

FURTHER INSIGTHS IN THE DESIGN OF POTENT UROPATHOGENIC *E. COLI* FIMH ANTAGONISTS

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Materials and Methods

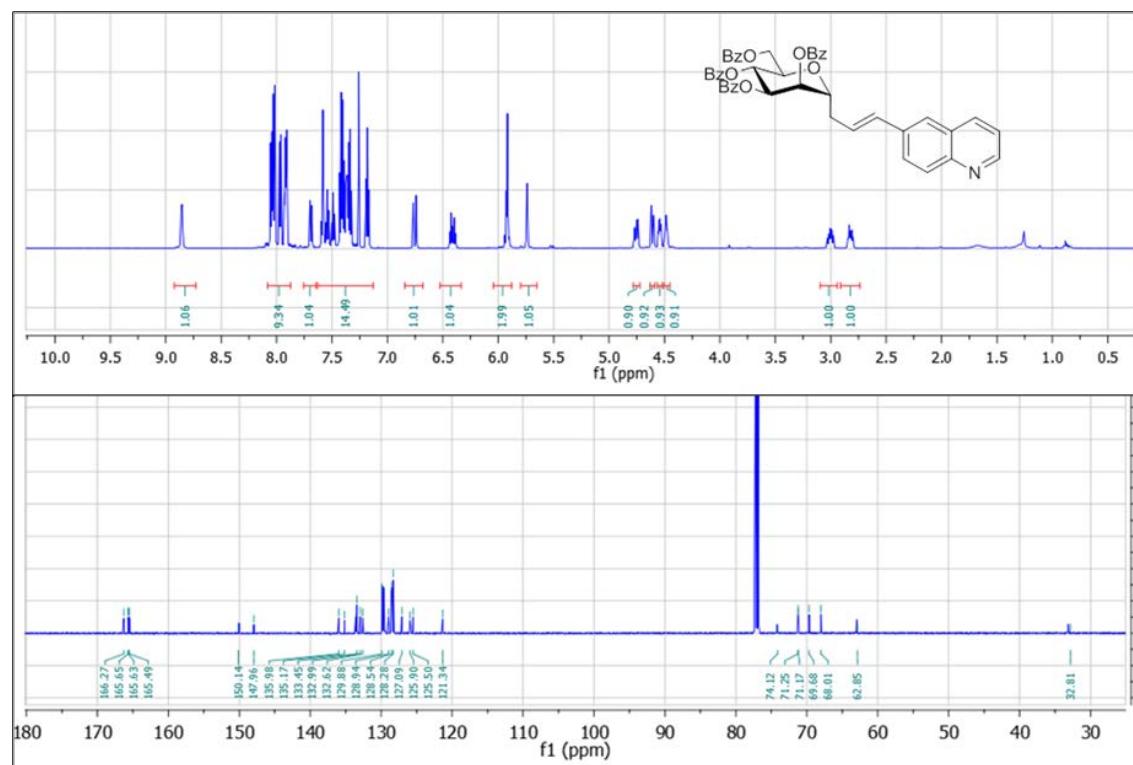


Figure S1. ¹HNMR and ¹³CNMR of compound 2 (CDCl₃, 600 and 151 MHz, respectively)

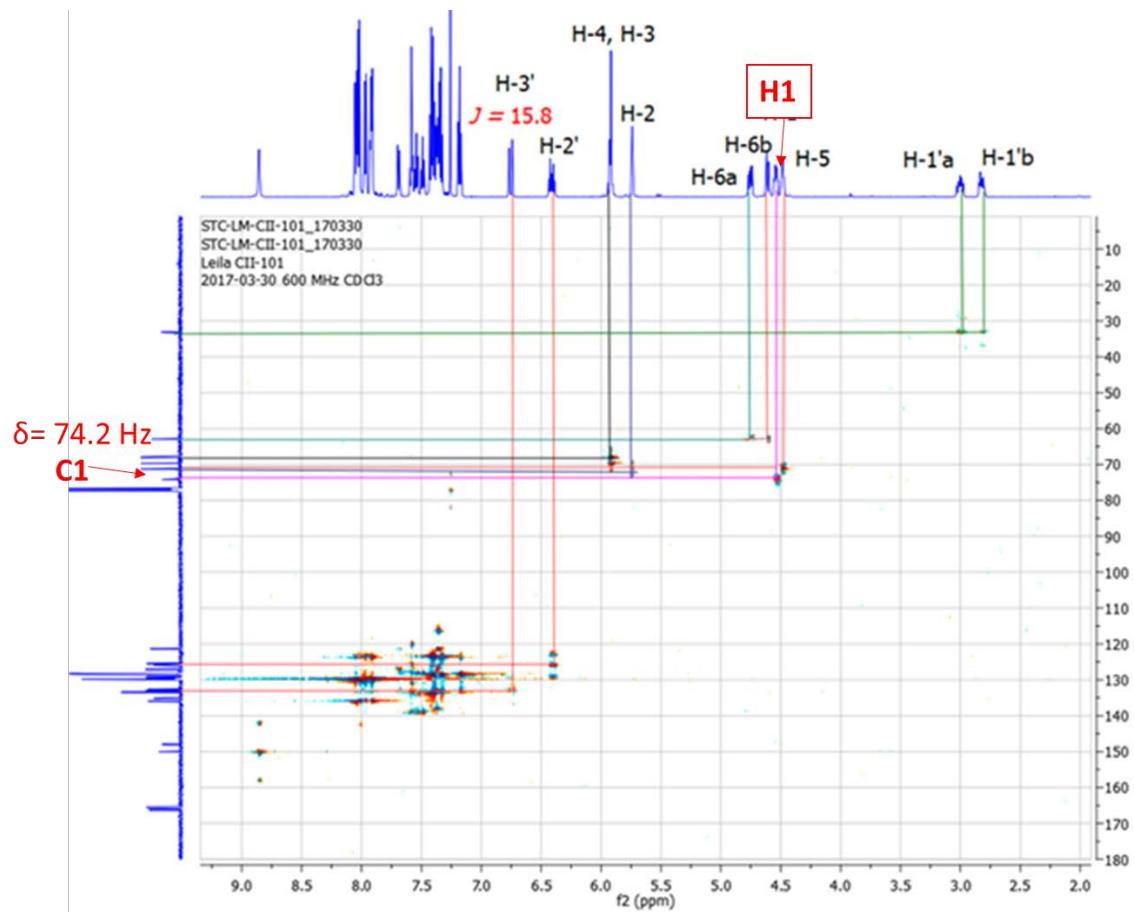


Figure S2. HSQC of purified compound 2. The coupling constant of H-3' ($J_{2',3'}=15.8$ Hz) indicates the presence of the trans stereoisomer. (CDCl_3 , 600 MHz)

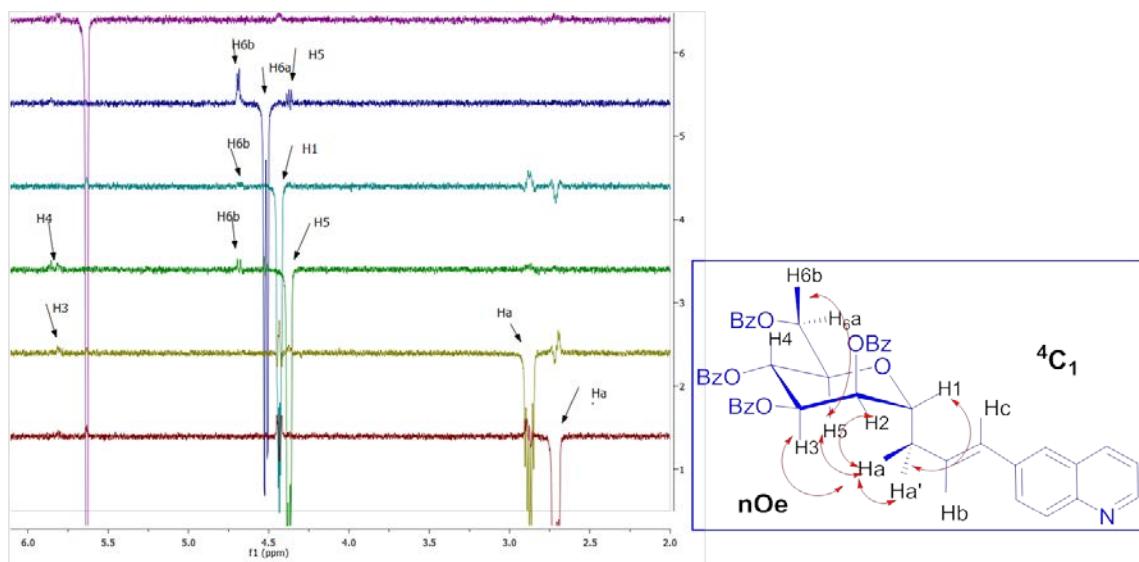


Figure S3. Conformational studies of compound 2 in solution using NOESY technique 600 MHz.

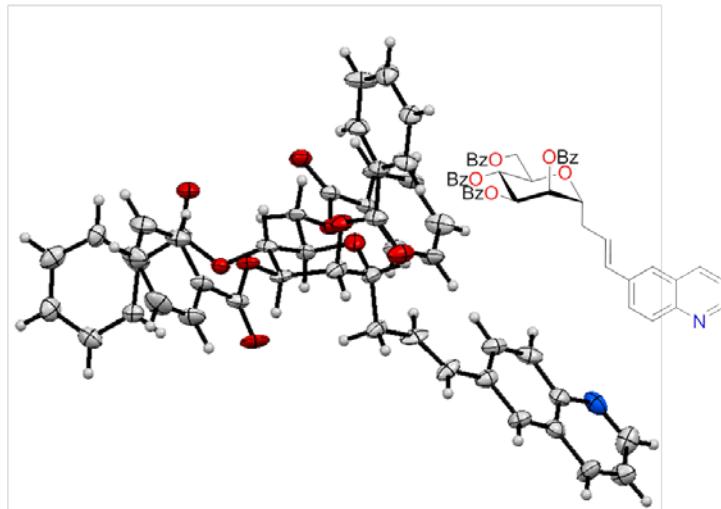


Figure S4. The ORTEP diagram of compound 2 clearly shows it to exist as the proper chair conformer 4C1. X-Ray crystal-structure and refinement data: formula, $(\text{C}_{46}\text{H}_{37}\text{NO}_9)$, monoclinic, space group P21, $a = 24.6459(12)$ Å, $b = 6.3450(3)$ Å, $c = 25.5657(12)$ Å, $\beta = 108.871(2)^\circ$, $V = 3783.0(3)$ Å 3 , $Z = 4$, $T = 150$ K, $D_{\text{calcd}} = 1.313$ g/cm 3 . Crystallographic data for the structure reported in this paper has been deposited at the Cambridge Crystallographic Data Centre (CCDC) with deposition no: (CCDC:1560371). Supplementary data can be obtained free of charge from CCDC, 12 Union Road, Cambridge CB2 1EZ,

UK (fax: (+44)1223-336-033; e-mail: deposit@ccdc.cam.ac.uk

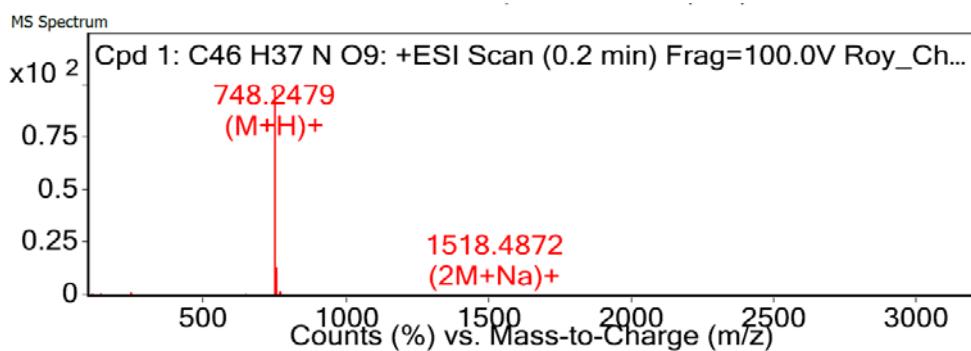


Figure S5. ESI+-HRMS spectrum of compound 2

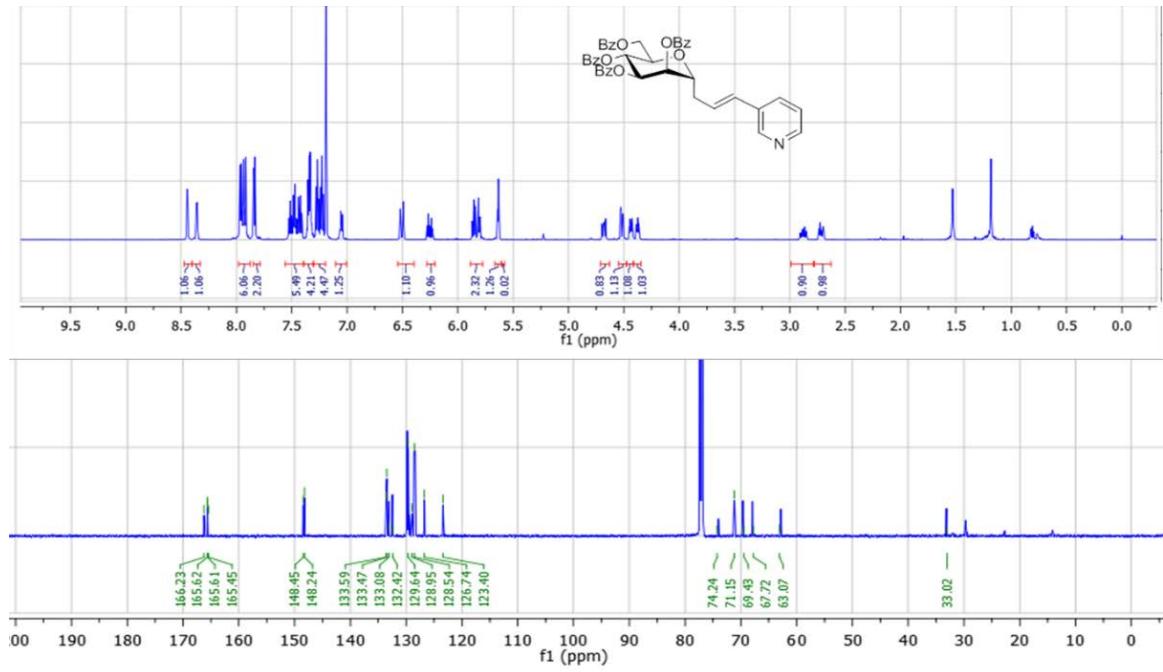


Figure S6. ^1H NMR and ^{13}C NMR of compound 3(CDCl_3 , 600 and 151 MHz, respectively)

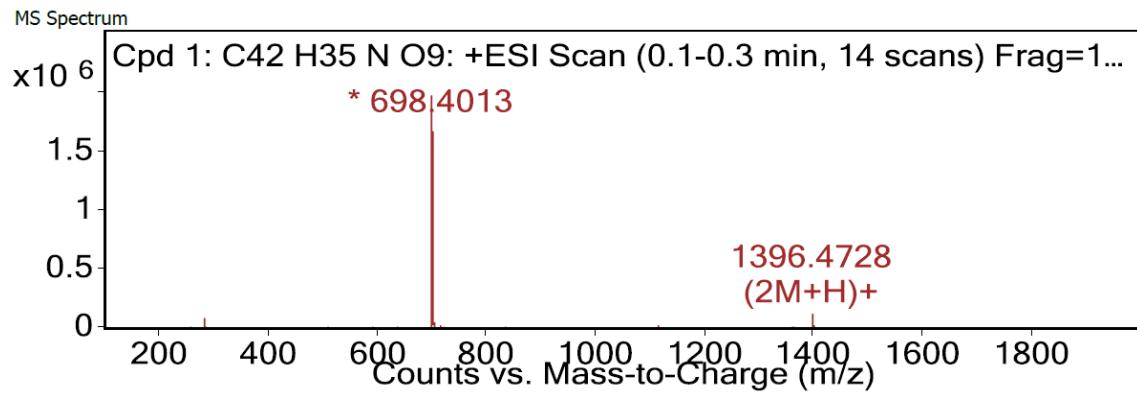


Figure S7. ESI+HRMS spectrum of compound 3

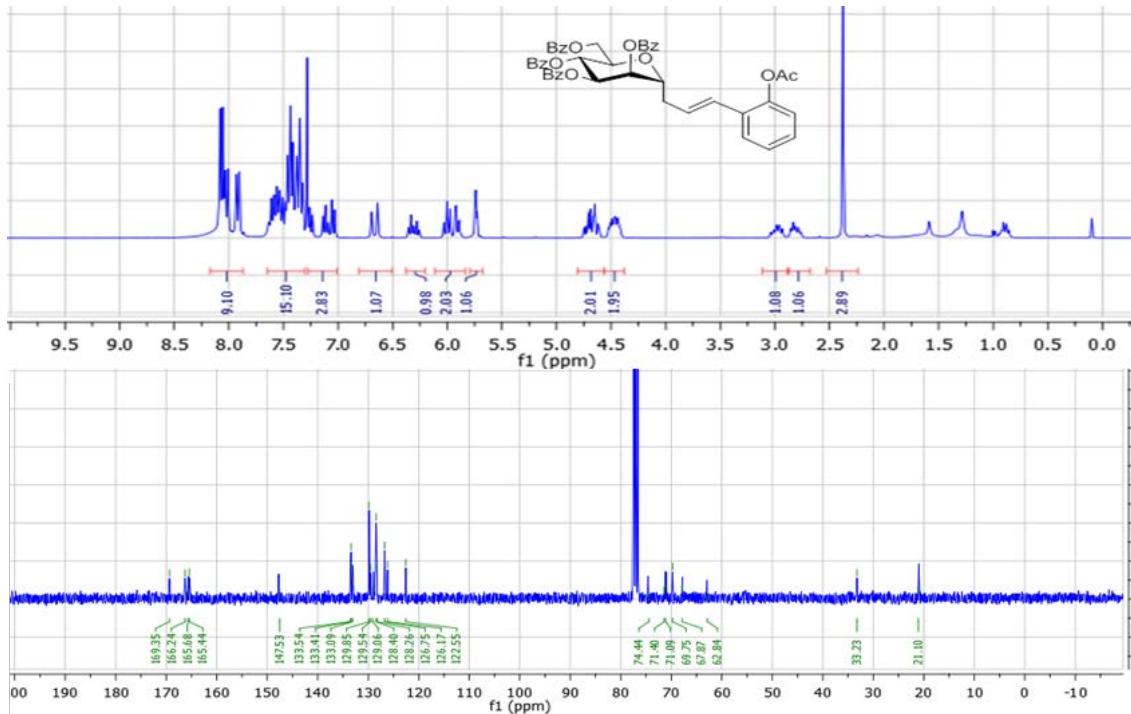


Figure S8. ^1H NMR and ^{13}C NMR of compound 4(CDCl_3 , 300 and 75 MHz respectively)

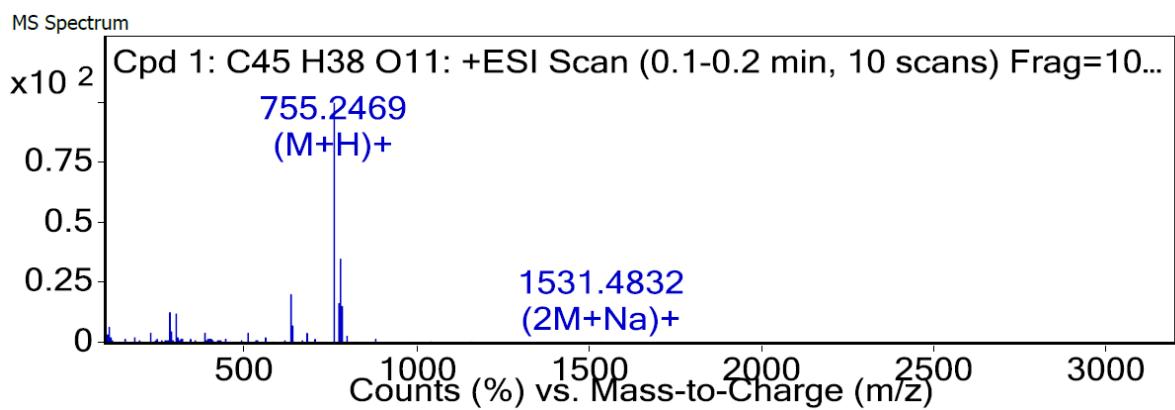


Figure S9. ESI+-HRMS spectrum of compound 4

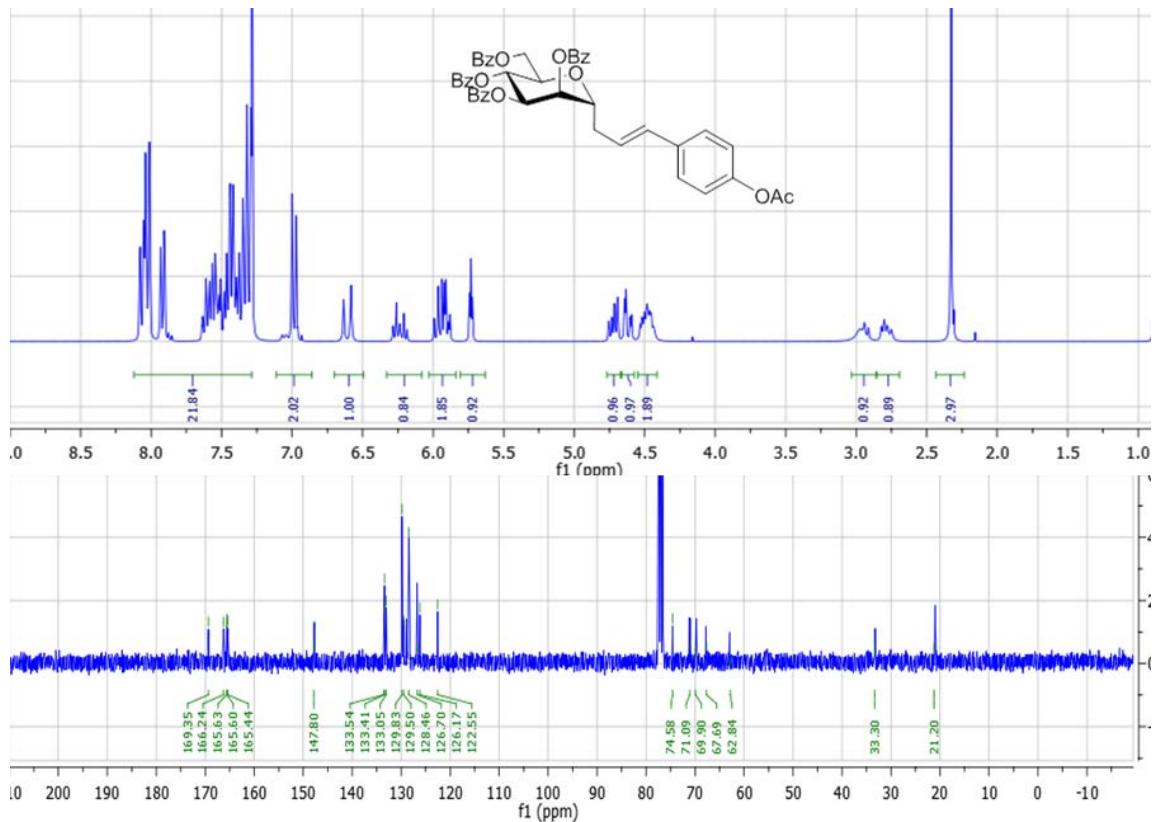


Figure S10. ¹HNMR and ¹³CNMR of compound 5(CDCl₃, 300 and 75 MHz respectively)

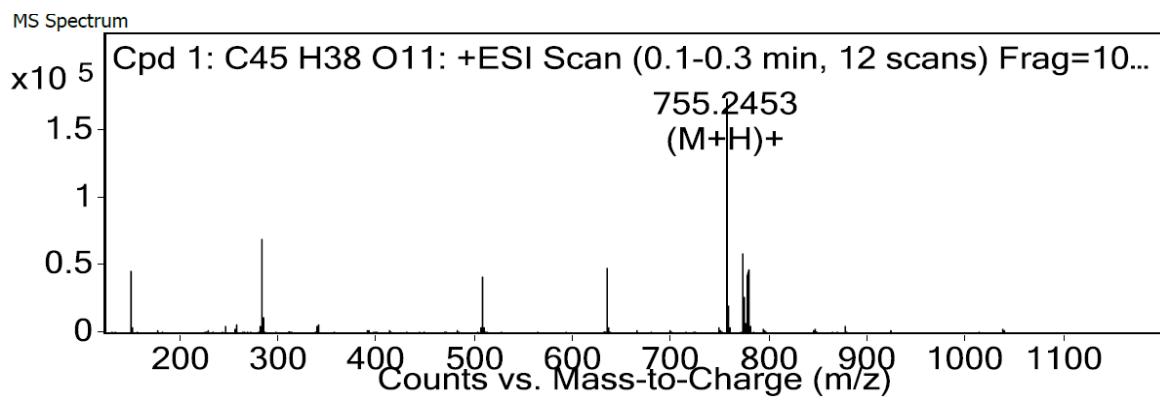


Figure S11. ESI+HRMS spectrum of compound 5

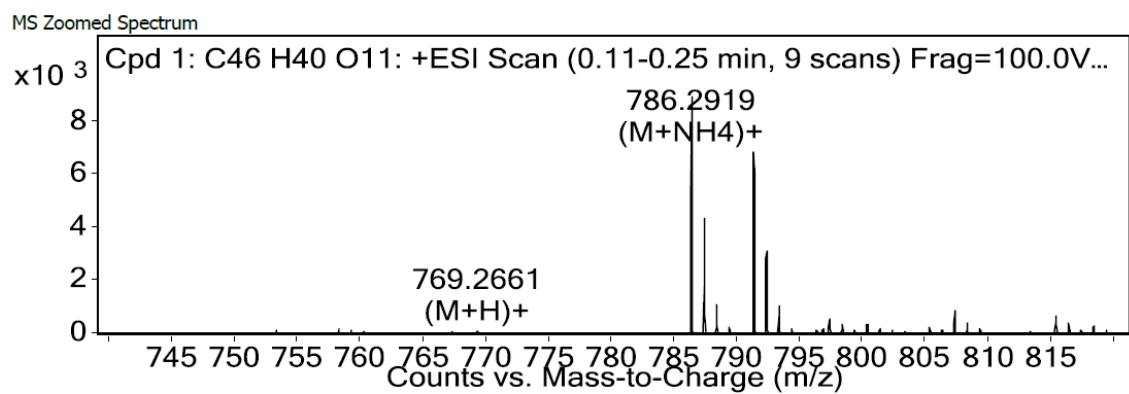
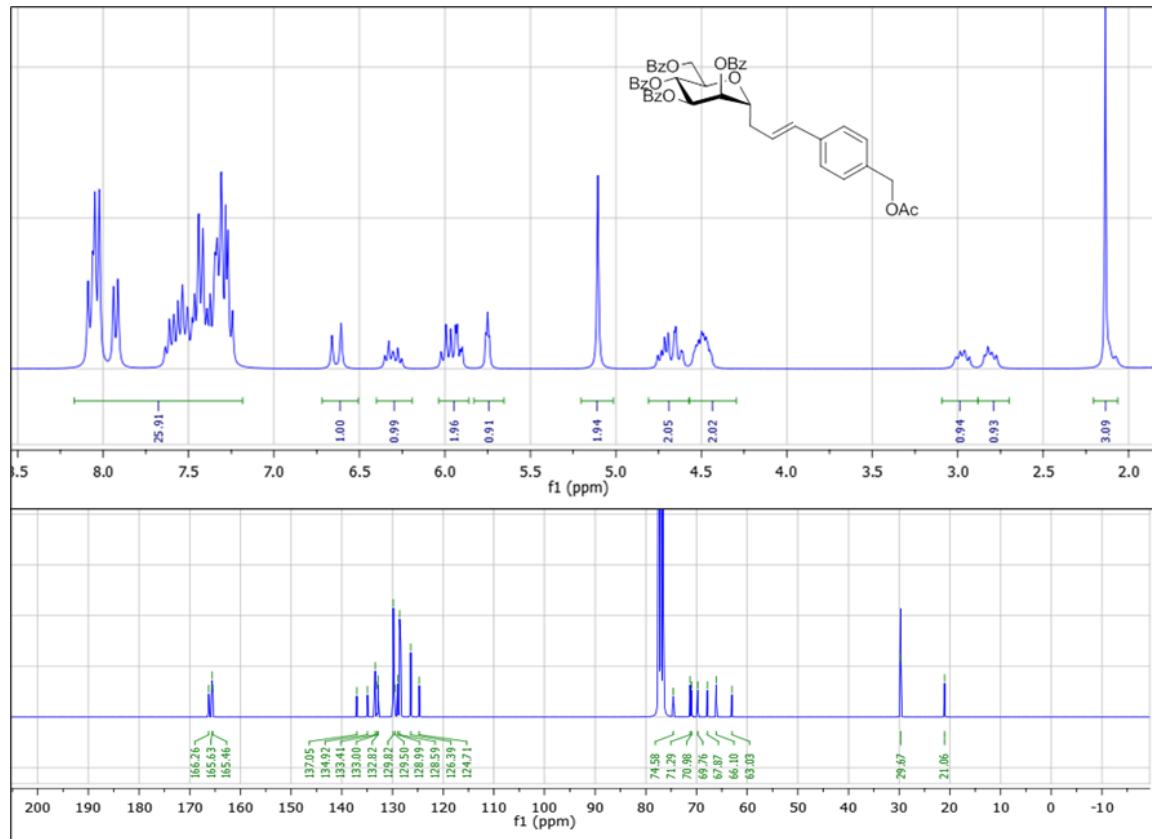


Figure S13. ESI+HRMS spectrum of compound 6

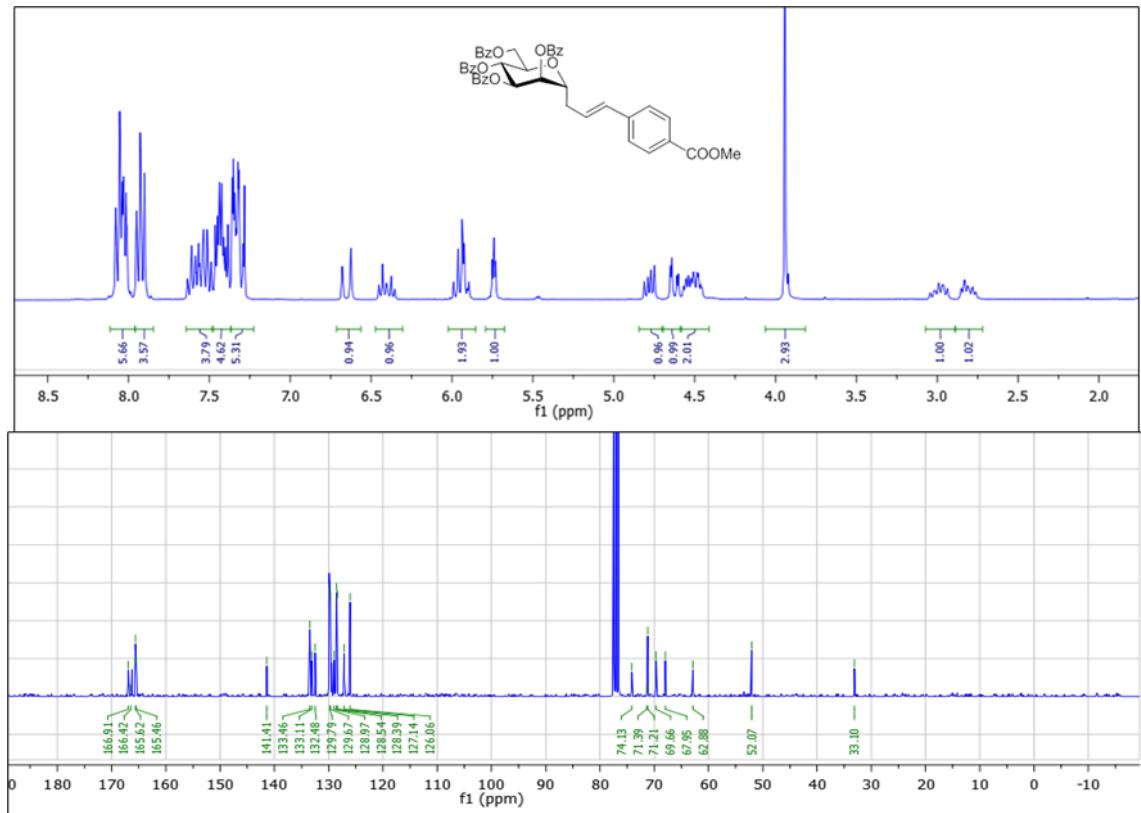


Figure S14. ^1H NMR and ^{13}C NMR of compound 7(CDCl_3 , 300 and 75 MHz respectively)

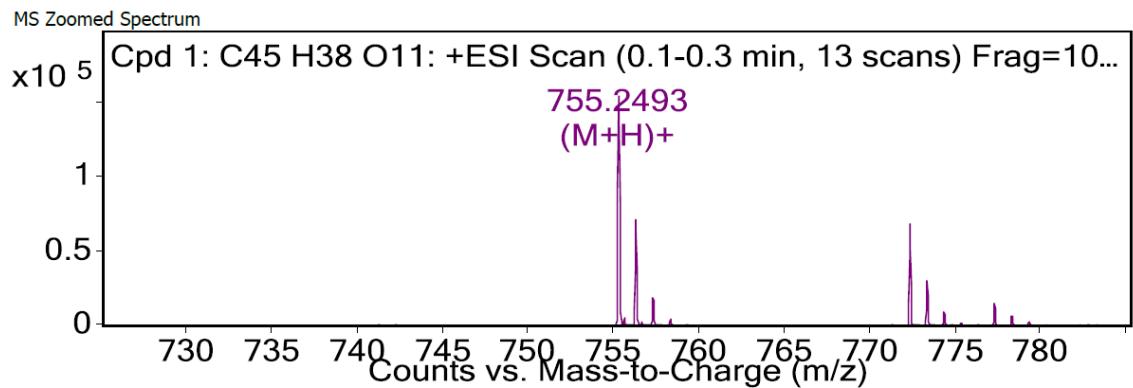


Figure S15. ESI+HRMS spectrum of compound 7

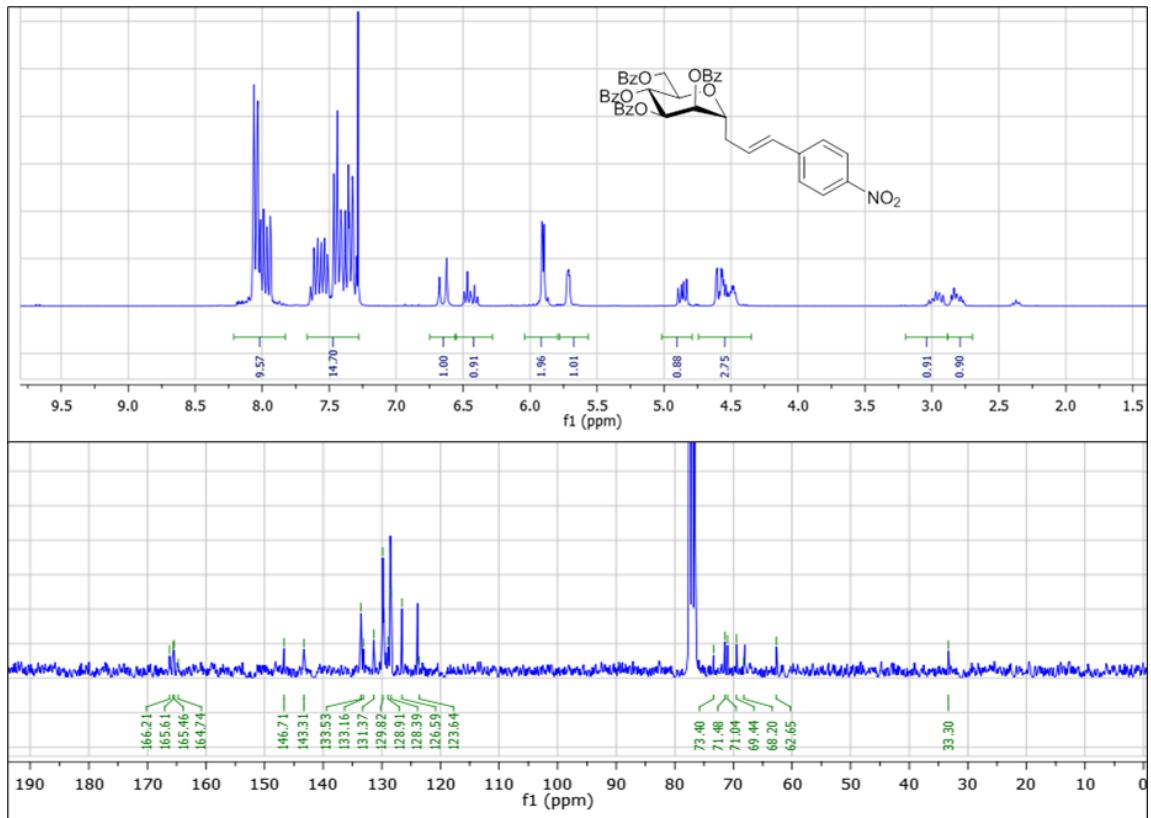


Figure S16. ^1H NMR and ^{13}C NMR of compound 8(CDCl_3 , 300 and 75 MHz respectively)

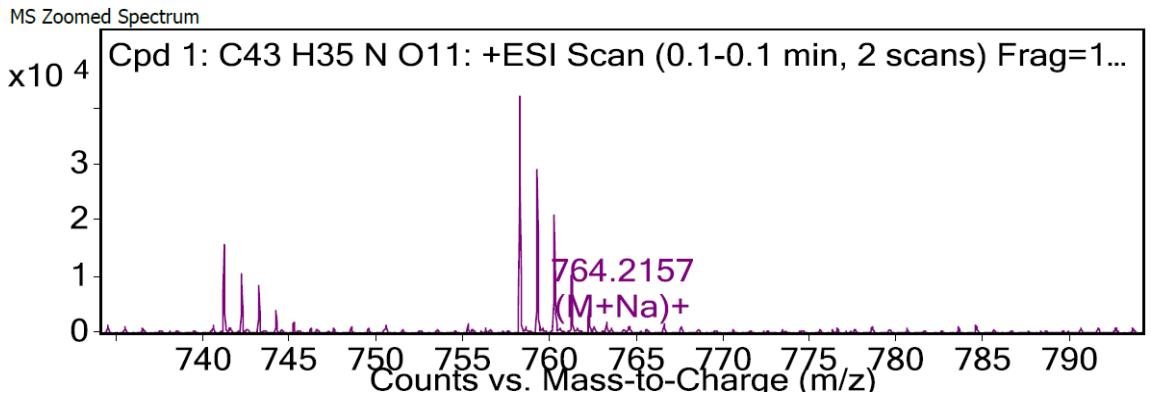


Figure S17. ESI+-HRMS spectrum of compound 8

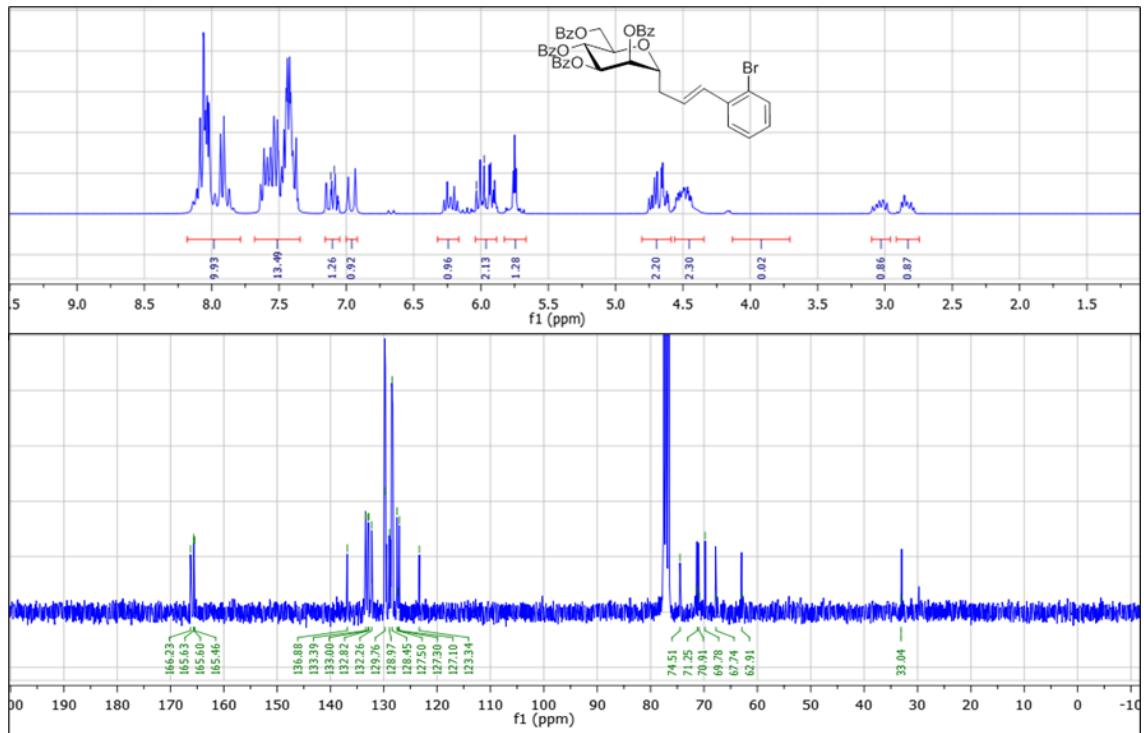


Figure S18. ¹HNMR and ¹³CNMR of compound 10(CDCl₃, 300 and 75 MHz respectively)

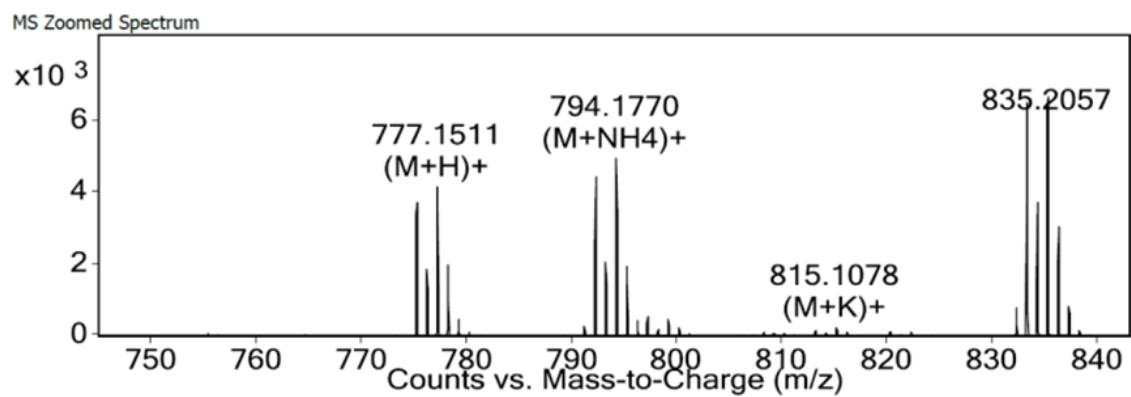


Figure S19. ESI+-HRMS spectrum of compound 10

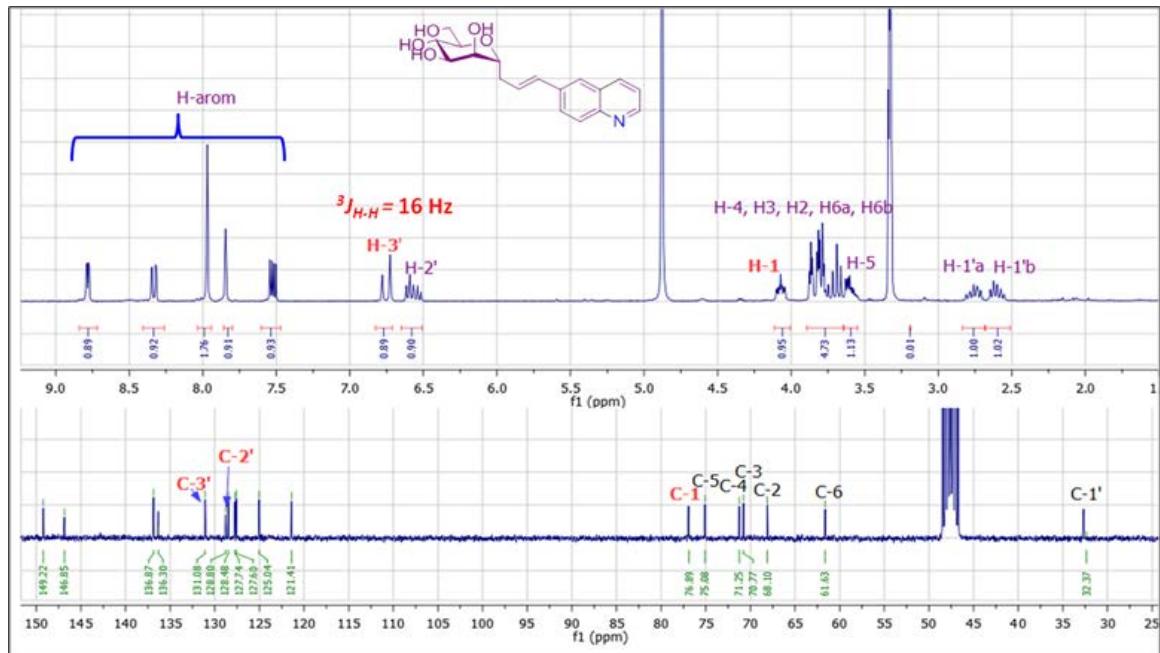


Figure S20. Fully assigned ^1H NMR and ^{13}C NMR of compound 11(CD₃OD, 600 and 151MHz respectively)

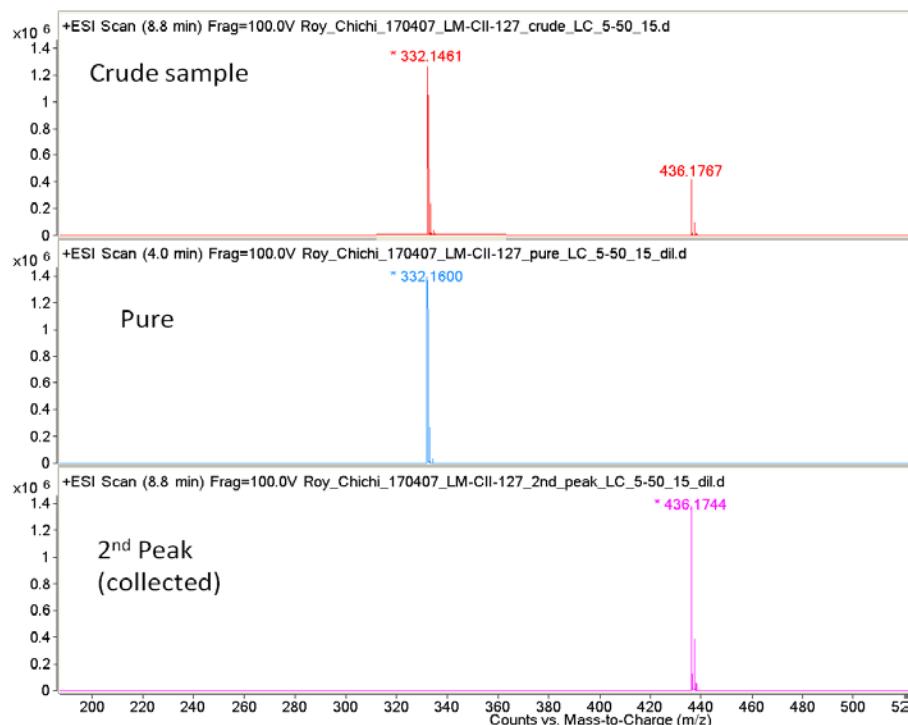


Figure S21. HPLC-TOF-MAS analysis spectrum of compound 11

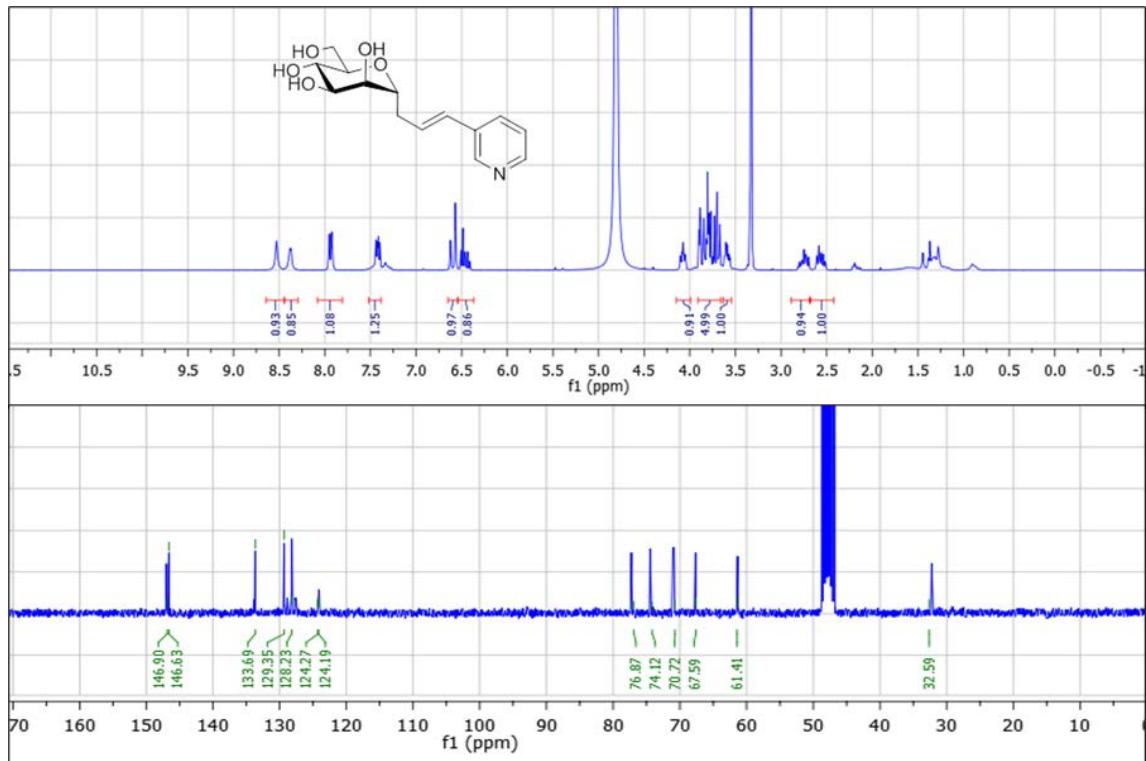


Figure S22. ¹H NMR and ¹³C NMR spectrum of compound 12 (CD₃OD, 300 and 75MHz respectively)

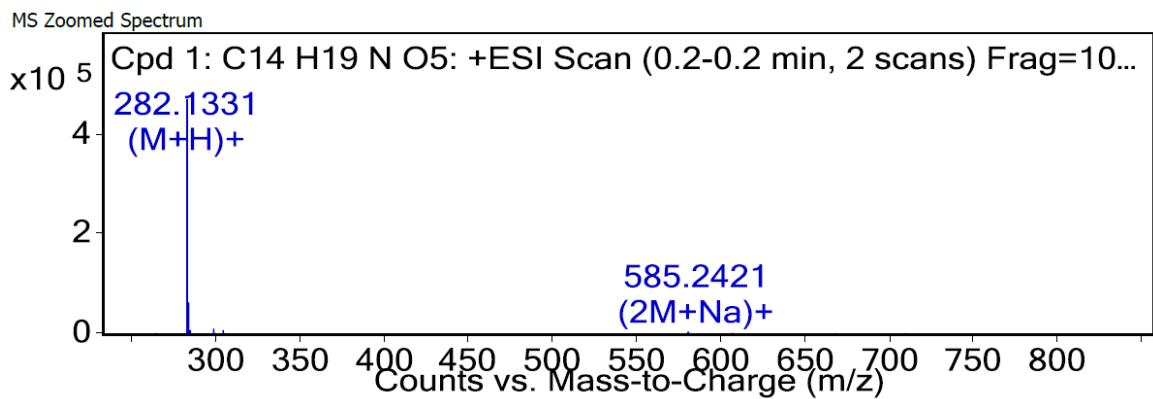


Figure S23. ESI+-HRMS spectrum of compound 12

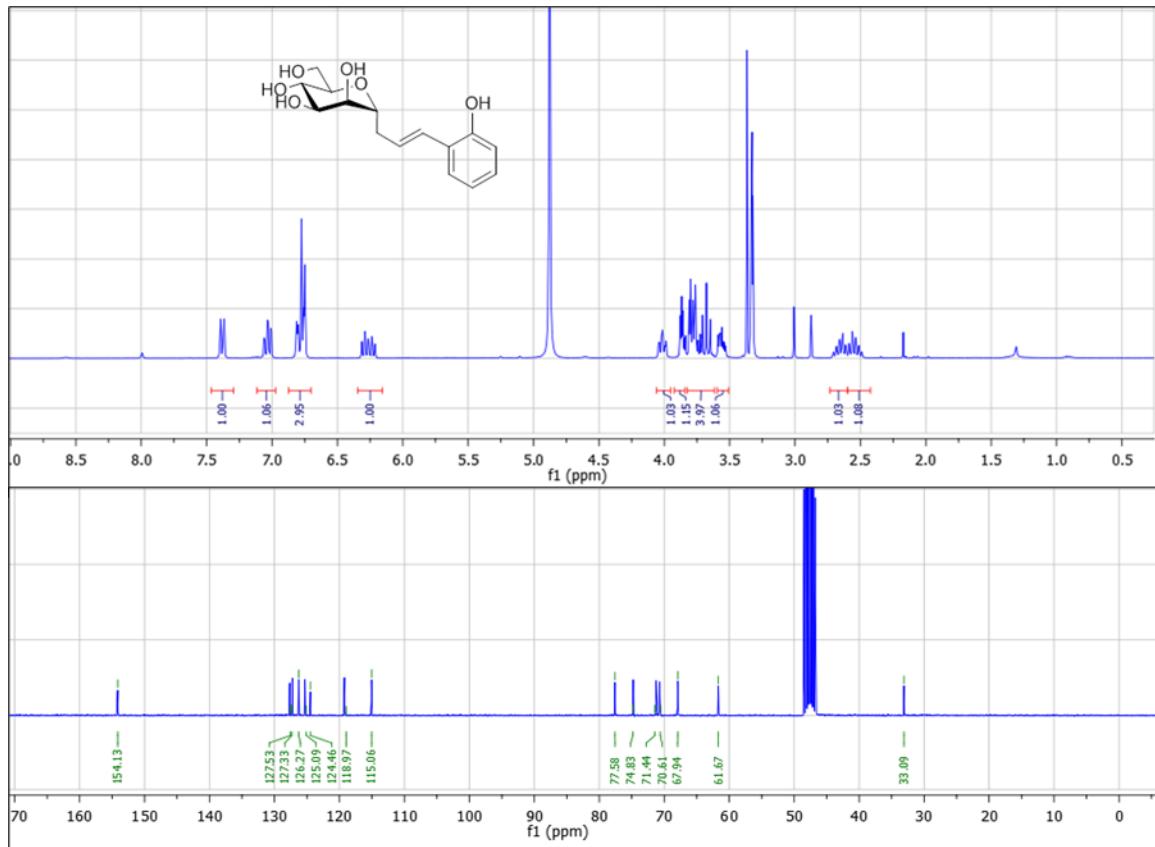


Figure S24. ^1H NMR and ^{13}C NMR spectrum of compound 13(CD_3OD , 300 and 75MHz respectively

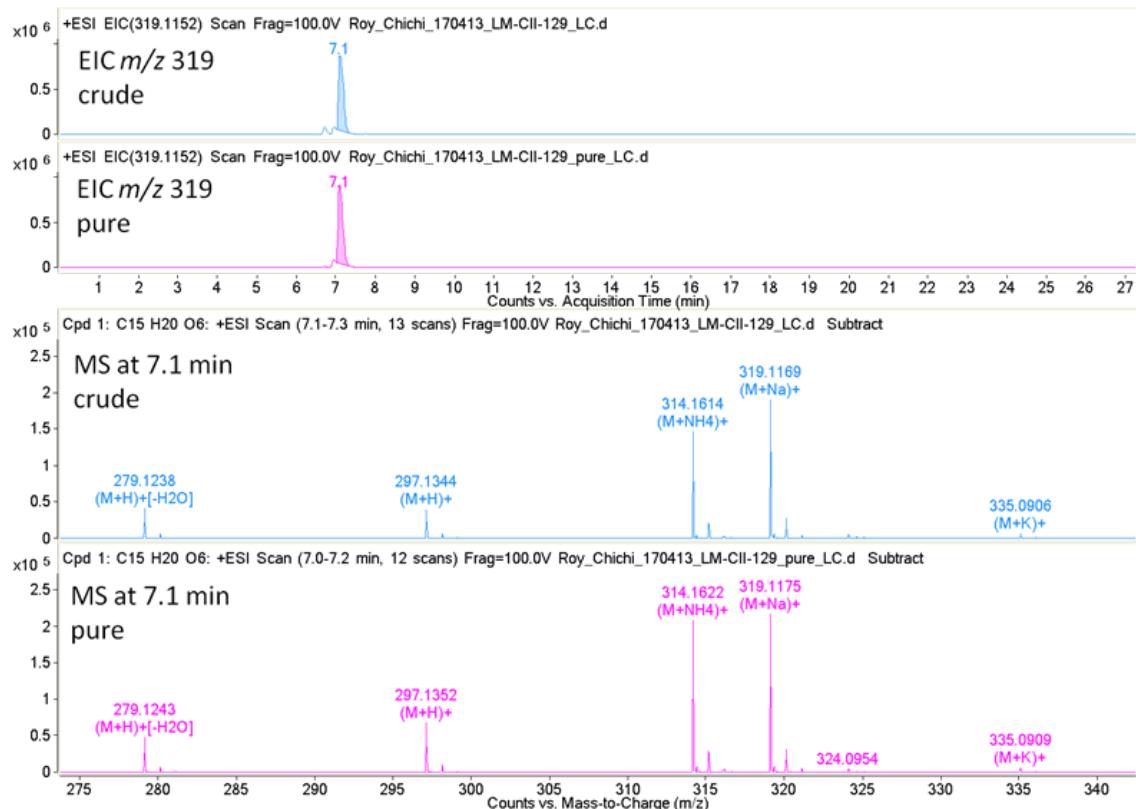


Figure S25. HPLC-TOF-MAS analysis spectrum of compound 13

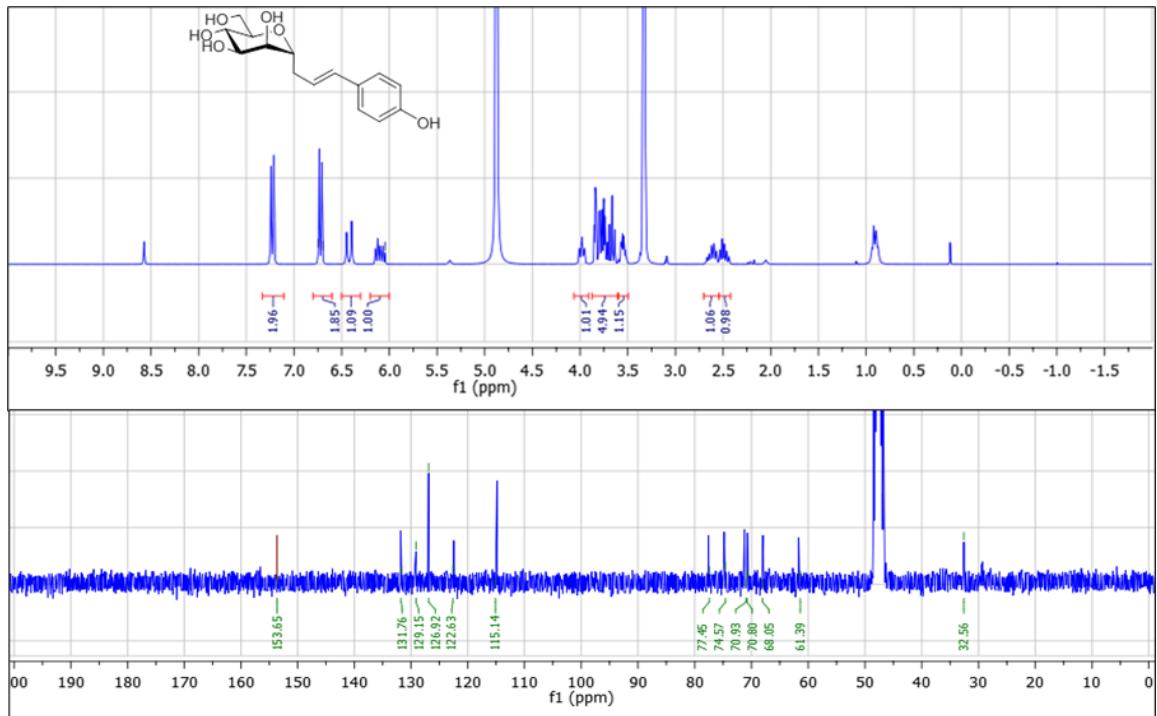


Figure S26. ^1H NMR and ^{13}C NMR spectrum of compound 14 (CD_3OD , 300 and 75MHz respectively)

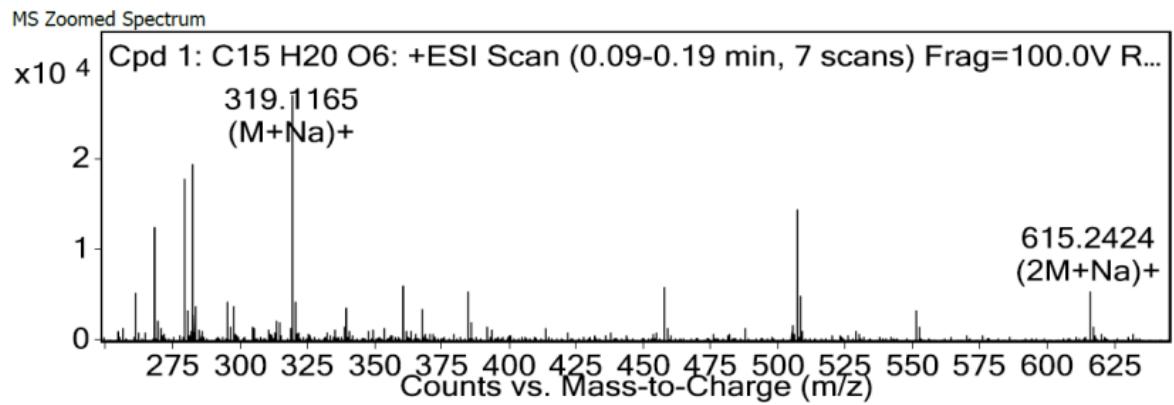


Figure S27. ESI+-HRMS spectrum of compound 14

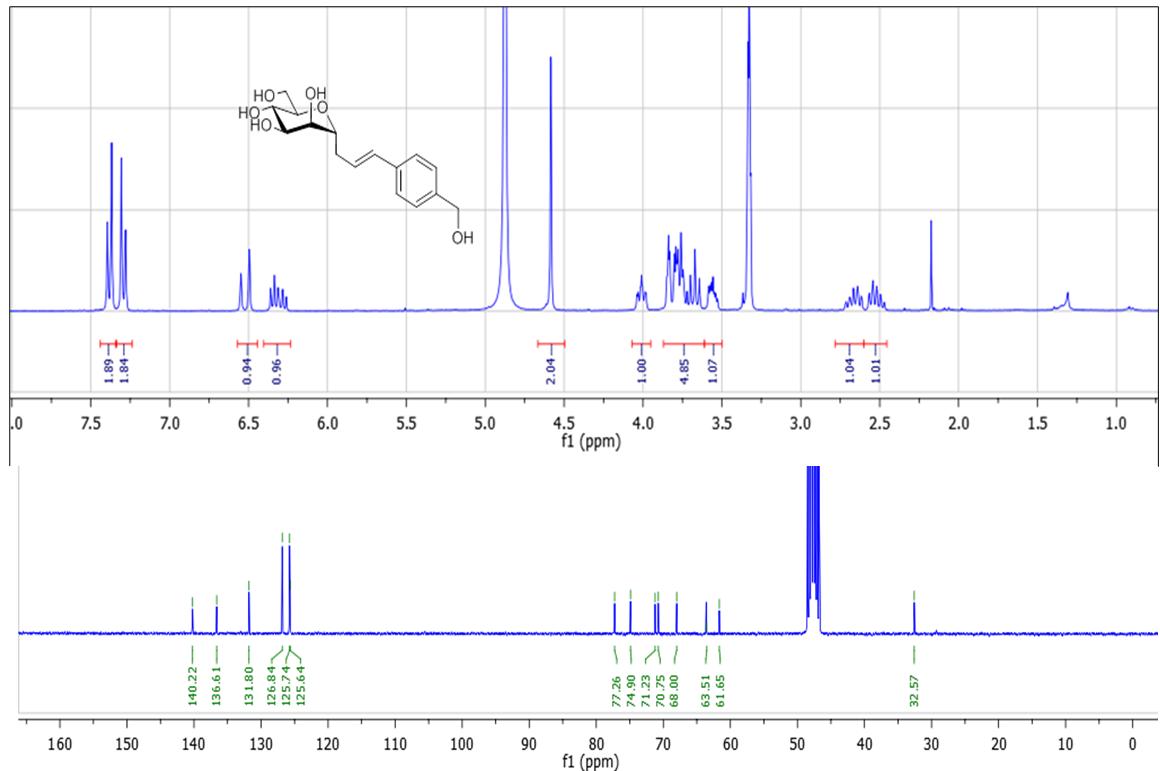


Figure S28. ^1H NMR and ^{13}C NMR spectrum of compound 15 (CD₃OD, 300 and 75MHz respectively)

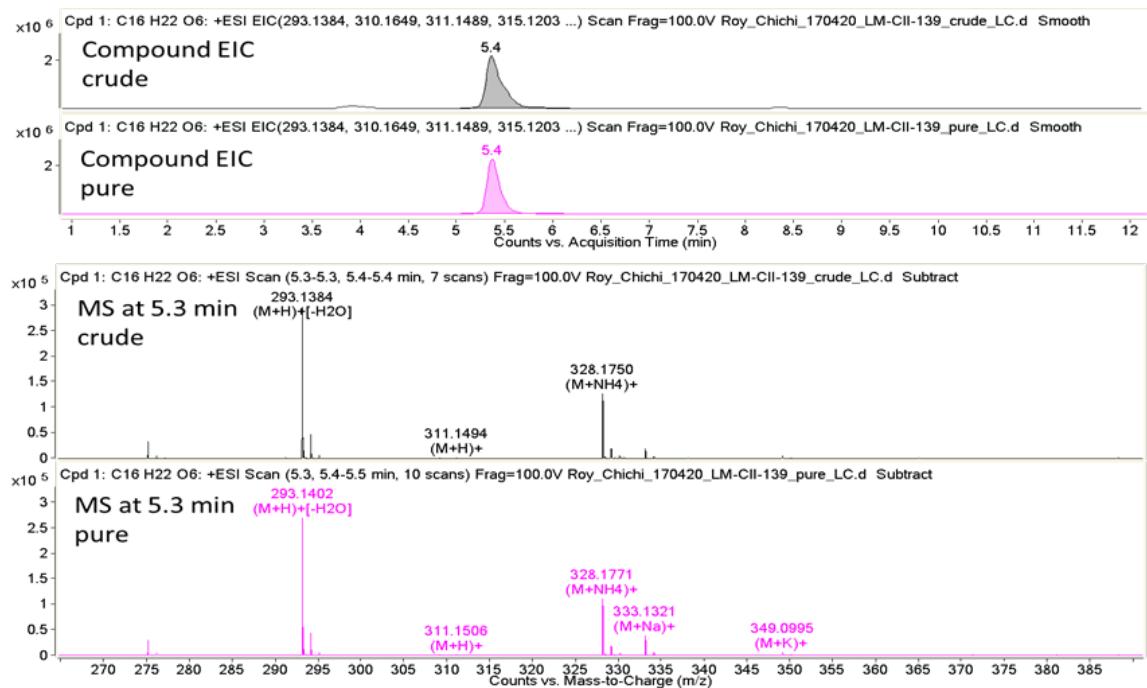


Figure S29. HPLC-TOF-MAS analysis spectrum of compound 15

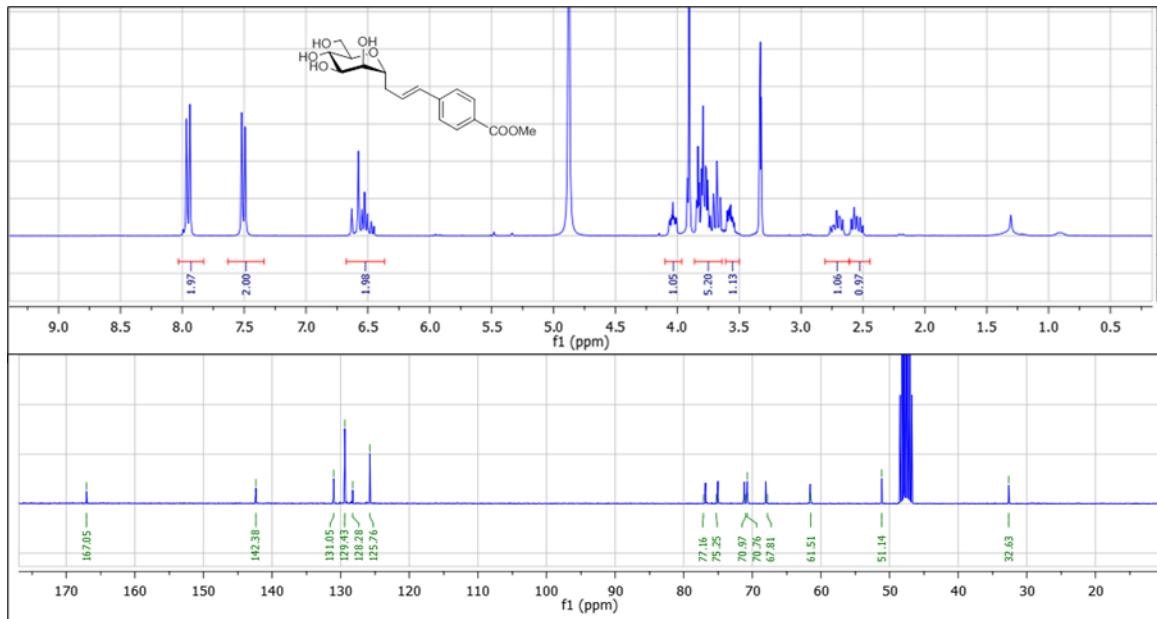


Figure S30. ^1H NMR and ^{13}C NMR spectrum of compound 16(CD_3OD , 300 and 75MHz respectively

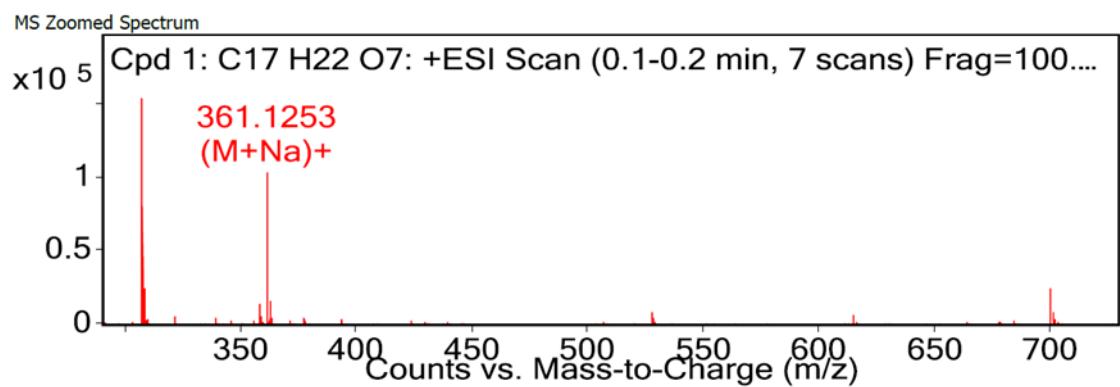


Figure S31. ESI+-HRMS spectrum of compound 16

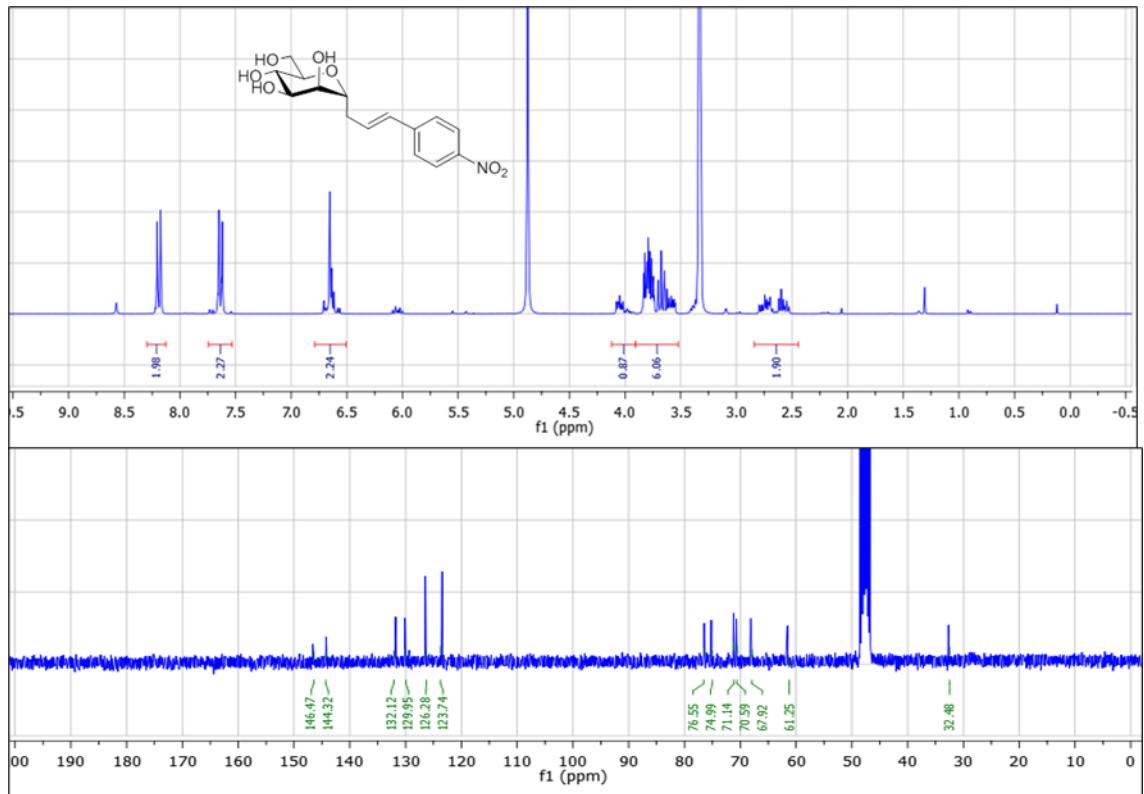


Figure S32. ¹H NMR and ¹³CNMR spectrum of compound 17 (CD₃OD, 300 and 75MHz respectively

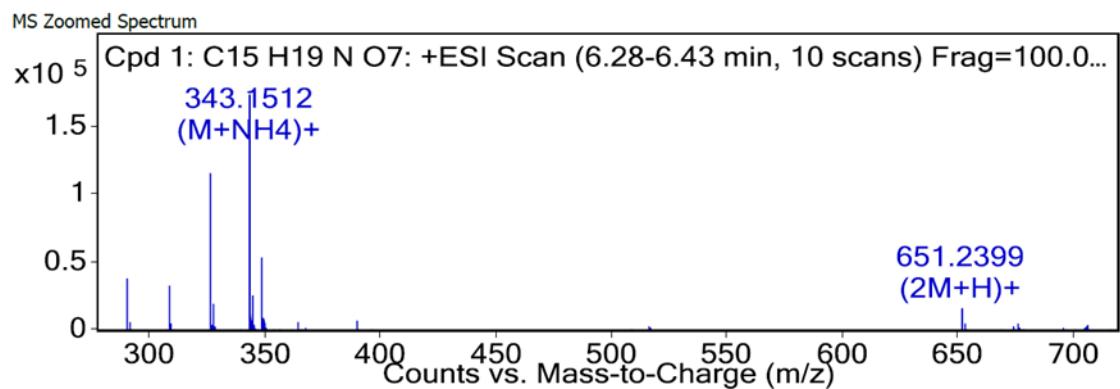


Figure S33. ESI+-HRMS spectrum of compound 17

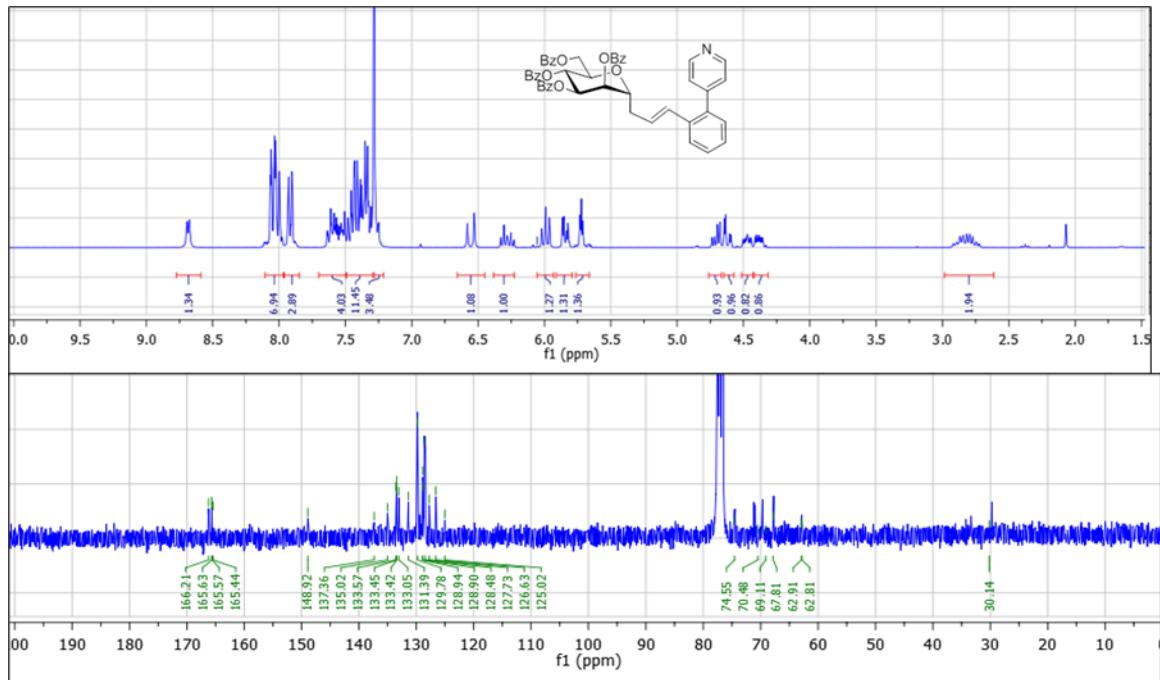


Figure S34. ¹H NMR and ¹³C NMR spectrum of compound 19 (CD₃OD, 300 and 75MHz respectively)

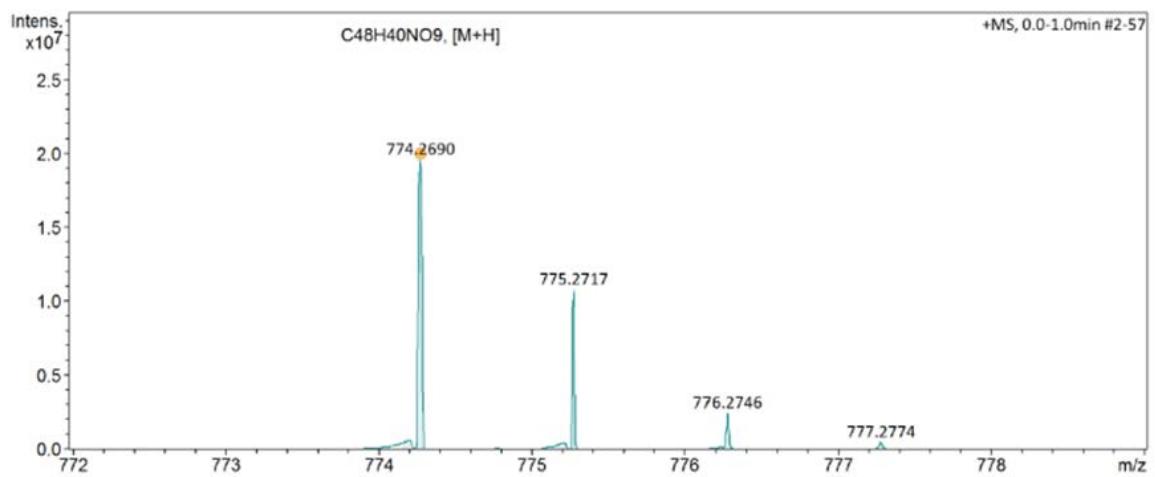


Figure S35. ESI+-HRMS spectrum of compound 19

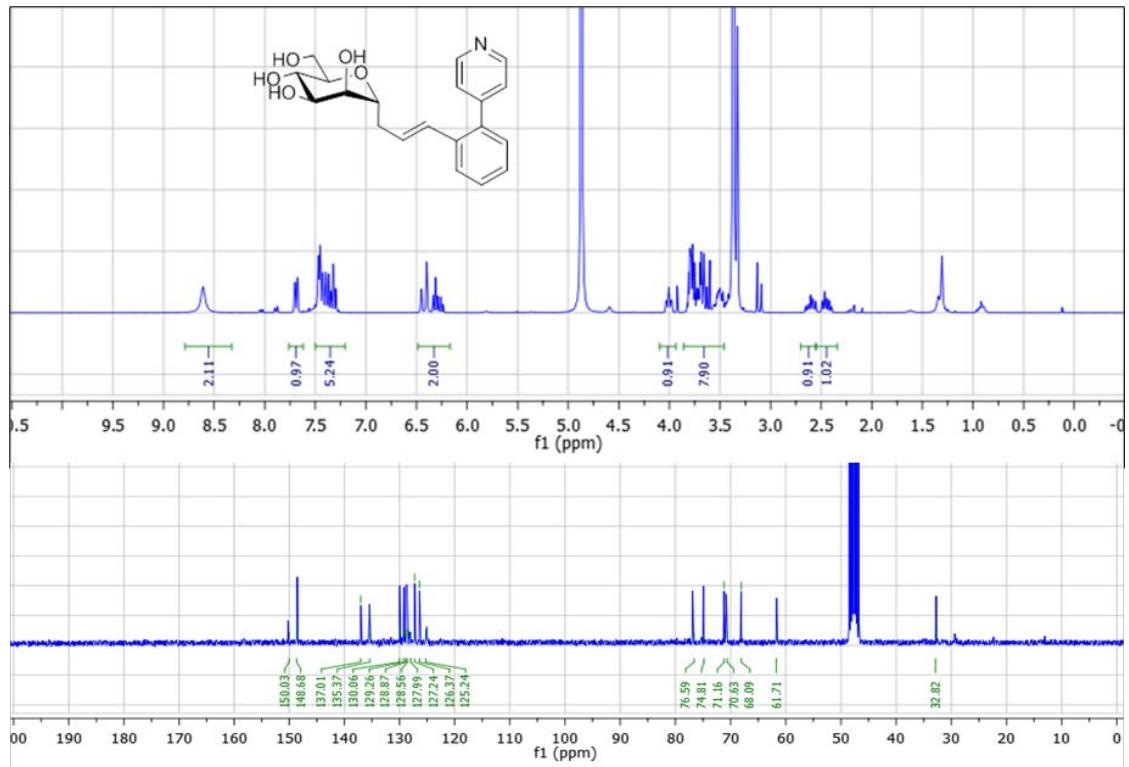
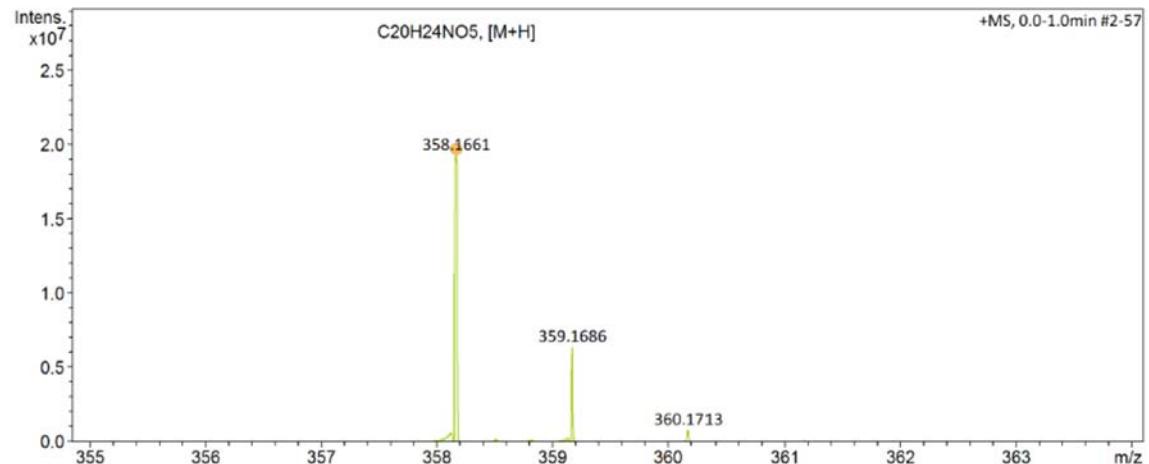


Figure S36. ^1H NMR and ^{13}C NMR spectrum of compound 20 (CD_3OD , 300 and 75MHz respectively)



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Figure S37. ESI+HRMS spectrum of compound 20

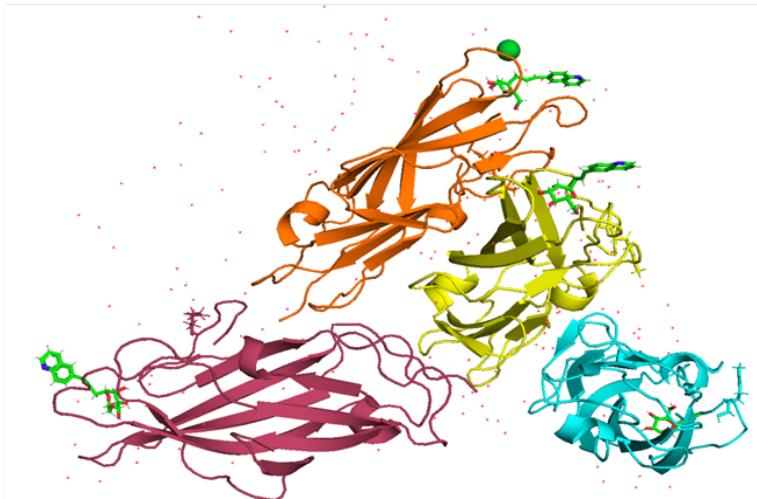


Figure S38. Single crystal of FimH bound with ligand 11

Table S1. X-ray data collection and refinement statistics.

	FimH – (cpd 11)
PDB entry code	8BVD
Resolution range*	49.73 – 3.00 (3.18 - 3.00)
Space group	P 61 2 2
Unit cell constants	a, b = 151.784 Å, c = 225.034 Å α, β = 90.00°, γ = 120.00°
Total number of reflections	1224300 (196951)
Number of unique reflections	31057 (4900)
Multiplicity	39.4 (40.2)
Completeness (%)	98.6 (98.3)
< I/σ(I) >	12.41 (1.37)
Wilson B-factor (Å²)	46.9
R-meas.	0.464 (3.422)
CC1/2	0.996 (0.615)
Resolution range* for refinement	(131.45 – 3.00)
Number of reflections used in refinement	27492 (34)
Reflections used for R-free	1439 (768)
R-work	0.2526 (0.301)
R-free	0.2906 (0.344)
Coordinate error (Å) (maximum-likelihood based)	0.328
Protein residues	632
RMS (bonds)	0.011
RMS (angles)	1.668
Ramachandran favoured (%)	95.03
Ramachandran allowed (%)	5.81
Ramachandran outliers (%)	0.16
Rotamer outliers (%)	0.20
Clash score	11.84
Average B-factor (Å²)	53.00
macromolecules	51.0
ligands	36.5

* Statistics for the highest-resolution shell are shown between parentheses.

	MeaDMan (23)				PNPMan (22)				HM (21)			
	1	2	3	4	5	6	7	8	9	10	11	12
A	200000	781.25			20000		78.13		2000		7.81	
B	100000	390.63			10000		39.06		1000		3.91	
C	50000	195.31			5000		19.53		500		1.95	
D	25000	97.66			2500		9.77		250		0.98	
E	12500	48.83			1250		4.88		125		0.49	
F	6250	24.41			625		2.44		62.5		0.24	
G	3125	12.21			312.5		1.22		31.25		PBS + FimH traceur	
H	1562.5	6.10			156.25		0.61		15.63		PBS seul	

	Inhibitor 11				Inhibitor 20				Inhibitor 18			
	1	2	3	4	5	6	7	8	9	10	11	12
A	614,2		2,4		581,63		2.27		3481.6		13.6	
B	307,2		1,2		290.82		1,14		1740.8		6.8	
C	153,6		0,6		145.41		0.57		870.4		3.4	
D	76,8		0,3		72.70		0.28		435.2		1.7	
E	38,4		0,15		36.35		0.14		217.6		0.85	
F	19,2		0,075		18.18		0.07		108.8		0,43	
G	9,6		0,0375		9.09		0.035		54.4		PBS + FimH traceur	
H	4,8		0,0188		4.54		0,018		27.2		PBS seul	

Figure S39. A perform serial dilutiono f fimH with ligands