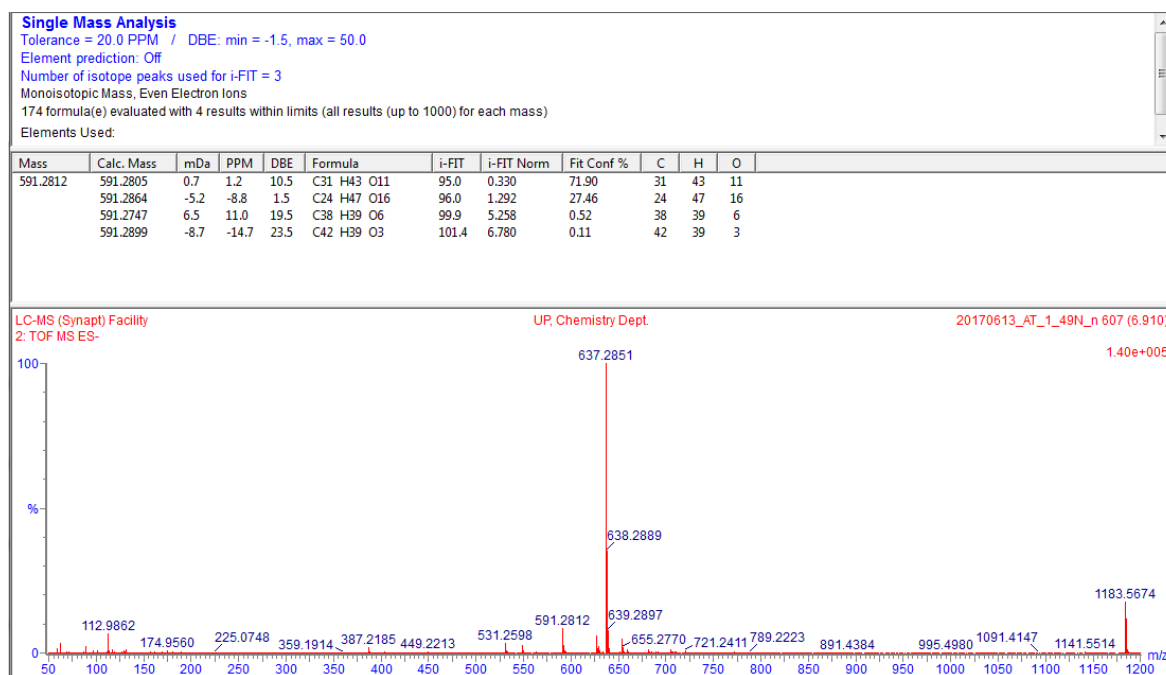
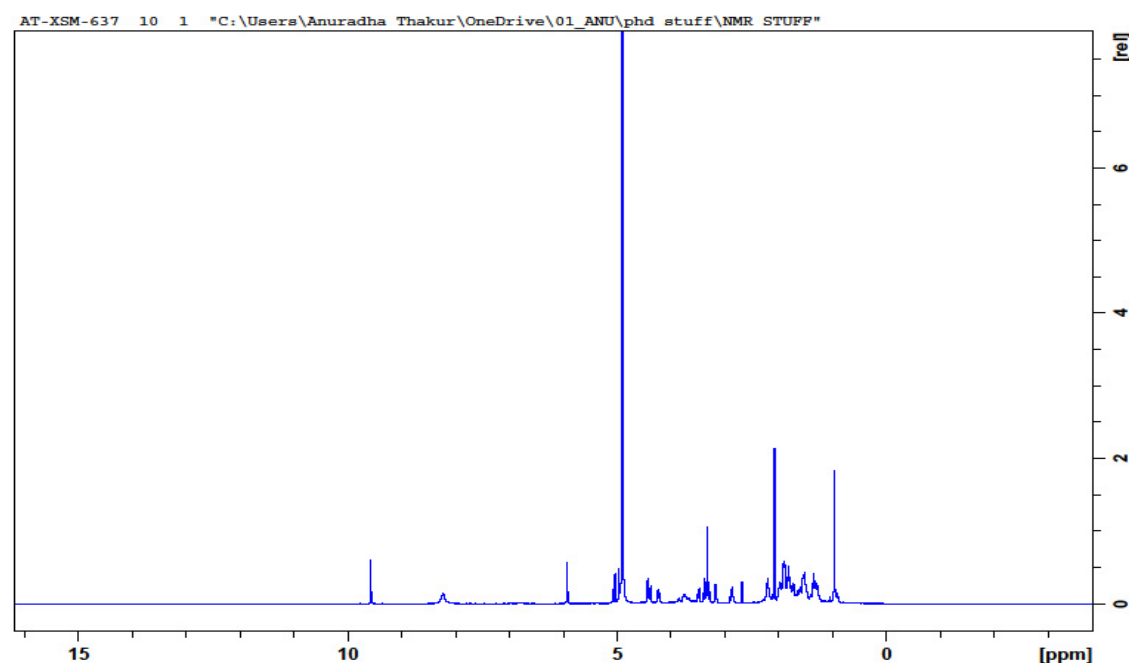


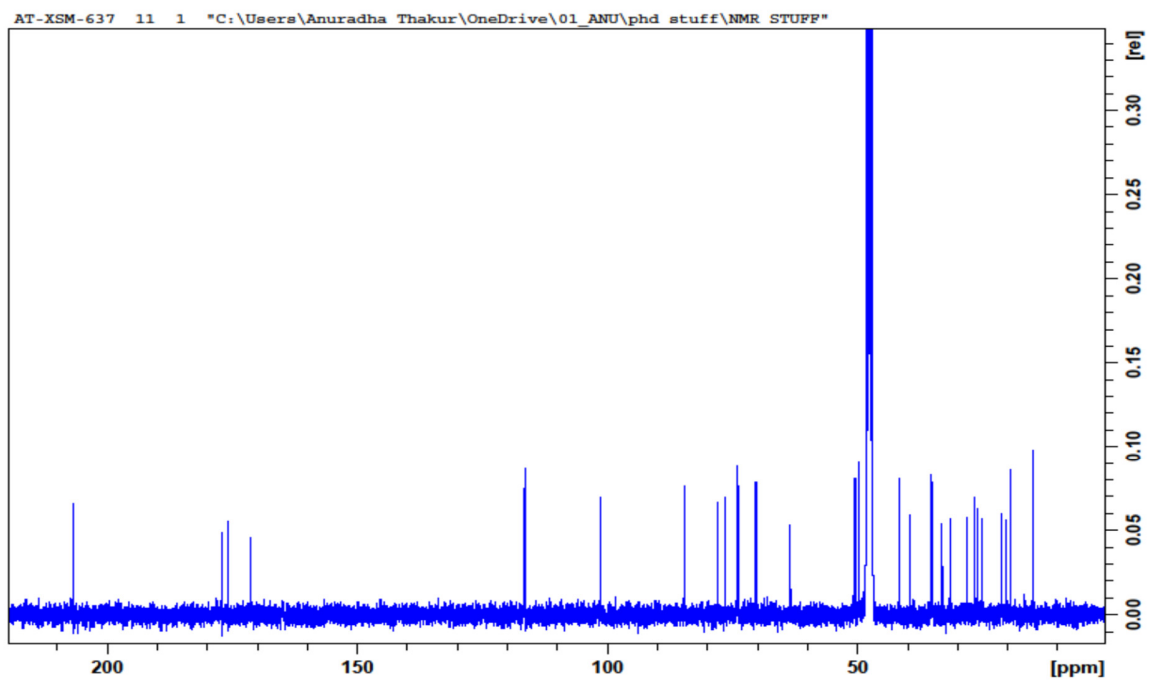
Supplementary Materials:



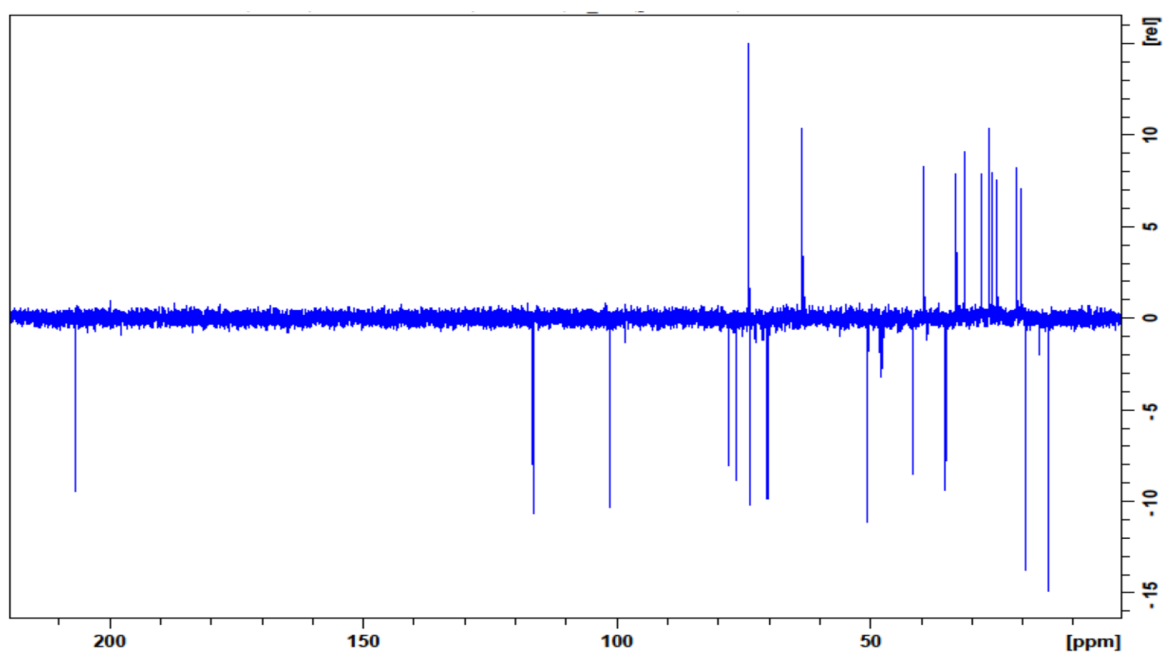
Supplementary Figure S1. iFit value of acetylated glycosydated crotoxigenin **1** in *X. undulatum* leaves extracted with DCM:MeOH



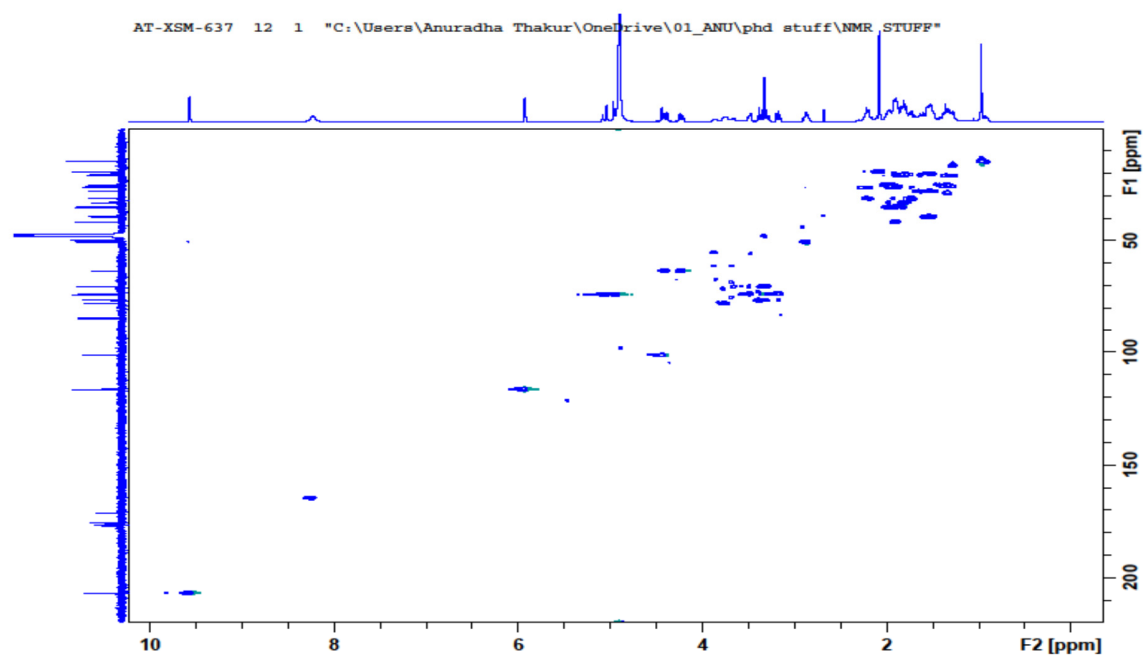
Supplementary Figure S2. ^1H NMR for acetylated glycosydated crotoxigenin **1** in MeOD- d_4



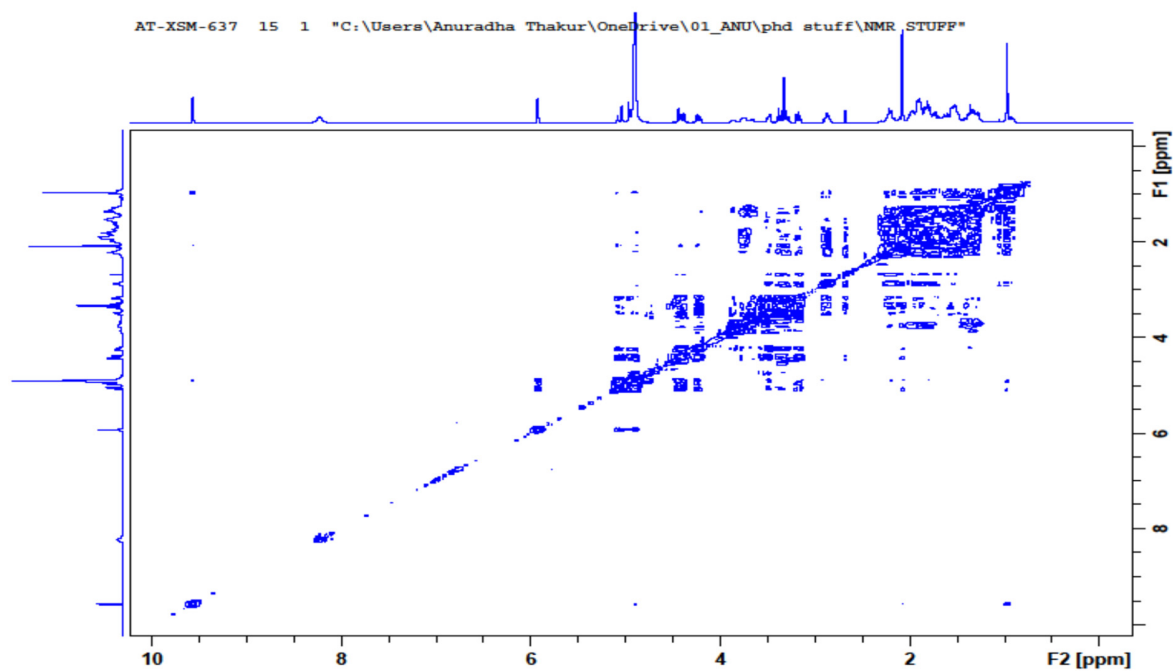
Supplementary Figure S3. ^{13}C NMR for acetylated glycosydated crotoxigenin **1** in $\text{MeOD-}d_4$



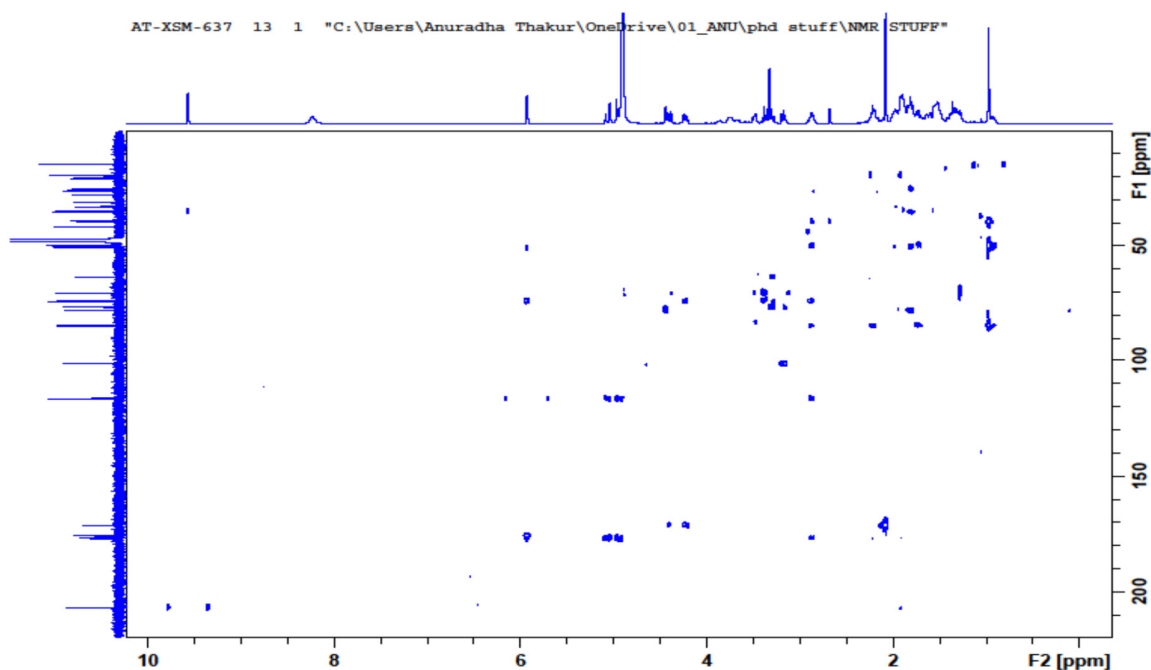
Supplementary Figure S4. DEPT 135 for acetylated glycosydated crotoxigenin **1** in $\text{MeOD-}d_4$



Supplementary Figure S5. HSQC spectrum for acetylated glycosydated crotoxogenin **1** in MeOD-*d*₄



Supplementary Figure S6. COSY spectrum for acetylated glycosydated crotoxogenin **1** in MeOD-*d*₄



Supplementary Figure S7. HMBC for acetylated glycosyated crotoxigenin 1 in MeOD-*d*₄

Single Mass Analysis

Tolerance = 20.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

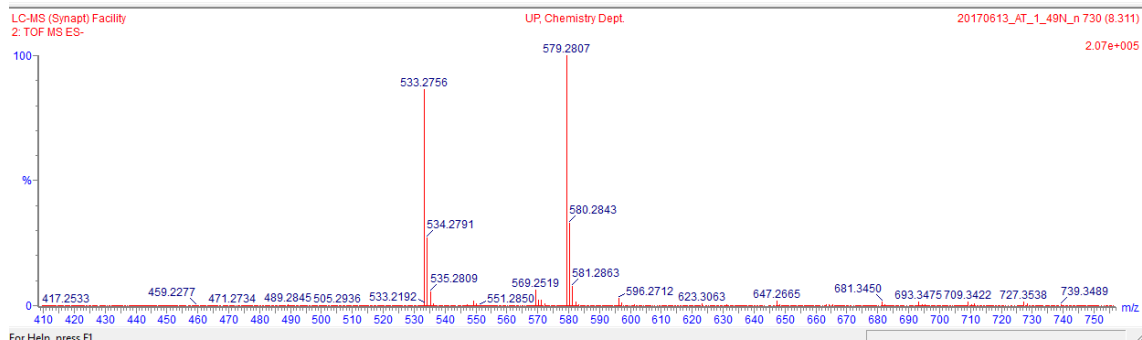
Number of isotope peaks used for iFIT = 3

Monoisotopic Mass, Even Electron Ions

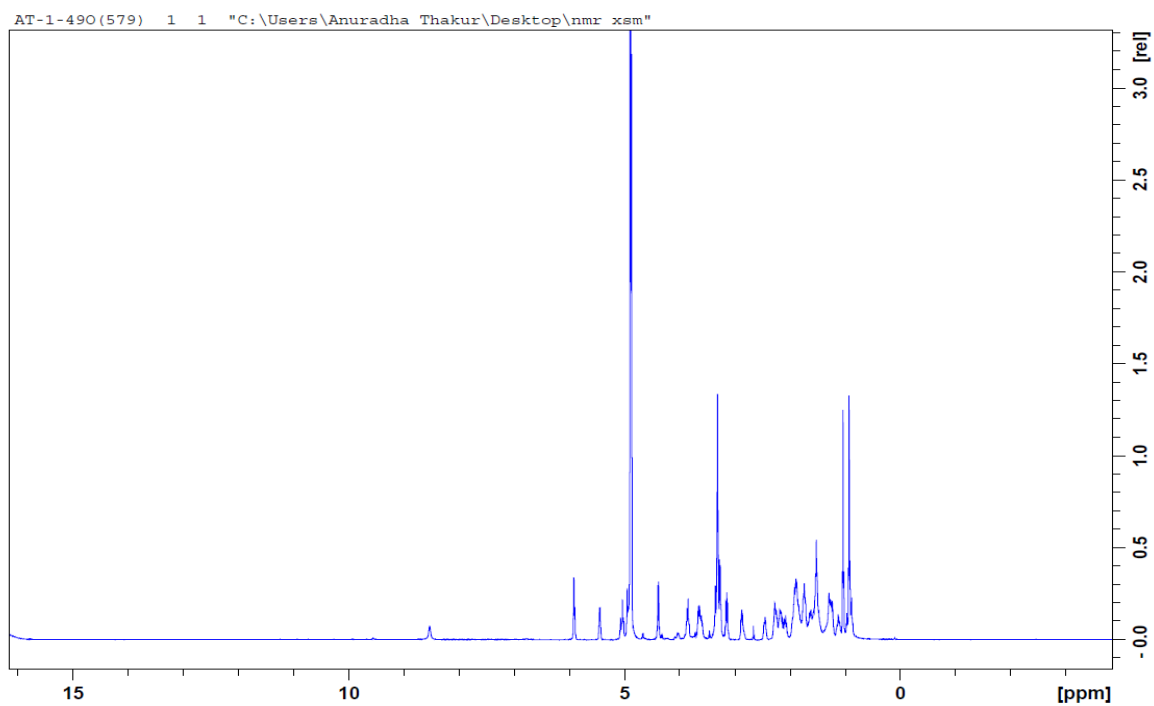
141 formula(e) evaluated with 4 results within limits (all results (up to 1000) for each mass)

Elements Used:

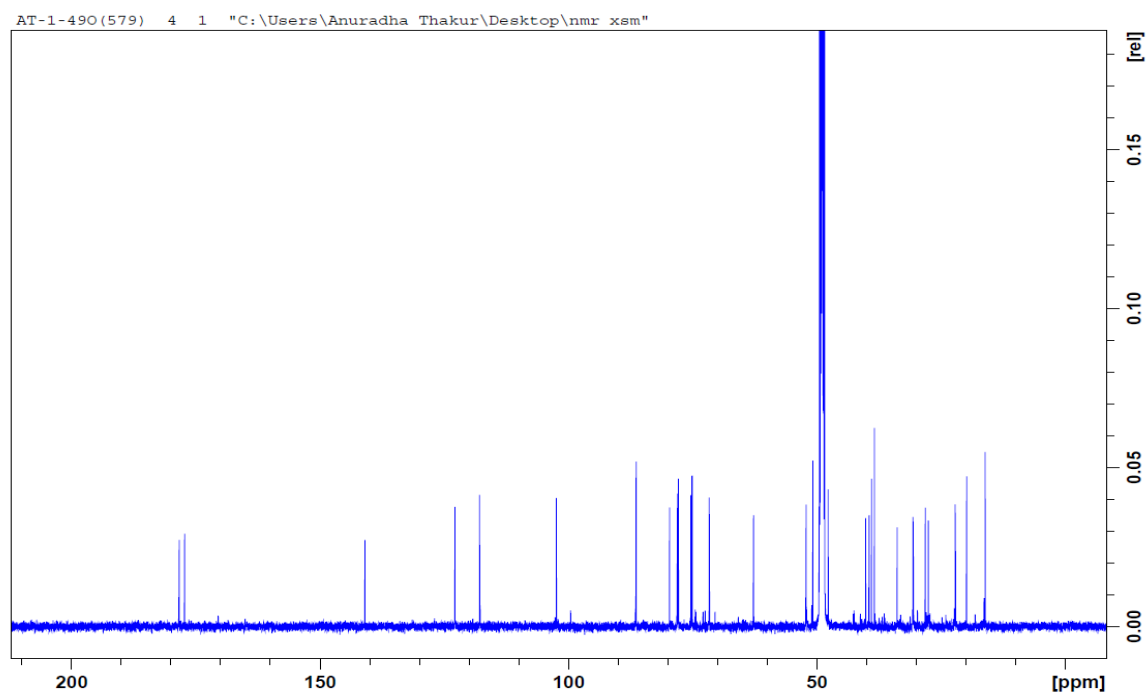
Mass	Calc. Mass	mDa	PPM	DBE	Formula	i-FIT	i-FIT Norm	Fit Conf %	C	H	O
533.2756	533.2751	0.5	0.9	9.5	C ₂₉ H ₄₁ O ₉	286.5	0.001	99.94	29	41	9
	533.2809	-5.3	-9.9	0.5	C ₂₂ H ₄₅ O ₁₄	294.0	7.476	0.06	22	45	14
	533.2692	6.4	12.0	18.5	C ₃₆ H ₃₇ O ₄	296.5	9.998	0.00	36	37	4
	533.2844	-8.8	-16.5	22.5	C ₄₀ H ₃₇ O	299.0	12.519	0.00	40	37	1



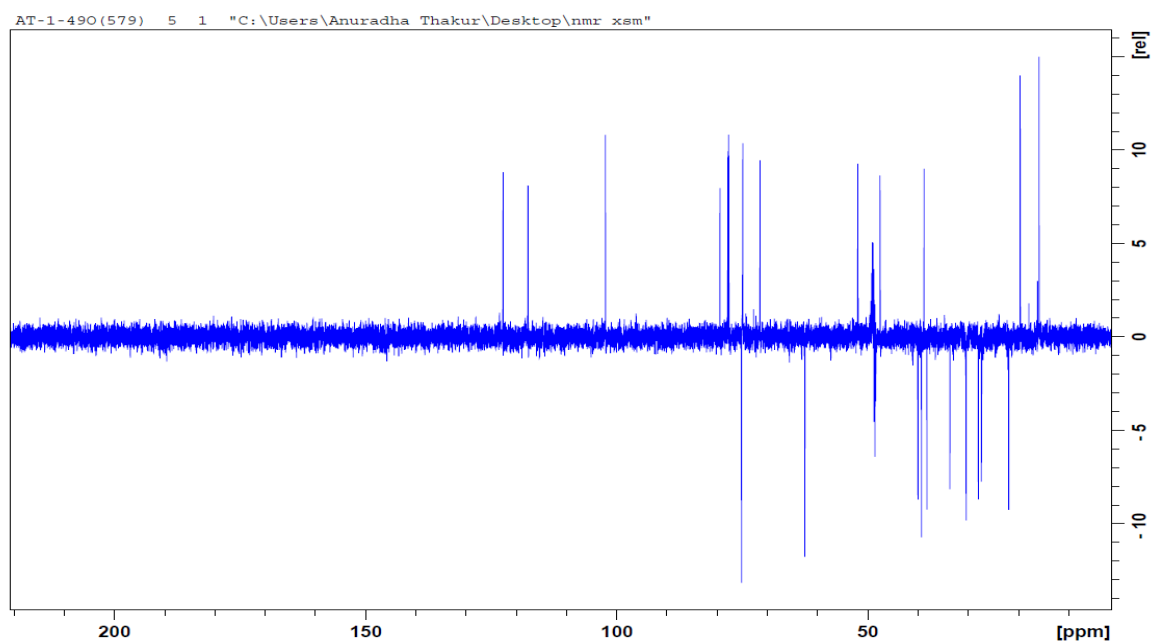
Supplementary Figure S8. iFit value of Xysmalogenin-3, β -D-glucopyranoside 2 in *X. undulatum* leaves extracted with DCM:MeOH



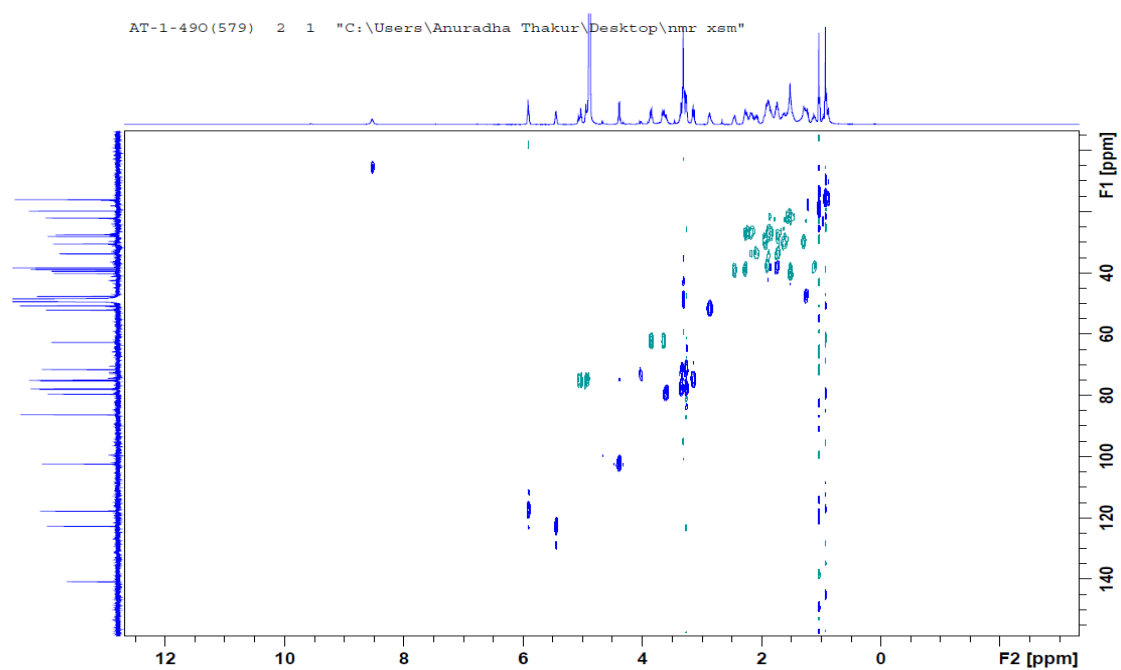
Supplementary Figure S9. ^1H NMR for Xysmalogenin-3, β -D-glucopyranoside **2** in $\text{MeOD-}d_4$



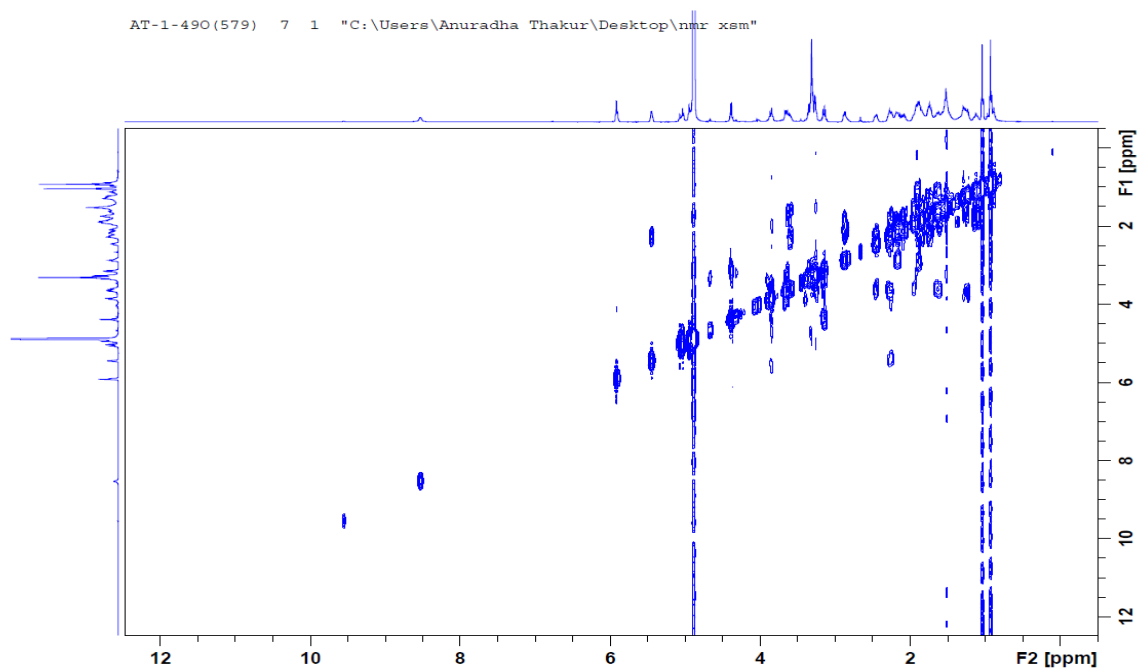
Supplementary Figure S10. ^{13}C NMR for Xysmalogenin-3, β -D-glucopyranoside **2** in $\text{MeOD-}d_4$



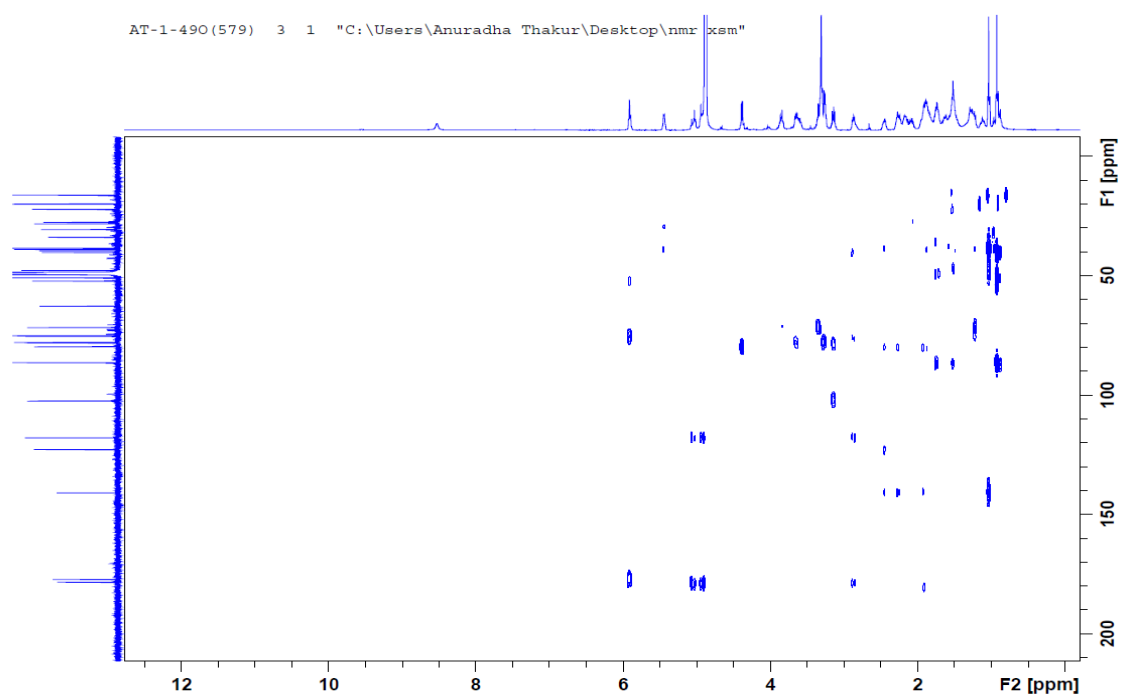
Supplementary Figure S11. DEPT 135 for Xysmalogenin-3, β -D-glucopyranoside **2** in MeOD-*d*₄



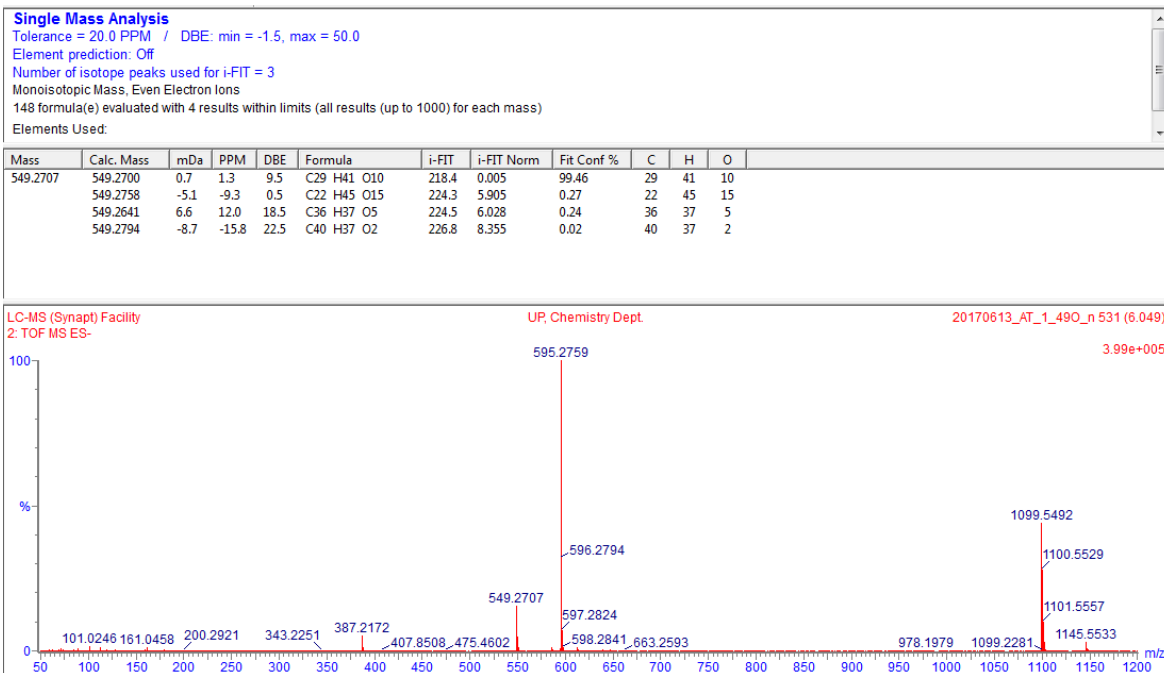
Supplementary Figure S12. HSQC for Xysmalogenin-3, β -D-glucopyranoside **2** in MeOD-*d*₄



Supplementary Figure S13. COSY spectrum for Xysmalogenin-3, β -D-glucopyranoside **2** in MeOD- d_4



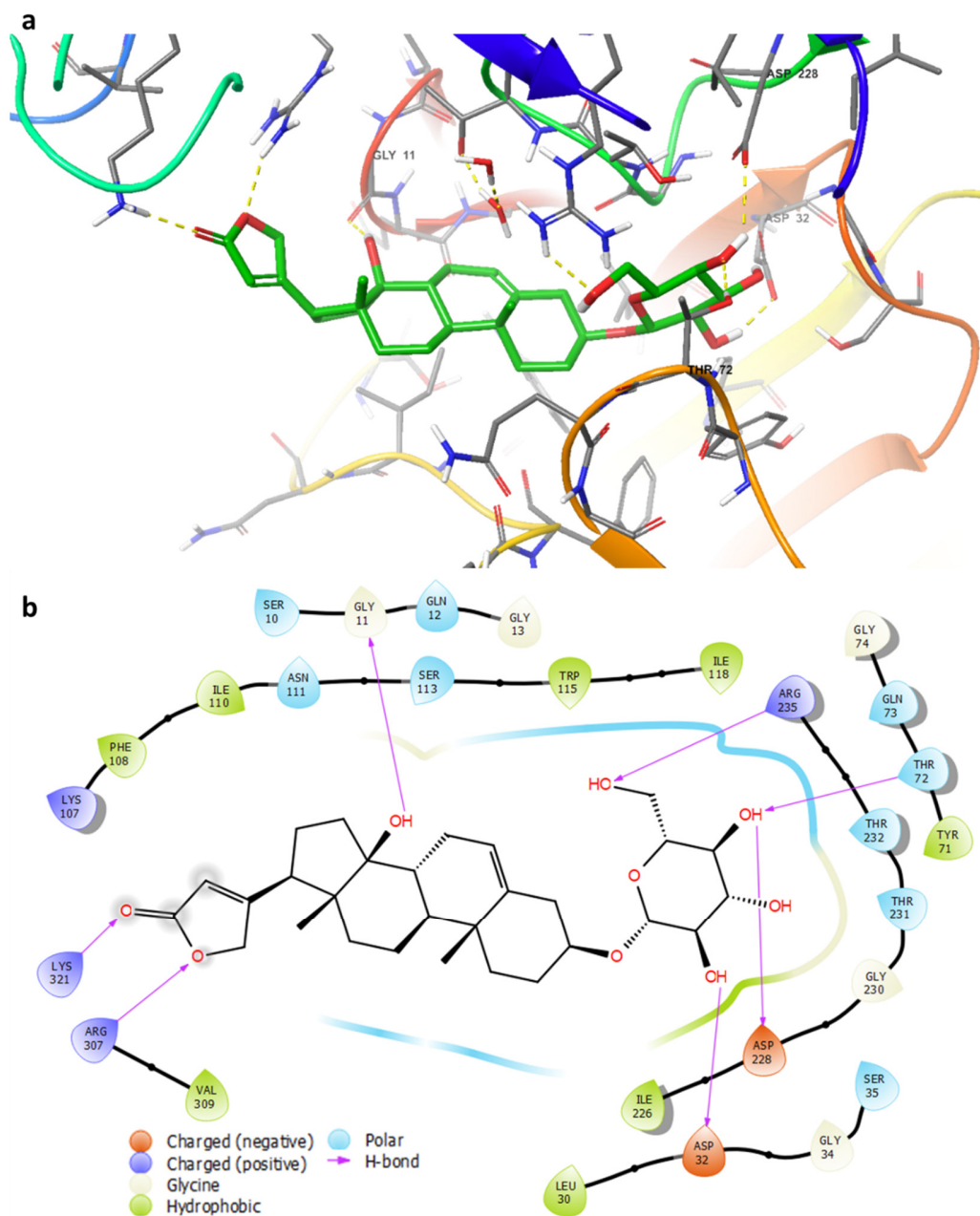
Supplementary Figure S14. HMBC spectrum for Xysmalogenin-3, β -D-glucopyranoside **2** in MeOD- d_4



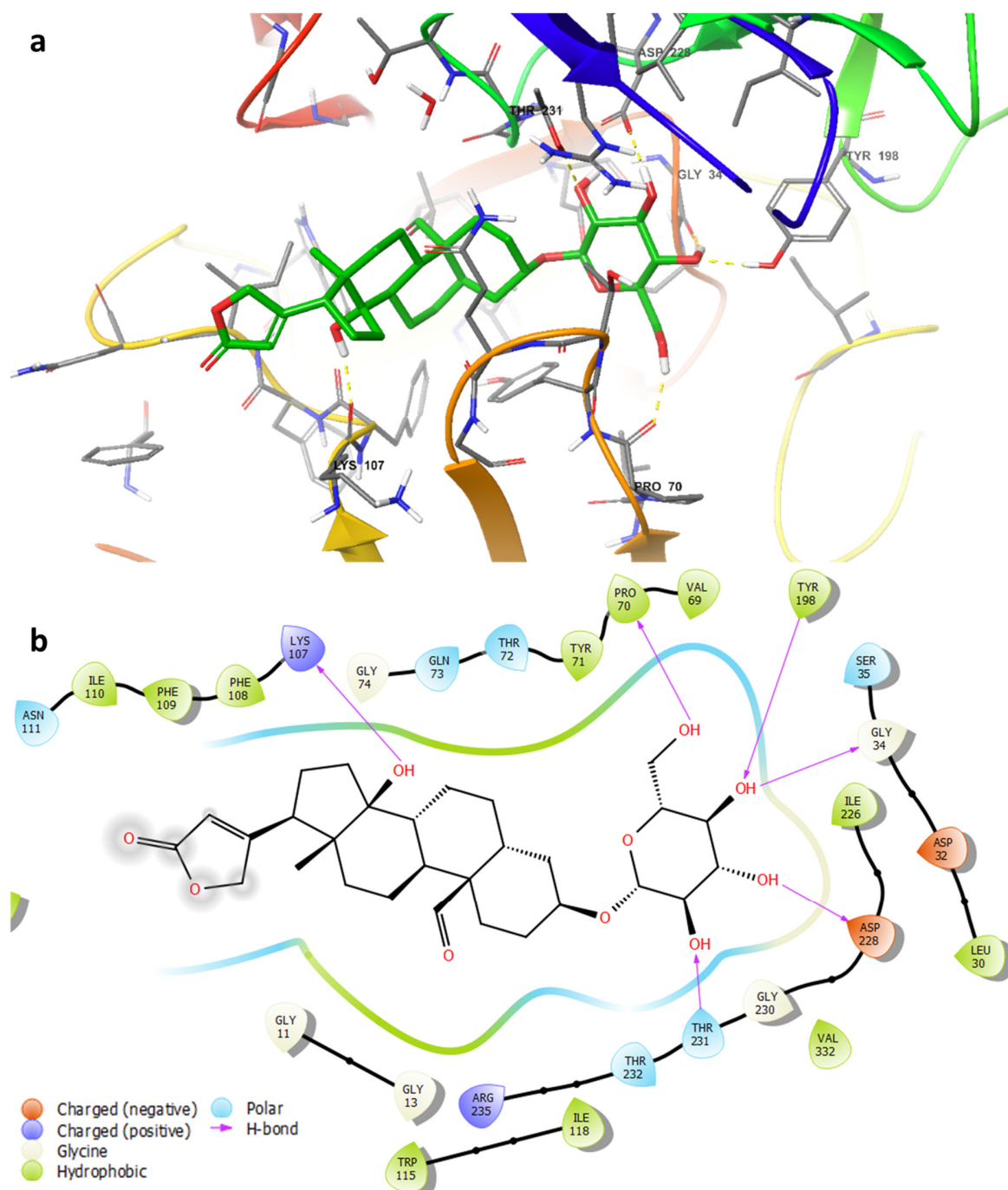
Supplementary Figure S15. iFit value of crotoxinigenin 3-O-glucopyranoside **3** in *X. undulatum* leaves extracted with DCM:MeOH

Supplementary Table S1. ¹H NMR and ¹³C NMR data for crotoxogenin 3-*O*-glucopyranoside **3** in MeOD-*d*4 compared to literature reports [1] for the compound in DMSO-*d*6.

Isolated Crotoxogenin 3- <i>O</i> -glucopyranoside (18)			¹ H NMR and ¹³ C NMR literature data for compound (59) in DMSO- <i>d</i> 6	
Position	¹³ C (ppm)	¹ H (ppm), <i>J</i> (Hz)	¹³ C (ppm)	¹ H (ppm), <i>J</i> (Hz)
1	28.0	1.83 m, 1.54 m	28.03	1.85 m, 1.50 m
2	31.3	1.85 m	30.36	1.80 m
3	77.0	3.82 m	76.78	3.05 m
4	25.9	1.85 m, 1.33 m	30.20	1.82 m, 1.14 m
5	41.6	1.89 m	42.03	1.45 m
6	24.9	1.76 m	26.27	1.74 m
7	20.3	1.48 m	21.41	1.60 m
8	35.2	1.96 m	47.57	1.43 m
9	35.0	1.79 m	35.49	1.81 m
10	50.4	-	49.99	-
11	21.2	1.82 m, 1.32 m	21.41	1.60 m
12	39.4	1.51 m	42.18	1.44 m
13	49.6	-	49.19	-
14	84.5	-	83.37	-
15	32.9	1.71 m	31.59	1.94 m, 1.55 m
16	26.5	2.16 m	27.17	2.13 m
17	50.5	2.8 (q, 14.6)	51.23	2.72 m
18	14.8	0.95 s	15.54	0.69 s
19	206.8	9.54 s	209.66	9.94 s
20	175.8	-	173.96	-
21	73.7	5.04 (d, 1.65), 4.92 (dd, 18.5, 1.74)	73.48	4.90 dd
22	116.4	5.91 s	116.38	5.89 s
23	176.9	-	176.31	-
1'	100.8	4.42 d (7.80)	100.71	4.20 d
2'	73.9	3.14 (dd, 9.03, 1.07)	75.75	3.59 m
3'	76.5	3.26	73.23	2.85 m
4'	70.3	3.27 m	70.11	3.02 m
5'	76.7	3.35m	76.78	3.06 m
6'	61.4	3.87 m, 3.65 m	61.14	3.64m, 3.40m



Supplementary Figure S16. (a) Predicted binding pose from induced fit docking for **2**, shown in green, bound to BACE1. Hydrogen bonds are shown in yellow dashed lines. Only selected residues are shown for clarity. (b) 2D ligand interaction diagram for the binding pose of **2**.



Supplementary Figure S17. (a) Predicted binding pose from induced fit docking for **3**, shown in green, bound to BACE1. Hydrogen bonds are shown in yellow dashed lines. Only selected residues are shown for clarity. (b) 2D ligand interaction diagram for the binding pose of **3**.

References:

- [1] L. Rascón-Valenzuela, C. Velázquez, A. Garibay-Escobar, L. A. Medina-Juárez, W. Vilegas, and R. E. Robles-Zepeda, “Antiproliferative activity of cardenolide glycosides from *Asclepias subulata*,” *J. Ethnopharmacol.*, vol. 171, pp. 280–286, 2015, doi: 10.1016/j.jep.2015.05.057.