

Supplementary Figure 1. ESI/MS/MS analyses of native and de-acetylated S2A. *Panel A*, Positive ESI mass spectra of native (N) and de-acetylated (DA) non-active S2A were obtained, and demonstrated a shift of approximately two acetyl groups (84 amu) when comparing major intact GPL ions (i.e. m/z 1595 vs. m/z 1511, and m/z 1561 vs. m/z 1477). *Panel B*, Major intact GPL ions were selected for collisionally activated dissociation (CAD). At a collision energy of 20 eV, product ions were produced, signifying the GPL ion with a loss of sugar moieties.

Supplementary Figure 2. Partial 1D ^1H NMR spectrum of S2A. Four dominant $-\text{OCH}_3$ signals and two dominant signals from the acetyl CH_3 groups could be observed for native S2A.

Supplementary Figure 3. Positive ESI mass spectra analyses of native and de-acetylated NS3.

When comparing the m/z values for native (N) and de-acetylated (DA) intact GPL ions, a shift of approximately one acetyl group (42 amu) was observed (m/z 1233 vs. m/z 1191, and m/z 1246 vs. m/z 1205). There was another ion (m/z 1219) that was observed in both the N and DA spectra indicating the non-acetylated nsGPL that also comprises NS3. All relevant ions are circled.

Supplementary Figure 4. 1D ^1H NMR spectrum of NS2 in CDCl_3 . *Panel A*, Full spectrum. *Panels B-F*, Expansions of select regions in (A) showing signal assignments for the 6-deoxytalose (dT) and rhamnose (R) residues.

Supplementary Figure 5. 2D ^1H - ^1H NMR spectrum of NS2 in CDCl_3 . *Panel A*, The H1-H2 cross-peaks for the 6-deoxytalose (dT) and the rhamnose (R) were used to initiate signal assignments in each residue (traces shown). *Panel B*, 2D ^1H - ^1H ROESY spectrum of NS2 showing ROEs for H1_R, H1_{dT}, and H4_{dT}. SC denotes correlations to side-chain nuclei. R: rhamnose; dT: 6-deoxytalose.

Assignment of ^1H chemical shifts, anomeric configuration and substitution pattern for native NS2.

The 1D ^1H NMR spectrum of NS2 contains two anomeric proton signals at 5.051 ppm and 4.798 ppm (Supplementary Table 1; Supplementary Fig. 5A, 5B). These signals gave identical integrations, and both were split by a small $^3J_{\text{H1,H2}}$ coupling (1.7 Hz and 1.8 Hz, respectively). Analysis of the 2D ^1H - ^1H COSY spectrum (Supplementary Fig. 6A) provided chemical shift assignments of all non-anomeric protons in each residue (Supplementary Fig. 5B-F), and more detailed interpretation of these signals in the 1D spectrum yielded a complete set of $^3J_{\text{HH}}$ values in each residue (Supplementary Table 2). Absolute configurations of the residues were not confirmed; it is assumed they were in the L-configuration. Anomeric configurations were confirmed by comparing $^3J_{\text{H1,H2}}$ values in the NS2 6-deoxytalose and rhamnose with those observed in standard talo- and mannopyranosyl rings and were in agreement with the α -anomers.

The 1D ^1H NMR spectrum of NS2 contained four intense singlets (Supplementary Fig. 5A, 5C) that gave identical integrations (three protons each), consistent with CH_3 groups. Three of these signals were observed at 3.4-3.5 ppm and were assigned to methyl ether ($-\text{OCH}_3$) protons (Supplementary Table 1). The remaining singlet at 2.177 was assigned to the CH_3 of an acetyl group. Analysis of the H2_{dT} and H2_R multiplets revealed the presence of $^3J_{\text{HCOH}}$ couplings in both cases (7.9 Hz, and 2.3 Hz, respectively; CDCl_3 solvent), showing that OH2 in both residues is unsubstituted (Supplementary Table 2). Since both residues are 6-deoxyhexopyranosyl rings, the remaining four OH groups (OH3 and OH4 of both residues) must therefore be substituted, since four CH_3 signals were observed.

To determine the substitution pattern, two approaches were taken. ^1H chemical shift differences were calculated for the NS2 6-deoxytalose and rhamnose residues, and for standard unsubstituted talo- and mannopyranosyl residues, using H1 as the reference signal. Since OH2 in both residues is unsubstituted, $\delta_{\text{H1}}-\delta_{\text{H2}}$ serves as a reasonable internal control, which is confirmed by the good agreement

observed between the NS2 6-deoxytalose and the α -talo standard, and between the NS2 rhamnose and the α -manno standard (Supplementary Table 3) (these differences also confirm the anomeric configuration of the NS2 6-deoxytalose and rhamnose). Values of $\delta_{H1}-\delta_{H3}$ and $\delta_{H1}-\delta_{H4}$ in the NS2 6-deoxytalose and rhamnose are *more positive* in three cases than the *positive* difference calculated in the reference compound, whereas in one case ($\delta_{H1}-\delta_{H4}$ of the NS2 6-deoxytalose), the difference is *negative* compared to a *positive* difference calculated in the reference. These results support the conclusion that OH3 and OH4 of NS2 rhamnose, and OH3 of NS2 6-deoxytalose are substituted as methyl ethers, and the remaining site, OH4 of 6-deoxytalose, is acetylated. *O*-Methylation of a CH-OH fragment thus induces a modest upfield shift of the CH proton, whereas *O*-acetylation induces a significant downfield shift.

The above substitution pattern was supported by an analysis of the ^1H - ^1H 2D ROESY spectrum of NS2 (Supplementary Fig. 6B). Inspection of the H4_{dT} cross-peaks revealed strong correlations to H3 and H5, a moderate correlation to the most upfield ether $-\text{OCH}_3$ (presumably at O3), and a very weak correlation to the acetyl CH_3 . This pattern is consistent with the proposed substitutions in the NS2 6-deoxytalose. In this interpretation, it is assumed that the acetyl CH_3 is, on average, relatively far removed from H4 due to conformation about the C4-O4 bond, and that conformation about the C3-O3 bond results in an average orientation of the CH_3 that places it relatively close to H4. The ROESY data also confirm free $-\text{OH}$ groups in the NS2 6-deoxytalose and rhamnose, since H1 in both residues exhibits an NOE to the corresponding OH2 proton.

Supplementary Table 1

Table 1. ^1H chemical shifts of sample NS2

residue	chemical shift (ppm)									
	H1	H2	H3	H4	H5	H6	OH2	CH ₃ (1) (ether)	CH ₃ (2) (ether)	CH ₃ (acetyl)
deoxyTalo (dT)	5.051	3.802	3.562	~5.307	3.950	1.158	2.790	3.418		2.177
Rhamno (R)	4.798	4.002	3.452	3.057	3.619	~1.26	2.633	3.525	3.468	

^aIn CDCl₃; 22 °C; chemical shifts referenced to the internal residual CHCl₃ signal (7.270 ppm).

Supplementary Table 2

Table 2. ^1H - ^1H spin-couplings in sample NS2

residue	^1H - ^1H spin coupling (Hz) ^a						
	$^3J_{\text{H1,H2}}$	$^3J_{\text{H2,H3}}$	$^3J_{\text{H3,H4}}$	$^3J_{\text{H4,H5}}$	$^3J_{\text{H5,H6}}$	$^4J_{\text{H2,H4}}$	$^3J_{\text{H2,OH2}}$
deoxyTalo (dT)	1.7	~3.1	~3.1	~1.2	5.7	1.2	7.9
Rhamno (R)	1.8	3.4	9.1	~9.5	5.8		2.3
α -Talo ^b	1.9	3.2	3.2	1.3		1.4	
β -Talo ^b	1.2	3.2	3.3	1.2		1.2	
α -Talo ^c	~1.9	3.4	~3.5	~1.3			
α -Manno ^c	1.8	3.5	9.5				
β -Manno ^c	0.9	3.2	9.6				

^aIn Hz \pm 0.1 unless otherwise noted; in CDCl₃; 22° C. ^bData for α - and β -D-talopyranoses taken from Ref. 1. ^cData for methyl α -D-talopyranosides and methyl α - and β -D-mannopyranosides, taken from Ref. 2.

Ref. 1: Snyder, J. R., Johnston, E. R., and Serianni, A. S. (1989) *J Am Chem Soc* **111**, 2681-2687

Ref. 2: Podlasek, C. A., Wu, J., Stripe, W. A., Bondo, P. B., and Serianni, A. S. (1995) *J Am Chem Soc* **117**, 8635-8644

Supplementary Table 3

Table 3. ^1H chemical shift differences for residues 6-deoxytalose (dT) and rhamnose (R) in NS2, and for unsubstituted manno- and talopyranosyl rings.

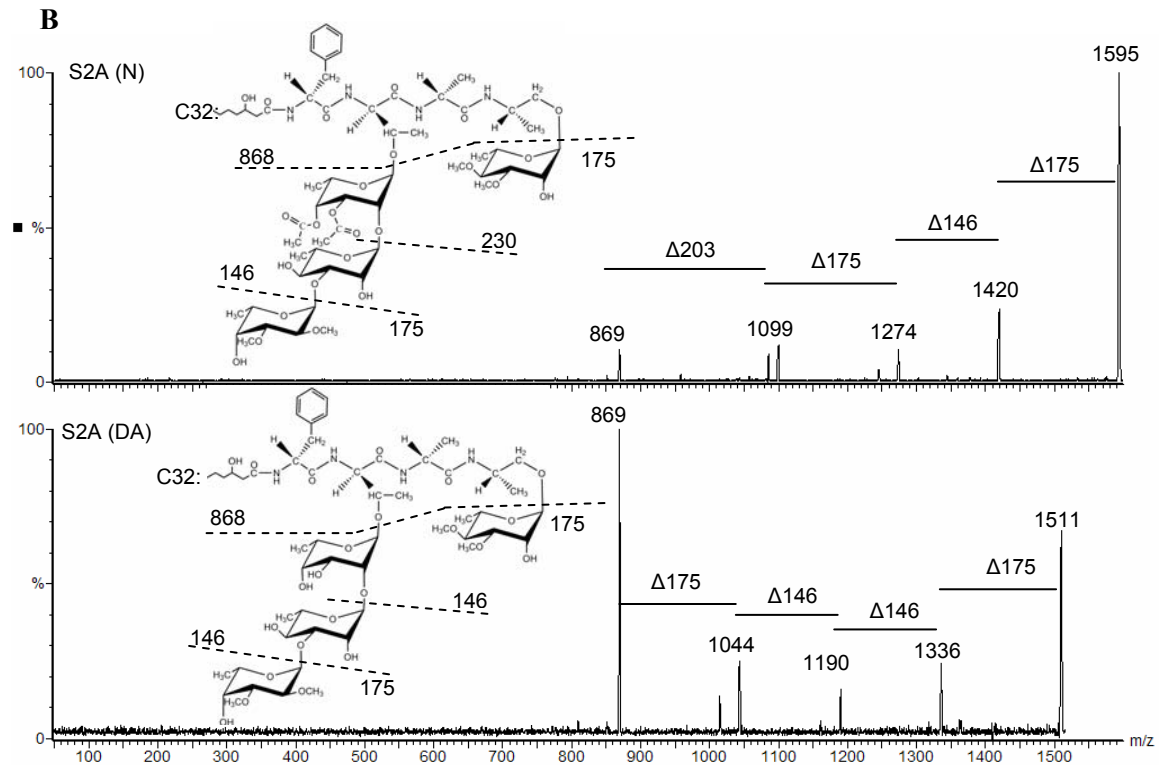
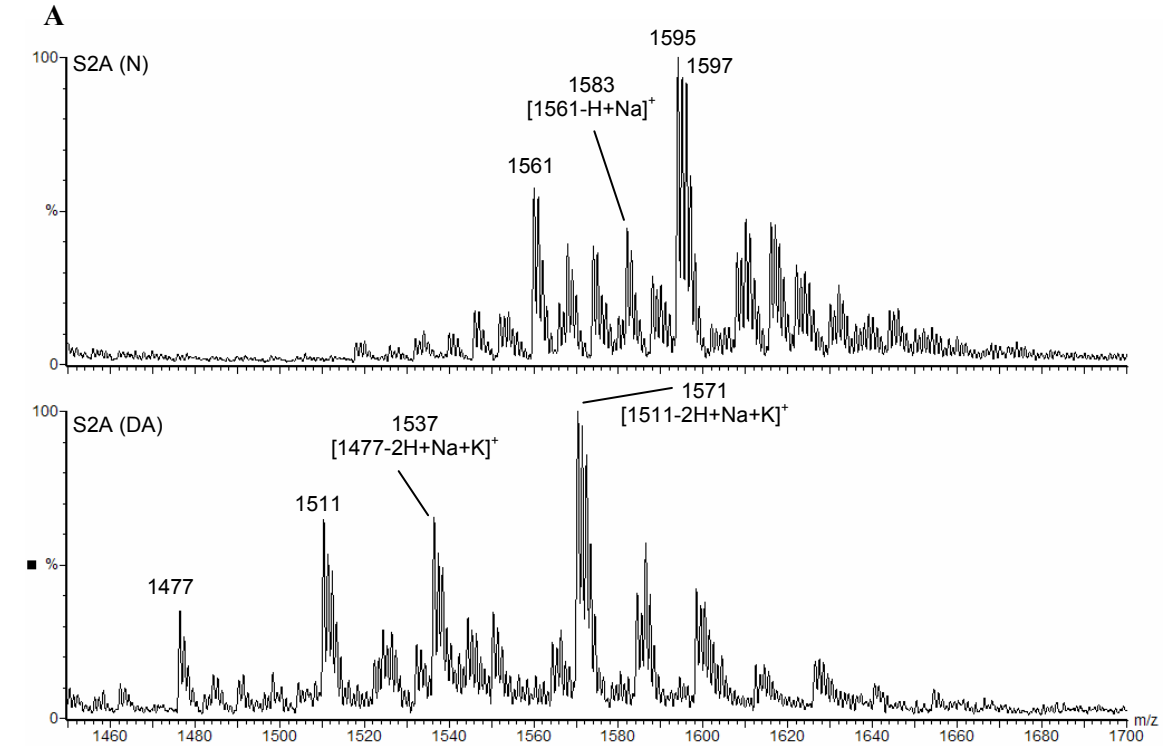
residue	^1H chemical shift difference (ppm)		
	$\delta\text{H1}-\delta\text{H2}$	$\delta\text{H1}-\delta\text{H3}$	$\delta\text{H1}-\delta\text{H4}$
dT	1.25	1.49	-0.26
α -Talo ^a	1.41	1.32	~1.34
β -Talo ^a	0.87	1.00	~-0.95
R	0.80	1.35	1.74
α -Manno ^b	0.83	1.00	1.12
β -Manno ^b	0.59	0.94	1.01

^aData for α - and β -D-talopyranoses taken from Ref. 1. ^bData for methyl α - and β -D-mannopyranosides taken from Ref. 2

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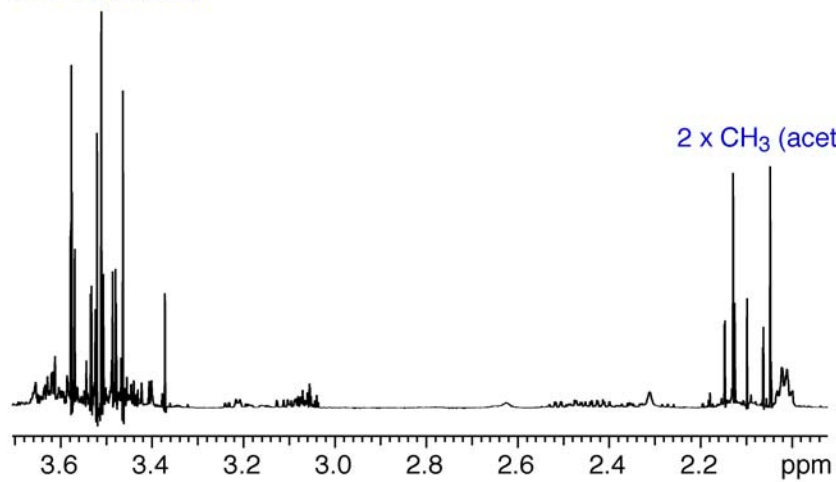
Ref. 2: Podlasek, C. A., Wu, J., Stripe, W. A., Bondo, P. B., and Serianni, A. S. (1995) *J Am Chem Soc* **117**, 8635-8644

Supplementary Figure 1

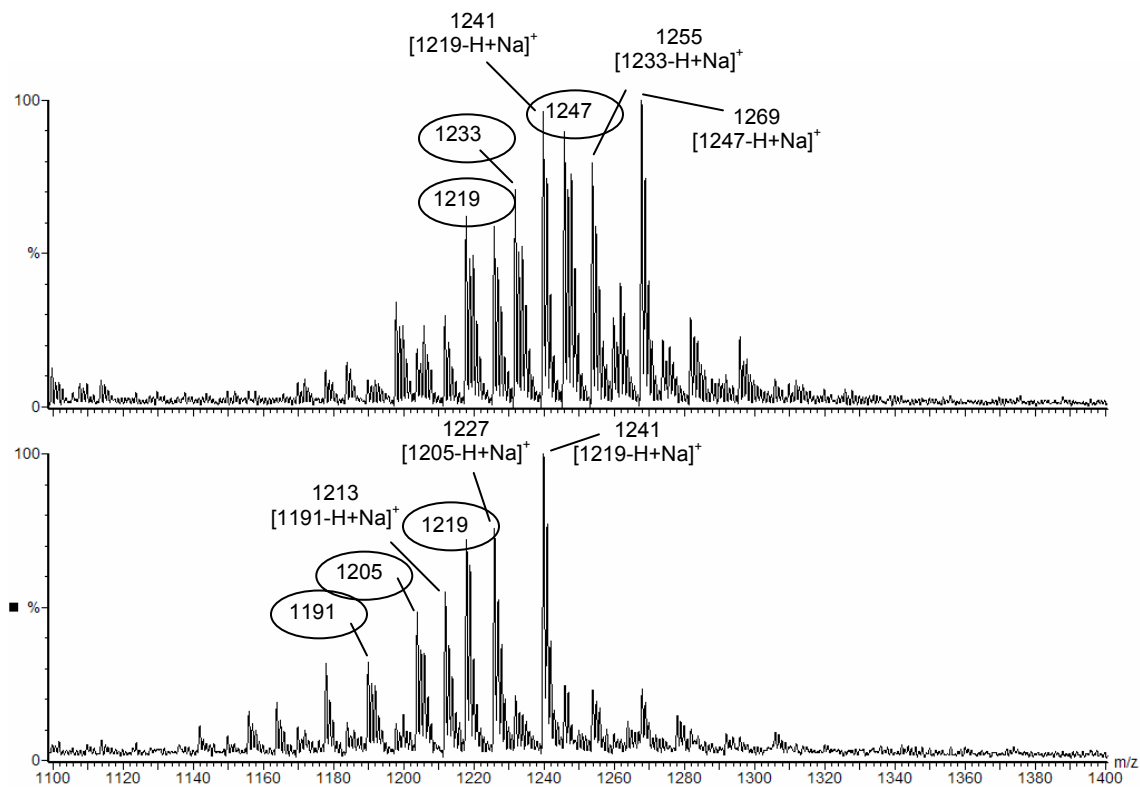


Supplementary Figure 2

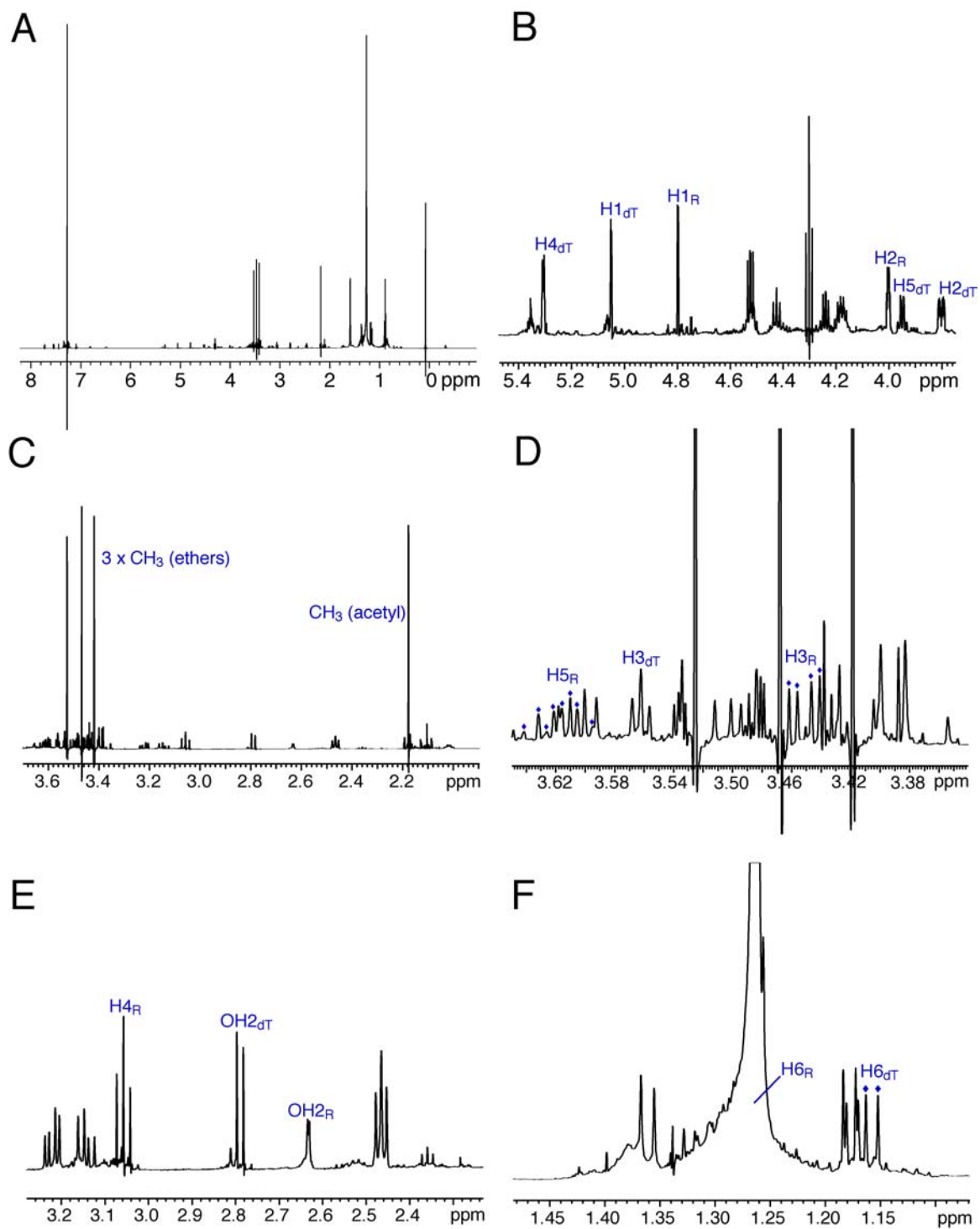
4 x CH₃ (ethers)



Supplementary Figure 3

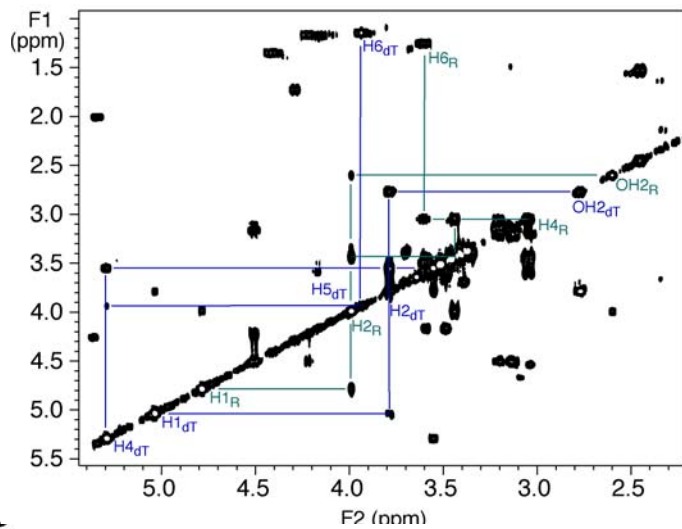


Supplementary Figure 4



Supplementary Figure 5

A



B

