Typical Preparation of *Francisella tularensis O*-antigen Yields a Mixture of Three Types of Saccharides[†]

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Supporting Information

Supplemental Table 1 Observed fragment ions for [4,2,2][0,0,0] from multi-stage tandem

Product ion	Charge	Saccharide	Mass	Fragmentation				
(m/z)	_	Composition ¹	Error	_				
			(ppm)					
MS ² Precursor ion	MS ² Precursor ion <i>m/z</i> 824.2961 (2+)							
412.1321	1	[1,0,1][0,0,0]º+Na⁺	-1.4	B ₂				
426.1476	1	[1,1,0][0,0,0]º+Na⁺	-1.7	Z ₂				
430.1426	1	[1,0,1][0,0,0] +Na ⁺	-1.5	C ₂				
444.1581	1	[1,1,0][0,0,0] +Na ⁺	-1.8	Y ₂				
455.1378	1	[2,0,0][0,0,0]º+Na⁺	-1.5	Y ₇ /B ₃ ; Z ₃ /C ₇ ; Y ₃ /B ₇ ; Z ₇ /C ₃				
473.1482	1	[2,0,0][0,0,0] +Na ⁺	-1.8	Y ₇ /C ₃ ; Y ₃ /C ₇				
628.2063	1	[2,0,1][0,0,0]º+Na+	-1.6	B ₃				
629.7244	2	[3,2,1][0,0,0] +2Na ⁺	-1.5	Y ₆				
635.2143	2	[4,1,1][0,0,0]º+2Na⁺	-1.3	Y ₇ /B ₇ ; Z ₇ /C ₇				
642.2221	1	[2,1,0][0,0,0]º+Na⁺	-1.3	Z ₃				
644.2194	2	[4,1,1][0,0,0] +2Na ⁺	-1.6	Y ₇ /C ₇				
646.2168	1	[2,0,1][0,0,0] +Na ⁺	-1.7	C ₃				
660.2324	1	[2,1,0][0,0,0] +Na ⁺	-1.7	Y ₃				
721.7491	2	[4,1,2][0,0,0] ^o +2Na ⁺	-0.6	B ₇				
730.7540	2	[4,1,2][0,0,0] +2Na ⁺	-1.1	C ₇				
737.7610	2	[4,2,1][0,0,0] +2Na ⁺	-2.2	Y ₇				
				B ₄ ; Z ₄ ; Y ₇ /B ₅ ; Z ₇ /C ₅ ; Y ₆ /B ₆ ; Z ₆ /C ₆ ;				
815.2906	1	[2,1,1][0,0,0]º+Na⁺	-1.4	Y ₅ /B ₇ ; Z ₅ /C ₇				
833.3009	1	[2,1,1][0,0,0] +Na ⁺	-1.7	C ₄ ; Y ₄ ; Y ₇ /C ₅ ; Y ₆ /C ₆ ; Y ₅ /C ₇				
988.3593	1	[2,1,2][0,0,0] ^o +Na ⁺	-1.3	B ₅				
1002.3752	1	[2,2,1][0,0,0] ^o +Na ⁺	-1.0	Z ₅				
1020.3858	1	[2,2,1][0,0,0] +Na ⁺	-1.0	Y ₅				
1031.3651	1	[3,1,1][0,0,0]º+Na+	-1.2	Y ₇ /B ₆ ; Z ₇ /C ₆ ; Y ₆ /B ₇ ; Z ₆ /C ₇				
1049.3757	1	[3,1,1][0,0,0] +Na ⁺	-1.2	Y ₇ /C ₆ ; Y ₆ /C ₇				
1204.4340	1	[3,1,2][0,0,0]º+Na+	-1.0	B ₆				
1218.4495	1	[3,2,1][0,0,0]º+Na+	-1.1	Z ₆				
1222.4445	1	[3,1,2][0,0,0] +Na ⁺	-1.0	C_6				
1236.4599	1	[3,2,1][0,0,0] +Na ⁺	-1.2	Y ₆				
1438.5188	1	[4,1,2][0,0,0] +Na ⁺	-1.1	C ₇				
1452.5338	1	[4,2,1][0,0,0] +Na ⁺	-1.5	Y ₇				
1625.6023	1	[4,2,2][0,0,0] +Na ⁺	-1.6	[M+Na]⁺				
MS ³ Precursor ion	<i>m/z</i> 833.3	009 (1+), to simplify th	e nomencla	ature of the MS^3 fragment ions, Y_4				
is assumed to be t	he precurs	sor ion of MS ³ unless s	pecified.					
239.0634	1	[1,0,0][0,0,0]º+Na+	-2.0	Y ₃ /B ₆ ; Z ₃ /C ₆ ; Y ₂ /B ₇ ; Z ₂ /C ₇				
257.0738	1	[1,0,0][0,0,0] +Na ⁺	-2.6	Y ₃ /C ₆ ; Y ₂ /C ₇				
412.1316	1	[1,0,1][0,0,0]⁰+Na⁺	-2.7	Y ₄ /B ₆				
426.1472	1	[1,1,0][0,0,0]⁰+Na⁺	-2.7	Z ₂				
444.1576	1	[1,1,0][0,0,0] +Na ⁺	-3.0	Y ₂				
455.1372	1	[2,0,0][0,0,0]⁰+Na⁺	-2.9	Y ₃ /B ₇ ; Z ₃ /C ₇				
473.1476	1	[2,0,0][0,0,0] +Na ⁺	-3.2	Y ₃ /C ₇				
545.1685	1		-3.1	$^{0,2}A_3$ from C ₄				

MS in positive ion mode on an LTQ-Orbitrap XL mass spectrometer.

				Y_{6}/B_{5} ; Z_{6}/C_{5} from Y_{7}/C_{5} ; Y_{6}/C_{6} or
599.2156	1	[1,1,1][0,0,0]⁰+Na+	-2.6	Y_5/B_6 ; Z_5/C_6 from Y_6/C_6 ; Y_5/C_7
628.2053	1	[2,0,1][0,0,0]º+Na+	-3.3	Y ₄ /B ₇
642.2210	1	[2,1,0][0,0,0]º+Na+	-3.1	Z ₃
646.2158	1	[2,0,1][0,0,0] +Na ⁺	-3.3	Y ₄ /C ₇
660.2314	1	[2,1,0][0,0,0] +Na ⁺	-3.3	Y ₃
815.2895	1	[2,1,1][0,0,0]º+Na+	-2.8	Z ₄
MS ⁴ Precursor ion	<i>m/z</i> 646.2	158 (1+)		
239.0632	1	[1,0,0][0,0,0]º+Na⁺	-3.0	Y ₃ /B ₆ ; Z ₃ /C ₆ ; Y ₂ /B ₇ ; Z ₂ /C ₇
257.0739	1	[1,0,0][0,0,0] +Na ⁺	-2.2	Y ₃ /C ₆ ; Y ₂ /C ₇
412.1316	1	[1,0,1][0,0,0]º+Na⁺	-2.7	Y ₄ /B ₆ ; Z ₄ /C ₆
430.1421	1	[1,0,1][0,0,0] +Na⁺	-2.7	Y ₄ /C ₆
455.1372	1	[2,0,0][0,0,0]º+Na+	-2.9	Y ₃ /B ₇ ; Z ₃ /C ₇
473.1475	1	[2,0,0][0,0,0] +Na ⁺	-3.4	Y ₃ /C ₇
545.1684	1		-3.3	^{0,2} A ₃ from C ₄
628.2055	1	[2,0,1][0,0,0]º+Na+	-2.9	Y ₄ /B ₇ ; Z ₄ /C ₇
MS ⁴ Precursor ion	<i>m/z</i> 660.2	314 (1+)		
228.0836	1	[0,1,0][0,0,0] +Na ⁺	-3.1	Y ₁
239.0632	1	[1,0,0][0,0,0]º+Na+	-3	Y ₃ /B ₆ ; Z ₃ /C ₆ ; Y ₂ /B ₇ ; Z ₂ /C ₇
257.0738	1	[1,0,0][0,0,0] +Na ⁺	-2.6	Y ₃ /C ₆ ; Y ₂ /C ₇
426.1471	1	[1,1,0][0,0,0]º+Na⁺	-3.0	Z ₂
444.1575	1	[1,1,0][0,0,0] +Na⁺	-3.2	Y ₂
455.1371	1	[2,0,0][0,0,0] ^o +Na ⁺	-3.2	Y ₃ /B ₇ ; Z ₃ /C ₇
473.1474	1	[2,0,0][0,0,0] +Na ⁺	-3.6	Y ₃ /C ₇
642.2208	1	[2,1,0][0,0,0] ^o +Na ⁺	-3.4	Z ₃

¹Saccharide compositions are given as [GalNAcAN, QuiNAc, Qui4NFm] [HexNAc, Hex, Kdo].

 $^{\underline{o}}$ represents loss of water.

Supplemental Table 2 Observed fragment ions for ¹⁸O labeled [4,2,2][0,0,0] from multi-

Product ion	Charge	Saccharide Composition ¹	Mass	Fragmentation		
(m/z)			Error			
			(ppm)			
MS ² Precursor ion	<i>m/z</i> 825.2	987 (2+)				
428.1519	1	[1,1,0][0,0,0] ^o + ¹⁸ O- ¹⁶ O+Na ⁺	-1.6	Z ₂		
646.2168	1	[2,0,1][0,0,0]+Na⁺	-1.7	C ₃		
662.2367	1	[2,1,0][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.6	Y ₃		
730.7539	2	[4,1,2][0,0,0]+2Na ⁺	-1.2	C ₇		
738.7635	2	[4,2,1][0,0,0]+ ¹⁸ O- ¹⁶ O+2Na ⁺	-1.7	Y ₇		
				B ₄ ; Y ₇ /B ₅ ; Z ₇ /C ₅ ; Y ₆ /B ₆ ;		
815.2906	1	[2,1,1][0,0,0]º+Na⁺	-1.4	Z ₆ /C ₆ ; Y ₅ /B ₇ ; Z ₅ /C ₇		
835.3053	1	[2,1,1][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.5	Y ₄		
988.3594	1	[2,1,2][0,0,0]º+Na⁺	-1.2	B_5		
1004.3794	1	[2,2,1][0,0,0] ^º + ¹⁸ O- ¹⁶ O+Na⁺	-1.1	Z ₅		
1022.3901	1	[2,2,1][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-0.9	Y ₅		
1049.3758	1	[3,1,1][0,0,0]+Na⁺	-1.1	Y ₇ /C ₆ ; Y ₆ /C ₇		
1204.4342	1	[3,1,2][0,0,0]º+Na⁺	-0.8	B_6		
1222.4447	1	[3,1,2][0,0,0]+Na⁺	-0.9	C_6		
1238.4646	1	[3,2,1][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-0.8	Y ₆		
1438.5191	1	[4,1,2][0,0,0]+Na⁺	-0.9	C ₇		
1627.6078	1	[4,2,2][0,0,0]+ ¹⁸ O- ¹⁶ O+2Na ⁺	-0.8	[M+Na]⁺		
MS ³ Precursor ion <i>m/z</i> 835.3053 (1+)						
412.1317	1	[1,0,1][0,0,0]º+Na⁺	-2.4	Y_4/B_6		
428.1516	1	[1,1,0][0,0,0]º+ ¹⁸ O- ¹⁶ O+Na⁺	-2.3	Z ₂		
430.1422	1	[1,0,1][0,0,0] +Na⁺	-2.5	Y ₄ /C ₆		
446.1621	1	[1,1,0][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-2.4	Y ₂		
473.1479	1	[2,0,0][0,0,0] +Na ⁺	-2.5	Y ₃ /C ₇		
628.2057	1	[2,0,1][0,0,0]º+Na⁺	-2.6	Y ₄ /B ₇ ; Z ₄ /C ₇		
646.2163	1	[2,0,1][0,0,0]+Na ⁺	-2.5	Y ₄ /C ₇		
662.2362	1	[2,1,0][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-2.4	Y ₃		

stage tandem MS in positive ion mode on an LTQ-Orbitrap XL mass spectrometer.

¹Saccharide compositions are given as [GalNAcAN, QuiNAc, Qui4NFm] [HexNAc, Hex, Kdo].

^o represents loss of water.

Supplemental Table 3 Observed fragment ions for [4,2,2][0,0,0] and [6,3,3][0,0,0] from

Observed	Charge	Saccharide	Mass	Fragmentation
Peaks (m/z)	U	Composition ¹	Accuracy	C
. ,		•	(ppm)	
MS ² of Precurs	sor ion <i>m/z</i>	846.3049 (2-)		
215.0674	1	[1,0,0][0,0,0] ^o -[H ⁺]	0.3	Y ₇ /B ₂ ; Z ₇ /C ₂ ; Y ₆ /B ₃ ; Z ₆ /C ₃ ;
				Y ₃ /B ₆ ; Z ₃ /C ₆ ; Y ₂ /B ₇ ; Z ₂ /C ₇
232.0827	1	[0,1,0][0,0,0] ^{oַ,**} -[H ⁺]	0.2	Z ₁
236.0776	1	[0,0,1][0,0,0]** -[H ⁺]	0.1	C ₁
305.0991	1		0.2	^{0,2} A ₂
402.1518	1	[1,1,0][0,0,0]º-[H⁺]	0.0	Z ₂
431.1419	1	[2,0,0][0,0,0]º-[H⁺]	-0.2	Y ₇ /B ₃ ; Z ₇ /C ₃ ; Y ₃ /B ₇ ; Z ₃ /C ₇
448.1572	1	[1,1,0][0,0,0] ^{o,**} -[H ⁺]	-0.2	Z ₂
575.2204	1	[1,1,1][0,0,0]º-[H⁺]	-0.4	Y ₆ /B ₅ ; Z ₆ /C ₅ ; Y ₅ /B ₆ ; Z ₅ /C ₆
618.2261	1	[2,1,0][0,0,0]º-[H⁺]	-0.5	Z ₃
656.2338	2		-0.9	^{0,2} A ₇
664.2314	1	[2,1,0][0,0,0] ^{0,**} -[H ⁺]	-0.8	Z ₃
706.7577	2	[4,1,2][0,0,0] -2[H ⁺]	-0.8	C ₇
727.7631	2	[4,2,1][0,0,0] ^{o,**} -2[H ⁺]	-0.6	Z ₇
729.7605	2	[4,1,2][0,0,0]** -2[H ⁺]	-0.7	C ₇
791.2946	1	[2,1,1][0,0,0]º-[H⁺]	-0.8	B ₄ ; Z ₄
800.2995	2	[4,2,2][0,0,0] -2[H ⁺]	-1.3	[M-2H] ²⁻
823.3025	2	[4,2,2][0,0,0]** -2[H ⁺]	-0.9	[M+FA-2H] ²⁻
846.3053	2	[4,2,2][0,0,0] ² **-2[H ⁺]	-0.8	[M+2FA-2H] ²⁻
982.3734	1	[2,1,2][0,0,0] -[H⁺]	-1.2	C_5
1007.3688	1	[3,1,1][0,0,0]º-[H⁺]	-1.0	Y ₇ /B ₆ ; Z ₇ /C ₆ ; Y ₆ /B ₇ ; Z ₆ /C ₇
1024.3839	1		-1.2	^{2,4} A ₆
1028.3788	1	[2,1,2][0,0,0]** -[H ⁺]	-1.2	C_5
1097.4002	1		-1.2	^{0,2} A ₆
1180.4371	1	[3,1,2][0,0,0]º-[H⁺]	-1.3	B_6
1198.4477	1	[3,1,2][0,0,0] -[H⁺]	-1.3	C_6
1223.4428	1	[4,1,1][0,0,0]º-[H⁺]	-1.4	Y ₇ /B ₇ ; Z ₇ /C ₇
1240.4581	1		-1.4	^{2,4} A ₇
1244.4529	1	[3,1,2][0,0,0]** -[H ⁺]	-1.5	C_6
1295.4640	1		-1.2	^{0,2} A ₇ -H ₂ O
1313.4740	1		-1.6	^{0,2} A ₇
1396.5116	1	[4,1,2][0,0,0]º-[H⁺]	-1.2	B ₇
1410.5256	1	[4,2,1][0,0,0]º-[H⁺]	-2.4	Z ₇
1414.5215	1	[4,1,2][0,0,0] -[H⁺]	-1.7	C ₇
1460.5269	1	[4,1,2][0,0,0]** -[H ⁺]	-1.7	C ₇
MS ² of Precurs	sor ion <i>m/z</i>	1242.4535 (2-)		
402.1516	1	[1,1,0][0,0,0]º-[H⁺]	-0.5	Z ₂
431.1418	1	[2,0,0][0,0,0] ^o -[H ⁺]	-0.4	Y ₁₁ /B ₃ ; Z ₁₁ /C ₃ ; Y ₇ /B ₇ ; Z ₇ /C ₇ ; Y ₃ /B ₁₁ ; Z ₃ /C ₁₁
575.2202	1	[1,1,1][0,0,0]º-[H⁺]	-0.7	Y ₁₀ /B ₅ ; Z ₁₀ /C ₅ ; Y ₉ /B ₆ ; Z ₉ /C ₆ ; Y ₆ /B ₉ ; Z ₆ /C ₉ ; Y ₅ /B ₁₀ ; Z ₅ /C ₁₀

tandem MS in negative ion mode on an FT-ICR mass spectrometer.

618.2258	1	[2,1,0][0,0,0] ^o -[H ⁺]	-1.0	Z ₃
664.2311	1	[2,1,0][0,0,0] ^{o,**} -[H ⁺]	-1.2	Z ₃
791.2939	1	[2,1,1][0,0,0] ^o -[H ⁺]	-1.7	B ₄ ; Z ₄
837.3001	1	[2,1,1][0,0,0] ^{o,**} -[H ⁺]	-0.7	B ₄ ; Z ₄
944.3460	2		-2.5	^{0,2} A ₁₀
982.3728	1	[2,1,2][0,0,0] -[H ⁺]	-1.8	C ₅
1007.3685	1	[3,1,1][0,0,0]º-[H ⁺]	-1.3	Y ₁₁ /B ₆ ; Z ₁₁ /C ₆ ; Y ₁₀ /B ₇ ; Z ₁₀ /C ₇ ; Y ₇ /B ₁₀ ; Z ₇ /C ₁₀ ; Y ₆ /B ₁₁ ; Z ₆ /C ₁₁
1015.8758	2	[5,3,2][0,0,0] ^{<u>0</u>,** -2[H⁺]}	-1.7	Z ₁₀
1028.3780	1	[2,1,2][0,0,0]** -[H ⁺]	-2.0	C ₅
1052.3834	2		-2.2	^{0,2} A ₁₁
1093.9021	2	[6,2,3][0,0,0] ^o -2[H ⁺]	-2.0	B ₁₁
1097.3993	1		-2.0	^{0,2} A ₆
1100.9096	2	[6,3,2][0,0,0] ^o -2[H ⁺]	-2.3	Z ₁₁
1102.9063	2	[6,2,3][0,0,0] -2[H ⁺]	-2.9	C ₁₁
1123.9125	2	[6,3,2][0,0,0] ^{o,**} -2[H ⁺]	-2.1	Z ₁₁
1180.4363	1	[3,1,2][0,0,0]º-[H⁺]	-2.0	B ₆
1194.4516	1	[3,2,1][0,0,0]º-[H⁺]	-2.3	Z ₆
1196.4491	2	[6,3,3][0,0,0] -2[H ⁺]	-2.2	[M-2H] ²⁻
1205.4533	2	[6,3,3][0,0,0] ^o -2[H ⁺]	-3.1	[M-H ₂ O-2H] ²⁻
1219.4512	2	[6,3,3][0,0,0]** -2[H ⁺]	-2.7	[M+FA-2H] ²⁻
1223.4419	1	[4,1,1][0,0,0]º-[H⁺]	-2.1	Y ₁₁ /B ₇ ; Z ₁₁ /C ₇ ; Y ₇ /B ₁₁ ; Z ₇ /C ₁₁
1367.5196	1	[3,2,2][0,0,0] ^⁰ -[H ⁺]	-2.6	Y ₁₀ /B ₇ ; Z ₁₀ /C ₇ ; Y ₉ /B ₁₀ ; Z ₉ /C ₁₀
1410.5249	1	[4,2,1][0,0,0] ^⁰ -[H ⁺]	-2.9	Z ₇
1583.5931	1	[4,2,2][0,0,0]º-[H⁺]	-2.9	B ₈ ; Z ₈
1774.6715	1	[4,2,3][0,0,0] -[H ⁺]	-3.2	C ₉
1889.6987	1		-2.8	^{0,2} A ₁₀

¹Saccharide compositions are given as [GalNAcAN, QuiNAc, Qui4NFm] [HexNAc, Hex, Kdo].

 $^{\circ}$ represents loss of water; $^{+\circ}$ represents water adduct; ** stands for formic acid adduct; and *

stands for loss of formic acid.

Supplemental Table 4 Observed fragment ions for [6,3,3][0,0,0] from multi-stage tandem

Product ion	Charge	Saccharide	Mass	Fragmentation			
(m/z)		Composition ¹	Error				
			(ppm)				
MS ² Precursor ion	MS ² Precursor ion <i>m/z</i> 1220.4479 (2+)						
412.1320	1	[1,0,1][0,0,0]º+Na⁺	-1.7	B ₂			
426.1476	1	[1,1,0][0,0,0]º+Na⁺	-1.7	Z ₂			
430.1424	1	[1,0,1][0,0,0] +Na⁺	-2.0	C ₂			
444.1581	1	[1,1,0][0,0,0] +Na⁺	-1.8	Y ₂			
				Y ₁₁ /B ₃ ; Z ₁₁ /C ₃ ; Y ₇ /B ₇ ; Z ₇ /C ₇ ;			
455.1376	1	[2,0,0][0,0,0]º+Na⁺	-2.0	Y ₃ /B ₁₁ ; Z ₃ /C ₁₁			
473.1480	1	[2,0,0][0,0,0] +Na ⁺	-2.3	Y ₁₁ /C ₃ ; Y ₇ /C ₇ ; Y ₃ /C ₁₁			
628.2062	1	[2,0,1][0,0,0]º+Na⁺	-1.8	B ₃			
642.2219	1	[2,1,0][0,0,0]º+Na⁺	-1.6	Z_3			
646.2162	1	[2,0,1][0,0,0] +Na ⁺	-2.6	C_3			
660.2323	1	[2,1,0][0,0,0] +Na ⁺	-1.9	Y ₃			
815.2897	1	[2,1,1][0,0,0]º+Na⁺	-2.6	B ₄ ; Z ₄			
833.3003	1	[2,1,1][0,0,0] +Na ⁺	-2.5	C ₄ ; Y ₄			
901.8249	2	[4,2,3][0,0,0] ^o +2Na ⁺	-1.4	B ₉			
910.8305	2	[4,2,3][0,0,0]+2Na ⁺	-1.0	C ₉			
917.8388	2	[4,3,2][0,0,0]+2Na ⁺	-0.5	Y ₉			
923.3280	2	[5,2,2][0,0,0] ^o +2Na ⁺	-1.2	Y ₁₀ /B ₁₁ ; Z ₁₀ /C ₁₁ ; Y ₁₁ /B ₁₀ ; Z ₁₁ /C ₁₀			
932.3331	2	[5,2,2][0,0,0]º+2Na+	-1.3	Y ₁₁ /C ₁₀ ; Y ₁₀ /C ₁₁			
968.3439	2		-1.0	^{0,2} A ₁₀			
988.3593	1	[2,1,2][0,0,0]º+Na ⁺	-1.3	B ₅			
1002.3752	1	[2,2,1][0,0,0] ^o +Na ⁺	-1.0	Z ₅			
1006.3701	1	[2,1,2][0,0,0]+Na ⁺	-1.0	C ₅			
1009.8623	2	[5,2,3][0,0,0]º+2Na+	-1.2	B ₁₀			
1016.8703	2	[5,3,2][0,0,0]º+2Na+	-1.0	Z ₁₀			
1018.8675	2	[5,2,3][0,0,0]+2Na ⁺	-1.2	C ₁₀			
1025.8756	2	[5,3,2][0,0,0]+2Na ⁺	-0.9	Y ₁₀			
1031.3652	2	[6,2,2][0,0,0]º+Na ⁺	-1.1	Y ₁₁ /B ₁₁ ; Z ₁₁ /C ₁₁			
1040.3705	2	[6,2,2][0,0,0]+2Na ⁺	-1.1	Y ₁₁ /C ₁₁			
1049.3756	1	[3,1,1][0,0,0] +Na ⁺	-1.3	Y ₁₁ /C ₆ ; Y ₁₀ /C ₇ ; Y ₇ /C ₁₀ ; Y ₆ /C ₁₁			
1117.8993	2	[6,2,3][0,0,0] ^o +2Na ⁺	-1.3	B ₁₁			
1126.9037	2	[6,2,3][0,0,0]+2Na ⁺	-2.1	C ₁₁			
1204.4327	1	[3,1,2][0,0,0]º+Na ⁺	-2.1	B ₆			
1236.4599	1	[3,2,1][0,0,0] +Na ⁺	-1.2	Y ₆			
1247.4393	1	[4,1,1][0,0,0]⁰+Na⁺	-1.4	Y ₁₁ /B ₇ ; Z ₁₁ /C ₇ ; Y ₇ /B ₁₁ ; Z ₇ /C ₁₁			
1265.4499	1	[4,1,1][0,0,0]+Na ⁺	-1.3	Y ₁₁ /C ₇ ; Y ₇ /C ₁₁			
1337.4713	1		-1.0	^{0,2} A ₇			
1434.5242	1	[4,2,1][0,0,0]⁰+Na⁺	-0.9	Z ₇			
1438.5187	1	[4,1,2][0,0,0] +Na ⁺	-1.2	C ₇			
1452.5350	1	[4,2,1][0,0,0] +Na ⁺	-0.7	Y ₇			
1607.5910	1	[4,2,2][0,0,0]º+Na+	-2.0	B ₈ ; Z ₈			
1625.6009	1	[4,2,2][0,0,0] +Na ⁺	-2.4	C ₈ ; Y ₈			

MS in positive ion mode on an LTQ-Orbitrap XL mass spectrometer.

1780.6608	1	[4,2,3][0,0,0]º+Na+	-1.3	B ₉			
MS ³ Precursor ion	MS ³ Precursor ion m/z 833.3003 (1+), to simplify the nomenclature of the MS ³ fragment ions, Y ₄						
is assumed to be t	he precurs	sor ion of MS ³ unless s	pecified.				
412.1320	1	[1,0,1][0,0,0]º+Na⁺	-1.7	Y ₄ /B ₁₀			
426.1469	1	[1,1,0][0,0,0]º+Na⁺	-3.5	Z ₂			
430.1417	1	[1,0,1][0,0,0] +Na ⁺	-3.7	Y ₄ /C ₁₀			
444.1574	1	[1,1,0][0,0,0] +Na ⁺	-3.5	Y ₂			
473.1473	1	[2,0,0][0,0,0] +Na ⁺	-3.8	Y ₃ /C ₁₁			
628.2051	1	[2,0,1][0,0,0]º+Na⁺	-3.6	Y ₄ /B ₁₁			
646.2155	1	[2,0,1][0,0,0] +Na ⁺	-3.7	Y ₄ /C ₁₁			
660.2312	1	[2,1,0][0,0,0] +Na ⁺	-3.6	Y ₃			
815.2890	1	[2,1,1][0,0,0]º+Na⁺	-3.4	Z ₄			
833.2993	1	[2,1,1][0,0,0]+Na ⁺	-3.7	Y ₄			

¹Saccharide compositions are given as [GalNAcAN, QuiNAc, Qui4NFm] [HexNAc, Hex, Kdo].

^o represents loss of water.

Supplemental Table 5 Observed fragment ions for ¹⁸O labeled [6,3,3][0,0,0] from multi-

Product ion	Charge	Saccharide Composition ¹	Mass	Fragmentation				
(m/z)	-		Error	_				
			(ppm)					
MS ² Precursor ion	MS ² Precursor ion <i>m/z</i> 1221.4487 (2+)							
428.1518	1	[1,1,0][0,0,0]º+ ¹⁸ O- ¹⁶ O+Na⁺	-1.8	Z ₂				
446.1624	1	[1,1,0][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.7	Y ₂				
473.1480	1	[2,0,0][0,0,0]+Na ⁺	-2.3	Y ₁₁ /C ₃ ; Y ₇ /C ₇ ; Y ₃ /C ₁₁				
628.2063	1	[2,0,1][0,0,0]º+Na⁺	-1.6	B ₃				
646.2166	1	[2,0,1][0,0,0]+Na ⁺	-2.0	C ₃				
662.2365	1	[2,1,0][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.9	Y ₃				
815.2903	1	[2,1,1][0,0,0]º+Na⁺	-1.8	B ₄				
835.3052	1	[2,1,1][0,0,0]+ ¹⁸ O- ¹⁶ O +Na ⁺	-1.6	Y ₄				
988.3592	1	[2,1,2][0,0,0]º+Na⁺	-1.4	B ₅				
1004.3791	1	[2,2,1][0,0,0]º+ ¹⁸ O- ¹⁶ O+Na⁺	-1.4	Z ₅				
1018.8668	2	[5,2,3][0,0,0]+2Na ⁺	-1.9	C ₁₀				
1022.3896	1	[2,2,1][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.4	Y ₅				
1031.3651	2	[6,2,2][0,0,0]º+2Na⁺	-1.2	Y ₁₁ /B ₁₁ ; Z ₁₁ /C ₁₁				
1040.3703	2	[6,2,2][0,0,0]+2Na ⁺	-1.3	Y ₁₁ /C ₁₁				
1076.3807	2		-1.4	^{0,2} A ₁₁				
1117.8996	2	[6,2,3][0,0,0]º+2Na⁺	-1.1	B ₁₁				
1126.9044	2	[6,2,3][0,0,0]+2Na ⁺	-1.5	C ₁₁				
1204.4337	1	[3,1,2][0,0,0]º+Na⁺	-1.2	B ₆				
1438.5182	1	[4,1,2][0,0,0]+Na⁺	-1.5	C ₇				
1454.5386	1	[4,2,1][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.1	Y ₇				
1607.5918	1	[4,2,2][0,0,0] ^o +Na ⁺	-1.5	B ₈				
1627.6068	1	[4,2,2][0,0,0]+ ¹⁸ O- ¹⁶ O+Na ⁺	-1.4	Y ₈				
1780.6605	1	[4,2,3][0,0,0]⁰+ Na⁺	-1.4	B ₉				

stage tandem MS in positive ion mode on an LTQ-Orbitrap XL mass spectrometer.

¹Saccharide compositions are given as [GalNAcAN, QuiNAc, Qui4NFm] [HexNAc, Hex, Kdo].

^o represents loss of water.

Supplemental Figure 1 Multi-stage tandem MS reveals the structural linkage for

[4,2,2][0,0,0]. Tandem mass spectra of the SEC fraction enriched with [4,2,2][0,0,0] was recorded with an Oribtrap mass spectrometer in positive ion mode. CID with 20% normalized collision energy was applied to achieve fragmentation of the $[M + 2Na]^{2+}$ at m/z 824.2961 (A). The ion observed at m/z 833.3009 in the MS² spectrum was isolated and CID was applied to obtain the MS³ spectrum (B). Two dominant fragment ions at m/z 646.2158 and 660.2314 in the MS³ spectrum were isolated and fragmented to achieve MS⁴ spectra (C and D). The fragment ions are designated using the Domon and Costello nomenclature (*27*). "/" is used for internal cleavages, and alternative assignments are shown above one another.



Supplemental Figure 2 Multi-stage tandem mass spectra of the ¹⁸O labeled saccharide [4,2,2][0,0,0] reveals the reducing-end residue. After quadrupole ion trap isolation of the ¹⁸O labeled saccharide [4,2,2][0,0,0] at m/z 825.2987, CID with 20% normalized collision energy was applied to achieve fragmentation (A). The ion at m/z 835.3053 further isolated from the MS² spectrum and CID was applied to obtain MS³ spectrum (B). All fragments contain sodium.



Supplemental Figure 3 Similar fragment ions are observed for saccharides in negative

ion mode. CID MS^2 spectra of saccharide chains [4,2,2][0,0,0] at m/z 846.3049 (A) and [6,3,3][0,0,0] at m/z 1242.4535 (B) were obtained in negative ion mode using a Solarix FT-ICR mass spectrometer.



Supplemental Figure 4 Multi-stage tandem MS of the $[M + 2Na]^{2+}$ *m/z* 1220.4479 of the saccharide [6,3,3][0,0,0]. After quadrupole ion trap isolation *m/z* 1220.4479, CID with 20% normalized collision energy was applied to achieve fragment ions (A). The ion at *m/z* 833.3003 was further isolated from the MS² spectrum and CID was applied to obtain MS³ spectrum (B).



Supplemental Figure 5 Tandem MS of the $[M + 2Na]^{2+}$ *m/z* 1221.4487 of the ¹⁸O-labeled saccharide [6,3,3][0,0,0] reveals the reducing-end residue. After quadrupole isolation *m/z* 1221.4487, CID with 20% normalized collision energy was applied to achieve fragmentation of the ¹⁸O labeled saccharide [6,3,3][0,0,0].

