214.0435,189.0482,188.0389, 182.0715,157.0762,130.0655	$C_{13}H_{11}N_3O_2S$	-2.1	Pyrisulfoxin B [3-4]	Streptomyces californicus
15	18.22	273.0694	259.0545,243.0590,229.0441,204.0480,188.0540, 181.0770,156.0810,130.0658	$C_{14}H_{12}N_2O_2S$	-1.4	unknown	no
16	10.22	270.0701	263.0850,249.0693,235.0537,219.0597,203.0650,	CHNOS	0.4	unknown	<u>no</u>

			189.0484,157.0763,130.0652					
17	20.31	276.0797	258.0693,244.0537,214.0439,212.0824,189.0487,	$C_{13}H_{13}N_3O_2S$	-3.6	unknown	Strontomygog gn CS40	
			182.0712,157.0768,130.0655				Streptomyces sp. CS40	
18	20.43	276.0794	231.0583,217.0434,216.0348,189.0479,188.0388,	$C_{13}H_{13}N_3O_2S$	-4.7	unknown	Streptomyces sp. CS40	
10	20.43	270.0794	185.0707,157.0761,130.0652	$C_{13} 1_{13} 1_{3} 0_{2} 0_{2} 0_{3}$	+./	UIIKIIOWII	Sucptomyces sp. C340	
19	21.44	276.0795	258.0701,244.0536,214.0441,212.0821,189.0486,	C ₁₃ H ₁₃ N ₃ O ₂ S	-4.3	.3 SF2738A/B [2-3]	Streptomyces sp. CS40	
17	21.44	210.0195	182.0713,157.0765,130.0657	$C_{13} 1_{13} 1_{3} C_{2} S_{3}$	-4.5	51 ² /50A/D [2-5]	Sucptomyces sp. C340	
20a	21.74	284.2235	266.2128,232.1345,212.1166,173.1198,143.0851	$C_9H_{29}N_7OS$	0.7	unknown	no	
20b	21.74	284.1387	256.2277,219.1737,194.0954,160.1081,147.1065,133.0856	C ₁₆ H ₁₇ N ₃ O ₂	-4.2	unknown	Streptomyces sp. H7372 and	
200	21.74	204.1307	230.2277,217.1757,174.0754,100.1081,147.1005,155.0830	$C_{16} 1_{17} N_3 O_2$	-4.2	UIIKIIOWII	other Streptomyces species	
21	22.68	298.2400	280.2274,270.2439,239.1414,184.0747,160.1102,117.0930	$C_{10}H_{31}N_7OS$	4.0	unknown	no	
22	23.48	330.3370	312.3283,298.2385,266.2117,233.1741,209.1686,	C ₂₀ H ₄₃ NO ₂	-0.6	unknown	no	
			175.1346,131.0721					
	-							

[M+Na]+

References

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compound	IC ₅₀ (µM)						
	SGC7901	MDA-MB-231	A549	HepG2			
1	>40	>40	>40	>40			
2	>40	>40	>40	>40			
3	>40	>40	>40	>40			
4	>40	>40	>40	>40			
5	>40	>40	>40	>40			
6	>40	>40	>40	>40			
7	1.7 ± 0.5	6.3 ± 1.0	14.7 ± 3.2	5.8 ± 0.5			
8	>40	>40	>40	>40			
9	>40	>40	21.3 ± 2.9	>40			
10	>40	>40	>40	>40			
11	>40	34.5 ± 5.3	25.8 ± 0.7	7.2±1.1			
12	>40	>40	>40	>40			
13	>40	>40	>40	>40			

Table S2 Cytotoxic activities of **1-13** isolated from the $\Delta Srlsr2$ strain of *S* roseosporus

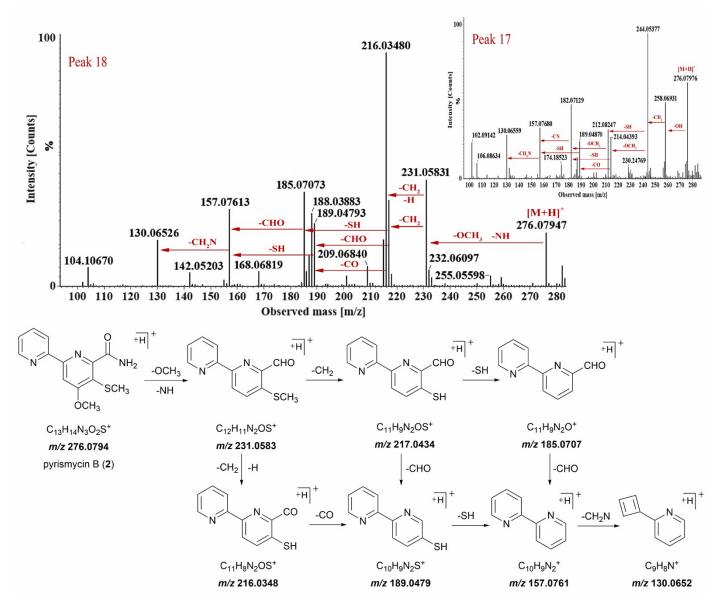


Figure S1 HR-MS/MS spectra and proposed fragmentation pathway of peaks 17 and 18 (1 and 2).

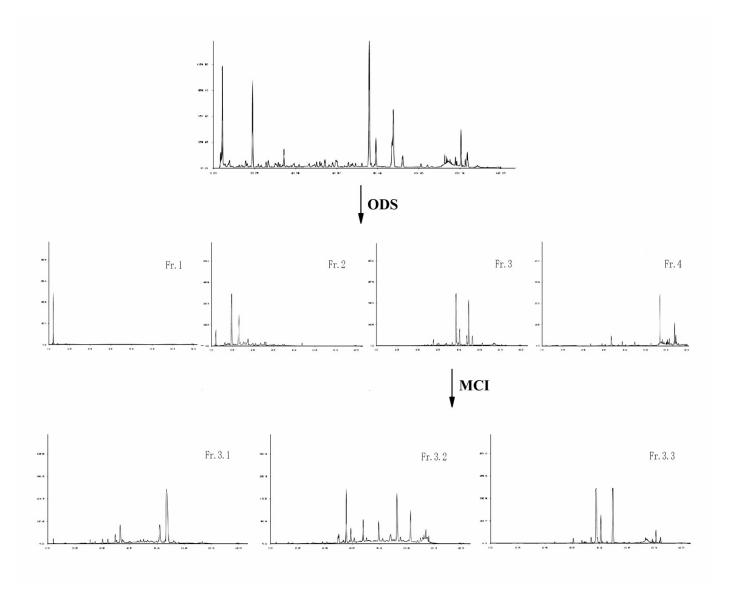


Figure S2. HPLC guided isolation of sulfur-containing metabolites.

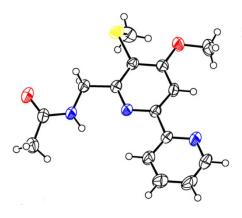
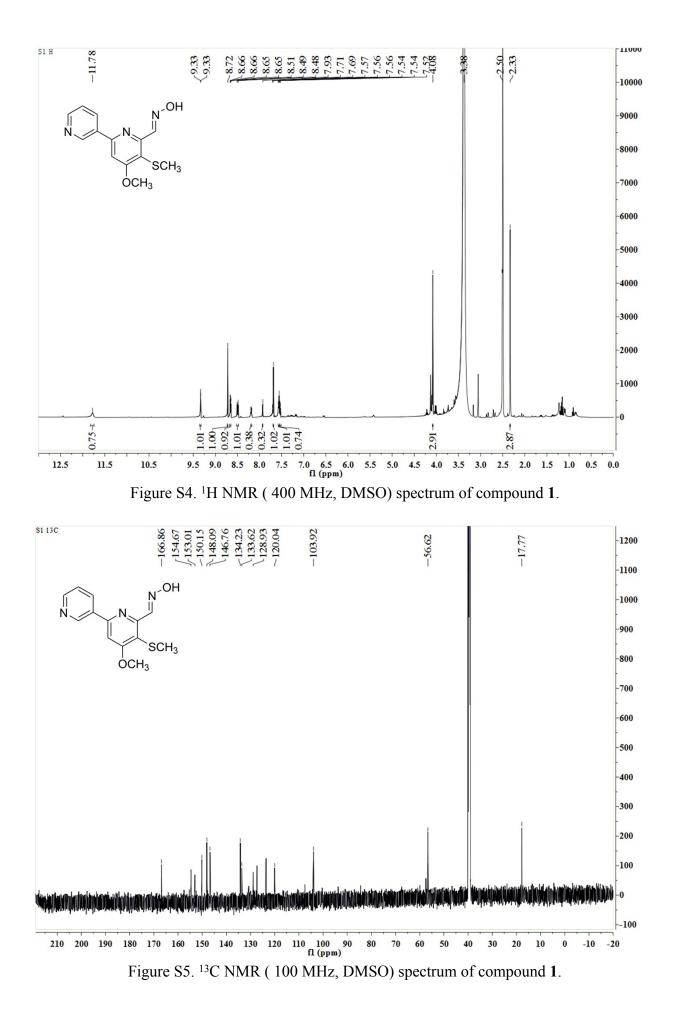


Figure S3. X-ray crystal structure of 10



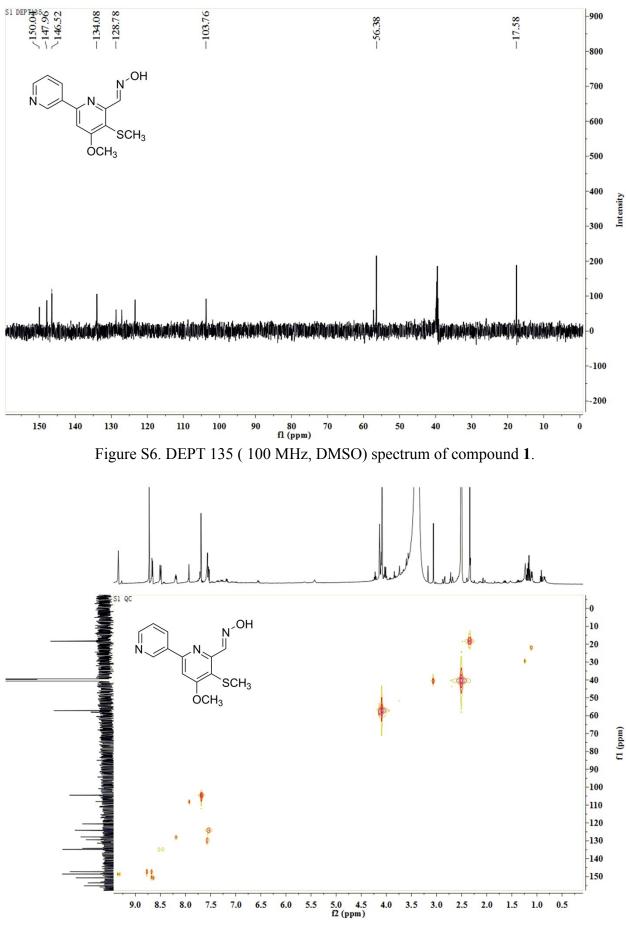


Figure S7. HSQC (400 MHz, DMSO) spectrum of compound 1.

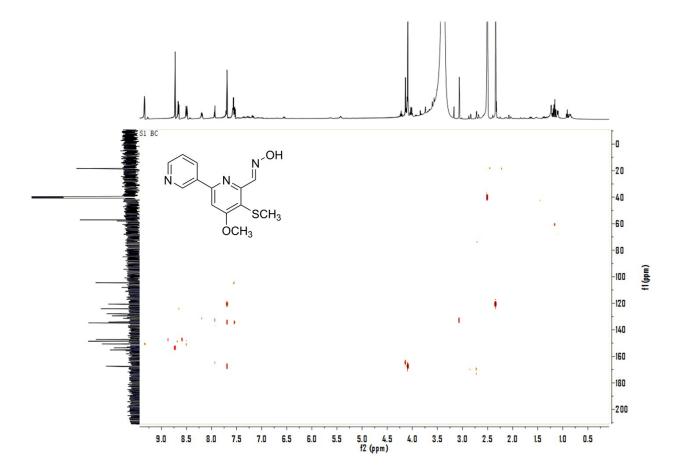


Figure S8. HMBC (400 MHz, DMSO) spectrum of compound 1.

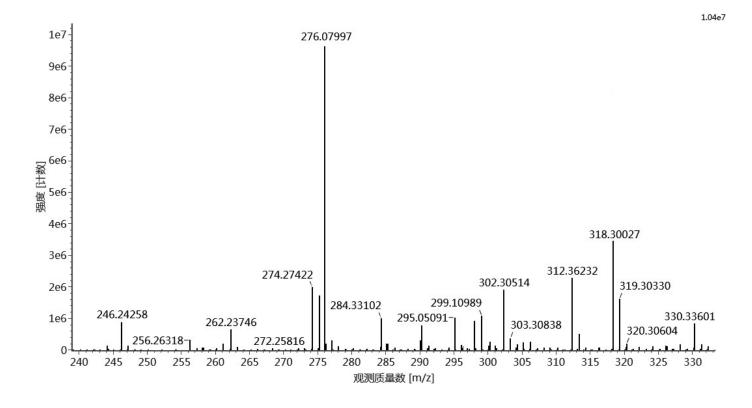


Figure S9. HRESIMS spectrum of compound 1.

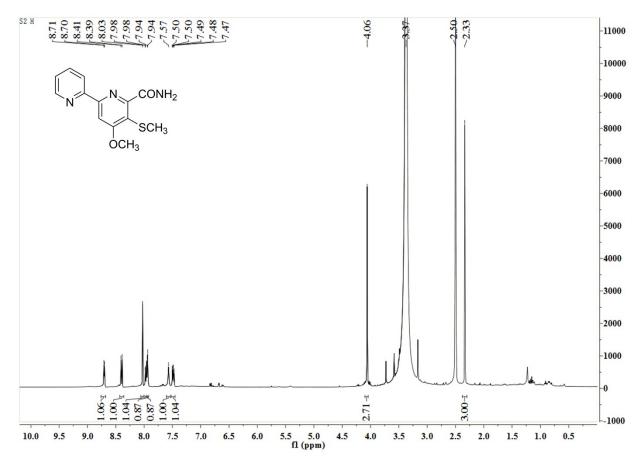


Figure S10. ¹H NMR (400 MHz, DMSO) spectrum of compound 2.

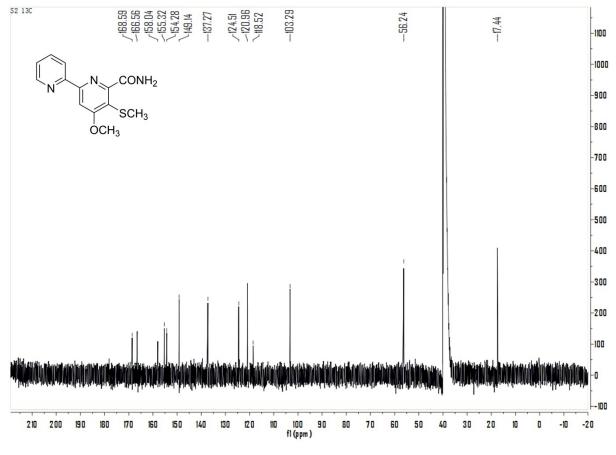
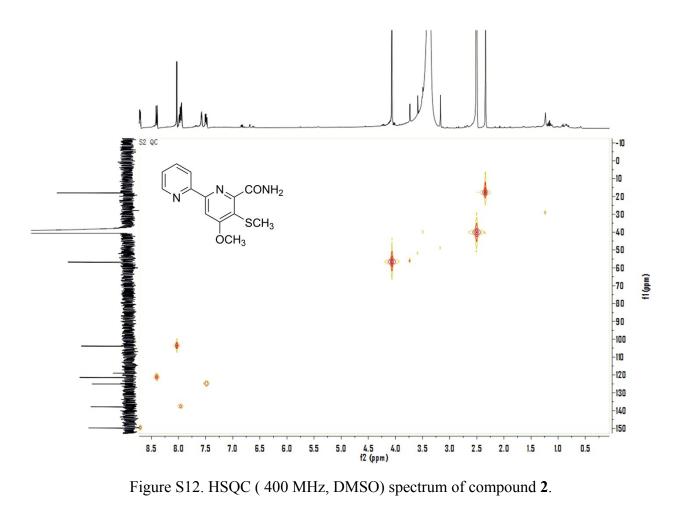


Figure S11. ¹³C NMR (100 MHz, DMSO) spectrum of compound **2**.



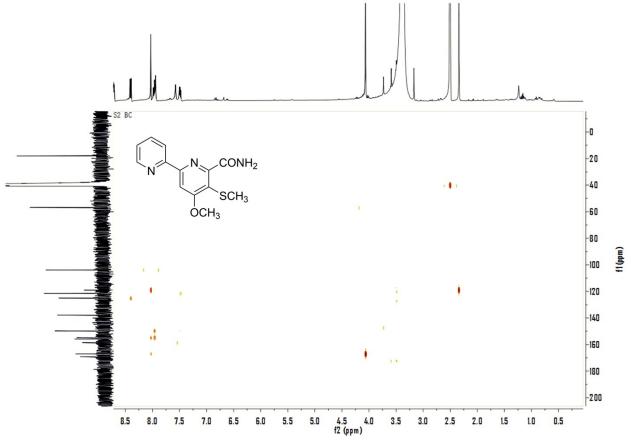


Figure S13. HMBC (400 MHz, DMSO) spectrum of compound 2.

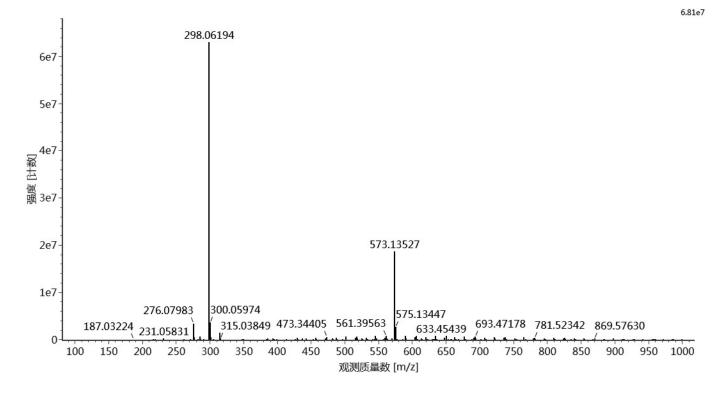


Figure S14. HRESIMS spectrum of compound 2.

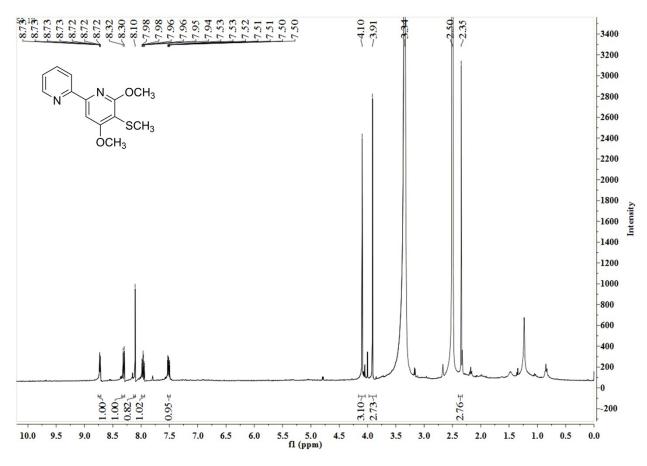
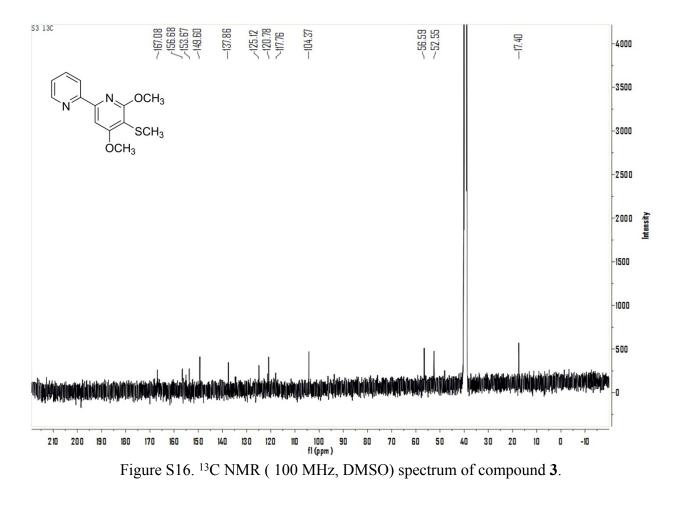


Figure S15. ¹H NMR (400 MHz, DMSO) spectrum of compound **3**.



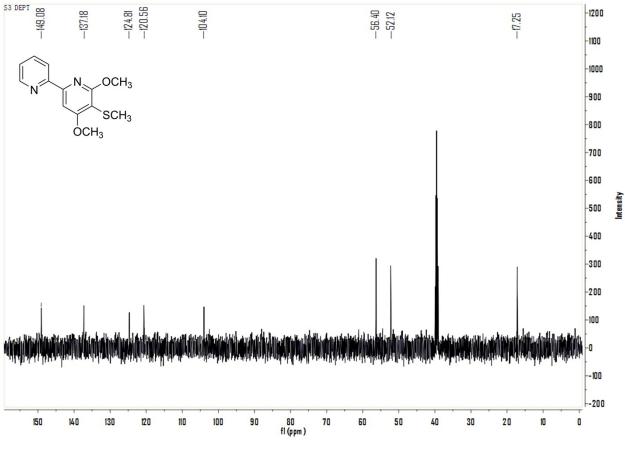


Figure S17. DEPT 135 (100 MHz, DMSO) spectrum of compound 3.

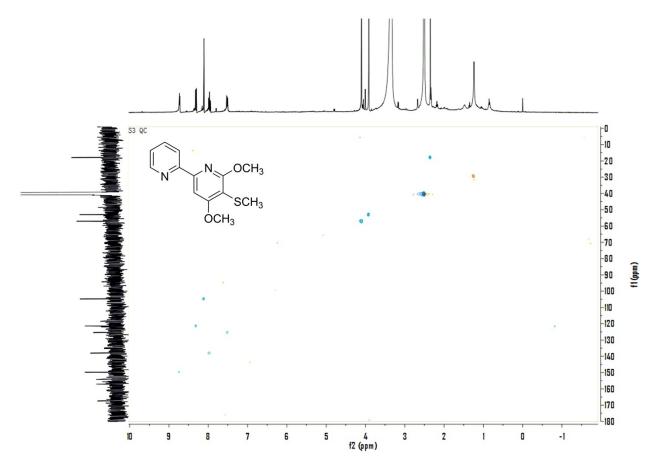


Figure S18. HSQC(400 MHz, DMSO) spectrum of compound 3.

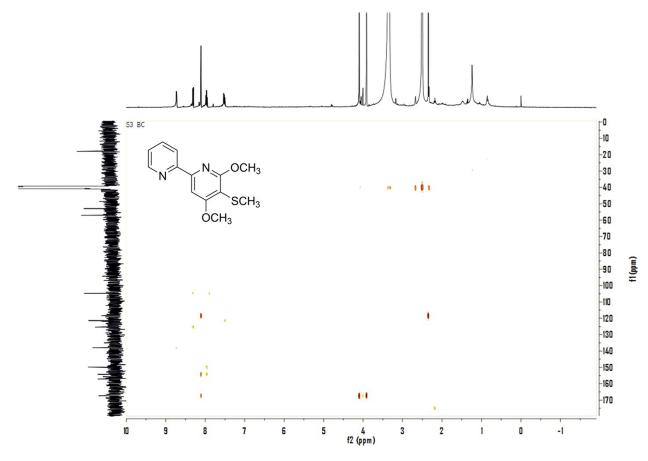
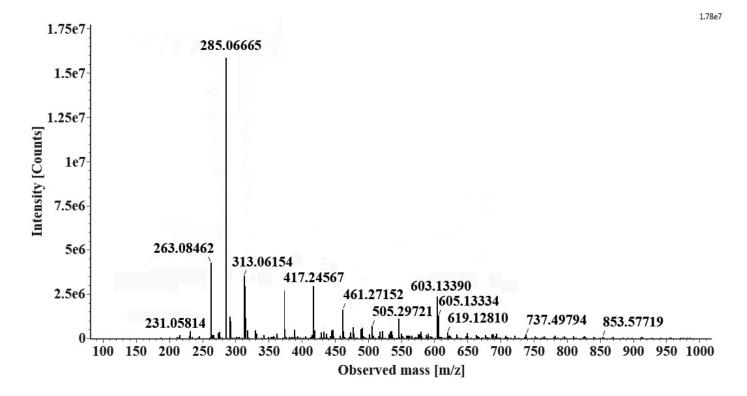


Figure S19. HMBC (400 MHz, DMSO) spectrum of compound 3.





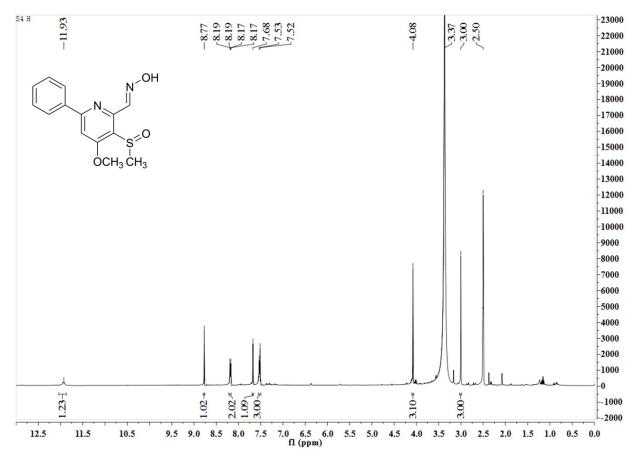


Figure S21. ¹H NMR (400 MHz, DMSO) spectrum of compound 4.

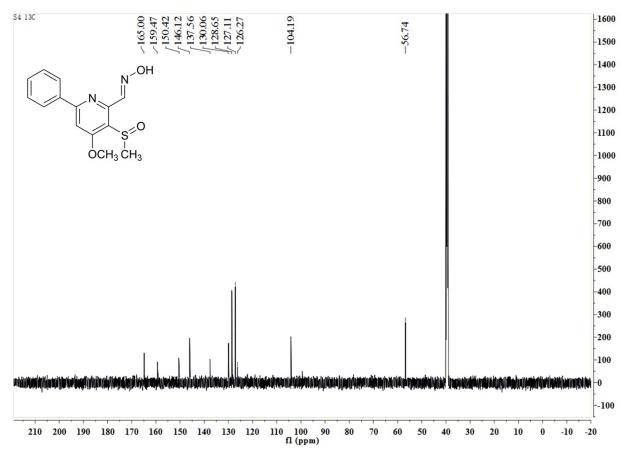


Figure S22. ¹³C NMR (100 MHz, DMSO) spectrum of compound 4.

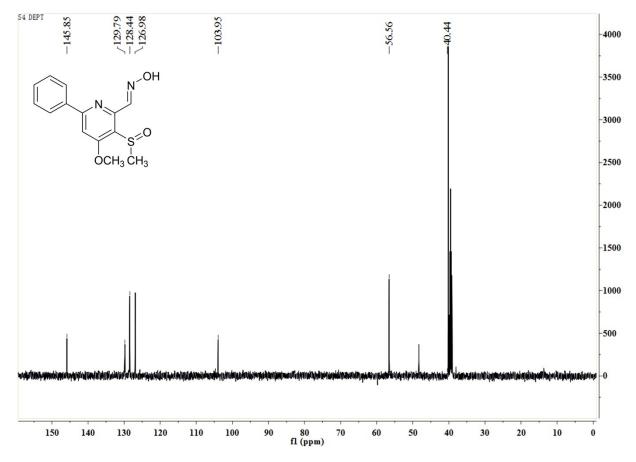


Figure S23. DEPT 135 (100 MHz, DMSO) spectrum of compound 4.

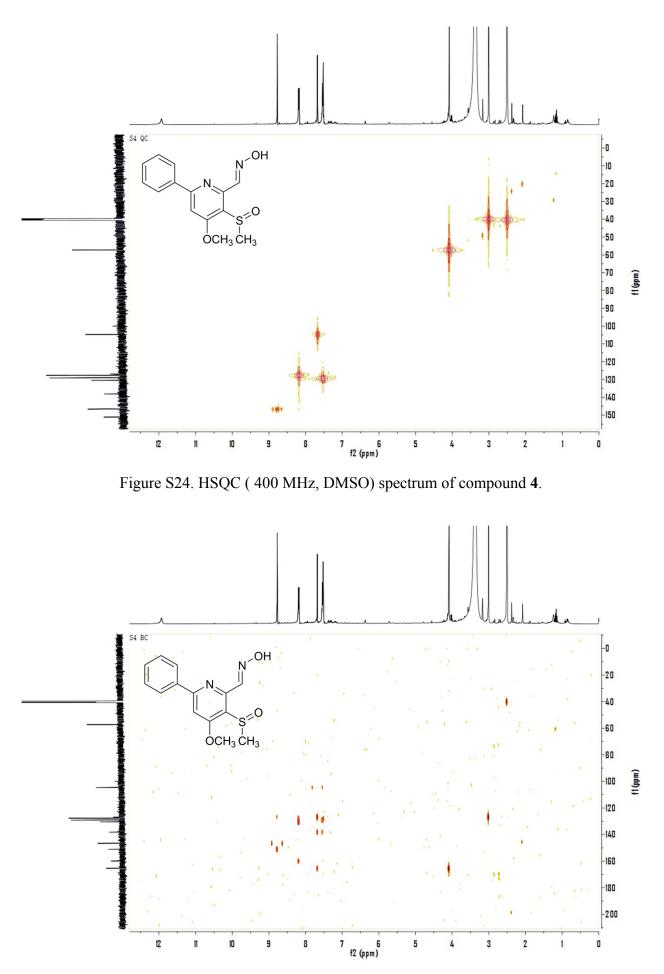
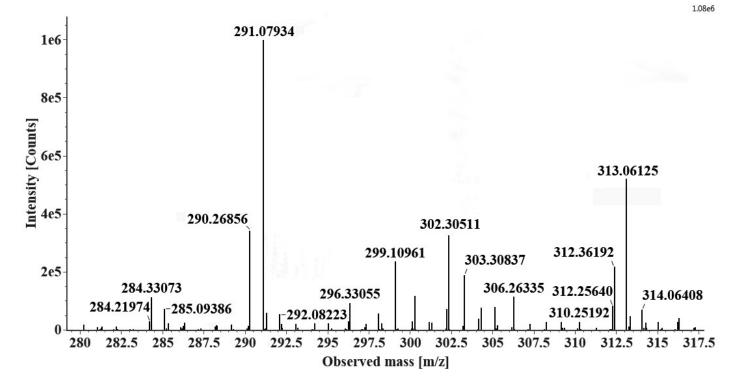
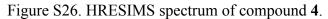


Figure S25. HMBC (400 MHz, DMSO) spectrum of compound 4.





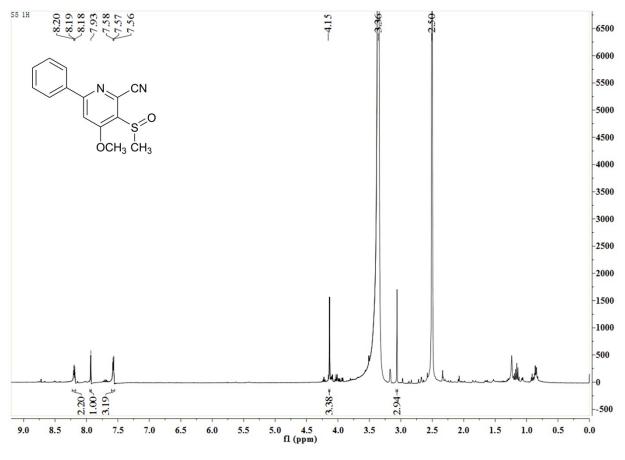


Figure S27. ¹H NMR (400 MHz, DMSO) spectrum of compound 5.

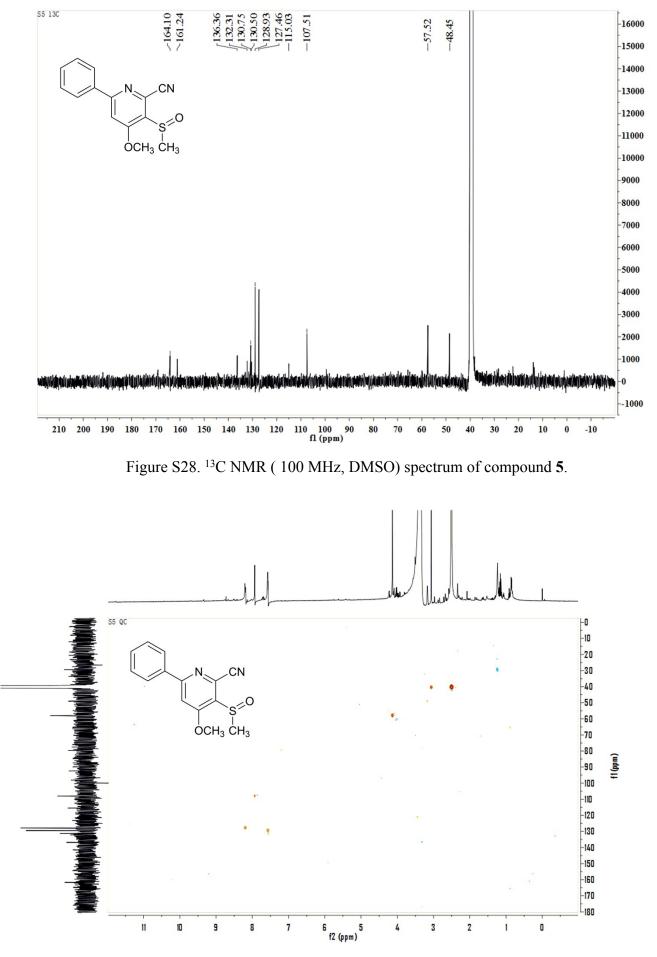


Figure S29. HSQC (400 MHz, DMSO) spectrum of compound 5.

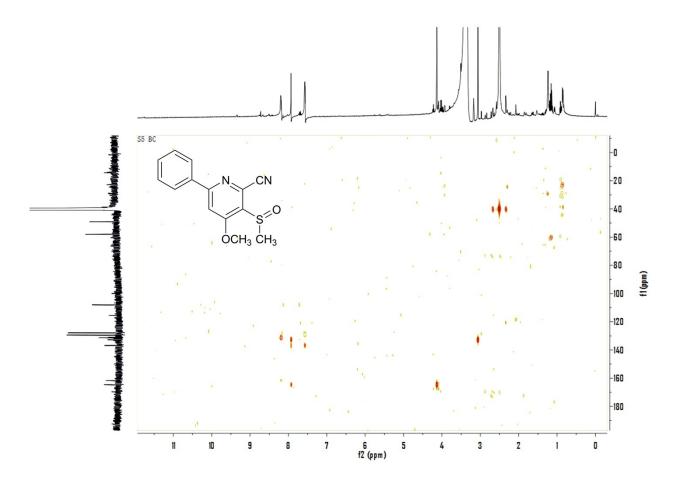


Figure S30. HMBC (400 MHz, DMSO) spectrum of compound 5.

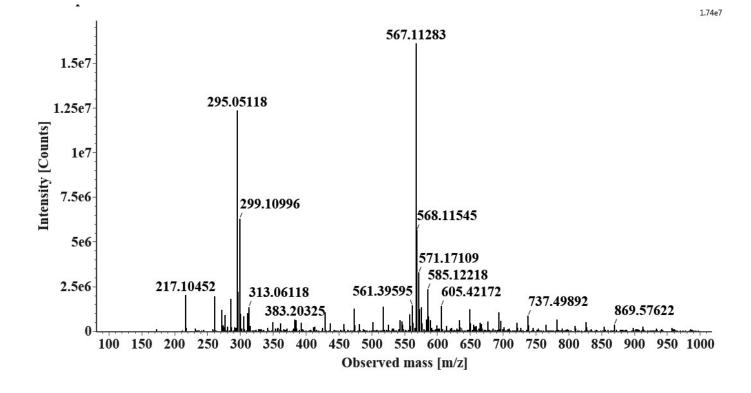


Figure S31. HRESIMS spectrum of compound 5

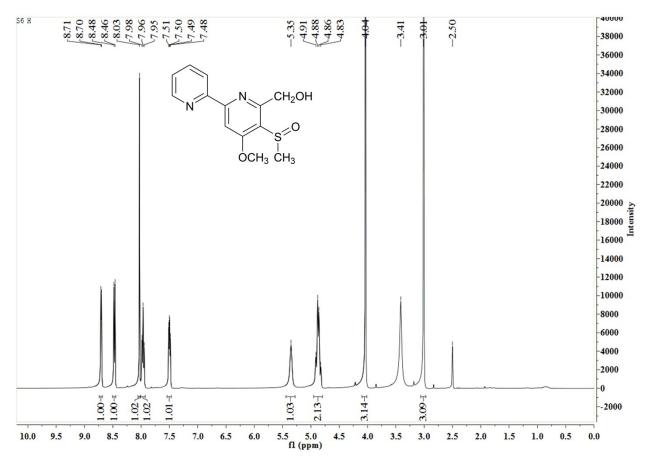


Figure S32. ¹H NMR (400 MHz, DMSO) spectrum of compound 6.

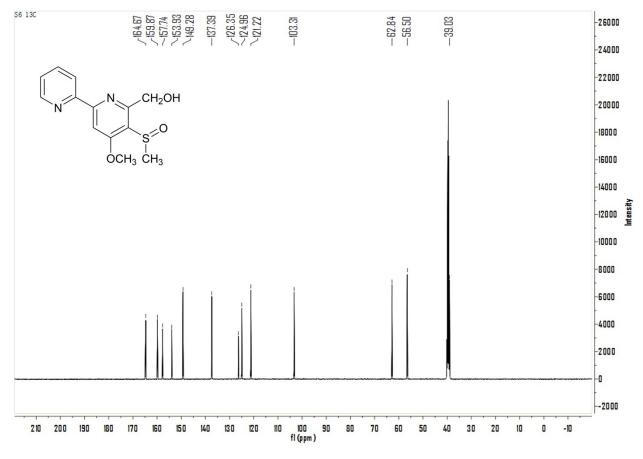
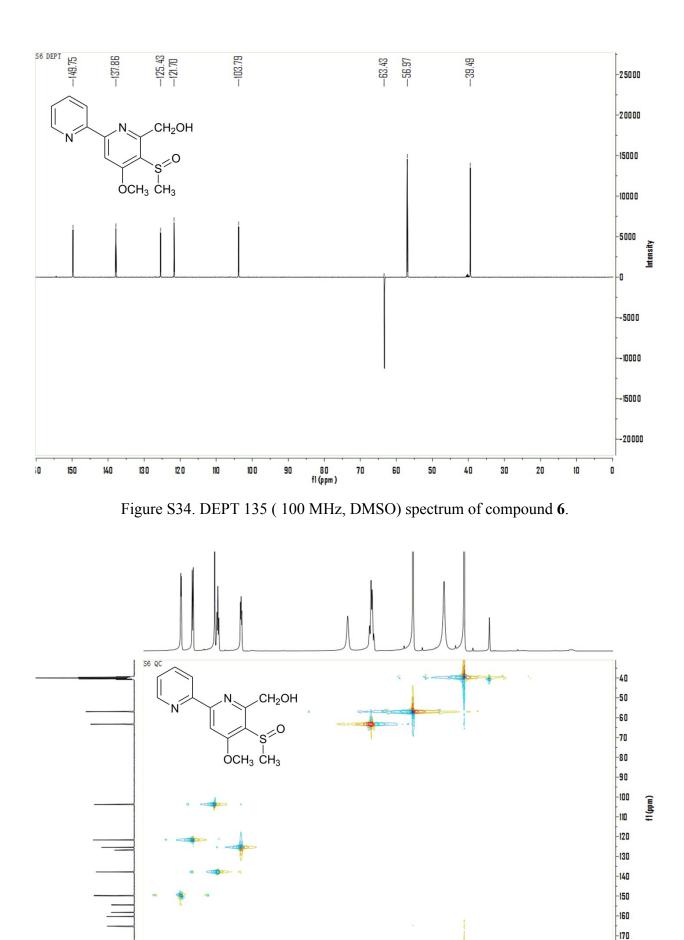


Figure S33. ¹³C NMR (100 MHz, DMSO) spectrum of compound 6.



5.0 4.5 f2 (ppm) 6.5 6.0 5.5 2.0 1.5 Figure S35. HSQC (400 MHz, DMSO) spectrum of compound 6.

4.0

3.5 3.0 2.5

1.0 0.5

9.0 8.5 8.0

7.5 7.0

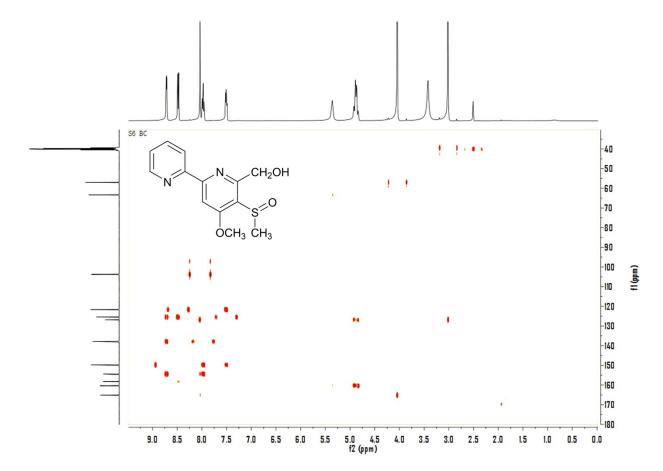


Figure S36. HMBC (400 MHz, DMSO) spectrum of compound 6.

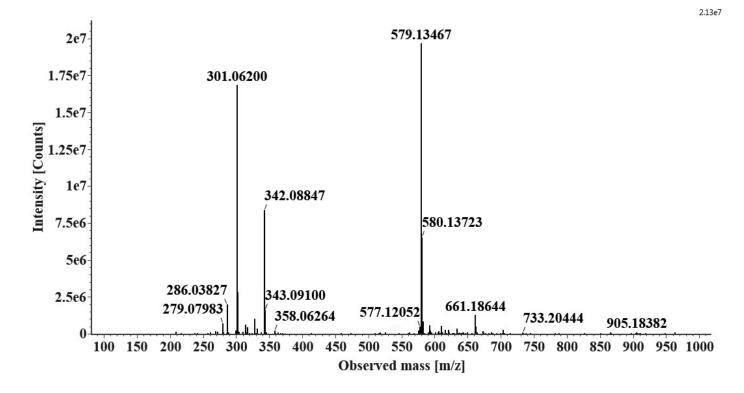


Figure S37. HRESIMS spectrum of compound 6.