

Supporting information

In vitro* anti-inflammatory, *in silico* molecular docking model and molecular dynamic study of oleanane-type triterpenes from aerial parts of *Mussaenda recurvata

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Table S2. NMR data of related compounds

No	Scutellaric acid (1C)			Hederagenin (1D)			Wilforol C (1A)			3 α , 24-dihydroxyolean-12-en-28-oic acid (1B)			3 β , 6 β , 24-trihydroxyolean-12-en-28-oic acid (3A)			Spathodic acid (2A)			Ilexosapogenin A (3B)			3 β , 6 β , 23-trihydroxyolean-12-en-28-oic acid (3)			3 β , 6 β , 19 α , 23-tetrahydroxyolean-12-en-28-oic acid (4)		
	[1]	[2]	[3]	[4]	[5]	[6]	[7]	[8]	[9]	[10]	[11]	[11]	[11]	[11]	[11]												
	DMSO- <i>d</i> ₆	Pyridine- <i>d</i> ₅	Pyridine- <i>d</i> ₅	Acetone- <i>d</i> ₆	DMSO- <i>d</i> ₆ +CDCl ₃	DMSO- <i>d</i> ₆	Pyridine- <i>d</i> ₅	DMSO- <i>d</i> ₆	Pyridine- <i>d</i> ₅	Pyridine- <i>d</i> ₅	Pyridine- <i>d</i> ₅	Pyridine- <i>d</i> ₅	Pyridine- <i>d</i> ₅	Pyridine- <i>d</i> ₅													
	¹ H	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C	¹ H	¹³ C		
3	3.55	4.23	73.5	3.94	75.7	3.88	76.5	3.77	68.8	4.29	72.6	4.23	78.4	4.30	73.3	4.31	73.3	4.04	73.3	4.04	73.4						
5			48.7		43.6		43.7				48.7		48.7		48.6		48.4		49.3		49.7						
6										5.06	66.9							4.85	67.5	4.83	67.9						
19												80.1	3.73	81.2	3.18	81.2		4.85	67.5	4.83	67.9						
CH ₂ -OH	3.43 3.19	4.18 3.73	67.8	3.88 3.68	71.3	3.55 3.44	71.3	3.65 3.37	64.4	4.41 4.05	66.4	4.18 3.73	63.0	4.29 3.82	64.5	4.28 3.82	67.8	4.18 3.83	67.1	4.19 3.85	67.3						
CH ₃ 24/23	0.68	1.05	13.3	0.79	18.2	0.69	18.3	0.89		1.73	14.1	1.05	22.9	1.10 1.17	13.6	1.10 1.16	13.0	1.51	14.7	1.53	14.6						
25	0.87	0.97	16.1	0.96	15.8	0.91	16.0	1.05		1.67	16.8	0.97	15.3	1.19	15.8	1.19	15.8	1.47	17.4	1.48	17.3						
26	1.10	1.04	17.7	1.07	17.5	0.97	17.7	0.76		1.63	17.9	1.04	16.6	1.22	17.5	1.22	17.5	1.42	18.6	1.46	18.4						
27	1.10	1.26	26.3	1.23	26.2	1.19	26.5	1.14		1.25	25.6	1.26	24.5	1.29	24.8	1.29	24.8	1.06	26.2	1.42	24.8						
29	0.87	0.95	33.4	0.92	33.3	0.82	33.5	0.89	33.1	0.91	32.6	0.95	28.1	1.72	28.9	1.71	28.9	0.80	33.2	0.99	28.8						
30	0.88	1.02	23.9	1.00	23.8	0.94	24.0	0.89	23.6	0.99	23.1	1.02	24.0		24.8		24.8	0.72	23.7	0.92	25.0						

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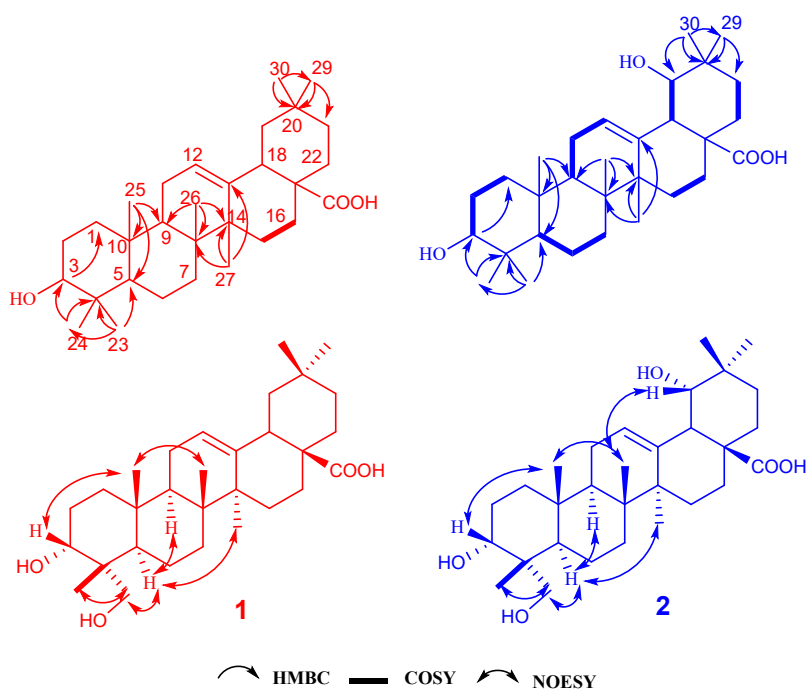


Figure S1. Selected HMBC and NOESY correlations of **1** and **2**

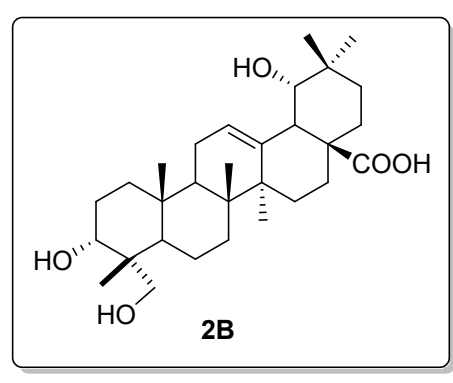
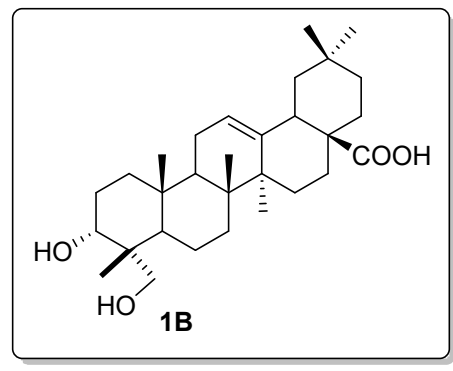
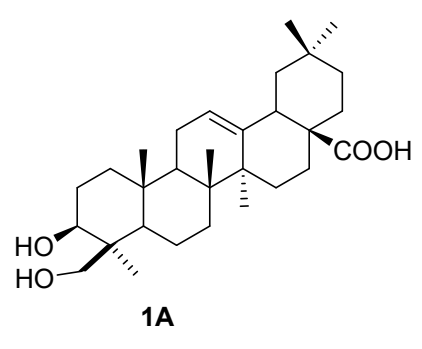
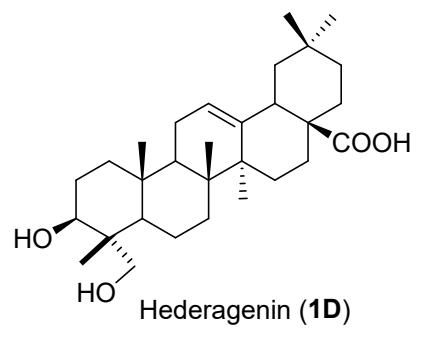
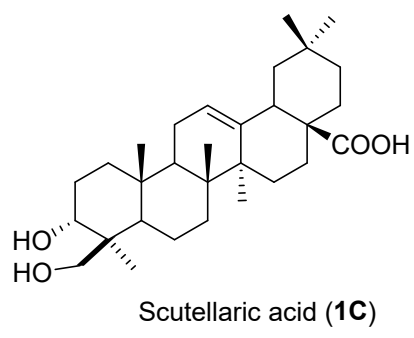


Figure S2. Four possible stereoisomers of 1

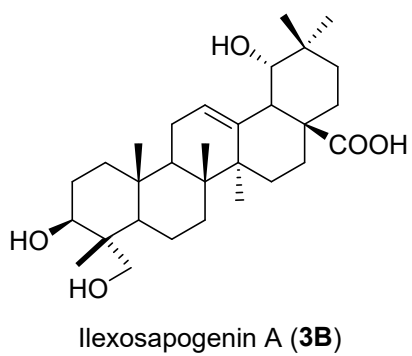
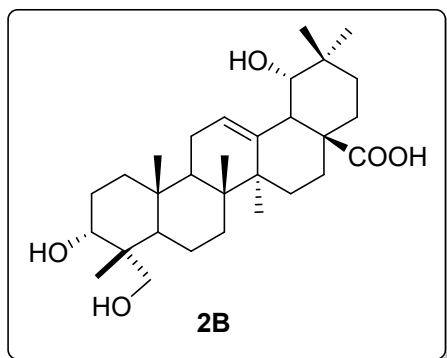
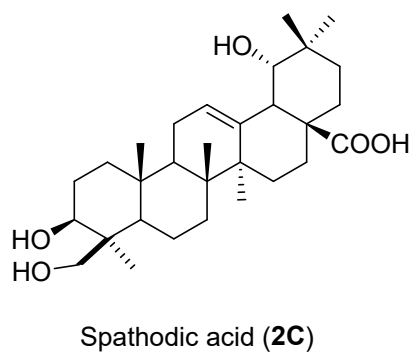
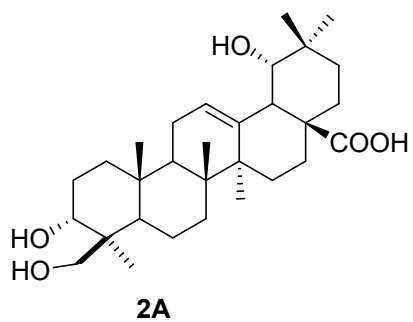


Figure S3. Four possible stereoisomers of **2**

Spectrum from TRI_MR36_(-)ESI.wiff2 (sample 1) - TRI_MR36_(-)ESI, -TOF MS (70 - 1500) from 0.190 min, noise filtered (noise multiplier = 1.5), Gaussian smoothed (0.5 points)

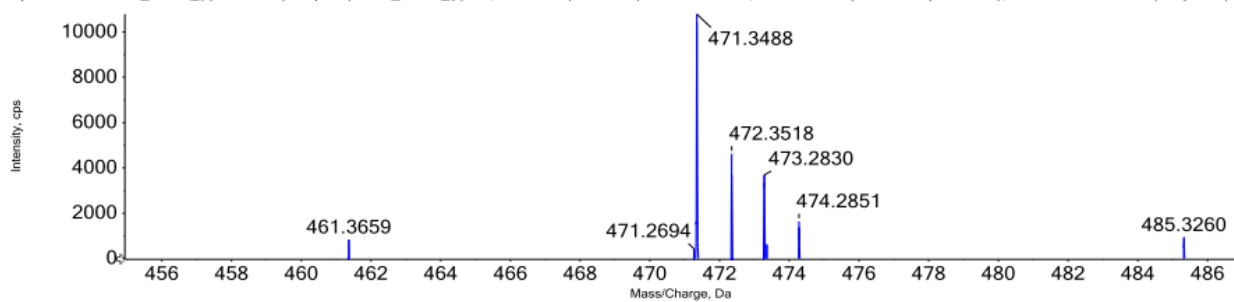


Figure S4. HRESIMS spectrum of **1**.

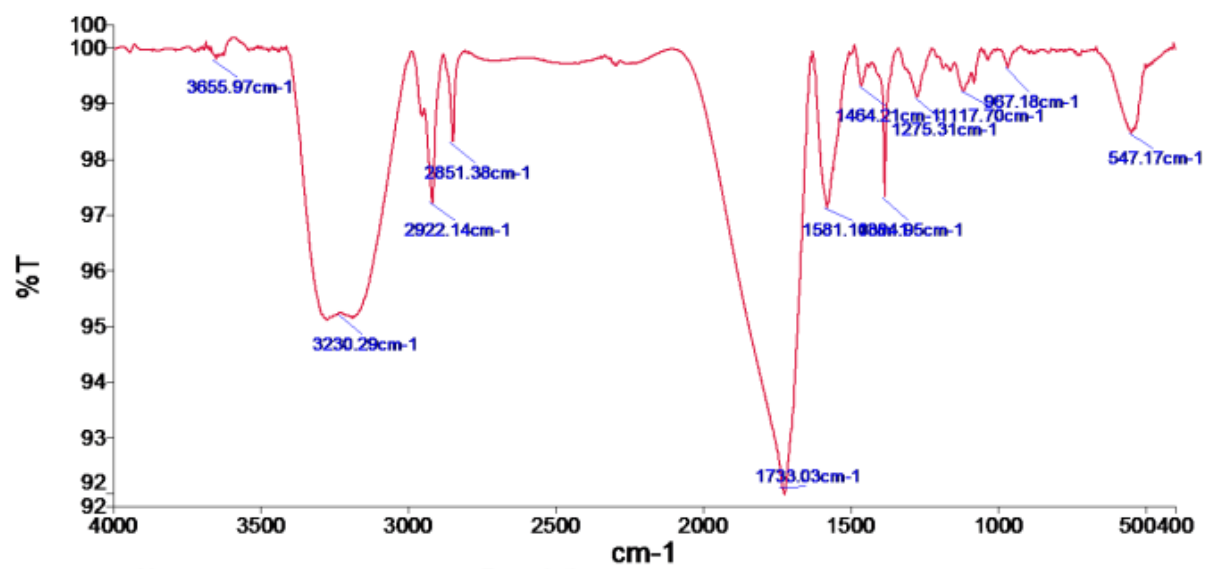


Figure S5. IR spectrum of 1.

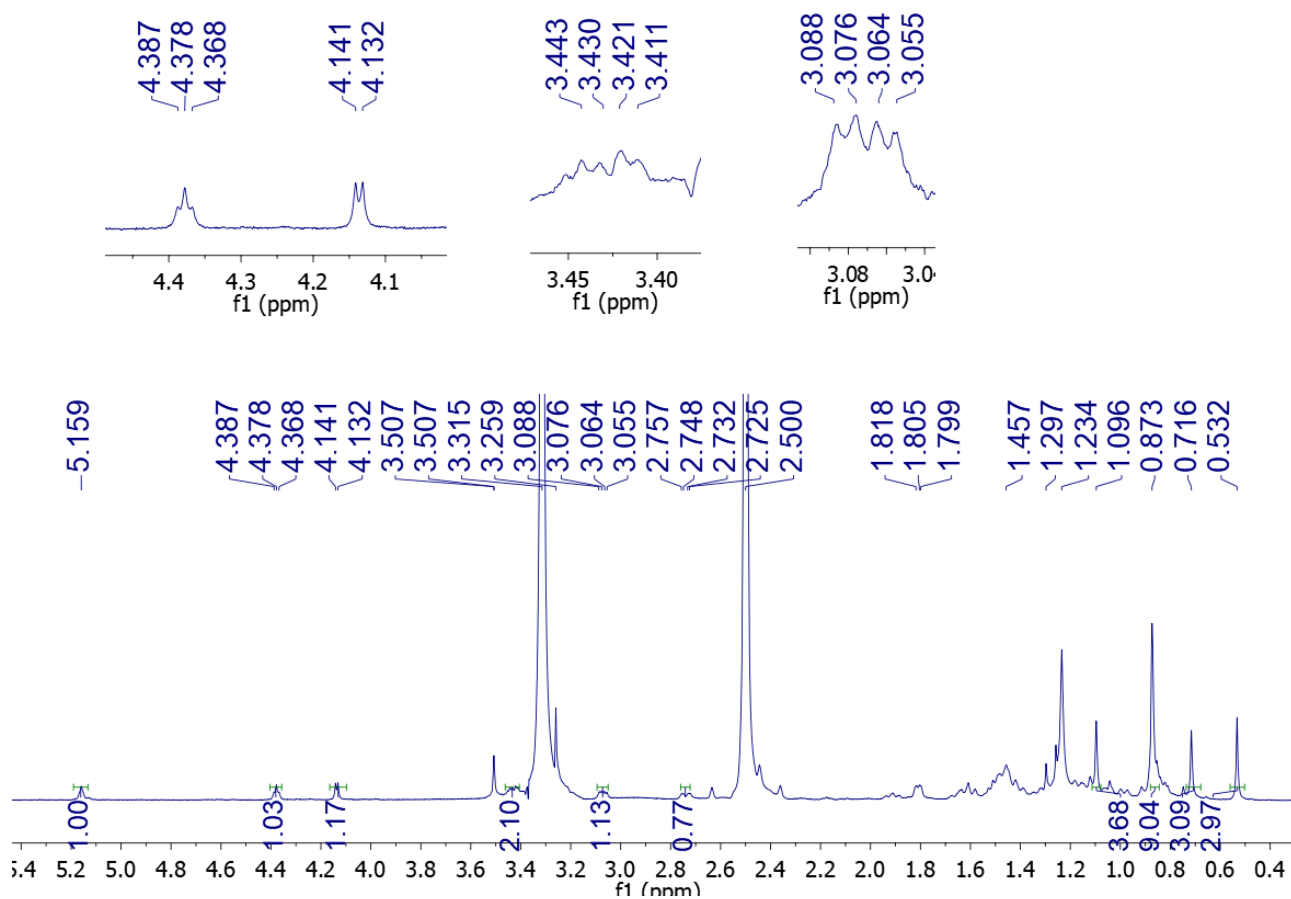


Figure S6. The $^1\text{H-NMR}$ ($\text{DMSO-}d_6$, 500 MHz) spectrum of **1**.

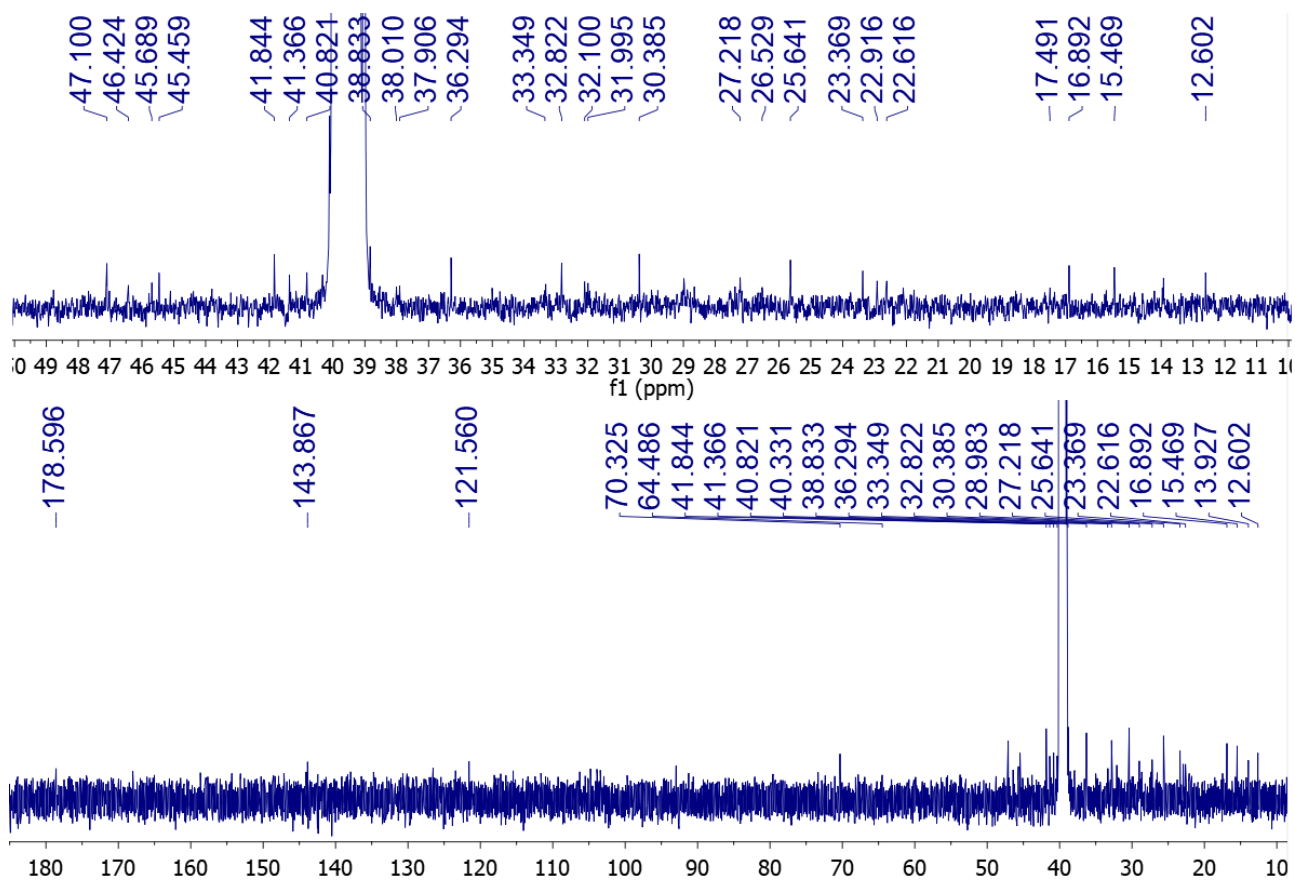


Figure S7. The ^{13}C -NMR ($\text{DMSO-}d_6$, 500 MHz) spectrum of **1**.

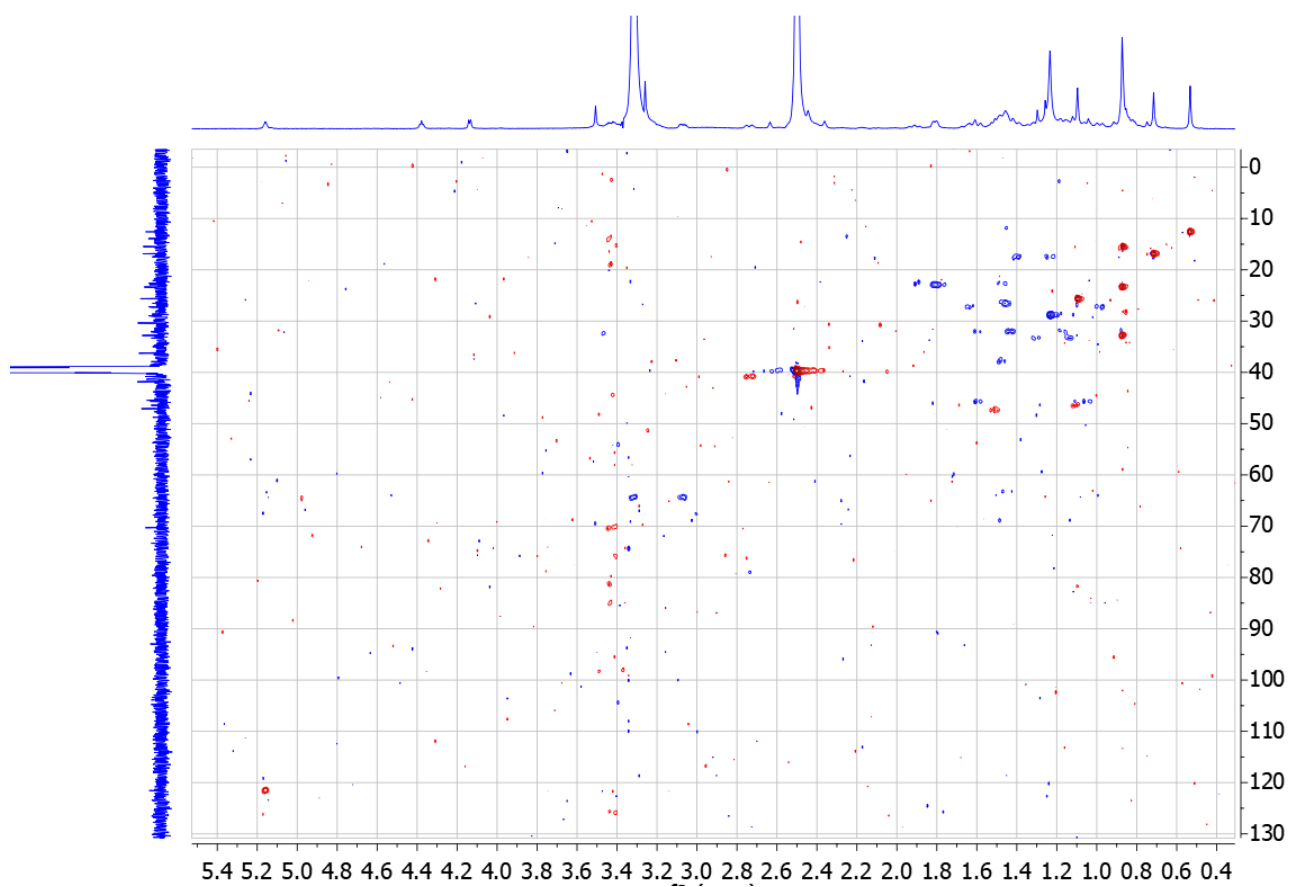


Figure S8. The HSQC (DMSO-*d*₆) spectrum of **1**

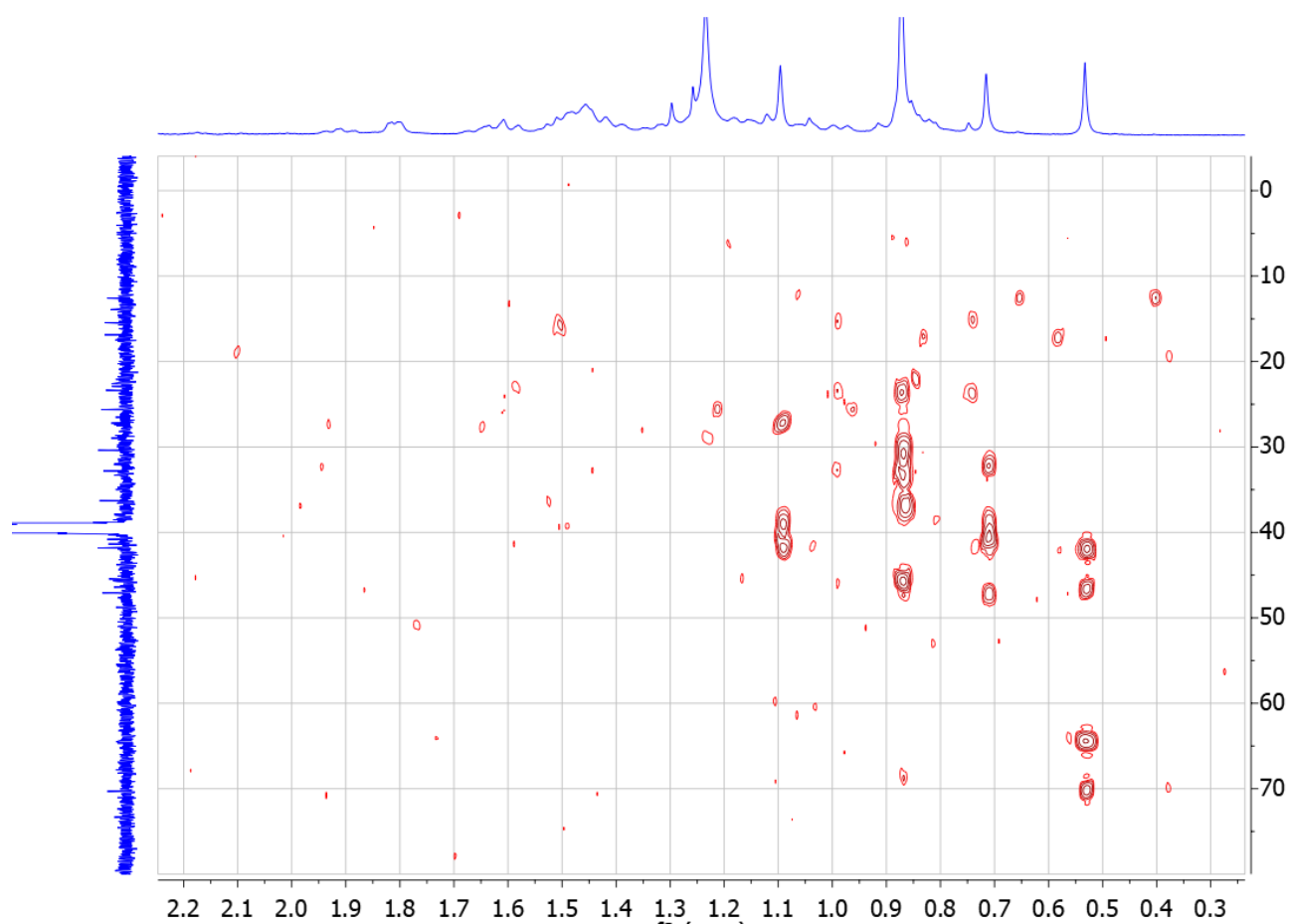
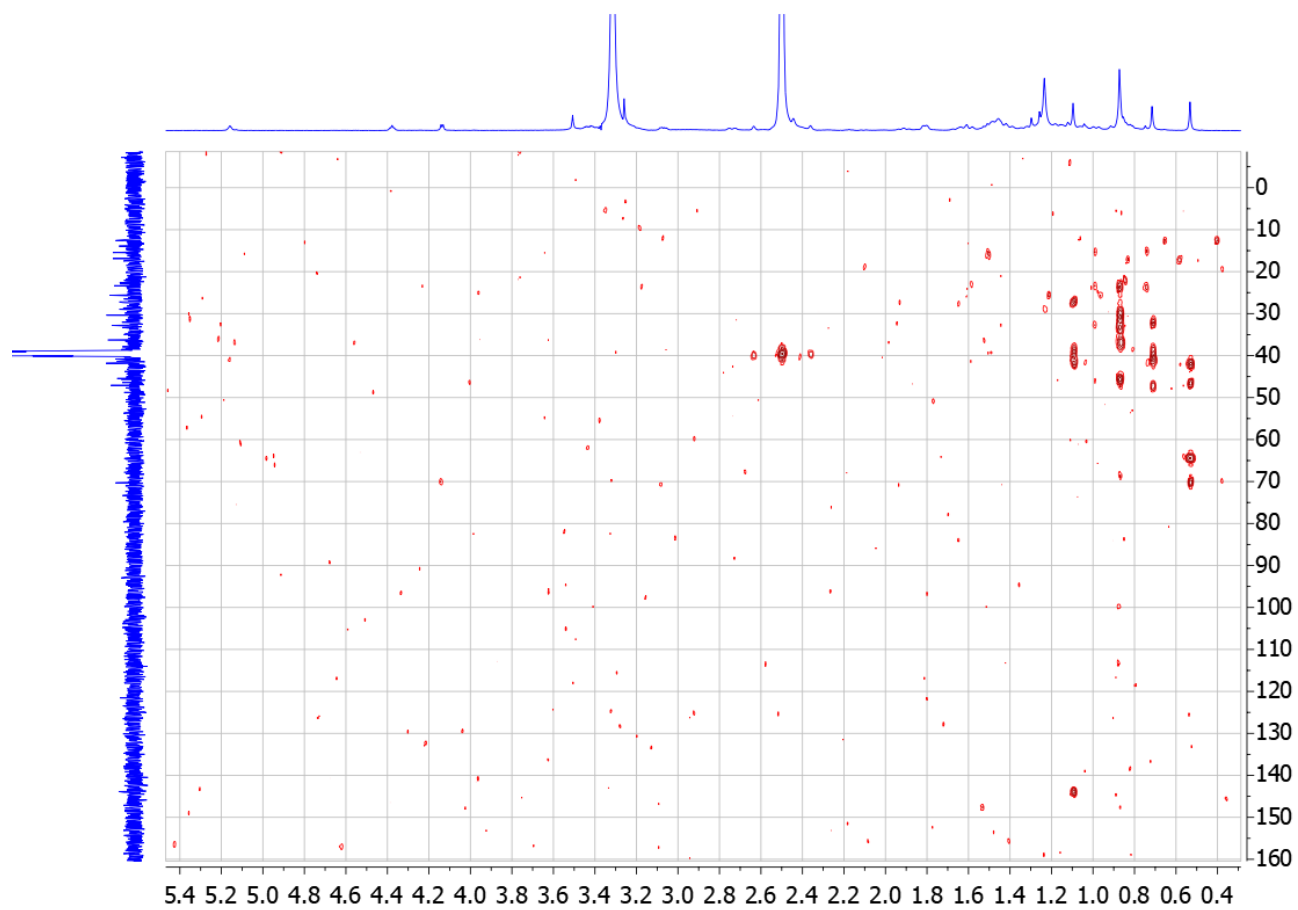


Figure S9. The HMBC ($\text{DMSO-}d_6$) spectrum of **1**.

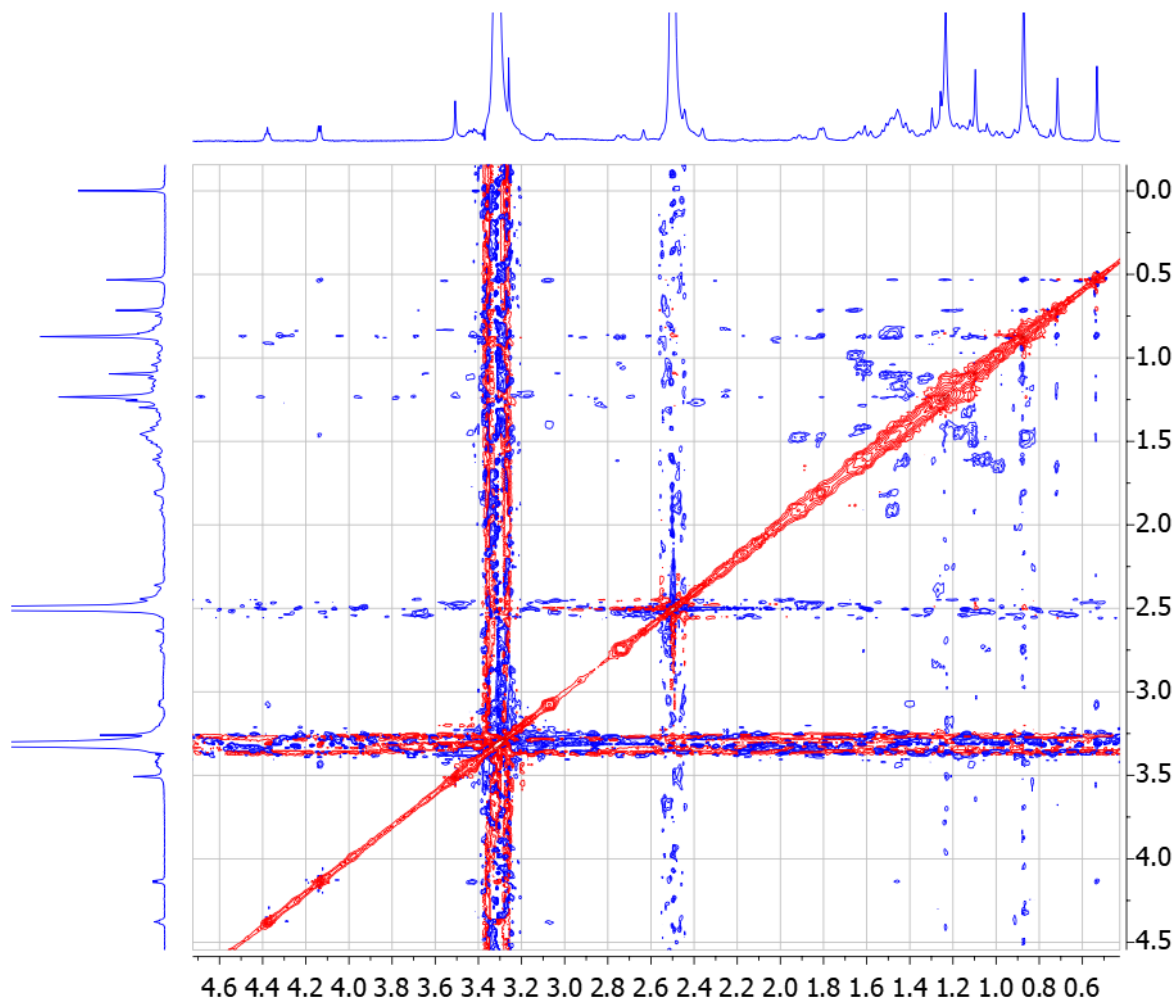


Figure S10. The NOESY (DMSO-*d*₆) spectrum of **1**.

Expanded spectrum

Spectrum from TRI_MR25_(-)ESI.wiff2 (sample 1) - TRI_MR25_(-)ESI, -TOF MS (70 - 1500) from 0.176 min, noise filtered (noise multiplier = 1.5), Gaussian smoothed (0.5 points)

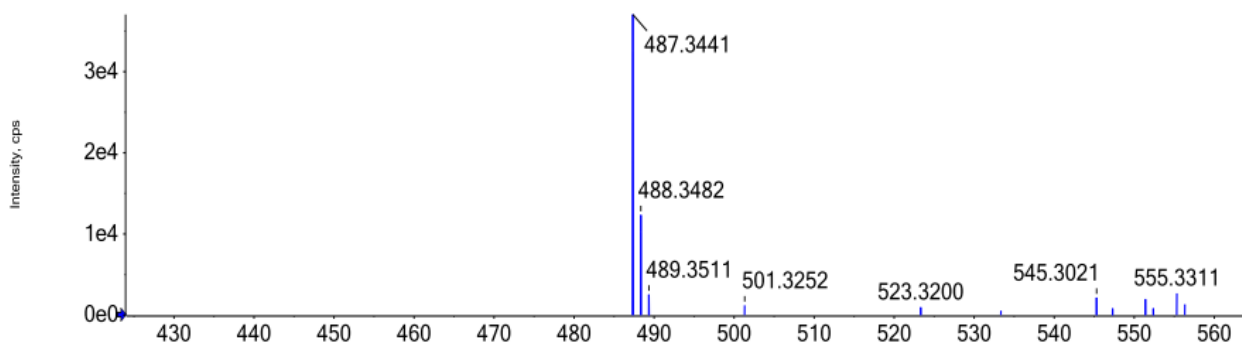


Figure S11. HRESIMS spectrum of **2**.

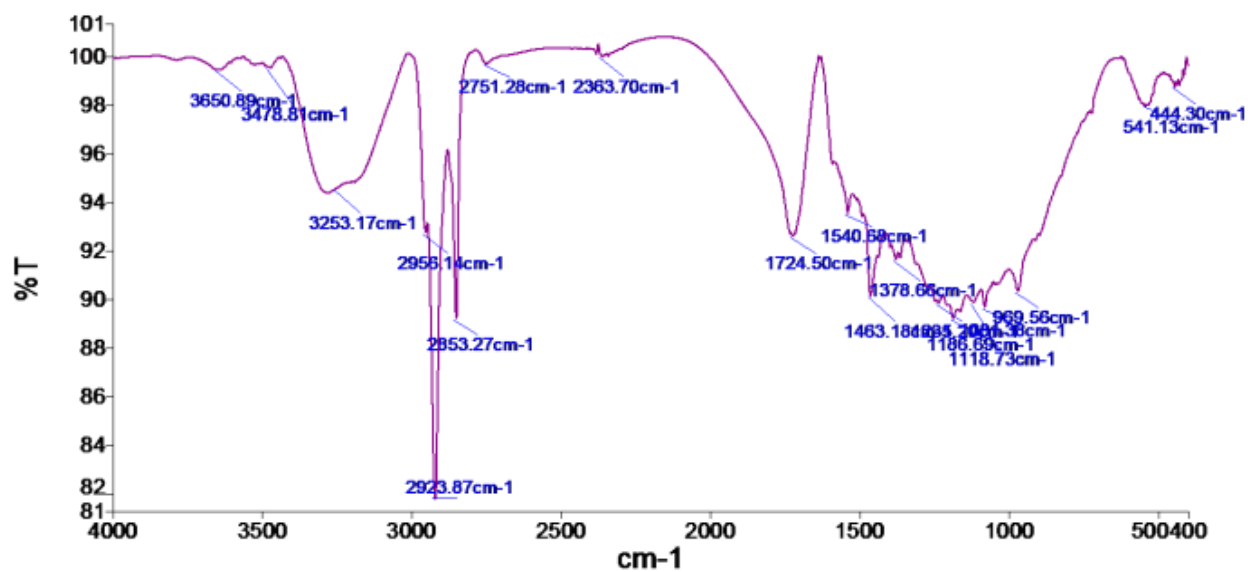


Figure S12. IR spectrum of 2.

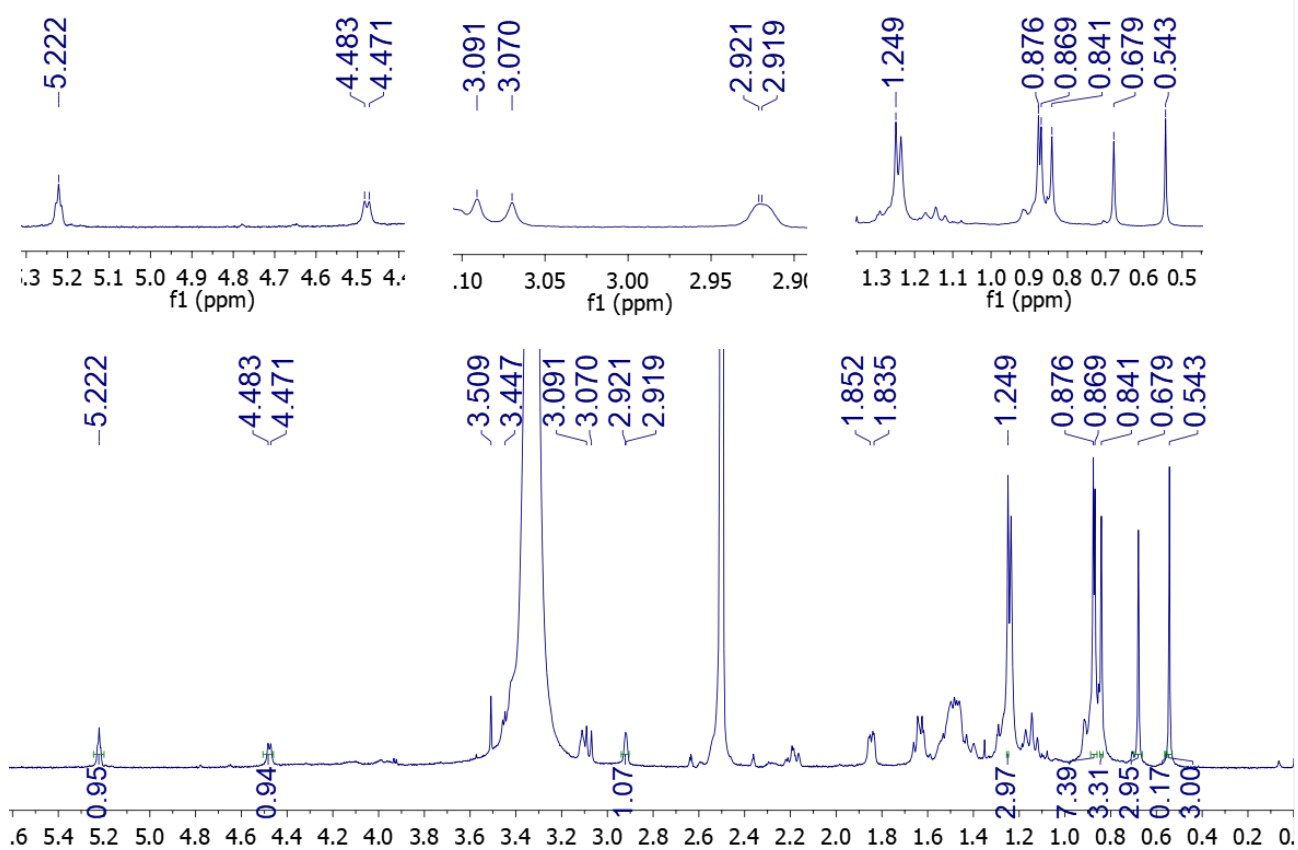


Figure S13. The $^1\text{H-NMR}$ ($\text{DMSO-}d_6$, 500 MHz) spectrum of **2**.

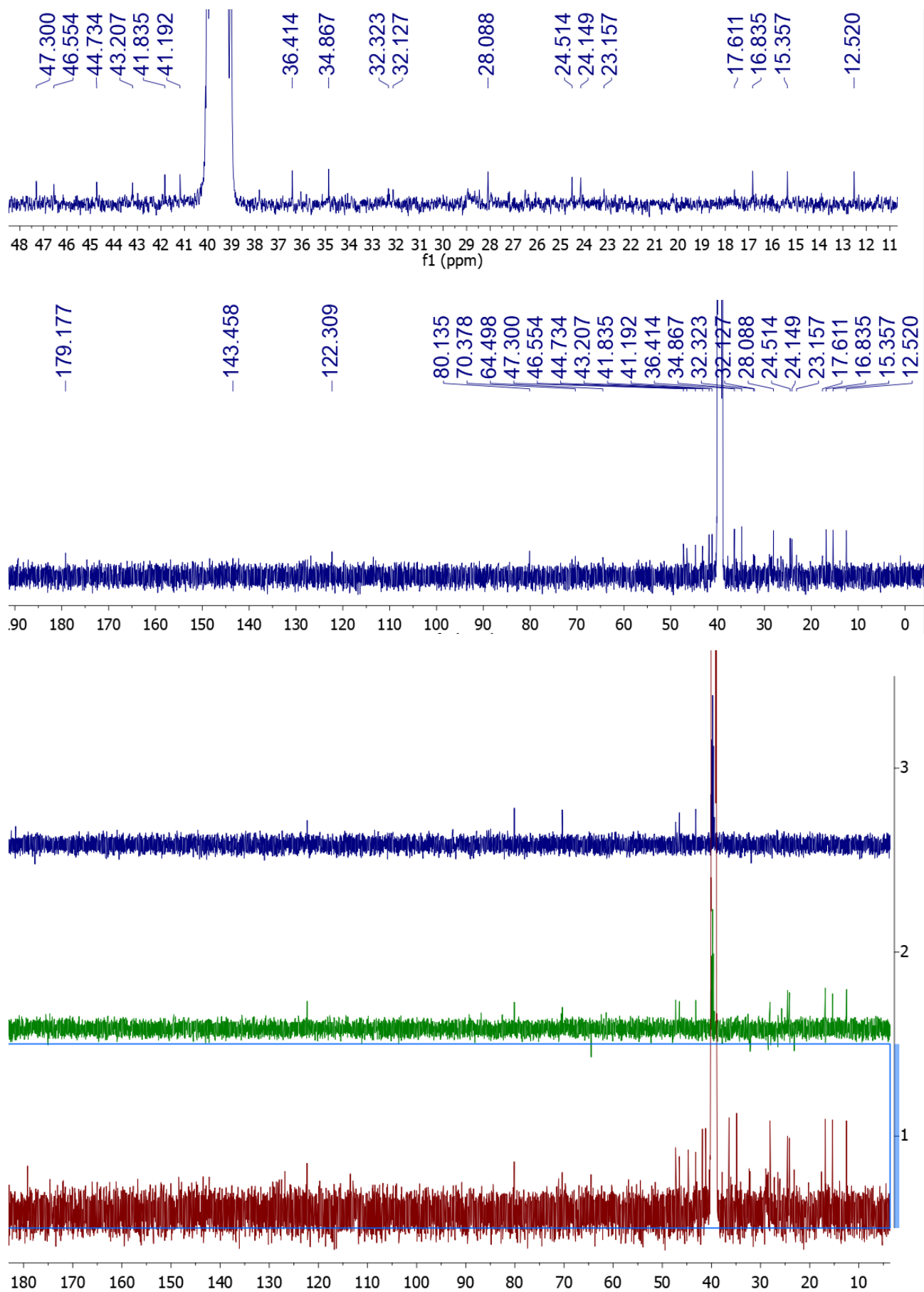


Figure S14. The ¹³C-NMR (DMSO-*d*₆, 500 MHz) spectrum of **2**.

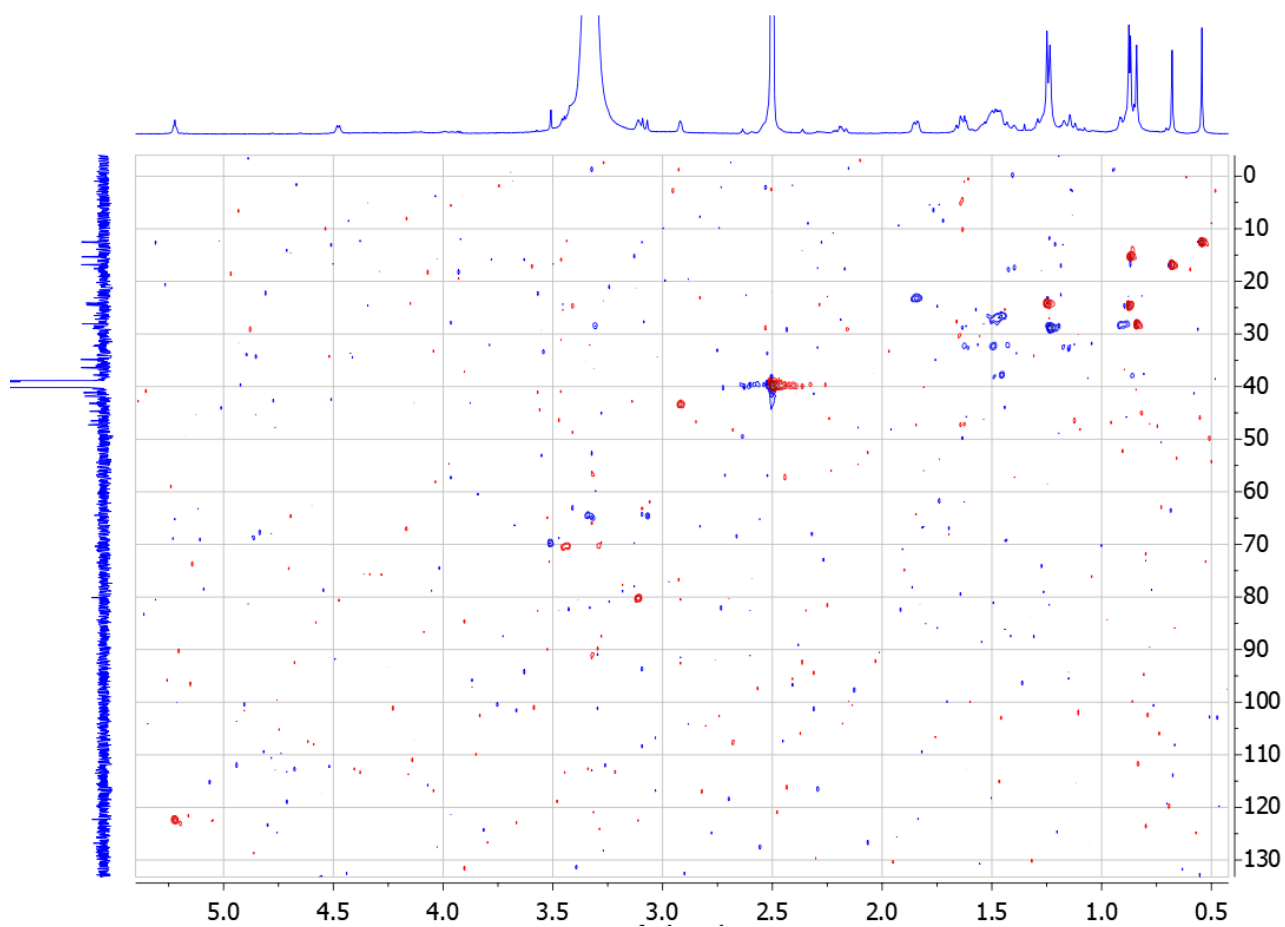


Figure S15. The HSQC (DMSO- d_6) spectrum of **2**

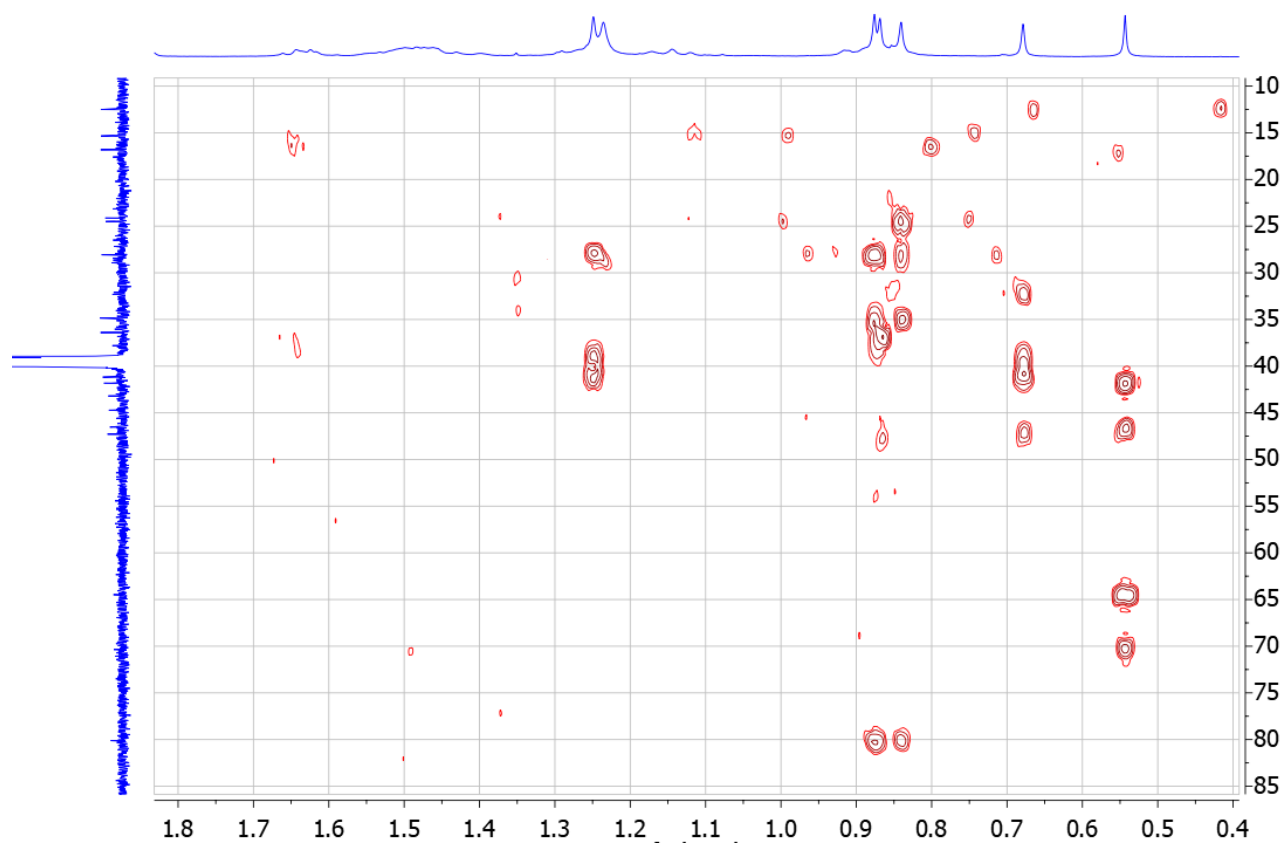
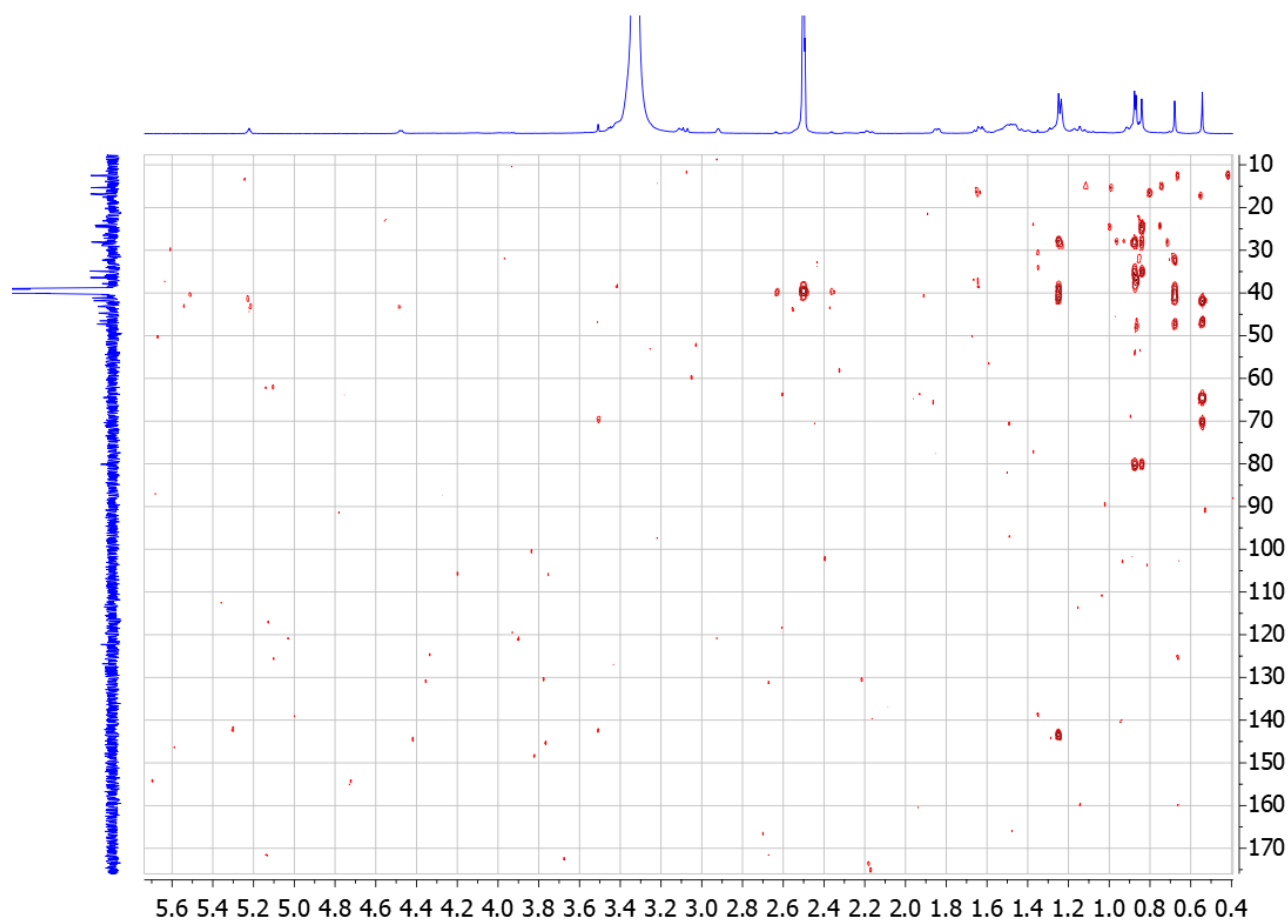


Figure S16. The HMBC (DMSO- d_6) spectrum of **2**.

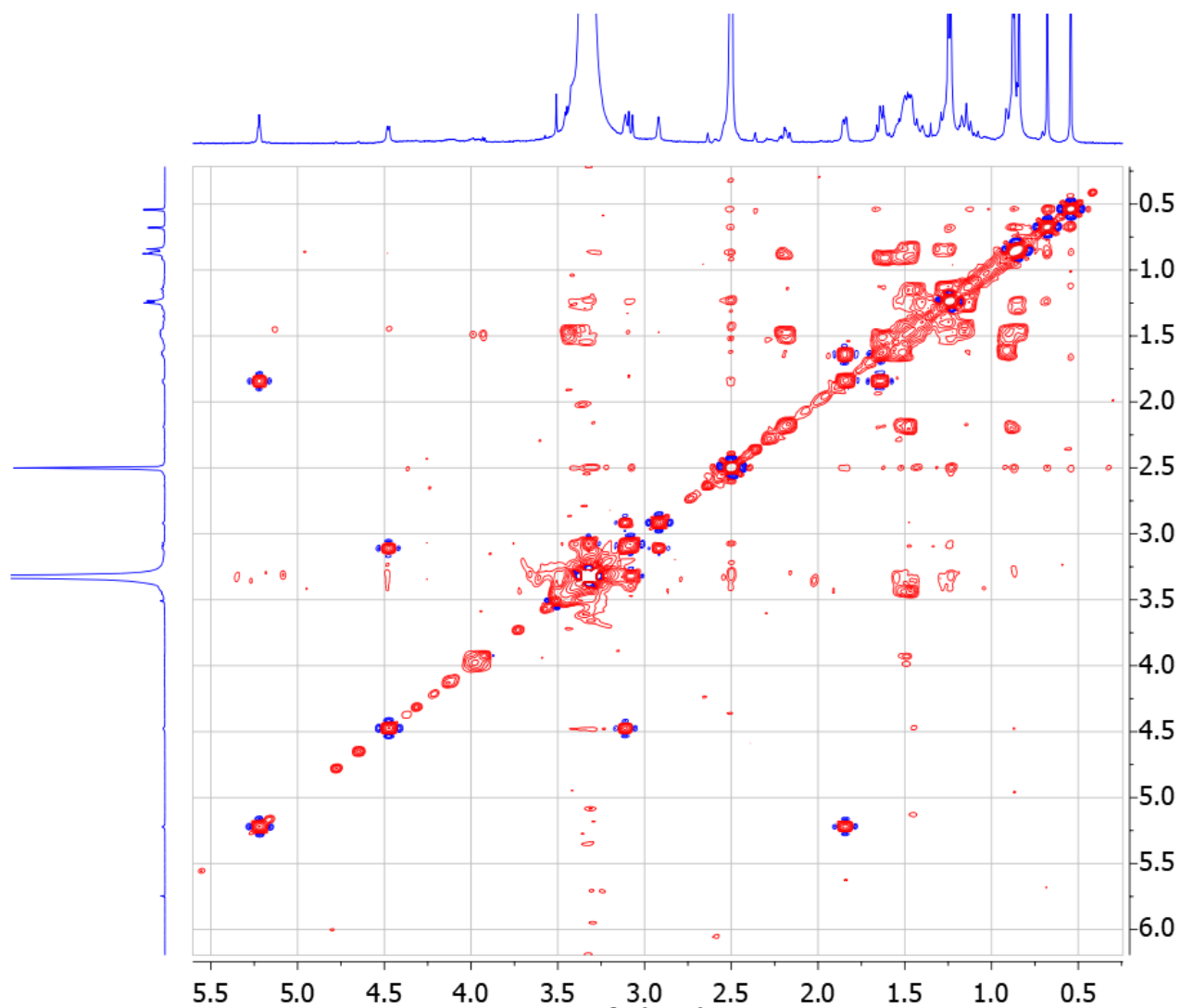


Figure S17. The COSY (DMSO- d_6) spectrum of **2**.

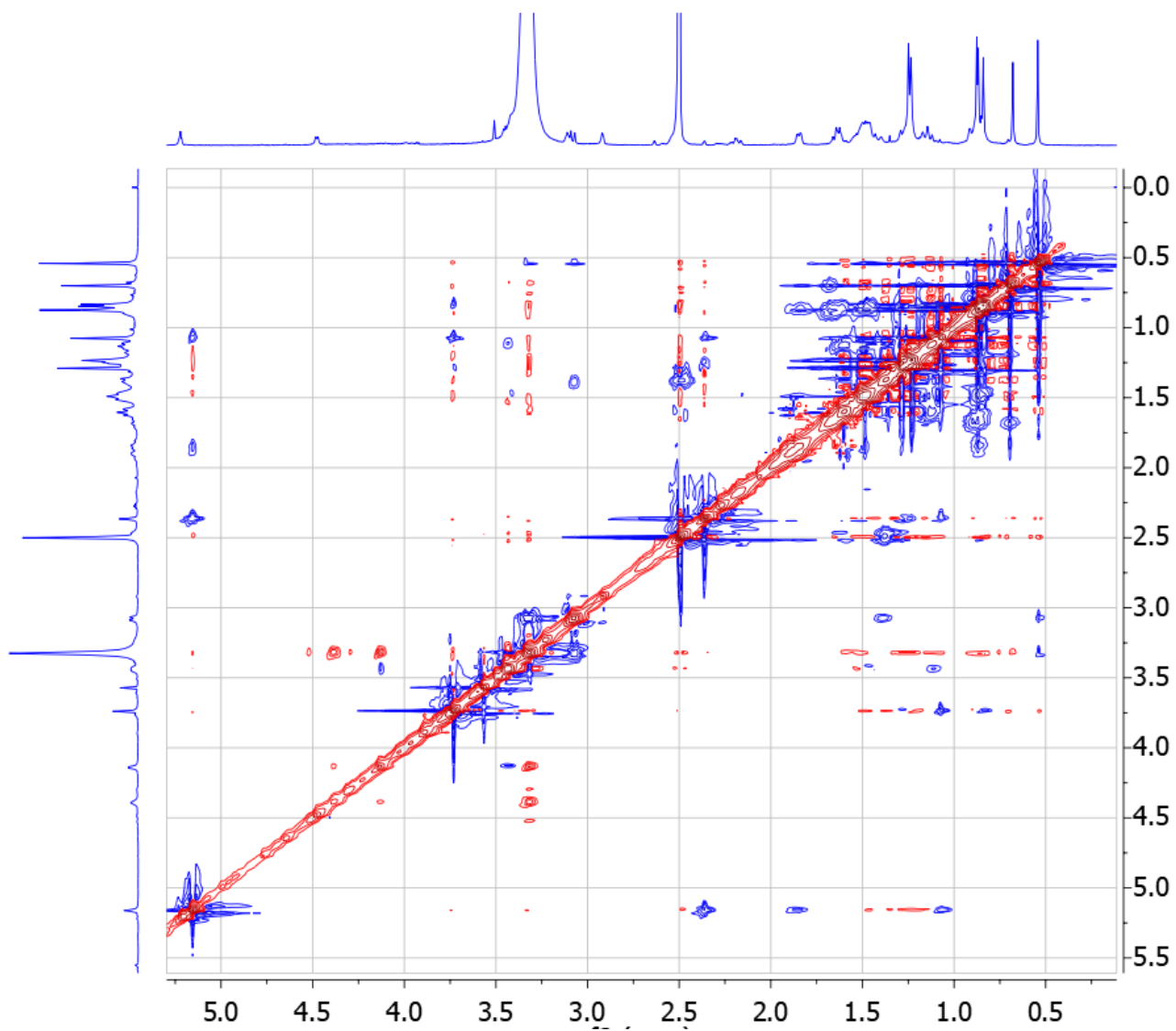


Figure S18. The NOESY (DMSO- d_6) spectrum of **2**.

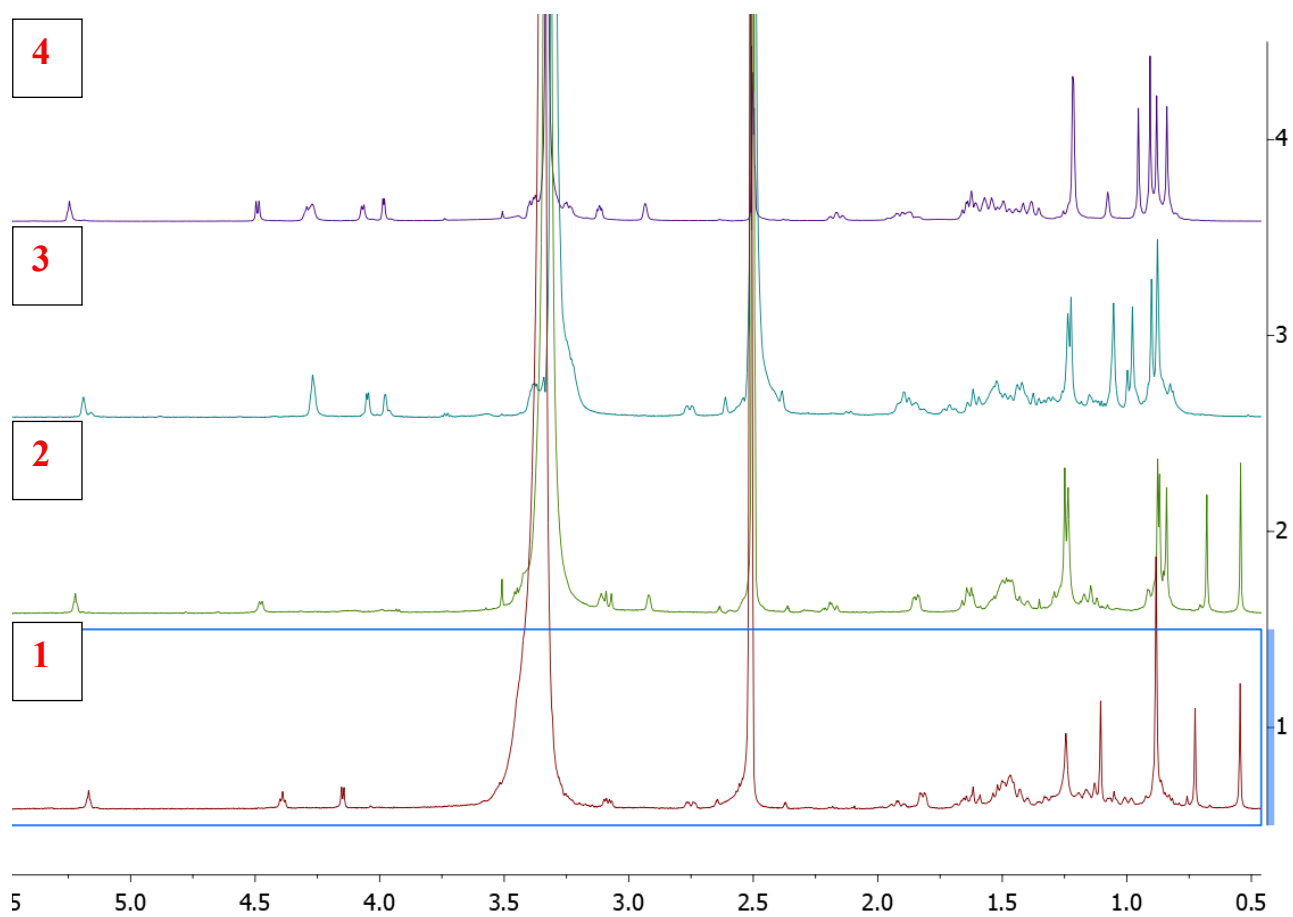


Figure S19. Comparison of $^1\text{H-NMR}$ spectra of **1-4**.

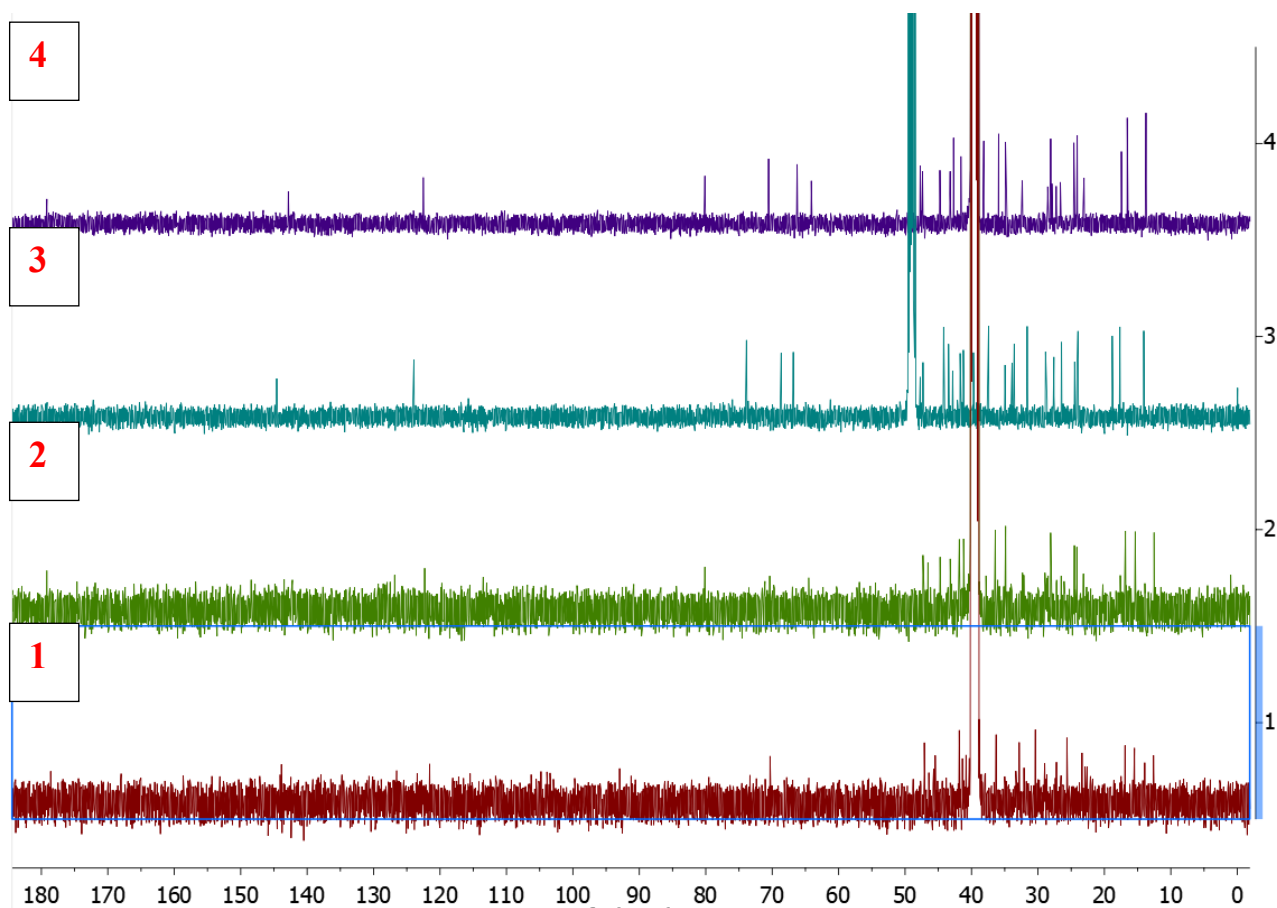


Figure S20. Comparison of ^{13}C -NMR spectra of 1-4.

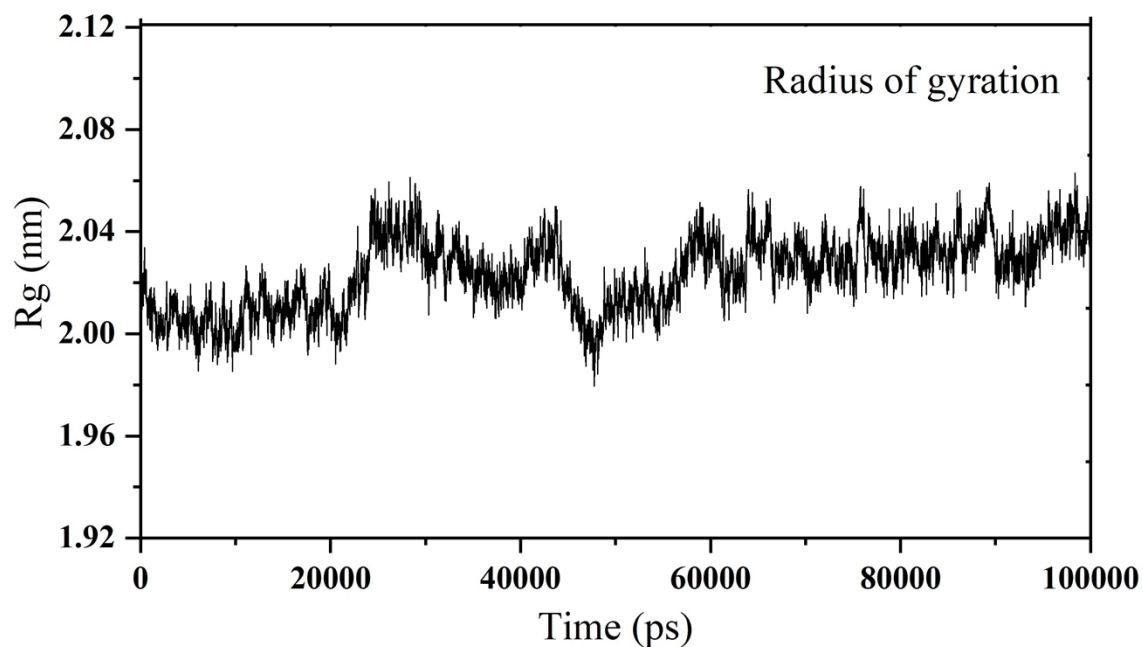
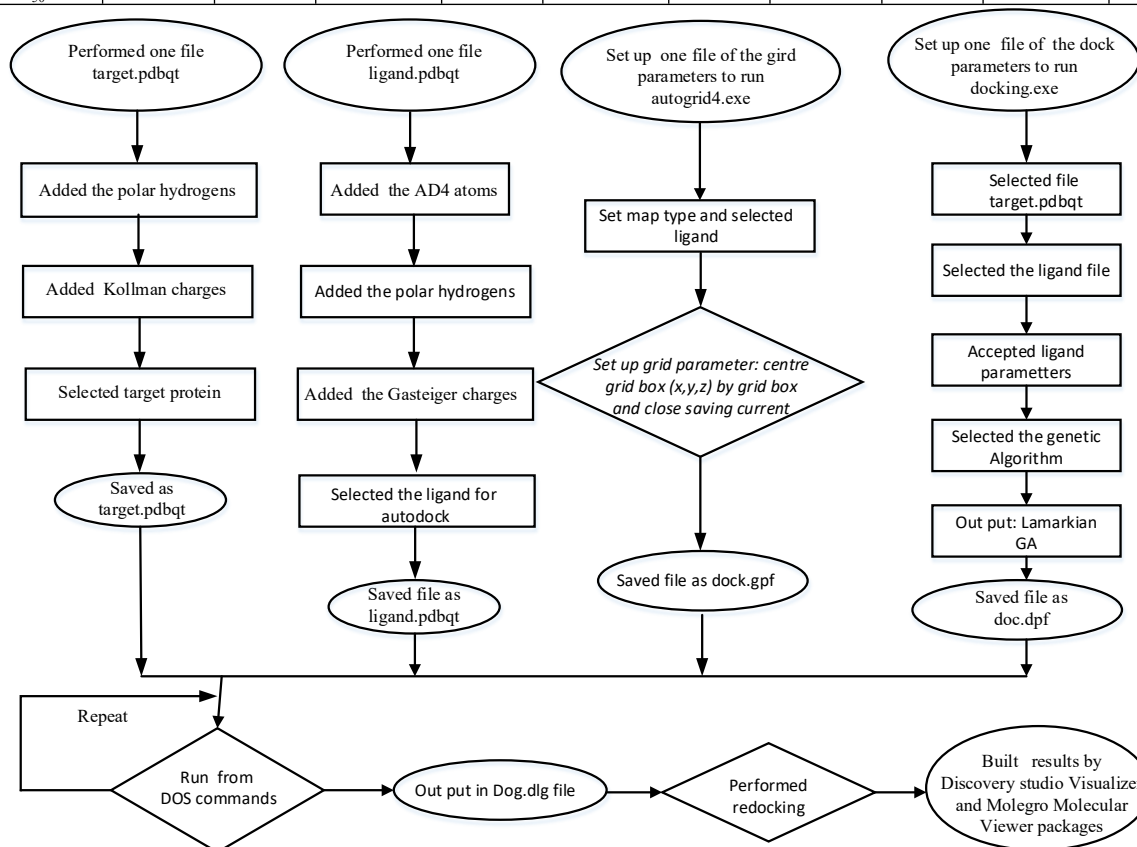


Figure S21. The radius of gyration of the protein-best pose was calculated using 100-ns MD simulations.

Table S3. Biological activity of tested compounds on the NO inhibitory activity

Concentration (µg/mL)	1		2		3		4		L-NMMA	
	% NO inhibition	% Survival	% NO inhibition	% Survival	% NO inhibition	% Survival	% NO inhibition	% Survival	% NO inhibition	% Survival
100	46.81	97.86	67.58	44.74	86.01	15.48	44.51	103.77	95.60	82.20
20	35.66	103.48	44.51	97.24	40.66	96.08	27.47	106.02	75.38	93.10
4	4.69		36.26		21.43		12.09		19.34	
0.8	-8.83		23.08		6.59		7.69		12.77	
IC ₅₀	>100	-	27.15±1.23	-	29.39±1.55	-	>100	-	10.24±1.65	-



Scheme S1. Procedure docking of ligand to receptor based on autodock