

Supplementary Materials: One-pot Bi-enzymatic Cascade Synthesis of Novel Ganoderma Triterpenoid Saponins

Te-Sheng Chang ^{1,+,*}, Chien-Min Chiang ^{2,+}, Tzi-Yuan Wang ³, Yu-Li Tsai ¹, Yu-Wei Wu ^{4,5}, Huei-Ju Ting ^{1,*}
and Jiumn-Yih Wu ^{6,*}

Table S1. NMR spectroscopic data for GAA-G2 (in pyridine-*d*₅; 700MHz).

Compound		δ_c ,	GAA-G2	HMBC
Position	type		δ_H (<i>J</i> in Hz)	
GAA moiety				
1	CH2	36.2	3.12, m	H-2, H-19
			1.52, m	
2	CH2	34.7	2.56, m	H-1
			2.52, m	
3	C	215.9		H-1, H-2, H-28, H-29
4	C	46.8		H-5, H-6, H-28, H-29
5	CH	49.1	1.71, dd (13.4, 2.1)	H-1, H-6, H-19, H-28, H-29
6	CH2	30.6	2.12, m	H-5, H-7
			1.97, m	
7	CH	67.1	5.17 dd (9.8, 7.5)	H-5, H-6
8	C	161.9		H-6, H-7, H-15, H-30
9	C	140.7		H-1, H-5, H-7, H-12, H-19
10	C	38.1		H-1, H-2, H-5, H-6, H-19
11	C	199.3		H-12
12	CH2	52.1	2.88, d (16.4)	H-18
			2.64, d (16.3)	
13	C	46.3		H-12, H-17, H-18, H-30
14	C	54.2		H-12, H-15, H-16, H-18, H-30
15	CH	82.7	5.34, dd (9.4, 6.3)	H-14, H-30, Glc1-H-1'
16	CH2	35.7	2.50, m	H-17
			1.96, m	
17	CH	49.2?	1.81, q (10.2)	H-12, H-16, H-18, H-21, H-22
18	CH3	17.7	1.09, s	H-12, H-17
19	CH3	18.9	1.43, s	H-1, H-5
20	CH	32.9	2.16, m	H-16, H-17, H-21, H-22
21	CH3	19.6	0.91, d (5.8)	H-22
22	CH2	49.9	2.51, m	H-17, H-21
			2.18, m	
23	C	209.0		H-22, H-24, H-25
24	CH2	47.0	3.03, dd (17.6, 8.4)	H-25, H-27
			2.54, m	
25	CH	35.6	3.26, m	H-24, H-27
26	C	178.2		H-24, H-25, H-27
27	CH3	17.6	1.33, d (7.2)	H-24, H-25
28	CH3	26.8	1.07, s	H-5, H-29
29	CH3	20.9	1.07, s	H-5, H-28
30	CH3	21.9	1.46, s	H-15
Glucose 1 moiety				
1'	CH	105.0	4.86, d (7.9)	H-15 , H-2', H-3', H-5'
2'	CH	75.2	4.01, t (8.2)	H-3' , H-5'
3'	CH	78.1	4.29, m	H-1', H-2' , H-4', H-5'
4'	CH	81.2	4.30, m	H2', H-3' , H-5', H-6', Glc2-H-1''
5'	CH	76.7	3.77, m	H-1', H-2', H-4', H-6'
6'	CH2	61.9	4.42 d (3.2), 2H	H-4', H-5'
Glucose 2 moiety				
1''	CH	103.0	5.88, d(3.9)	Glc1-H-4' , H-3''
2''	CH	74.3	4.16, dd (9.6, 3.8)	H-3''
3''	CH	75.29	4.55, m	H-1'', H-2'', H-4''
4''	CH	71.8	4.13, t (9.4)	H-3'', H-6''
5''	CH	75.30	4.55, m	H-2'', H-4'', H-6''
6''	CH2	62.6	4.52, m	H-4'', H-5''
			4.31, m	

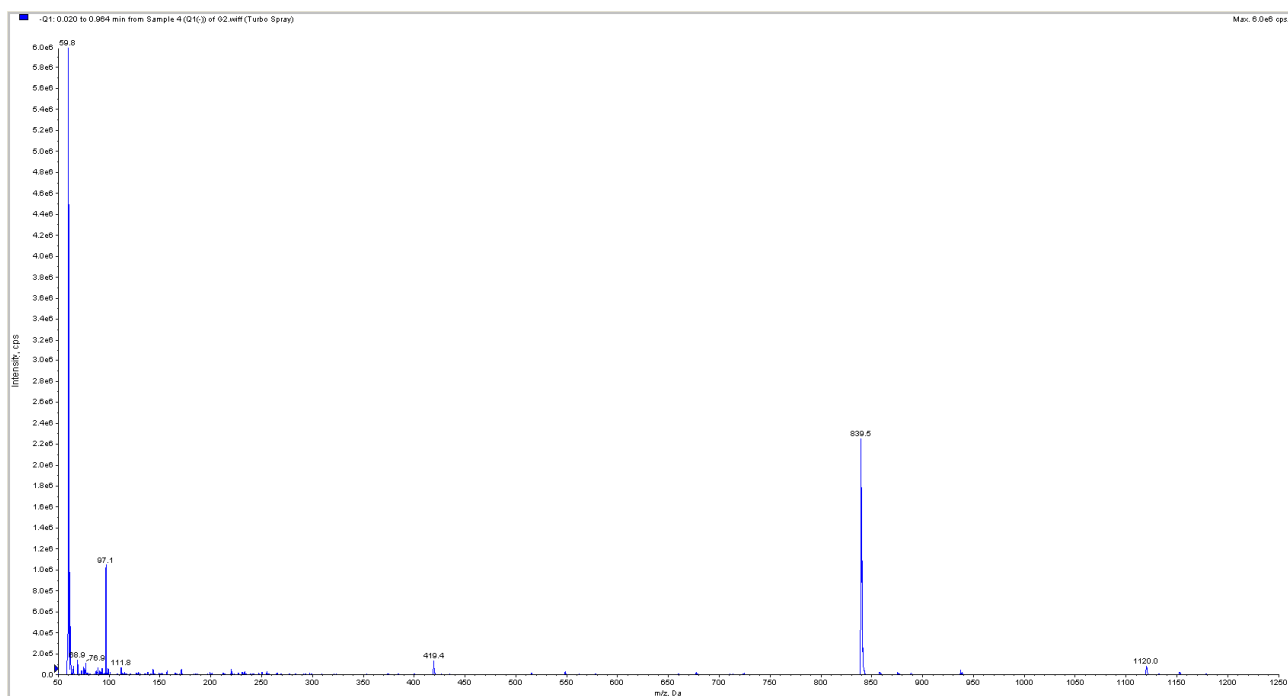


Figure S1. Mass analysis of GAA-G2 at the negative mode. A significant signal at m/z 839.5 showed the corresponding m/z signal of molecular weight 840 of GAA-diglucoside ($516 + 180 \times 2 - 18 \times 2$) at the negative mode $[GAA-G2 - H]^-$.

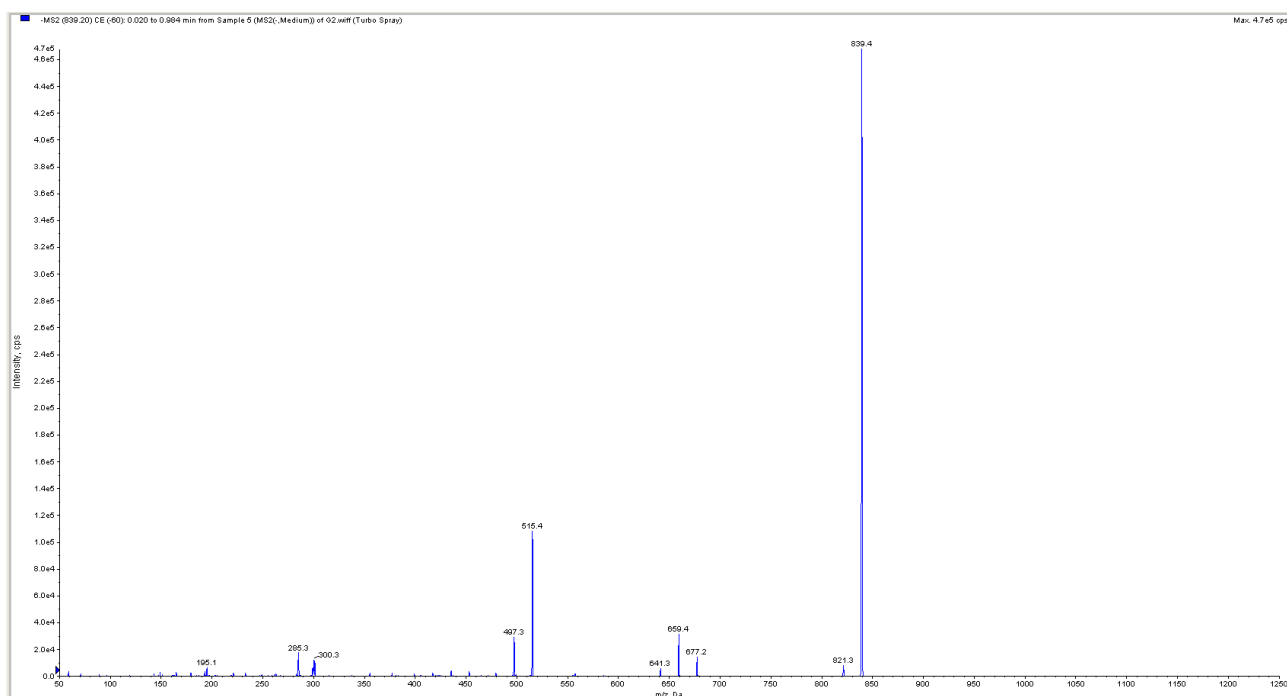


Figure S2. Mass-mass analysis of the m/z 839.5 peak of GAA-G2 at the negative mode. Two significant peaks at m/z 677.2 and 515.4 showed the corresponding m/z signals of molecular weights 678 and 516 of GAA-glucoside (516+180-18) and GAA (516) at the negative mode [GAA-G – H]⁻ and [GAA – H]⁻.

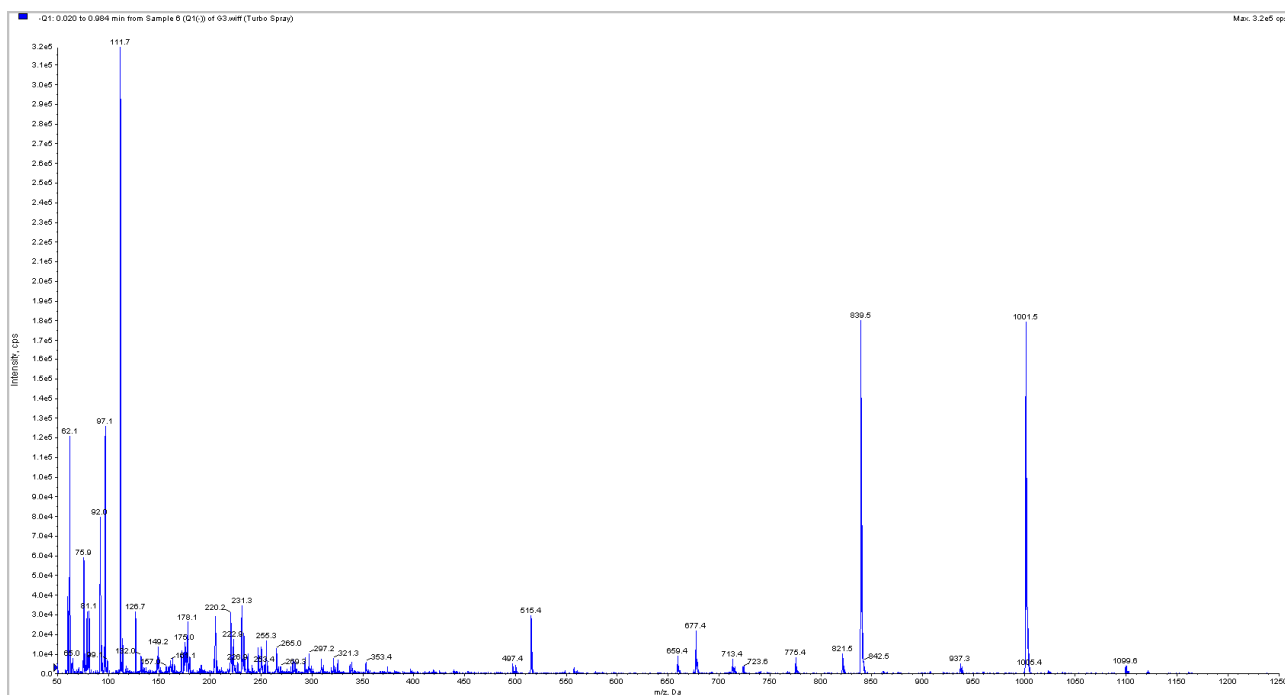


Figure S3. Mass analysis of GAA-G3 at the negative mode. A significant signal at m/z 1001.5 showed the corresponding m/z signal of molecular weight 1002 of GAA-triglucoside ($516+180 \times 3-18 \times 3$) at the negative mode $[\text{GAA-G3} - \text{H}]^-$.

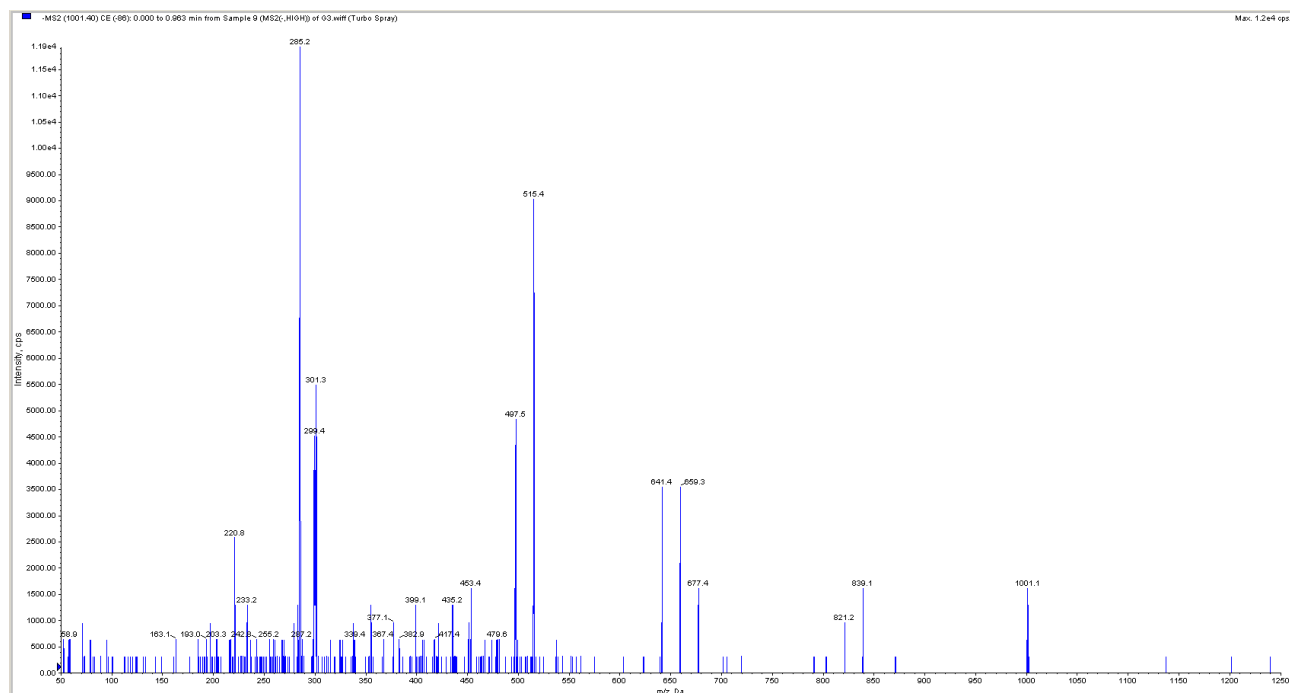


Figure S4. Mass-mass analysis of the m/z 1001.5 peak of GAA-G3 at the negative mode. Three significant peaks at m/z 839.1, 677.4 and 515.4 showed the corresponding m/z signals of molecular weights 840, 678 and 516 of GAA-diglucoside ($516+180 \times 2-18 \times 2$), GAA-glucoside ($516+180-18$) and GAA (516) at the negative mode $[GAA-2G-H]^-$, $[GAA-G-H]^-$ and $[GAA-H]^-$.

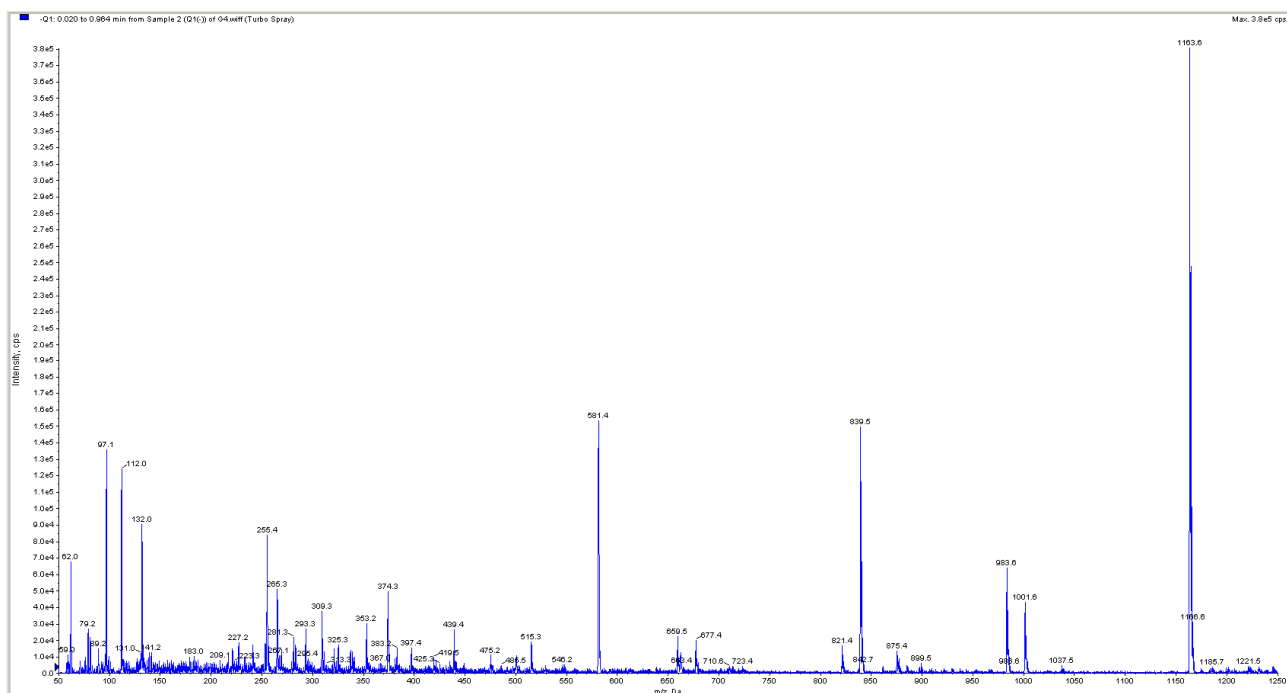


Figure S5. Mass analysis of GAA-G4 at the negative mode. A significant signal at m/z 1163.6 showed the corresponding m/z signal of molecular weight 1164 of GAA-tetraglucoside ($516+180 \times 4-18 \times 4$) at the negative mode [GAA-G4 - H].

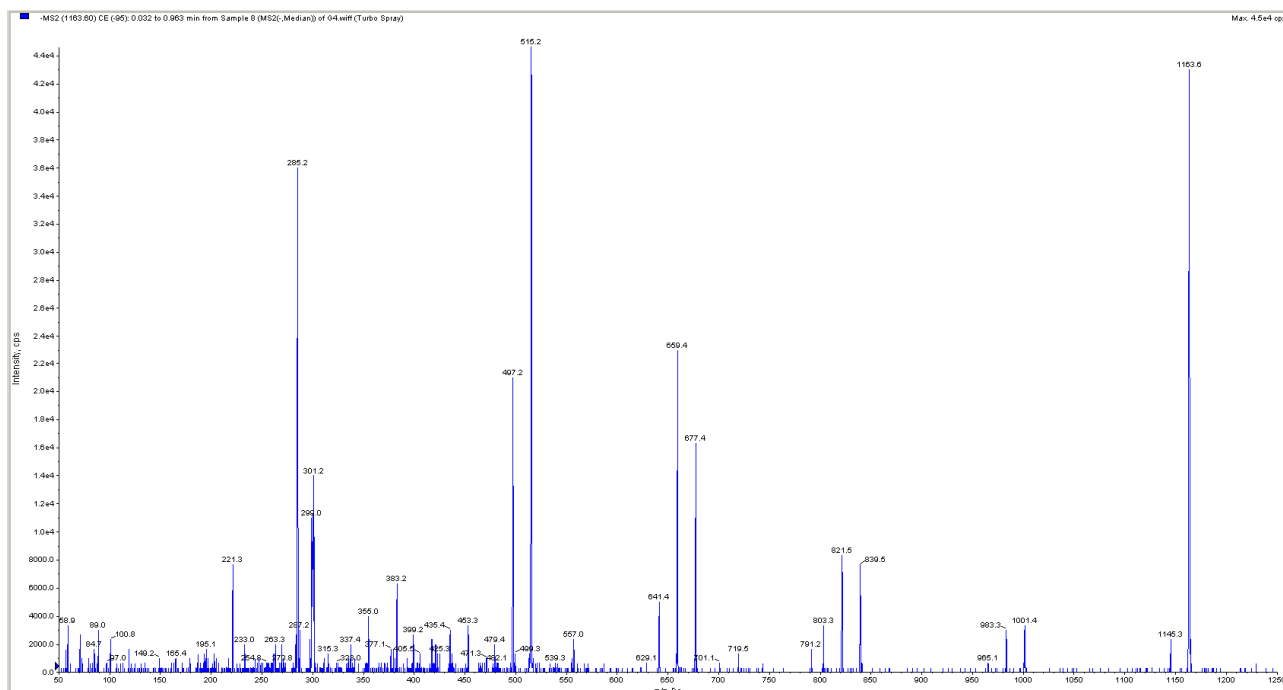


Figure S6. Mass-mass analysis of the m/z 1163.6 peak of GAA-G4 at the negative mode. Three significant peaks at m/z 1001.4, 839.5, 677.4 and 515.2 showed the corresponding m/z signals of molecular weights 1002, 840, 678 and 516 of GAA-triglucoside ($516+180 \times 3-18 \times 3$), GAA-diglucoside ($516+180 \times 2-18 \times 2$), GAA-glucoside ($516+180-18$) and GAA (516) at the negative mode $[GAA-3G-H]^-$, $[GAA-2G-H]^-$, $[GAA-G-H]^-$ and $[GAA-H]^-$.

GAA-G2, Pyr-d5, 1H

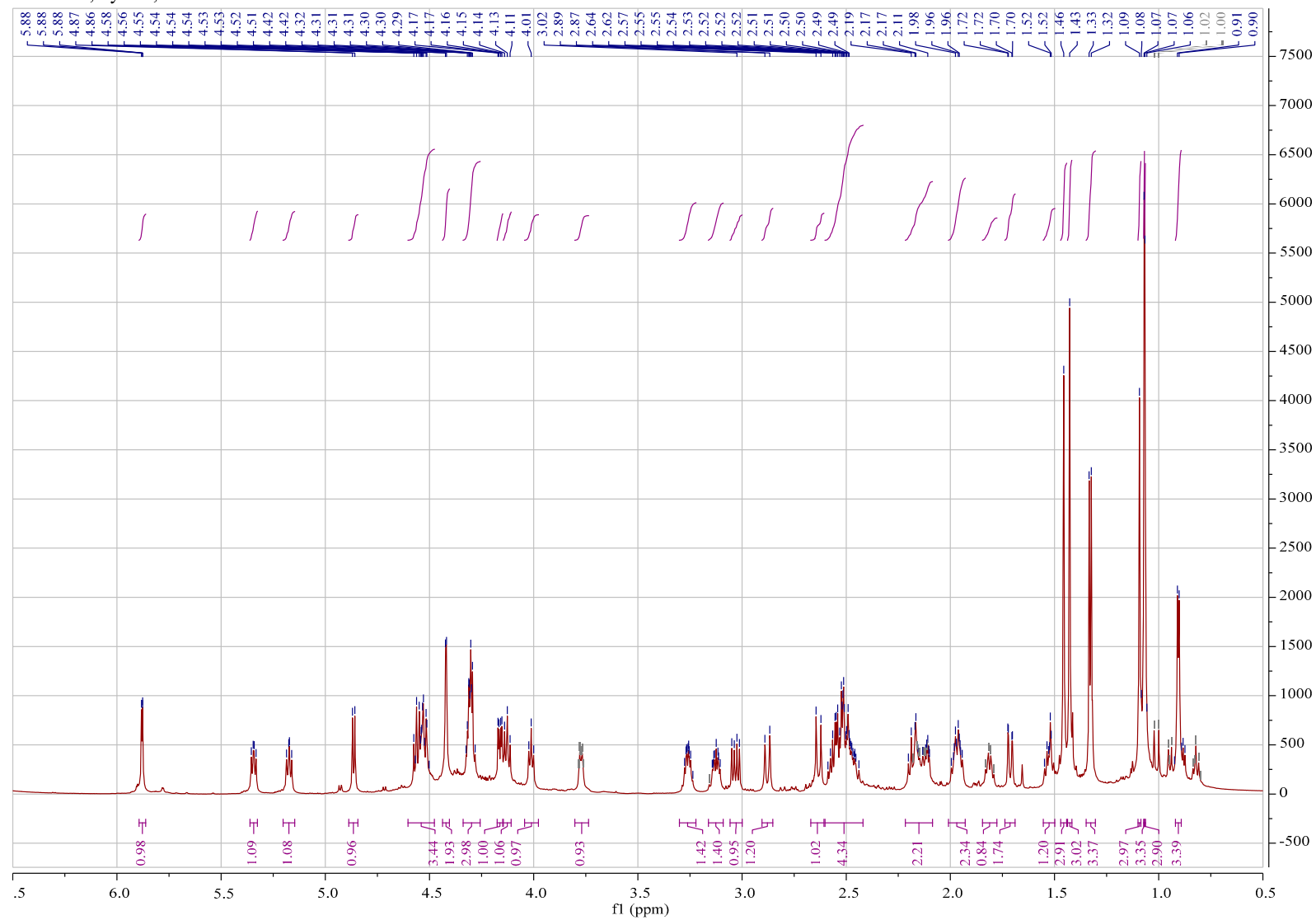


Figure S7. The ^1H -NMR (700 MHz) spectrum of GAA-G2 in pyridine- d_5 .

GAA-G2, pyridine-d₅, ¹³C

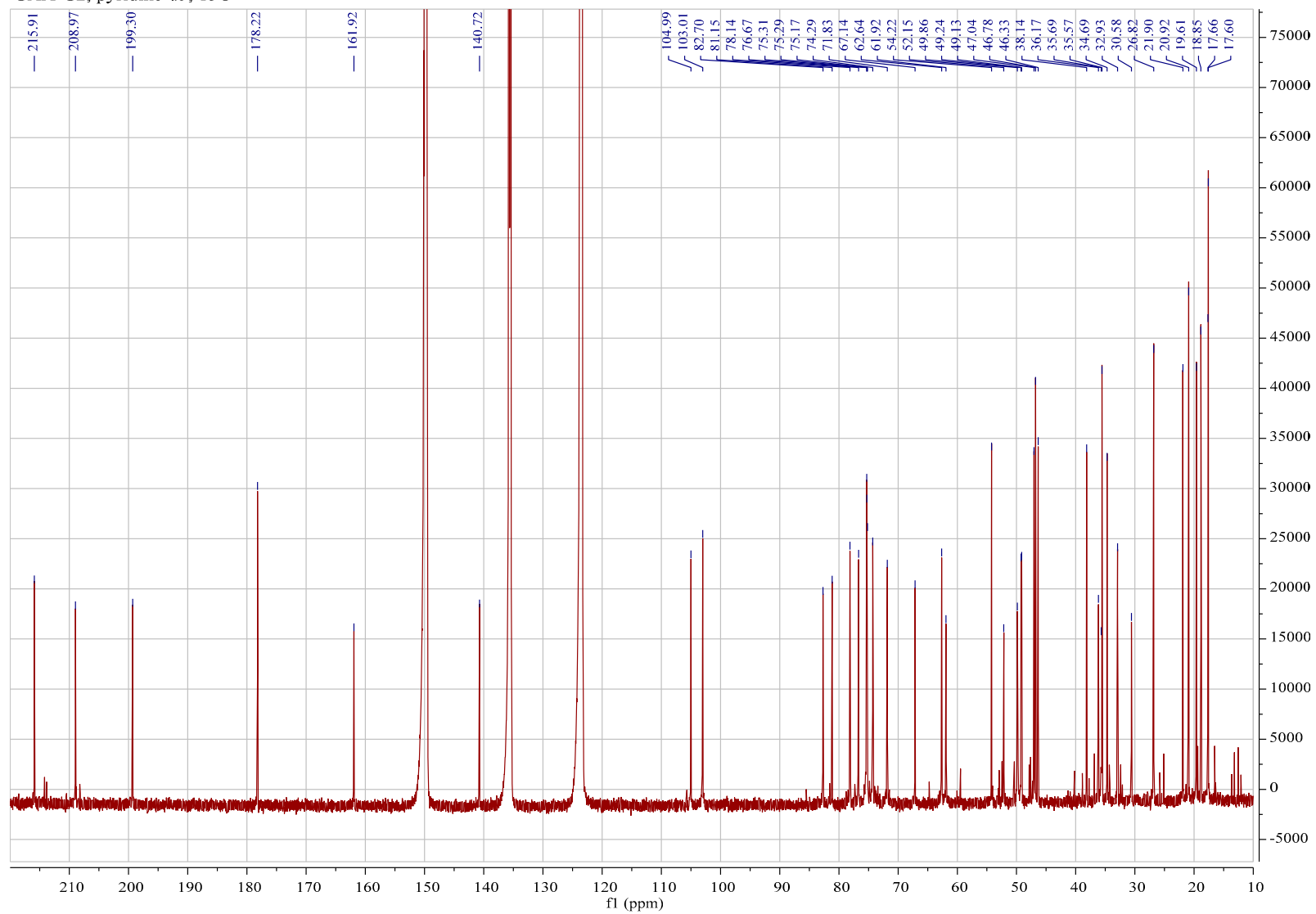


Figure S8. The ¹³C-NMR (176 MHz) spectrum of GAA-G2 in pyridine-d₅.

Dept-135, pyridine-d5



Figure S9. The DEPT-135 (176 MHz) spectrum of GAA-G2 in pyridined-5.

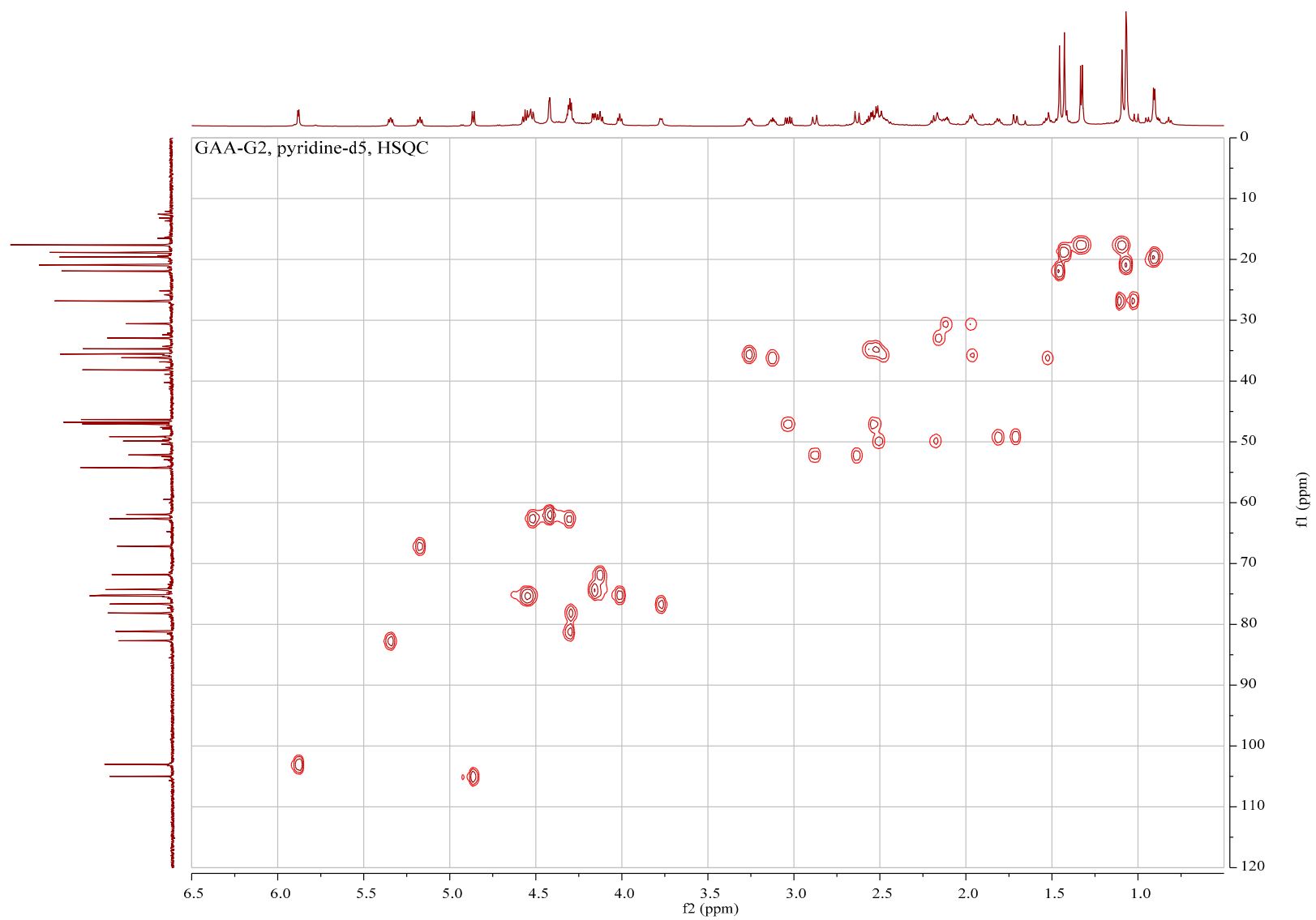


Figure S10. The HSQC (700 MHz) spectrum of GAA-G2 in pyridine-d₅.



Figure S11. The HMBC (700 MHz) spectrum of GAA-G2 in pyridine-d₅.

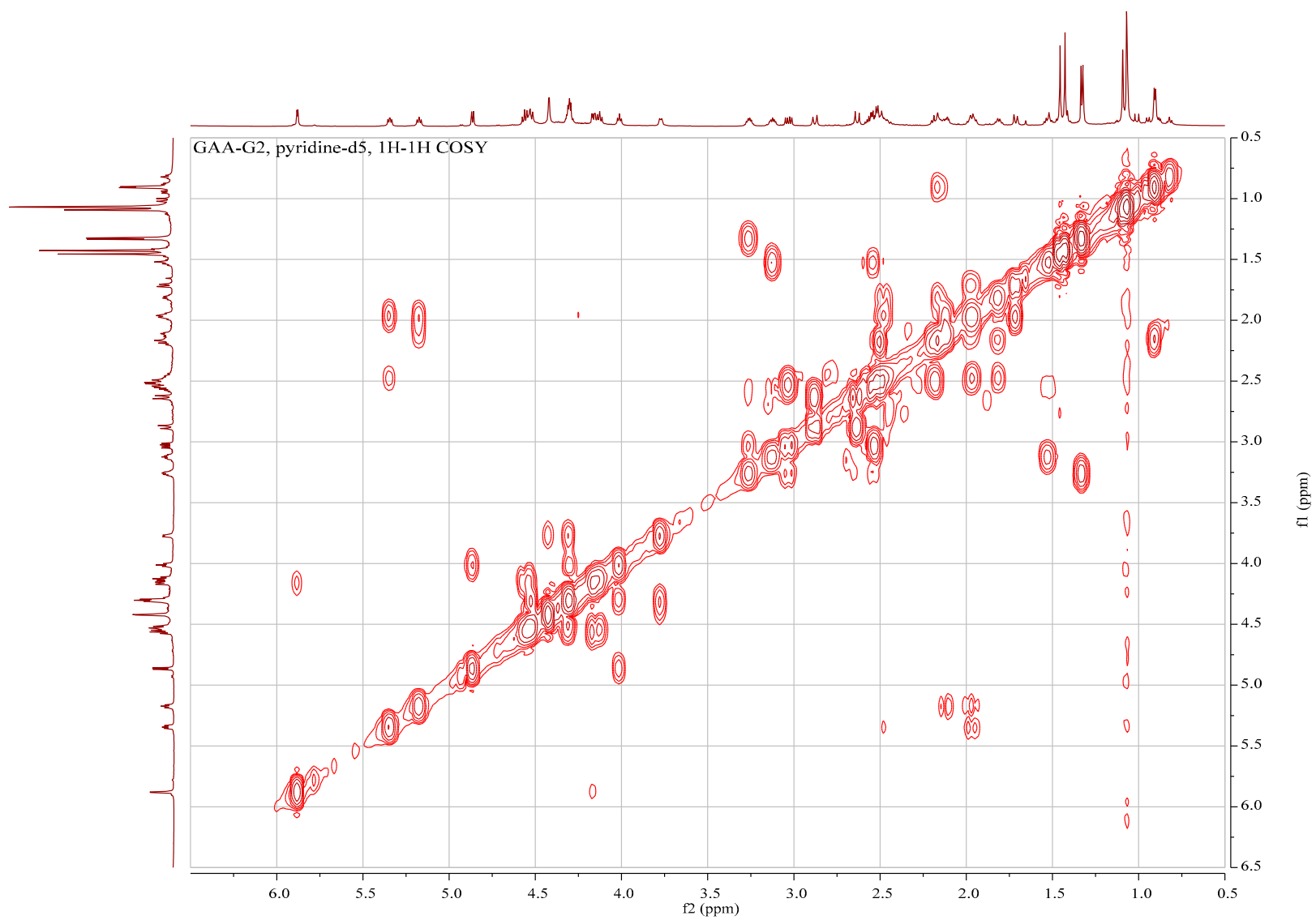


Figure S12. The ¹H-¹H COSY (700 MHz) spectrum of GAA-G2 in pyridine-d₅.

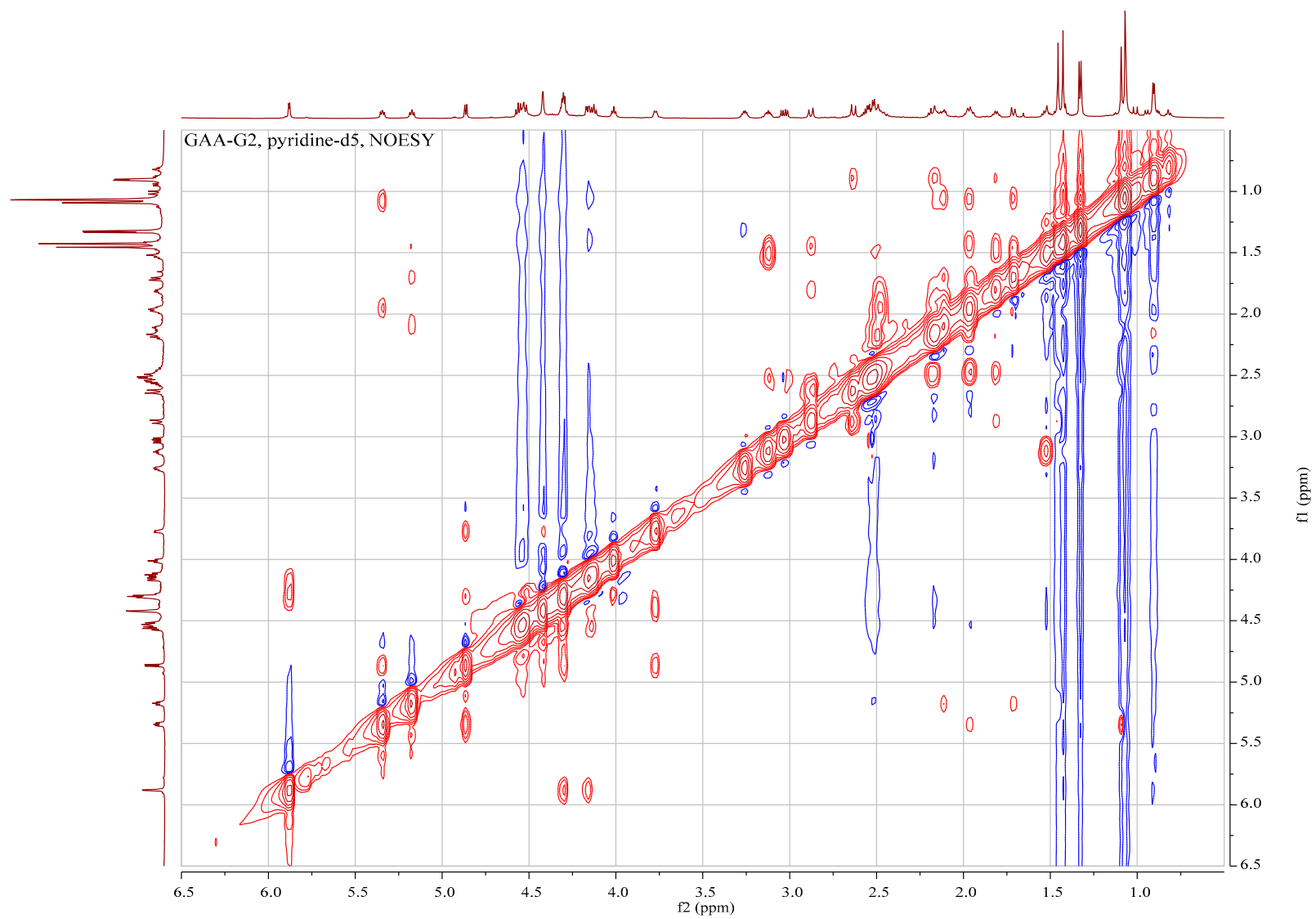


Figure S13. The NOESY (700 MHz) spectrum of GAA-G2 in pyridine-d₅.