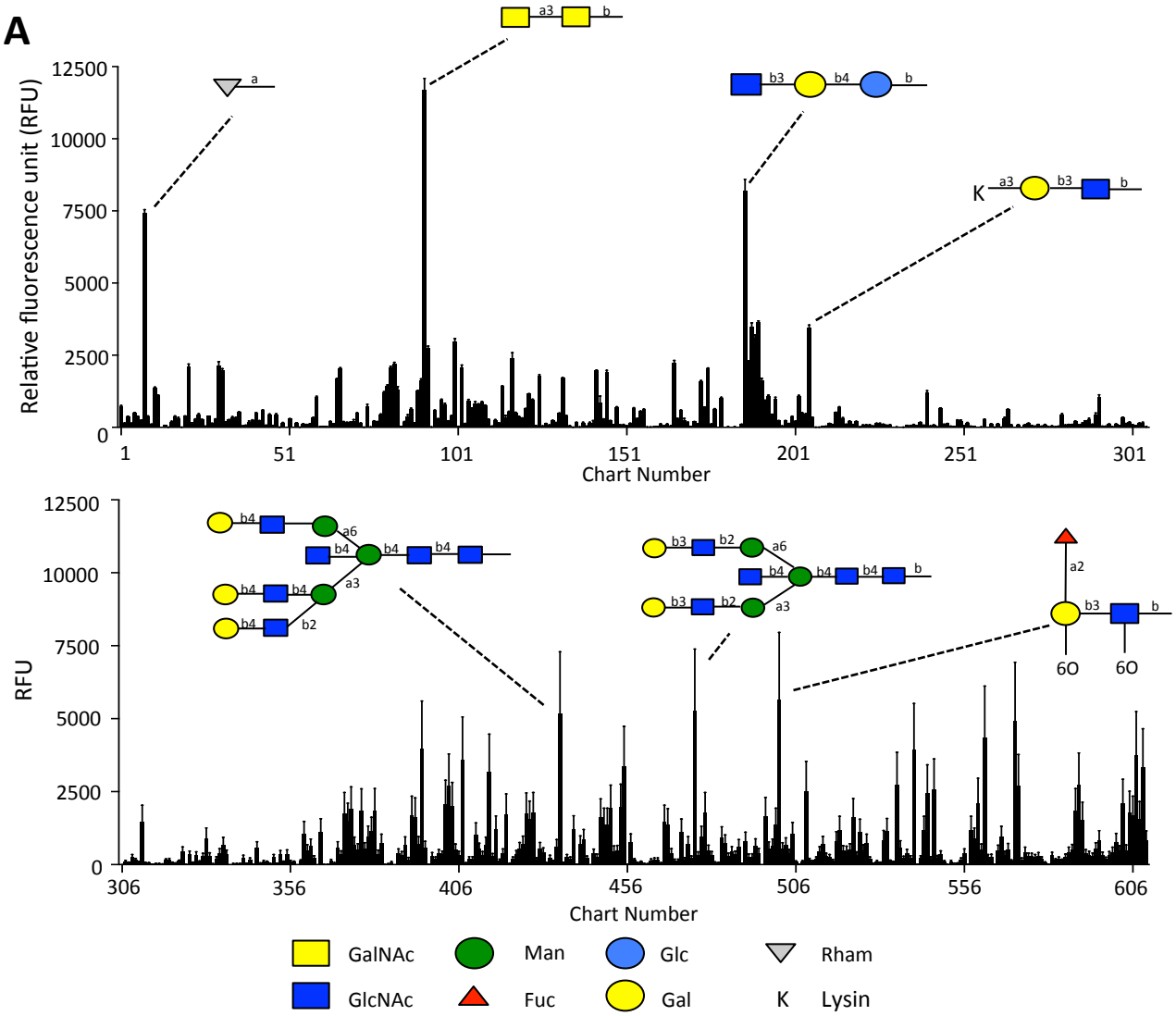
A



B

The top 20 glycans bound by Privigen.

Chart #	Glycan Name	Antibody binding ratio (ABR)
562	GalNAcb1-3GlcNAcb-Sp0	43449.31575
30	(3S)Galb1-3GlcNAcb-Sp0	21369.6875
377	Neu5Aca2-6Galb1-4GlcNAcb1-3GalNAc-Sp14	18457.11625
65	Fuca1-2Galb1-3GlcNAcb1-3Galb1-4Glcb-Sp8	16635.94575
90	GalNAca1-3(Fuca1-2)Galb-Sp18	16223.9325
114	Gala1-3Galb1-3GlcNAcb-Sp0	14177.7985
525	Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-6GalNAc-Sp14	11079.85325
123	Gala1-4GlcNAcb-Sp8	9381.1425
331	Neu5Aca2-3Galb1-3(Fuca1-4)GlcNAcb1-3Galb1-3(Fuca1-4)GlcNAcb-Sp0	8931.22675
107	Gala1-3(Fuca1-2)Galb-Sp8	8055.35775
491	Neu5Aca2-3Galb1-3GlcNAcb1-6GalNAca-Sp14	7876.8205
485	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb-Sp24	7270.845
514	GalNAcb1-4(6S)GlcNAc-Sp8	6894.9575
414	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-3GalNAca-Sp14	6734.65
494	Fuca1-2Galb1-4GlcNAcb1-6GalNAca-Sp14	6290.3545
105	Gala1-3(Fuca1-2)Galb1-4GlcNAc-Sp0	6245.08875
116	Gala1-3Galb1-4Glcb-Sp0	5418.57625
569	(3S)GlcAb1-3Galb1-4GlcNAcb1-3Galb1-4Glc-Sp0	5133.6885
561	GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-3)GalNAca-Sp14	5031.56975
487	Neu5Aca2-6Galb1-4GlcNAcb1-6(Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-3)Galb1-4Glc-Sp21	4362.745

Figure S1. Recognition of distinct carbohydrate-structures (glycans) by IgG antibodies in Privigen IVIG. **(A)** A total of 610 glycans were screened for binding of IgG in IgGPrepB by the glycan array CFG 5.1 as outlined in the Materials and Methods section. The carbohydrate structures of several of the top Privigen bound glycans are indicated as illustrative examples. **(B)** Results of top 20 glycans sorted by IgGPrepB ABR. RFU below the cut-off of the 95th percentile of the myeloma control were considered as negative and have been excluded from the table. Error bars represent s.e.m.

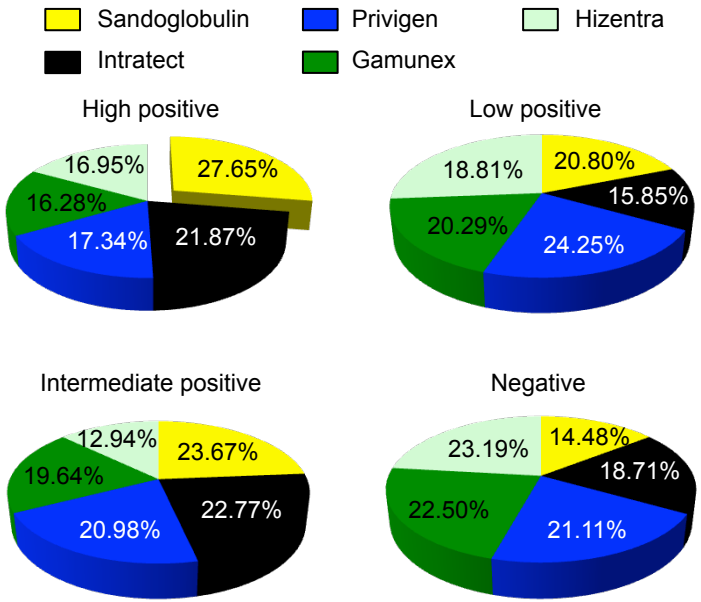


Figure S2. Relative distribution of negative, high-, low and intermediate positive bound glycans. Pie charts depicting the relative distribution of Sandoglobulin, Privigen, Hizentra, Intratect and Gamunex for negative, high-, intermediate-, and low positive bound glycans revealing a predominance of Sandoglobulin for high bound glycans. Categorized binding intensity was computed, as outlined in the Materials and methods section.

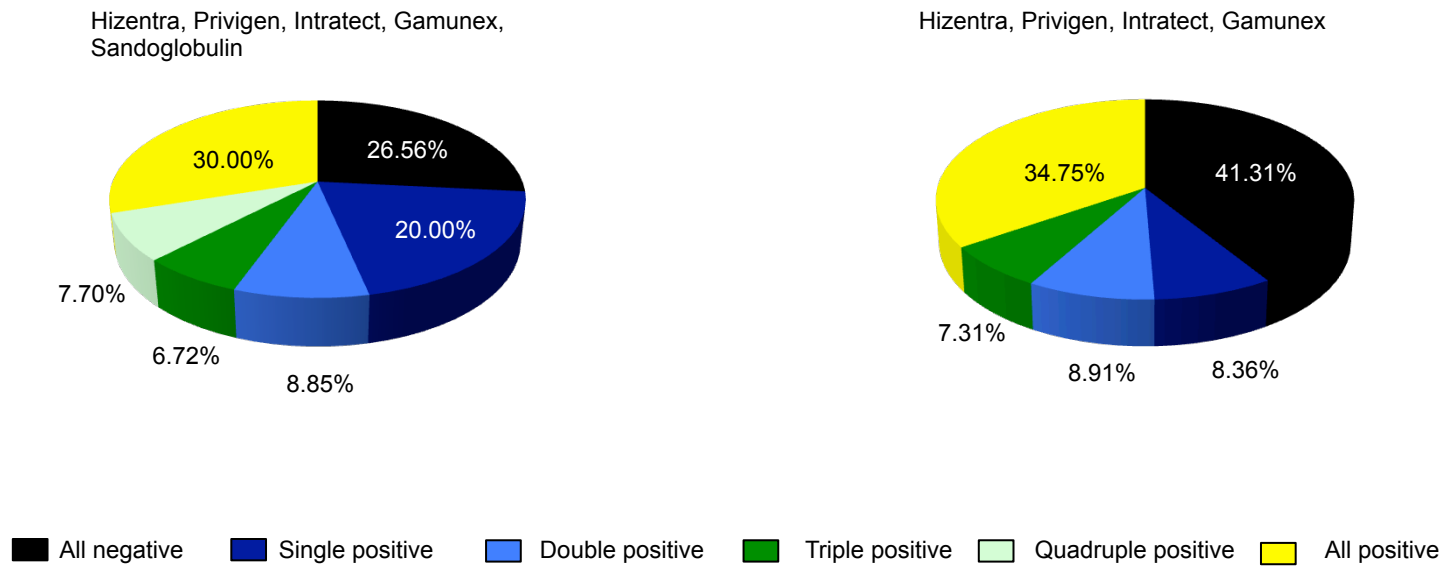


Figure S3. Similarities in glycan recognition among different IVIG preparations. Pie charts depicting the percentages of glycans recognized either by one (single positive), several or all preparations, including (left) or without (right) Sandoglobulin.

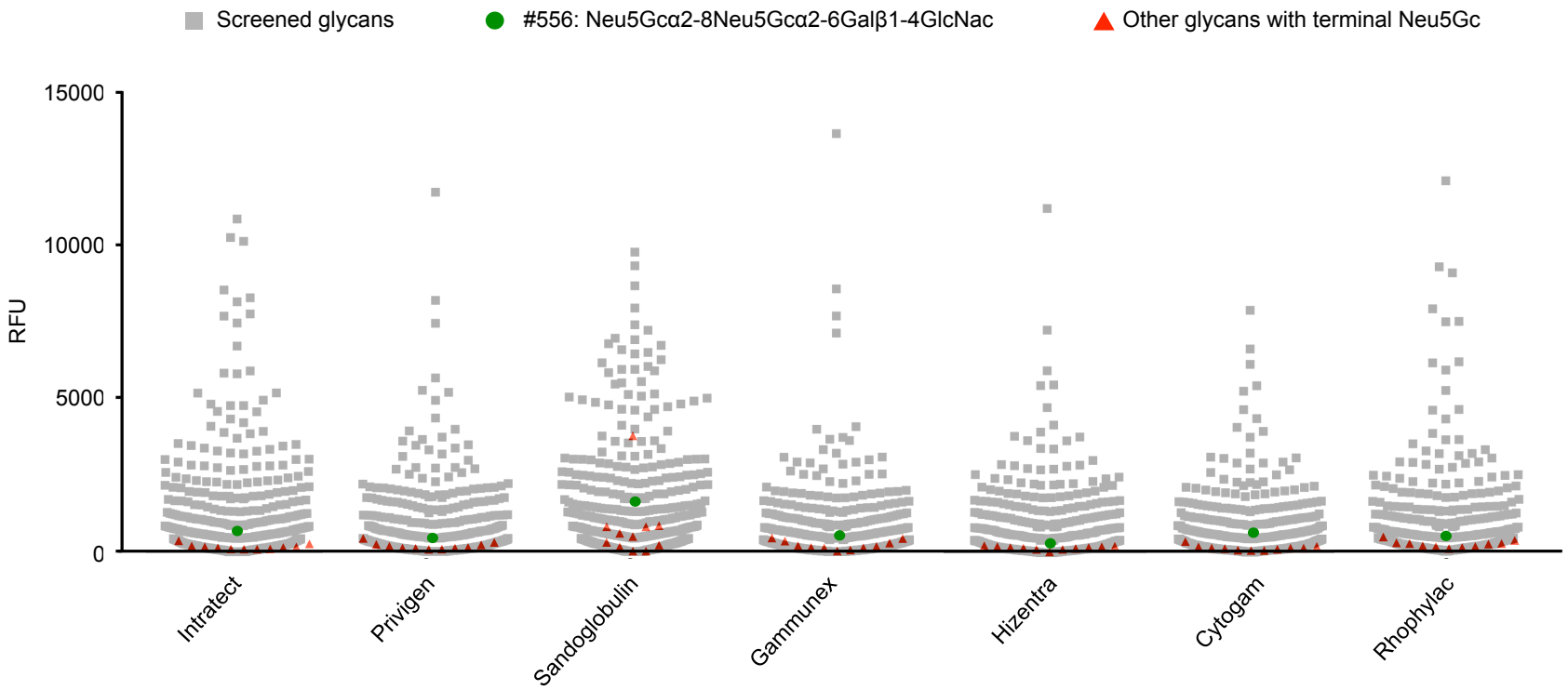
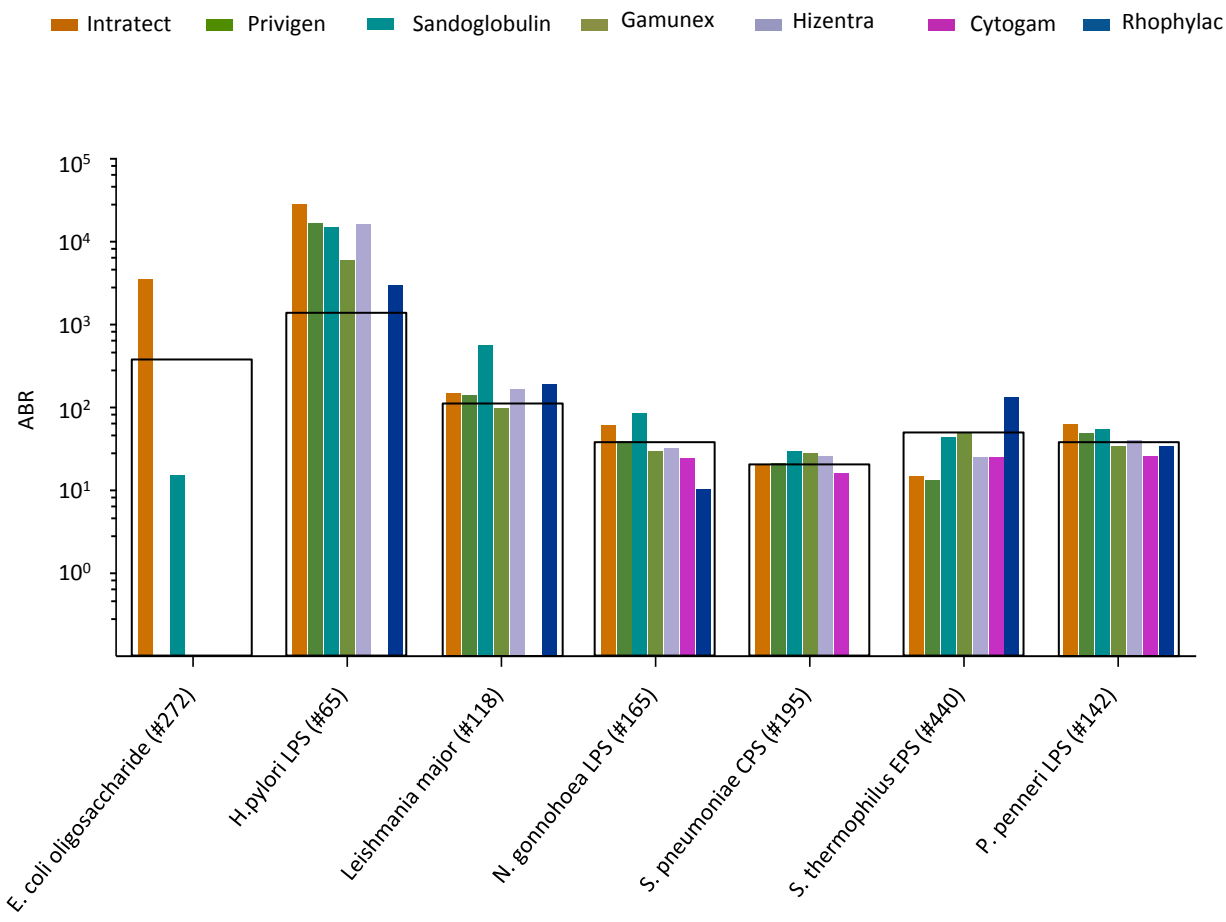


Figure S4. Immunogenicity of Neu5Gc-terminated glycans. Binding reactivities (RFU) of Sandoglobulin, Gammunex, Hizentra, Intratect, Privigen, Cytogam and Rhophylac to Neu5Gc-terminated carbohydrate structures (red or green).

Fig. S5

A



B

The top 20 microorganism associated glycans bound by IVIG / SCIG																					
Chart #	Glycan Name	LPS / LOS		LTA		CPS				EPS				Other				Eukaryotic		Predominant Ab Isotype	
		gram -		gram +		gram +		gram -		gram +		gram -		gram +		gram -					
		Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Pathogen	Comm./Environ.	Non-IgG2	IgG2
562	GalNAcβ1-3GlcNAcβ-Sp0		x						x											x	
65	Fucα1-2Galβ1-3GlcNAcβ1-3Galβ1-4Glcβ-Sp8	x																			x
107	Galα1-3(Fucα1-2)Galb-Sp8															x					x
190	GlcNAcβ1-4GlcNAcβ1-4GlcNAcβ1-4GlcNAcβ1-4GlcNAcβ1-Sp8																x			x	
8	Rhaa-Sp8		x																	x	
145	Galβ1-3GalNAcβ1-4Galβ1-4Glcβ-Sp8	x																		x	
118	Galα1-3Galb-Sp8									x								x			x
122	Galα1-4Galβ1-4Glcβ-Sp0	x	x																		x
66	Fucα1-2Galβ1-3GlcNAcβ1-3Galβ1-4Glcβ-Sp10	x																			
79	Fucα1-3GlcNAcβ-Sp8	x																			
108	Galα1-3(Fucα1-2)Galb-Sp18									x				x							x
142	Galβ1-3GalNAcβ-Sp8	x	x																	x	
165	Galβ1-4GlcNAcβ1-3Galβ1-4Glcβ-Sp8	x	x			x														x	
100	Galα1-2Galb-Sp8																	x			
115	Gala1-3Galb1-4GlcNAcb-Sp8		x											x							x
264	Neu5Aca2-3Galb1-4Glcb-Sp8	x	x														x				x
102	Galα1-3(Fucα1-2)Galβ1-3GlcNAcβ-Sp8	x																		x	
468	Gala1-3(Fuca1-2)Galb1-3GalNAcb-Sp8	x																		x	
179	GlcNAcβ1-3GalNAcα-Sp8	x																			x
448	Galβ1-4GlcNAcβ1-2Mana-Sp0		x																	x	

Figure S5. IgG binding to microorganism-associated carbohydrate structures. **(A)** Exemplary examples of ABR values for seven different glycans, associated with surface expression on microorganisms, in a grouped box plot for different IVIG and SCIG preparations. **(B)** The top 20 microorganism-associated glycans bound by IVIG/SCIG. Bacterial glycans were identified using the Bacterial Carbohydrate Structure Data Base (BCSDB). Abbreviations: CPS = capsular polysaccharides, EPS = exopolysaccharides, LPS = lipopolysaccharides, LOS = lipooligosaccharides, LTA = lipoteichoic acid, Comm. = commensal, Environ. = environmental, Ab = antibody

Supplemental Table 1: Characteristics of immunoglobulin products used in the study

Product	**IgG2 content**[1)]	**Manufacturing processes: major purification steps**	Excipients	**Route Form**	**Characteristics**	**Plasma origin; Donors per lot**	**Manufacturer**
Intratect	37%	Cold ethanol fractionation, cation-exchange chromatography	Glycine	i.v. liquid	Total serum IgG	na	Biotest
Privigen	28.7%	Cold ethanol fractionation, octanoic acid precipitation, anion-exchange chromatography	Proline	i.v. liquid	Total serum IgG	EU[2)], USA; 1000 - 60'000	CSL Behring
Hizentra	22-34%	Cold ethanol fractionation, octanoic acid precipitation, anion-exchange chromatography	Proline, polysorbate 80	s.c. liquid	Total serum IgG	EU[2)], USA; 1000 - 60'000	CSL Behring
Gamunex	29.7%	Cold ethanol fractionation, capyrlate precipitation, anion exchange chromatography	Glycine	i.v. liquid	Total serum IgG	na	Grifols/Talecris
CytoGam	28%	Cold ethanol fractionation	Sucrose, albumin	i.v. liquid	Total IgG from anti-CMV enriched serum	USA 1500-5000	CSL Behring
Rhophylac	7.6%	Cation-exchange chromatography	Glycine, albumin, NaCl	i.v.; i.m. liquid	Hyperimmune IgG anti-RhD blood group antigen	USA; <400	CSL Behring
Sandoglobulin	30.2%	Cold ethanol fractionation	Sucrose, NaCl	i.v. lyophil.	Total serum IgG	USA; >16'000	CSL Behring

[1)] Average IgG2 content as reported in Summary of Product Characteristics (SPC) documents or Product Insert leaflets of the various products
[2)] Austria, Switzerland, Germany
na; information not available

Supplemental Table 2. Comprehensive list of glycans bound by antibodies contained in IVIG. Results for each of the 610 glycans on the array and for 8 different IVIg preparations (Intratect, Privigen, Gamunex, Hizentra, Cytogam, Rhophylac, Sandoglobulin, IgG2-depleted Sandoglobulin) and 1 control mix IgG are shown. Displayed are average RFU and S.D. for each preparation (n=6 per preparation).

Chart #	Masterlist Name	Intratect		Privigen		Gamunex		Hizentra		Cytogam		Rhophylac		Sandoglobulin		IgG2 depl.		Control Mix	
		RFU	S.D.	RFU	S.D.	RFU	S.D.	RFU	S.D.	RFU	S.D.	RFU	S.D.	RFU	S.D.	RFU	S.D.	RFU	S.D.
1	Gala	1253.4	108.8	728.3	71.3	667.8	134.8	752.7	117.0	1062.6	58.2	1039.0	106.3	2273.6	392.5	177.7	80.8	136.7	18.7
2	Glca	446.6	61.0	126.5	64.1	75.3	16.1	112.5	28.2	150.6	70.0	366.4	120.0	129.6	56.6	73.4	17.0	47.3	28.1
3	Mana	292.9	25.0	374.1	50.0	108.5	61.9	314.6	90.3	236.4	55.0	278.4	5.6	315.7	240.7	154.0	24.6	71.6	24.2
4	GalNAca	392.4	119.0	144.6	14.0	302.3	101.6	106.5	24.4	163.2	53.0	230.4	90.7	194.0	73.1	66.6	15.5	26.3	14.2
5	GalNAca	613.7	20.2	501.6	45.9	289.1	71.1	142.1	91.4	396.7	114.9	731.4	221.7	336.2	52.2	263.5	70.4	191.1	102.5
6	Fuca	633.7	33.3	354.7	79.6	510.9	49.4	290.8	110.6	576.6	54.1	1038.9	73.8	283.3	97.3	161.0	37.1	43.3	30.9
7	Fuca	239.5	73.5	244.3	25.9	81.7	30.3	150.7	76.2	368.3	93.2	269.1	82.8	352.4	64.3	162.4	84.4	20.4	38.7
8	Rhaa	8546.4	458.5	7429.8	284.4	7684.5	270.7	7270.2	234.3	6094.0	399.2	7458.4	792.0	6022.8	611.7	3392.7	638.4	45.4	39.8
9	Neu5Aca	609.6	133.1	374.0	68.6	199.8	26.5	331.9	58.3	280.9	91.9	271.5	42.3	84.9	36.0	208.1	17.4	131.8	43.6
10	Neu5Aca	87.7	15.5	111.1	14.5	110.1	54.5	75.0	38.0	109.6	25.0	109.0	25.7	43.9	67.5	28.4	29.6	2.9	19.0
11	Neu5Acb	1160.1	103.8	1349.8	126.9	516.1	38.1	1060.8	66.6	939.9	119.2	503.4	25.7	1335.3	247.1	518.9	131.2	75.6	7.0
12	Galb	1631.4	91.3	1109.2	28.3	666.1	59.1	742.6	223.3	1511.3	173.9	1748.4	289.0	1324.6	23.1	299.5	66.8	117.6	35.7
13	Glcb	125.4	75.3	0.0	8.8	0.0	16.1	60.5	7.9	41.1	14.3	31.8	18.2	0.0	53.9	20.3	14.1	3.3	38.6
14	Manb	535.7	43.7	63.9	23.3	5.7	12.9	156.3	46.7	160.9	25.7	376.2	139.1	8.2	26.2	58.8	38.6	75.3	21.7
15	GalNAcb	615.7	71.3	0.0	40.6	39.0	34.1	201.8	58.5	77.9	35.7	115.6	50.5	0.0	43.6	67.2	20.2	10.5	35.7
16	GlcNAcb	711.6	59.2	176.5	93.3	23.6	7.5	0.0	202.2	157.2	45.9	829.9	285.2	287.6	65.4	115.2	48.9	114.4	44.8
17	GlcNAcb	758.9	122.7	355.9	95.7	455.1	67.0	299.3	179.4	491.5	69.9	1593.7	321.3	396.3	62.6	113.2	37.0	11.2	11.6
18	GlcN(Gc)b	430.5	125.0	317.2	78.9	73.2	21.4	13.6	48.6	290.6	28.8	582.0	124.5	335.1	71.2	108.1	26.3	274.4	45.0
19	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3)GalNAca	24.7	11.1	0.0	23.4	48.8	32.0	92.6	99.6	83.9	61.6	82.7	35.7	20.6	40.3	24.6	23.0	63.4	58.9
20	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3)GalNAc	559.8	150.5	383.7	40.2	420.9	85.3	287.9	32.5	624.9	42.9	503.6	36.6	558.4	83.0	86.1	30.3	153.4	16.1
21	GlcNAcb1-6(GlcNAcb1-4)(GlcNAcb1-3)GlcNAc	5813.1	496.0	2087.9	257.1	1964.2	233.2	2244.4	85.2	1429.4	83.6	971.4	31.6	3103.2	769.0	372.3	100.0	16.1	8.4
22	6S(3S)Galb1-4(6S)GlcNAcb	0.0	13.1	106.4	137.7	73.1	13.1	107.3	51.2	99.8	43.1	350.6	338.8	392.2	85.3	44.5	21.2	118.1	34.3
23	6S(3S)Galb1-4GlcNAcb	262.6	66.2	275.3	51.8	319.6	36.9	173.2	14.2	359.1	42.7	799.1	120.6	175.1	48.1	109.8	26.3	86.9	12.2
24	(3S)Galb1-4(Fuca1-3)(6S)Glc	369.5	53.0	441.0	91.9	391.2	21.6	399.6	32.7	347.8	37.8	432.8	47.5	629.2	526.6	128.9	18.8	156.2	23.7
25	(3S)Galb1-4Glcb	607.8	72.0	259.6	49.5	173.6	59.9	206.4	41.5	358.9	76.5	386.0	81.9	354.0	71.3	103.2	20.0	124.1	14.1
26	(3S)Galb1-4(6S)Glcb	584.0	64.5	29.2	37.0	19.7	20.0	33.2	68.7	43.8	19.8	352.4	77.4	15.5	39.5	41.3	15.3	25.0	10.6
27	(3S)Galb1-4(6S)Glcb	879.9	55.1	380.0	28.9	472.1	55.2	483.0	100.0	583.4	36.4	639.2	17.4	727.6	63.8	241.6	42.7	70.2	19.2
28	(3S)Galb1-3(Fuca1-4)GlcNAcb	157.7	30.3	139.2	13.3	199.4	100.2	28.8	174.1	152.7	25.4	348.7	111.2	27.2	19.1	102.6	9.2	23.1	11.7
29	(3S)Galb1-3GalNAca	267.3	39.5	262.6	79.5	122.5	15.1	173.0	53.4	194.8	30.4	218.4	59.5	193.1	21.5	119.5	42.6	64.4	16.2
30	(3S)Galb1-3GlcNAcb	4773.0	621.7	2137.0	355.6	1786.4	221.7	1606.7	482.3	1430.0	157.6	3300.6	508.5	1851.8	249.6	1823.1	311.3	0.1	15.8
31	(3S)Galb1-3GlcNAcb	2829.8	1525.1	1971.0	172.4	1234.4	41.5	2212.0	235.2	2099.4	73.3	2902.4	289.3	3590.0	754.7	2301.5	110.9	29.8	26.6
32	(3S)Galb1-4(Fuca1-3)GlcNAc	220.0	37.2	163.1	74.6	214.9	35.6	47.9	34.8	295.5	58.2	367.6	59.7	91.0	59.2	95.9	47.5	126.7	43.6
33	(3S)Galb1-4(Fuca1-3)GlcNAc	302.0	49.6	260.8	38.9	182.2	65.7	176.4	60.9	177.7	84.0	352.0	63.1	232.8	156.4	114.3	27.6	1.1	9.0
34	(3S)Galb1-4(6S)GlcNAcb	400.9	35.2	351.9	59.8	288.1	48.5	386.6	107.5	432.5	37.3	615.7	61.3	324.2	103.2	117.8	48.7	18.1	42.8
35	(3S)Galb1-4(6S)GlcNAcb	397.2	111.8	327.7	56.0	198.0	41.0	264.8	21.3	215.7	165.9	550.8	88.0	414.4	82.0	113.4	22.0	73.2	12.4
36	(3S)Galb1-4GlcNAcb	667.5	66.0	519.5	63.8	507.9	63.1	533.8	116.0	718.2	37.7	605.1	45.3	390.1	96.8	112.7	76.6	4.3	24.6
37	(3S)Galb1-4GlcNAcb	73.2	2.4	46.1	17.0	4.6	14.2	71.2	32.3	53.3	12.7	30.7	38.1	33.9	12.2	39.8	37.6	13.7	18.2
38	(3S)Galb	228.3	47.7	74.2	23.6	48.8	20.7	79.7	39.7	153.8	39.6	430.9	87.6	0.0	8.8	73.0	36.1	117.6	20.2
39	(6S)(4S)Galb1-4GlcNAcb	2435.1	300.9	188.0	75.0	43.0	36.5	368.0	70.6	492.4	118.8	347.7	129.3	15.5	38.7	40.9	20.1	0.1	13.5
40	(4S)Galb1-4GlcNAcb	1727.8	34.4	249.1	29.2	20.2	8.9	86.2	35.7	285.3	42.6	319.7	7.7	21.5	35.8	59.0	13.4	175.8	35.7
41	(6P)Mana	742.1	128.8	485.7	72.3	467.6	44.3	238.8	172.1	380.4	70.1	556.8	59.3	663.3	21.2	142.8	22.1	67.1	18.9
42	(6S)Galb1-4Glcb	386.1	98.8	209.3	70.1	104.2	11.5	128.4	56.8	306.3	82.2	643.2	482.3	286.6	34.6	38.1	31.9	215.0	54.4
43	(6S)Galb1-4Glcb	722.0	80.7	591.8	37.7	866.0	141.9	404.7	55.6	692.5	108.0	652.0	46.7	479.6	82.1	134.4	18.5	79.3	10.0
44	(6S)Galb1-4GlcNAcb	121.1	21.6	73.2	18.9	112.1	55.4	121.0	41.0	210.9	35.9	283.6	160.0	244.2	72.2	114.9	15.7	128.3	36.2
45	(6S)Galb1-4(6S)Glcb	540.4	80.4	439.7	59.2	331.5	65.0	798.9	219.2	503.9	31.8	720.3	70.9	489.9	115.4	137.2	50.8	61.8	23.2
46	Neu5Aca2-3(6S)Galb1-4GlcNAcb	33.2	68.3	50.1	60.6	110.9	2.1	52.8	33.9	112.9	45.1	216.9	18.5	82.2	99.7	39.6	16.7	70.1	10.0
47	(6S)GlcNAcb	831.3	137.3	444.0	73.5	457.6	40.4	572.1	188.9	561.4	74.6	1193.7	148.3	687.1	60.2	118.8	41.5	88.0	26.1
48	Neu5,9Ac2a	58.6	25.0	0.0	34.8	111.5	39.2	26.5	32.6	238.3	144.4	243.9	45.5	1002.4	742.8	32.4	18.5	4.9	8.6
49	Neu5,9Ac2a2-6Galb1-4GlcNAcb	168.6	30.3	159.5	27.4	91.4	41.6	402.5	93.7	66.1	16.6	113.5	38.2	14.0	21.5	63.6	13.9	37.1	12.0
50	Mana1-6(Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	266.8	80.5	0.0	16.9	0.0	7.7	0.0	66.9	186.0	50.1	341.5	59.7	374.2	61.4	77.5	68.4	0.1	14.5
51	Mana1-6(Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	335.4	61.5	298.7	19.4	189.0	45.2	864.6	732.4	237.5	80.2	218.2	38.7	255.3	132.2	171.1	0.8	165.6	51.7
52	GlcNAcb1-2Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	92.6	46.0	0.0	9.3	55.3	51.2	73.0	37.9	91.8	39.4	137.0	26.6	67.2	41.0	46.6	14.9	26.8	17.9
53	GlcNAcb1-2Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	180.8	58.1	156.6	15.9	91.9	29.4	244.1	89.1	94.0	14.9	128.7	24.2	93.0	32.2	139.3	40.0	71.5	18.7
54	Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	111.3	58.9	0.8	34.7	150.2	34.2	99.1	47.3	221.4	96.6	223.8	23.9	120.1	44.9	81.0	51.9	45.9	15.5
55	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	66.5	30.6	111.7	51.6	139.5	38.5	129.4	62.9	80.5	34.0	118.4	69.8	267.3	321.8	6.6	50.3	58.5	21.6
56	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Man-a1-3)Manb1-4GlcNAcb1-4GlcNAcb	170.8	12.8	98.3	24.4	344.7	248.0	126.9	13.7	100.9	32.9	205.2	46.1	82.6	57.0	41.6	29.1	33.0	9.4
57	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	28.7	42.6	111.1	51.4	123.4	13.5	384.3	277.0	99.0	24.0	149.0	82.8	290.5	89.8	84.6	34.9	128.3	18.7
58	Fuca1-2Galb1-3GalNAcb1-3Gala	464.3	41.4	326.7	62.1	163.6	22.3	172.0	56.9	225.0	67.6	374.4	42.1	472.2	32.1	125.1	54.4	30.0	19.0
59	Fuca1-2Galb1-3GalNAcb1-3Gala1-4Galb1-4Glcb	1664.9	41.8	1039.7	106.8	924.0	198.8	1250.6	64.3	851.8	121.2	1858.3	74.8	975.1	210.7	165.4	38.0	50.2	13.6
60	Fuca1-2Galb1-3(Fuca1-4)GlcNAcb	86.5	59.5	24.3	18.3	30.8	5.9	68.5	23.4	68.9	32.3	17.0	21.2	49.3	32.3	64.4	28.7	30.9	18.4
61	Fuca1-2Galb1-3GalNAca	97.0	38.0	30.3	12.4	31.6	18.3	57.4	14.8	49.1	20.3	68.2	212.5	6.4	16.2	64.1	40.0	110.5	13.2
62	Fuca1-2Galb1-3GalNAca	211.2	48.0	19.2	17.3	33.1	25.3	85.8	19.2	38.7	48.3	0.0	26.2	117.1	28.8	6.2	12.2	0.1	71.7
63	Fuca1-2Galb1-3GalNAcb1-4(Neu5Aca2-3)Galb1-4Glcb	203.8	90.3	207.3	26.0	51.1	7.3	145.9	44.2	132.1	34.3	150.0	26.0	53.6	10.2	108.1	19.1	137.2	27.8
64	Fuca1-2Galb1-3GalNAcb1-4(Neu5Aca2-3)Galb1-4Glcb	123.5	49.2	115.3	29.0	154.0	48.7	100.2	27.0	148.1	14.7	171.0	94.3	63.2	38.3	99.6	51.3	33.1	23.1
65	Fuca1-2Galb1-3GlcNAcb1-3Galb1-4Glcb	2807.2	288.7	1663.6	77.0	606.2	122.6	1652.5	173.0	1123.5	102.9	303.3	26.4	1509.8	123.0	37.7	14.2	0.1	5.7
66	Fuca1-2Galb1-3GlcNAcb1-3Galb1-4Glcb	2830.7	64.6	2033.5	94.6	2626.3	105.1	1678.8	99.3	1585.9	152.7	396.2	56.1	2222.2	290.3	413.2	78.2	73.4	31.4
67	Fuca1-2Galb1-3GlcNAcb	268.0	79.4	178.8	55.5	101.6	46.0	202.5	32.9	186.4	51.9	161.7	49.1	138.5	80.6	23.7	16.7	0.1	5.9
68	Fuca1-2Galb1-3GlcNAcb	258.7	61.5	151.0	79.6	210.6	52.6	266.1	48.8	304.2	52.3	421.4	64.8	134.2	46.3	40.1	21.0	18.2	21.2
69	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	109.1	64.0	170.9	37.5	146.9	34.4	164.2	55.0	114.7	22.5	302.2	54.5	8.9	14.0	37.6	7.8	18.4	3.8
70	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	422.6	41.5	217.6	126.2	354.3	21.6	297.4	38.9	431.9	49.6	307.0	42.1	331.5	39.0	128.6	21.2	123.8	18.5
71	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb	472.1	15.8	484.0	67.5	404.4	40.5	498.7	46.1	384.6	34.4	732.3	42.2	768.3	175.2	124.3	38.4	89.7	16.0
72	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb	158.0	17.2	149.0	26.1	0.0	153.7	68.0	13.8	89.9	17.3	145.2	19.5	49.5	17.5	115.4	37.9	43.1	8.5
73	Fuca1-2Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	423.6	56.1	0.0	27.3	0.0	8.4	0.0	60.3	7.0	18.7	53.3	14.7	0.0	21.9	4.0	14.0	0.1	13.5
74	Fuca1-2Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	582.6	21.1	719.1	212.4	867.8	39.4	1216.6	257.1	341.6	49.7	268.9	34.3	368.1	60.3	213.3	18.6	125.0	25.7
75	Fuca1-2Galb1-4GlcNAcb	643.7	89.1	10.3	31.4	190.5	118.1	160.4	23.6	82.5	19.9	119.1	77.5	78.7	15.4	39.2	19.9	0.1	12.5
76	Fuca1-2Galb1-4GlcNAcb	197.2	61.4	212.3	38.6	133.3	43.2	187.1	66.1	123.6	60.5	261.2	54.9	175.5	86.7	144.7	77.1	42.7	15.1
77	Fuca1-2Galb1-4Glcb	410.6	35.6	183.3	65.1	481.9	31.7	312.3	54.6	342.1	58.3	396.1	51.2	127.9	119.9	88.9	43.9	19.7	13.5
78	Fuca1-2Galb	659.0	68.5	436.9	70.5	312.6	61.5	457.2	46.2	478.1	62.9	445.0	77.7	486.4	15.3	112.9	29.2	76.8	33.2
79	Fuca1-3GlcNAcb	2415.1	118.9	1187.9	130.5	1333.1	83.3	1408.3	68.3	669.3	113.5	504.5	62.1	893.4	70.2	338.5	72.8	15.8	2.4
80	Fuca1-4GlcNAcb	2656.7	115.1	1411.4	152.0	1181.5	138.4	1563.1	50.4	1017.8	66.3	399.0	176.0	1804.6	421.8	792.5	94.9	115.2	65.7
81	Fucb1-3GlcNAcb	3292.0	66.0	2074.1	123.5	1913.0	266.6	2254.7	488.4	2105.6	85.7	3176.0	269.9	2861.5	388.8	815.2	120.3	123.5	22.2
82	GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb	2289.8	141.7	2194.7	154.9	1094.9	148.5	1792.6	288.9	593.9	33.9	822.0	104.5	948.5	123.6	415.4	64.6	103.9	39.2
83	GalNAca1-3(Fuca1-2)Galb1-4(Fuca1-3)GlcNAcb	1458.8	45.4	1299.1	277.4	899.9	186.3	1343.7	121.0	1082.7	30.0	1104.8	90.7	1103.1	93.4	516.8	43.1	351.6	43.4
84	(3S)Galb1-4(Fuca1-3)Glcb	157.3	52.1	55.2	8.2	44.5	16.1	115.6	51.6	94.9	15.5	46.4	32.5	165.9	68.7	49.8	14.7	56.9	16.2
85	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb	401.6	66.9	188.2	40.9	3.4	14.1	406.8	55.1	90.4	18.9	208.8	60.8	0.0	50.2	85.6	10.7	132.9	45.5
86	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb	961.0	43.7	372.7	161.2	46.0	7.5	856.2	101.6	39.2	32.5	0.0	52.5	54.8	17.0	0.0	30.6	0.1	20.0
87	GalNAca1-3(Fuca1-2)Galb1-4Glcb	565.7	17.9	629.3	92.0	28.1	18.3	516.5	94.3	75.7	11.9	147.8	12.1	64.2	25.8	121.0	25.2	122.5	9.5
88	GlcNAcb1-3Galb1-3GalNAca	338.3	30.2	156.9	16.6	193.1	39.1	117.4	38.3	190.9	46.8	40.6	53.3	169.7	26.5	63.0	19.9	0.1	54.8
89	GalNAca1-3(Fuca1-2)Galb	1324.0	223.2	1261.7	44.6	348.4	96.7	1123.4	63.0	380.1	98.5	304.3	45.2	332.4	18.7	140.9	45.6	55.8	9.6
90	GalNAca1-3(Fuca1-2)Galb	1876.8	86.6	1622.4	144.7	1661.5	188.0	1441.4	97.2	507.2	46.9	378.6	39.5	605.2	141.0	135.2	41.3	0.1	43.0
91	GalNAca1-3GalNAcb	10868.2	1298.6	11697.0	962.5	13628.3	1786.8	11238.7	1188.5	7850.7	805.8	29278.7	2276.7	1268.4	1251.6	1280.5	122.1	749.1	513.8
92	GalNAca1-3Galb	3856.0	127.1	2737.2	190.0	2531.0	202.1	2424.2	101.1	1928.5	161.8	1461.7	29.7	508.4	695.0	205.2	34.1	43.1	34.8
93	GalNAca1-4(Fuca1-2)Galb1-4GlcNAcb	0.0	134.2	0.0	46.5	183.8	70.0	0.0	89.2	0.0	69.0	230.0	62.5	0.0	381.5	0.0	17.8	0.1	29.3
94	GalNAcb1-3GalNAca	1349.0	57.9	579.7	44.1	820.0	94.6	567.6	122.1	686.3	31.6	0.0	142.3	1378.6	39.0	45.3	21.2	58.0	66.5
95	GalNAcb1-3(Fuca1-2)Galb	223.3	45.3	299.1	27.0	199.0	62.6	276.1	67.8	33.4	39.8	239.0	14.6	393.4	155.2	22.8	30.7	85.0	46.2
96	GalNAcb1-3Gala1-4Galb1-4GlcNAcb	2197.7	140.2	943.1	86.8	721.8	29.0	1152.1	42.8	584.2	25.2	1024.3	22.5	794.0	560.9	687.3	37.1	65.2	13.1
97	GalNAcb1-4(Fuca1-3)GlcNAcb	1416.2	26.2	782.7	134.0	485.7	57.7	1023.6	246.2	726.5	72.3	504.5	142.2	1060.1	103.2	259.2	21.6	59.3	10.4
98	GalNAcb1-4GlcNAcb	232.2	38.3	203.2	72.8	90.5	13.7	107.1	23.7	245.7	27.7	186.7	73.0	164.8	104.3	97.5	15.9	22.5	16.6
99	GalNAcb1-4GlcNAcb	237.2	69.3	405.4	26.3	290.2	25.3	406.2	179.8	377.9	103.7	353.8	46.1	241.3	155.6	67.7	18.1	17.1	4.2
100	Gala1-2Galb	8296.2	749.0	2971.8	263.2	1418.8	325.9	2748.2	378.1	1950.2	310.0	9034.8	1030.1	4599.9	702.8	476.6	153.6	72.3	19.6
101	Gala1-3(Fuca1-2)Galb1-3GlcNAcb	1665.4	107.2	34.5	16.1	0.0	7.7	486.1	84.5	84.1	21.9	354.2	96.5	0.0	100.6	57.0	23.5	0.1	16.2
102	Gala1-3(Fuca1-2)Galb1-3GlcNAcb	2931.0	164.3	2075.2	206.3	1588.9	136.1	2059.9	97.2	1055.5	30.9	1176.3	202.9	1397.0	154.2	1061.7	194.4	626.4	25.8
103	Gala1-3(Fuca1-2)Galb1-4(Fuca1-3)GlcNAcb	716.3	118.1	105.7	81.4	287.1	149.9	530.5	60.7	391.2	72.3	557.2	67.4	535.3	164.7	197.8	52.5	14.4	15.8
104	Gala1-3(Fuca1-2)Galb1-4(Fuca1-3)GlcNAcb	1069.9	84.2	918.7	117.6	748.4	48.7	821.9	54.0	573.6	58.0	736.5	97.2	603.5	173.2	231.0	20.0	78.2	8.8
105	Gala1-3(Fuca1-2)Galb1-4GlcNAc	804.4	26.0	624.5	96.8	824.1	82.4	643.6	205.2	158.2	38.1	157.3	83.2	259.6	41.6	112.2	36.2	0.1	7.9
106	Gala1-3(Fuca1-2)Galb1-4Glcb	1030.3	55.8	826.4	161.4	761.5	93.1	660.2	30.0	478.6	98.5	670.5	76.9	631.5	65.2	214.2	47.8	87.8	10.3
107	Gala1-3(Fuca1-2)Galb	1068.2	89.2	805.5	25.8	924.6	50.1	952.0	115.3	494.4	49.3	492.5	87.9	747.7	100.7	105.9	25.2	0.1	16.7
108	Gala1-3(Fuca1-2)Galb	771.9	76.5	881.5	36.9	664.9	11.7	904.9	186.8	388.4	28.2	285.0	18.2	438.1	44.9	33.9	10.3	83.5	15.0
109	Gala1-4(Gala1-3)Galb1-4GlcNAcb	803.2	81.5	743.9	82.1	871.6	83.9	736.9	89.3	686.8	107.0	799.1	50.7	2396.9	254.0	89.4	71.6	47.0	41.8
110	Gala1-3GalNAca	277.8	131.1	260.8	66.6	274.6	99.9	329.4	97.2	400.4	60.1	218.9	92.1	605.4	35.8	97.8	88.2	114.6	26.0
111	Gala1-3GalNAca	322.6	92.0	43.3	17.2	29.2	33.8	380.8	92.4	445.2	49.9	124.6	49.9	374.7	72.8	133.0	49.6	73.0	24.7
112	Gala1-3GalNAcb	887.5	97.8	243.8	81.3	0.0	188.7	581.7	44.6	38.9	9.4	29.1	13.7	7.0	53.4	141.9	13.6	82.5	20.5
113	Gala1-3Galb1-4(Fuca1-3)GlcNAcb	716.9	57.6	139.1	61.1	21.8	7.1	390.1	57.1	299.3	79.4	50.3	25.2	125.0	63.2	0.0	97.3	7.0	15.6
114	Gala1-3Galb1-3GlcNAcb	3288.5	307.5	1417.8	57.6	52.8	19.5	1801.2	88.4	714.3	143.9	1960.8	103.4	2166.7	310.5	78.7	18.9	0.1	76.4
115	Gala1-3Galb1-4GlcNAcb	2109.1	208.9	328.7	219.1	811.0	227.8	805.4	100.4	1132.2	99.2	198.9	88.1	2597.1	379.6	119.5	15.9	30.5	31.6
116	Gala1-3Galb1-4Glcb	855.1	97.1	541.9	35.0	127.8	62.8	408.3	31.6	94.3	23.4	442.6	97.4	2175.3	117.5	74.0	10.9	0.1	12.8
117	Gala1-3Galb1-4Glc	3705.1	141.2	2382.6	516.4	3665.2	289.4	2471.1	298.0	2894.7	122.5	1802.8	158.4	9749.0	371.2	332.3	133.4	113.1	250.7
118	Gala1-3Galb	512.9	86.0	481.7	113.2	341.7	87.9	579.2	102.1	61.1	29.9	655.3	111.4	1940.9	61.6	149.9	64.2	3.5	4.2
119	Gala1-4(Fuca1-2)Galb1-4GlcNAcb	498.9	33.8	377.1	18.5	240.4	90.0	230.6	97.2	185.5	43.9	604.4	32.7	980.3	83.6	65.2	26.1	0.1	56.6
120	Gala1-4Galb1-4GlcNAcb	458.6	54.7	314.3	42.6	360.4	52.9	347.4	239.6	435.3	57.5	777.9	123.6	664.2	84.9	123.6	47.4	40.8	6.9
121	Gala1-4Galb1-4GlcNAcb	834.4	43.8	641.7	45.0	570.9	50.8	634.3	72.6	670.9	76.8	1342.4	199.7	1944.7	118.1	378.2	100.7	35.0	24.2
122	Gala1-4Galb1-4Glcb	1177.3	88.1	1158.3	35.5	932.6	12.4	1124.5	38.8	1157.1	18.1	2665.1	163.0	2424.1	237.6	265.4	72.2	48.1	17.3
123	Gala1-4GlcNAcb	2805.9	124.9	938.1	100.5	308.7	176.5	785.3	148.7	495.1	89.6	1428.6	130.2	3239.9	388.5	51.7	12.8	0.1	17.4
124	Gala1-6Glcb	2622.5	519.9	55.5	25.4	0.0	10.2	796.1	285.1	111.3	80.6	2150.3	152.6	329.1	75.8	34.7	18.0	6.1	14.5
125	Galb1-2Galb	3537.7	451.1	1775.0	138.0	1373.7	170.0	1688.4	99.8	1415.6	151.6	2268.3	550.9	4805.0	188.9	1127.2	140.1	32.8	12.9
126	Galb1-3(Fuca1-4)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	135.7	54.2	39.2	34.6	64.4	19.3	148.4	59.1	68.6	18.9	72.3	19.8	25.1	20.5	22.2	6.5	0.1	18.5
127	Galb1-3GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	146.5	14.3	129.0	23.4	123.9	46.3	110.0	48.6	71.3	36.4	155.5	44.2	108.9	40.4	87.9	23.7	25.3	29.5
128	Galb1-3(Fuca1-4)GlcNAc	181.8	82.8	70.2	42.8	137.5	53.4	130.3	115.0	54.6	44.1	64.5	18.0	143.9	62.5	28.7	18.3	0.1	29.4
129	Galb1-3(Fuca1-4)GlcNAc	200.1	40.4	219.7	63.7	182.0	54.2	210.7	119.9	135.4	43.1	160.7	12.2	210.9	48.0	73.4	51.0	54.9	33.7
130	Fuca1-4(Galb1-3)GlcNAcb	243.0	46.7	158.4	43.1	230.3	72.5	339.8	109.5	180.2	24.3	193.0	50.9	314.5	154.1	123.8	12.2	31.6	12.6
131	Galb1-4GlcNAcb1-6GalNAca	1255.4	84.0	484.0	49.2	580.2	60.5	476.4	52.0	629.8	106.1	473.8	157.6	1184.7	60.3	77.3	15.4	68.3	21.9
132	Galb1-4GlcNAcb1-6GalNAc	1397.2	98.1	1701.3	53.4	1276.0	59.5	1467.2	118.9	2023.5	207.9	3618.0	144.2	2750.7	81.8	492.8	42.9	171.6	44.2
133	GlcNAcb1-6(Galb1-3)GalNAca	378.0	52.5	395.4	50.8	522.0	73.4	256.6	90.4	907.2	81.8	771.0	30.8	555.9	30.5	167.9	30.2	36.7	9.3
134	GlcNAcb1-6(Galb1-3)GalNAca	20.8	37.1	56.4	7.8	62.1	39.2	39.6	16.9	113.0	55.2	163.3	106.8	47.6	43.8	57.3	36.1	60.4	41.2
135	Neu5Aca2-6(Galb1-3)GalNAca	0.0	41.5	0.0	22.7	0.0	7.9	0.0	47.6	12.2	8.0	128.4	28.5	0.0	52.7	94.2	17.5	68.9	21.7
136	Neu5Aca2-6(Galb1-3)GalNAca	426.1	75.7	173.5	21.9	50.4	51.1	172.5	28.1	240.3	34.9	65.9	26.3	521.3	87.6	97.8	64.3	29.3	16.4

(Continued)

Supplemental Table 2. (Continued)

137	Neu5Acb2-6(Galb1-3)GalNAca	11.9	19.0	15.3	10.0	34.1	13.0	62.1	21.2	38.2	26.0	55.1	25.4	0.0	56.3	44.2	27.1	5.0	15.2
138	Neu5Aca2-6(Galb1-3)GlcNAcb1-4Galb1-4Glcb	166.8	64.5	152.1	61.1	170.0	40.3	91.4	33.9	151.1	27.6	98.8	52.5	561.0	141.8	6.9	25.3	4.2	24.7
139	Galb1-3GalNAca	69.8	31.5	42.7	13.9	18.0	15.1	22.3	21.4	32.8	10.0	44.6	15.5	171.4	61.9	59.5	9.4	0.1	11.5
140	Galb1-3GalNAca	125.2	53.4	155.3	57.6	103.4	16.8	146.6	68.3	109.6	49.8	225.1	22.0	733.4	33.1	0.0	138.3	44.7	17.9
141	Galb1-3GalNAca	120.2	24.1	159.5	189.3	46.0	15.9	75.5	57.8	16.2	18.2	88.3	60.0	425.0	57.3	74.1	24.3	1.2	31.3
142	Galb1-3GalNAcb	2470.7	115.1	1971.1	35.6	1370.2	97.1	1596.7	21.9	1033.2	31.8	1355.3	109.4	2174.6	240.1	1477.8	119.3	39.9	21.3
143	Galb1-3GalNAcb1-3Gala1-4Galb1-4Glcb	1707.1	107.5	843.9	600.7	1298.1	93.1	1319.9	58.8	980.3	36.2	2368.8	193.7	3045.7	198.2	261.3	30.7	2.5	28.1
144	Galb1-3GalNAcb1-4(Neu5Aca2-3)Galb1-4Glcb	254.3	57.1	287.7	28.3	270.2	71.4	175.7	37.9	203.0	47.3	288.3	104.3	178.3	21.4	123.1	23.6	0.1	4.5
145	Galb1-3GalNAcb1-4Galb1-4Glcb	2325.4	87.5	1896.7	192.4	1647.4	109.1	2013.3	178.6	1283.2	99.8	970.3	124.1	2578.4	148.5	663.4	66.6	5.2	49.4
146	Galb1-3Galb	234.0	36.5	238.5	12.3	131.5	67.6	245.6	16.5	97.8	18.7	166.0	44.2	651.4	166.1	83.5	13.5	62.4	10.0
147	Galb1-3GlcNAcb1-3Galb1-4GlcNAcb	151.4	20.3	74.3	53.5	49.0	42.4	40.4	22.8	26.7	16.3	19.7	11.2	87.7	22.9	0.0	38.0	0.1	19.7
148	Galb1-3GlcNAcb1-3Galb1-4Glcb	1023.0	86.4	679.3	86.7	763.8	105.7	843.2	30.9	567.9	78.8	232.7	39.1	2721.5	230.6	376.3	31.9	63.5	11.6
149	Galb1-3GlcNAcb	101.9	39.3	78.7	54.0	101.7	32.4	114.7	45.3	75.7	29.4	54.5	17.9	1757.4	154.5	10.0	46.0	22.8	6.8
150	Galb1-3GlcNAcb	107.8	27.2	82.0	51.2	113.2	40.2	137.3	77.3	183.2	82.0	361.2	120.3	3162.7	191.6	68.8	8.3	45.8	12.7
151	Galb1-4(Fuca1-3)GlcNAcb	101.0	16.6	53.2	14.3	140.8	84.3	13.0	28.4	71.8	24.5	133.5	35.9	148.7	81.0	48.8	9.1	0.1	15.6
152	Galb1-4(Fuca1-3)GlcNAcb	93.7	90.5	135.4	22.7	244.6	104.6	166.4	69.9	138.3	79.0	199.3	75.6	0.0	147.1	72.8	12.9	22.6	19.0
153	Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	885.6	55.4	671.2	38.3	831.9	74.2	414.6	60.6	787.9	32.2	641.2	61.5	744.9	58.8	470.3	85.5	54.7	26.5
154	Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	128.1	60.2	297.4	42.2	328.8	29.9	240.8	74.5	482.3	87.0	180.1	35.1	463.6	81.9	87.0	5.1	31.0	33.0
155	Galb1-4(6S)Glcb	380.3	73.8	550.2	48.3	758.6	126.0	478.7	88.3	705.0	91.1	1119.4	56.9	1989.3	203.1	90.3	26.7	92.5	22.7
156	Galb1-4(6S)Glcb	545.8	72.0	612.3	63.1	680.1	31.3	375.3	113.9	969.3	15.6	713.4	50.2	1310.9	216.8	128.1	59.4	95.4	51.8
157	Galb1-4GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb	0.0	23.4	0.0	47.9	0.0	13.0	44.1	20.4	3.5	6.3	0.0	22.4	20.5	24.3	0.0	8.7	17.0	11.8
158	Galb1-4GalNAcb1-3(Fuca1-2)Galb1-4GlcNAcb	44.7	151.3	0.0	48.9	0.0	15.7	3.7	8.5	35.1	16.7	163.6	81.8	0.0	13.7	51.7	17.8	65.5	16.7
159	Galb1-4GlcNAcb1-3GalNAca	31.3	14.2	21.4	10.2	14.7	11.6	6.4	26.9	34.9	7.4	56.5	37.4	9.9	21.1	33.0	26.1	28.2	27.2
160	Galb1-4GlcNAcb1-3GalNAc	0.0	10.9	25.9	16.9	16.4	15.8	19.0	14.2	83.5	28.1	107.6	45.2	92.8	22.7	88.0	50.7	137.0	58.3
161	Galb1-4GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	169.4	41.6	110.8	38.3	162.7	36.8	0.0	52.3	186.0	52.8	123.6	37.2	249.4	55.6	62.4	50.2	41.8	25.6
162	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	83.4	61.7	58.4	7.6	23.6	15.7	20.5	14.8	85.8	28.2	77.5	19.4	140.3	21.7	46.2	5.5	146.0	30.3
163	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	70.8	34.0	114.7	12.7	117.5	38.1	155.8	67.7	95.5	53.1	105.8	25.3	3006.4	3518.7	23.6	32.6	10.6	19.0
164	Galb1-4GlcNAcb1-3Galb1-4Glcb	102.0	33.8	47.0	35.5	83.9	27.5	69.4	50.1	87.3	1.9	116.5	31.4	679.5	132.1	55.1	8.0	64.3	11.2
165	Galb1-4GlcNAcb1-3Galb1-4Glcb	3452.1	180.2	2211.2	280.3	1640.4	155.6	1819.8	356.6	1356.3	175.2	573.6	73.8	4771.8	484.7	1059.2	55.8	55.7	27.9
166	Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	226.9	23.1	297.9	41.7	131.8	16.7	276.2	109.2	269.7	35.7	330.6	94.1	6244.6	1436.0	46.0	9.2	16.9	36.4
167	Galb1-4GlcNAcb1-6(Galb1-3)GalNAc	539.9	95.3	571.8	61.1	350.7	38.6	376.5	199.0	487.4	39.6	428.7	44.4	692.2	86.2	215.8	33.5	25.1	30.0
168	Galb1-4GlcNAcb	251.4	45.4	246.6	169.7	331.9	120.2	169.6	101.9	332.1	80.4	480.8	68.7	617.0	19.5	48.9	44.6	29.7	19.7
169	Galb1-4GlcNAcb	213.8	32.5	176.9	9.4	95.2	10.4	221.1	57.6	106.8	17.9	82.8	28.9	552.2	125.3	52.1	27.1	39.5	20.8
170	Galb1-4GlcNAcb	110.1	8.8	27.2	16.7	21.1	11.8	35.8	22.1	41.7	24.9	69.4	33.5	66.6	14.6	0.0	12.4	0.1	11.7
171	Galb1-4Glcb	239.4	71.6	197.4	25.7	133.8	64.2	221.2	88.7	91.0	33.4	182.2	41.0	606.2	109.9	77.2	17.0	74.3	14.0
172	Galb1-4Glcb	167.7	34.9	111.2	50.7	69.1	25.8	74.3	19.6	64.5	39.5	106.4	39.7	264.3	88.2	10.9	21.6	0.1	17.2
173	GlcNAca1-3Galb1-4GlcNAcb	2319.0	996.5	1573.0	132.8	1534.1	81.0	1137.5	234.1	2141.2	82.2	3827.7	442.2	6765.2	297.3	665.5	121.9	30.2	16.8
174	GlcNAca1-6Galb1-4GlcNAcb	846.8	171.9	680.7	74.9	867.8	126.6	500.1	244.6	650.8	89.3	6113.4	1208.7	2103.7	136.6	213.8	40.7	21.5	18.0
175	GlcNAcb1-2Galb1-3GalNAca	4952.9	1153.3	2039.4	53.3	1822.5	94.8	2139.8	184.4	1611.5	135.2	348.4	50.5	6435.6	1357.2	642.6	39.6	14.2	13.8
176	GlcNAcb1-6(GlcNAcb1-3)GalNAca	213.4	71.7	106.3	28.7	133.2	73.2	93.6	125.4	234.6	27.3	178.9	38.8	1567.5	95.1	70.2	22.0	25.4	5.2
177	GlcNAcb1-6(GlcNAcb1-3)GalNAca	736.3	48.7	629.8	45.2	717.6	47.9	600.5	54.8	1497.9	84.9	1304.9	90.1	811.7	118.0	142.7	15.4	121.8	60.3
178	GlcNAcb1-6(GlcNAcb1-3)Galb1-4GlcNAcb	0.0	27.2	0.0	11.0	203.4	162.6	0.0	23.2	77.8	21.8	228.2	84.0	18.1	16.5	9.5	20.2	0.1	14.7
179	GlcNAcb1-3GalNAca	1136.3	29.5	1020.7	62.4	926.6	73.8	400.8	105.6	1114.8	61.2	320.5	50.9	1878.2	248.4	148.8	21.2	60.1	20.9
180	GlcNAcb1-3GalNAca	0.0	34.8	0.0	18.4	0.0	18.0	46.9	15.1	14.0	26.8	26.0	20.1	0.5	48.0	13.1	7.6	0.1	8.2
181	GlcNAcb1-3Galb	259.5	68.5	22.3	12.4	37.1	11.7	0.0	25.5	104.9	17.0	367.2	51.6	0.0	32.2	65.9	9.1	93.0	8.7
182	GlcNAcb1-3Galb1-4GlcNAcb	0.0	26.7	0.0	7.2	0.0	19.2	0.0	6.5	13.3	30.7	0.0	8.8	7.2	29.1	13.5	39.8	25.5	5.3
183	GlcNAcb1-3Galb1-4GlcNAcb	2.7	34.5	25.0	6.2	11.5	4.4	29.0	8.6	63.9	10.0	146.0	35.3	0.0	20.8	63.8	20.4	149.7	51.3
184	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	70.1	34.8	0.0	19.4	63.4	13.3	64.8	52.5	33.0	33.4	34.1	20.4	127.9	57.0	25.5	36.9	39.2	27.8
185	GlcNAcb1-3Galb1-4Glcb	70.1	63.9	102.6	24.4	43.8	14.9	10.1	49.6	92.9	16.6	120.1	32.7	1087.6	128.2	81.2	39.0	168.5	46.0
186	GlcNAcb1-4-MDPLys	10259.3	720.8	8174.7	1041.6	8564.8	1899.9	5942.7	463.2	5399.9	367.8	3022.8	280.3	5488.4	2287.9	1796.2	210.5	25.6	22.6
187	GlcNAcb1-6(GlcNAcb1-4)GalNAca	4821.0	997.6	2272.9	75.2	2520.7	309.0	1827.3	115.2	2698.9	217.0	1206.3	200.0	6895.9	2066.7	62.9	26.8	73.8	67.1
188	GlcNAcb1-4Galb1-4GlcNAcb	7470.6	242.9	3478.3	362.2	2286.3	142.8	3659.5	178.5	1551.3	191.7	1185.5	38.5	5533.7	635.9	1419.1	97.3	54.5	38.9
189	GlcNAcb1-4GlcNAcb1-4GlcNAcb1-4GlcNAcb1-4GlcNAcb1-4GlcNAcb	5834.9	221.5	3095.1	272.8	3083.3	208.7	3010.8	110.9	2585.2	176.6	1067.4	74.5	2308.5	204.6	812.6	30.7	39.5	46.9
190	GlcNAcb1-4GlcNAcb1-4GlcNAcb1-4GlcNAcb1-4GlcNAcb	5904.1	216.7	3650.3	111.4	2523.8	91.6	2726.6	123.2	1388.1	822.3	1689.3	54.4	5030.3	64.6	1468.5	107.0	5.5	23.0
191	GlcNAcb1-4GlcNAcb1-4GlcNAcb	2284.1	107.0	1623.7	183.3	2105.6	87.3	1301.5	75.8	1535.0	83.9	911.7	96.3	2199.5	281.5	486.1	49.1	26.2	15.7
192	GlcNAcb1-6GalNAca	1828.1	44.8	917.2	69.1	624.8	28.9	737.8	74.8	614.0	33.5	1881.5	132.6	1691.3	359.8	256.6	23.9	4.1	13.2
193	GlcNAcb1-6GalNAca	2173.1	39.8	1062.6	136.8	963.1	118.1	982.2	64.2	1416.7	152.5	2259.4	173.6	2583.2	133.9	285.0	73.9	58.5	34.7
194	GlcNAcb1-6Galb1-4GlcNAcb	305.6	92.1	434.5	52.3	241.0	96.3	296.5	86.5	463.0	21.1	5217.9	609.3	570.6	98.7	85.6	63.2	43.8	83.7
195	Glca1-4Glcb	928.8	56.9	974.1	169.8	1275.0	201.4	1200.9	108.2	737.1	85.0	284.4	88.0	1372.4	62.0	134.6	43.9	45.9	29.1
196	Glca1-4Glca	264.1	51.5	229.7	30.8	161.7	13.9	338.8	171.3	124.3	38.8	96.4	39.6	577.0	46.3	25.7	16.6	0.1	30.7
197	Glca1-6Glca1-6Glcb	192.9	28.8	48.7	17.4	16.1	2.4	74.8	24.0	33.0	17.2	80.4	17.9	150.2	29.9	10.7	9.5	17.2	14.4
198	Glcb1-4Glcb	256.7	30.4	228.1	36.6	213.3	20.8	401.6	211.9	158.9	73.7	427.9	186.9	1093.4	121.1	61.6	48.8	43.3	3.8
199	Glcb1-6Glcb	461.9	118.8	115.8	47.7	450.6	139.3	230.8	122.6	569.7	136.9	2902.2	963.7	1277.1	286.7	163.3	44.6	12.6	13.5
200	G-ol	469.7	29.3	373.7	50.9	314.2	21.9	192.6	73.0	532.4	80.4	620.1	97.1	908.8	182.7	106.7	33.6	96.9	21.1
201	GlcAa	94.4	39.8	202.0	52.2	234.5	65.1	128.1	36.0	5.5	30.2	534.5	99.7	471.2	50.2	40.5	12.4	50.8	14.0
202	GlcAb	1347.3	21.7	1077.0	97.5	713.1	86.0	962.1	98.7	912.6	80.6	979.7	65.6	2885.9	430.2	658.2	42.7	1043.5	64.4
203	GlcAb1-3Galb	228.6	55.8	497.4	31.4	2993.2	200.5	383.7	138.4	1895.9	426.3	452.6	37.3	739.7	136.2	72.5	15.8	22.7	13.4
204	GlcAb1-6Galb	500.0	38.0	458.0	47.6	360.1	135.6	273.9	22.3	626.3	42.6	693.9	19.5	1470.2	146.8	80.7	27.9	93.3	32.2
205	KDNa2-3Galb1-3GlcNAcb	2342.8	45.9	3457.8	217.6	2477.6	321.7	2856.8	107.2	4050.3	401.6	7448.6	138.8	6477.9	528.6	146.8	21.9	97.8	19.5
206	KDNa2-3Galb1-4GlcNAcb	185.9	36.1	330.6	64.2	72.7	127.1	62.9	18.7	319.3	44.8	381.2	142.5	431.5	67.7	210.6	206.8	59.5	24.1
207	Mana1-2Mana1-2Mana1-3Mana	4.9	49.9	6.5	20.1	5.9	12.2	15.8	7.2	14.7	22.2	15.2	20.2	76.6	23.1	5.8	23.0	11.4	16.5
208	Mana1-2Mana1-6(Mana1-2Mana1-3)Mana	0.0	38.3	0.0	23.2	0.0	17.5	19.6	21.4	39.3	6.6	223.9	115.3	0.0	35.6	76.8	18.0	99.2	16.7
209	Mana1-2Mana1-3Mana	146.8	45.9	34.3	17.4	24.5	22.5	46.7	63.5	163.3	22.1	54.8	53.5	260.9	99.2	24.3	21.2	31.1	11.9
210	Mana1-6(Mana1-2Mana1-3)Mana1-6(Mana1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	191.9	94.1	23.2	54.2	0.1	27.9	56.7	12.6	78.9	23.9	160.2	4.3	297.1	65.7	71.6	32.3	144.5	19.6
211	Mana1-2Mana1-6(Mana1-3)Mana1-6(Mana1-2Mana1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	204.6	44.7	134.7	14.8	115.7	69.7	300.8	281.5	135.4	24.8	161.1	34.7	354.2	107.8	97.7	26.0	51.0	8.9
212	Mana1-2Mana1-6(Mana1-2Mana1-3)Mana1-6(Mana1-2Mana1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	119.6	113.0	77.1	59.8	152.1	59.1	54.6	36.0	207.4	16.9	168.2	103.3	686.9	38.3	110.3	103.5	176.7	82.9
213	Mana1-6(Mana1-3)Mana	821.0	101.5	518.2	20.1	439.5	105.3	333.4	174.7	229.5	5.7	83.7	33.0	1498.4	93.0	128.9	48.9	40.2	59.2
214	Mana1-2Mana1-2Mana1-6(Mana1-3)Mana	1007.5	73.3	673.0	114.1	1109.5	244.0	592.9	210.3	863.2	77.9	2098.1	819.5	6942.3	931.9	271.7	38.0	12.8	57.0
215	Mana1-6(Mana1-3)Mana1-6(Mana1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	332.7	101.9	296.1	108.1	320.0	94.9	215.5	115.4	309.8	67.3	392.5	85.0	1392.6	270.5	254.4	59.6	365.2	119.3
216	Mana1-6(Mana1-3)Mana1-6(Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	173.4	40.7	99.7	34.4	158.4	54.0	129.1	31.7	170.2	41.7	213.1	93.2	537.8	89.8	27.9	27.0	31.5	6.6
217	Manb1-4GlcNAcb	123.3	10.4	209.1	26.2	133.1	59.5	98.2	19.8	107.6	18.2	143.5	78.0	981.4	48.9	29.8	26.8	16.1	28.8
218	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	56.2	26.9	93.2	24.4	36.7	67.1	38.2	36.1	58.9	18.3	104.1	18.9	133.7	25.5	0.0	14.2	50.7	30.0
219	(3S)Galb1-4(Fuca1-3)(6S)GlcNAcb	200.9	61.3	195.7	39.0	65.8	44.9	125.8	6.6	127.8	34.9	176.7	61.3	245.0	106.9	47.5	24.6	0.1	25.4
220	Fuca1-2(6S)Galb1-4GlcNAcb	0.0	67.5	0.0	49.5	0.0	70.9	0.0	47.1	0.0	48.7	0.0	153.7	0.0	66.3	0.0	33.9	7.4	17.6
221	Fuca1-2Galb1-4(6S)GlcNAcb	33.6	52.6	57.3	71.4	0.0	43.5	683.6	230.5	64.0	54.6	6.2	19.7	3.2	128.0	0.0	10.9	60.8	28.1
222	Fuca1-2(6S)Galb1-4(6S)Glcb	111.5	35.3	89.1	19.5	52.2	18.1	95.5	24.9	114.8	31.7	130.8	52.6	199.7	96.6	71.5	42.7	35.9	14.9
223	Neu5Aca2-3Galb1-3GalNAca	164.4	81.3	75.3	46.9	60.6	14.6	129.1	47.8	188.8	30.4	141.0	47.2	1240.2	644.4	42.8	18.6	72.9	41.9
224	Neu5Aca2-3Galb1-3GalNAca	114.3	46.9	42.1	36.3	54.8	11.5	87.7	36.1	16.5	9.5	216.2	52.1	320.6	58.0	78.0	52.3	0.1	5.8
225	GalNAcb1-4(Neu5Aca2-8Neu5Aca2-8Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	80.5	44.8	139.7	12.9	139.5	30.9	115.2	38.7	210.2	37.6	167.0	30.1	223.5	32.0	152.1	90.7	80.4	11.4
226	GalNAcb1-4(Neu5Aca2-8Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	181.3	102.8	109.4	83.9	793.5	215.8	124.0	65.4	230.1	54.9	444.5	129.5	208.2	76.1	105.1	60.2	182.0	71.0
227	Neu5Aca2-8Neu5Aca2-8Neu5Aca2-3Galb1-4Glcb	47.2	49.4	104.9	25.4	123.1	50.5	73.1	79.5	198.0	83.3	191.5	50.2	137.8	61.6	152.9	110.7	46.2	21.3
228	GalNAcb1-4(Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	0.0	31.6	65.2	26.5	47.5	44.3	18.1	61.8	123.7	47.8	342.5	62.1	85.7	18.9	12.7	52.9	47.7	34.7
229	Neu5Aca2-8Neu5Aca2-8Neu5Aca	54.7	30.2	74.2	49.9	83.3	29.4	0.0	20.0	190.5	24.7	178.2	106.6	461.1	256.2	27.9	40.9	84.8	26.6
230	Neu5Aca2-3(6S)Galb1-4(Fuca1-3)GlcNAcb	0.0	12.1	0.0	17.2	0.6	15.3	15.1	9.5	24.2	8.7	34.4	27.8	0.0	16.8	32.5	47.0	35.2	16.9
231	GalNAcb1-4(Neu5Aca2-3)Galb1-4GlcNAcb	11.6	33.0	16.0	9.3	0.0	96.7	50.7	33.5	89.6	12.3	176.5	79.2	0.0	9.8	93.6	6.8	152.2	35.1
232	GalNAcb1-4(Neu5Aca2-3)Galb1-4GlcNAcb	0.0	17.0	0.0	3.0	0.0	52.1	0.0	71.4	55.0	15.5	12.0	2.2	0.0	67.5	9.3	21.2	5.4	23.4
233	GalNAcb1-4(Neu5Aca2-3)Galb1-4Glcb	0.0	19.7	6.5	14.8	0.0	141.0	24.6	18.9	42.9	10.1	114.9	30.7	92.7	167.4	78.9	32.4	112.8	31.0
234	Neu5Aca2-3Galb1-3GalNAcb1-4(Neu5Aca2-3)Galb1-4Glcb	51.3	14.3	68.5	9.9	37.2	25.6	90.7	13.3	66.8	27.7	66.3	36.2	113.2	67.8	56.5	34.7	41.3	12.9
235	Neu5Aca2-6(Neu5Aca2-3)GalNAca	51.4	19.5	56.2	35.8	54.9	45.3	72.1	9.2	74.6	23.6	75.8	12.3	63.1	61.7	63.5	41.4	20.0	18.1
236	Neu5Aca2-3GalNAca	56.7	32.7	85.1	40.8	18.3	17.8	53.6	38.7	60.5	58.1	73.3	4.5	89.4	212.2	24.0	8.8	12.4	17.6
237	Neu5Aca2-3GalNAcb1-4GlcNAcb	34.2	22.1	65.0	5.0	281.5	242.3	126.3	197.4	460.9	156.7	46.9	22.1	0.0	60.7	61.5	61.8	33.6	16.7
238	Neu5Aca2-3Galb1-3(6S)GlcNAc	248.8	32.1	159.9	48.1	91.7	21.5	106.3	58.8	129.5	56.7	175.7	123.3	1314.0	116.0	70.1	55.5	90.2	3.0
239	Neu5Aca2-3Galb1-3(Fuca1-4)GlcNAcb	102.3	21.0	73.8	15.3	67.8	26.2	579.2	254.6	120.5	42.5	22.6	37.2	151.6	43.9	29.9	48.6	16.4	19.2
240	Neu5Aca2-3Galb1-3(Fuca1-4)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	1155.4	195.4	1200.4	184.7	1201.6	60.2	946.3	75.9	1806.0	268.7	2436.7	125.3	2225.8	91.9	792.4	128.5	88.3	27.0
241	Neu5Aca2-3Galb1-4(Neu5Aca2-3Galb1-3)GlcNAcb	31.4	28.3	47.3	19.9	0.0	104.8	84.8	111.8	27.5	9.4	35.1	17.4	0.0	221.2	19.6	3.5	46.2	25.4
242	Neu5Aca2-3Galb1-3(6S)GalNAca	200.3	17.0	79.7	11.2	77.0	28.7	65.1	43.6	68.1	16.6	172.6	28.7	77.7	57.0	38.1	12.5	32.5	33.1
243	Neu5Aca2-6(Neu5Aca2-3Galb1-3)GalNAca	68.6	7.9	15.9	6.7	23.1	4.0	46.5	24.0	5.2	4.4	37.6	17.5	38.2	16.8	0.0	14.3	0.1	10.6
244	Neu5Aca2-6(Neu5Aca2-3Galb1-3)GalNAca	762.7	89.7	655.4	44.2	541.0	59.9	475.6	79.5	692.7	33.3	1292.1	106.2	1155.4	145.6	226.8	23.6	67.0	40.0
245	Neu5Aca2-3Galb	84.2	33.7	70.6	114.2	10.9	23.3	19.6	16.6	65.2	29.0	55.7	12.0	88.2	11.9	2.9	8.8	12.8	11.0
246	Neu5Aca2-3Galb1-3GalNAcb1-3Gala1-4Galb1-4Glcb	160.6	30.9	131.5	19.3	149.9	22.0	275.8	154.3	155.2	20.4	529.6	171.2	1407.3	265.8	39.5	6.5	17.5	19.7
247	Neu5Aca2-3Galb1-3GlcNAcb1-3Galb1-4GlcNAcb	36.8	15.4	41.2	18.2	60.9	24.9	57.2	18.3	57.1	14.5	42.2	13.8	126.7	9.4	49.8	20.9	64.9	24.9
248	Fuca1-2(6S)Galb1-4Glcb	56.4	32.3	30.8	18.1	22.7	73.6	0.0	175.0	48.7	8.6	19.0	25.9	540.8	87.5	65.0	32.6	48.3	14.3
249	Neu5Aca2-3Galb1-3GlcNAcb	272.8	117.8	231.4	76.6	358.2	31.5	175.1	14.5	207.8	17.1	396.0	208.6	7383.5	843.7	106.0	33.5	147.3	20.0
250	Neu5Aca2-3Galb1-3GlcNAcb	227.9	65.3	213.1	96.7	270.8	118.1	320.3	112.6	441.2	89.3	1088.9	223.7	5129.9	379.8	39.1	15.7	8.5	21.6
251	Neu5Aca2-3Galb1-4(6S)GlcNAcb	39.3	16.5	58.7	8.7	147.3	215.1	69.2	39.8	64.6	25.8	96.5	23.4	112.1	60.2	19.7	5.3	52.5	15.5
252	Neu5Aca2-3Galb1-4(Fuca1-3)(6S)GlcNAcb	122.5	15.9	146.9	59.9	144.1	69.9	230.2	100.9	156.8	34.1	150.1	77.2	254.4	63.0	76.7	23.2	50.6	35.6
253	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	156.7	69.9	36.5	14.0	0.0	87.2	128.7	39.9	170.3	63.8	67.6	19.2	252.7	60.5	90.8	27.3	25.8	14.7
254	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb	44.7	11.2	31.3	10.1	20.5	17.5	63.6	24.7	47.7	19.0	136.0	14.4	0.0	57.9	93.3	30.7	58.1	17.8
255	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb	0.0	34.9	0.0	16.4	0.0	14.2	0.0	209.2	0.0	30.8	0.0	10.4	0.0	51.2	0.0	14.4	1.8	15.7
256	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb	0.0	30.7	13.0	14.3	20.1	12.1	0.0	22.2	15.8	20.8	51.7	37.9	30.7	30.9	51.4	15.8	23.4	17.7
257	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4GlcNAcb	438.8	75.4	278.0	37.8	135.3	15.3	181.7	15.5	290.5	45.3	0.0	129.4	776.7	101.6	58.3	37.3	0.1	10.3
258	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	0.0	34.1	0.0	14.8	45.7	16.7	0.0	65.4	85.3	10.7	35.3	27.2	377.8	49.1	47.6	85.0	0.1	18.7
259	Neu5Aca2-3Galb1-4GlcNAcb	106.4	5.0	120.3	23.8	70.0	34.7	52.4	23.4	76.0	34.4	62.3	21.1	131.0	47.6	39.4	16.2	42.8	36.0
260	Neu5Aca2-3Galb1-4GlcNAcb	0.0	12.4	0.0	36.1	216.8	92.4	15.2	235.8	71.1	57.3	0.0	46.7	0.0	86.0	6.7	19.9	0.1	14.9
261	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	132.6	20.4	119.1	27.6	54.0	56.8	73.4	24.2	53.7	23.8	39.1	27.0	331.1	135.3	54.3	21.6	78.2	10.5
262	Fuca1-2Galb1-4(6S)Glcb	49.5	18.0	0.0	35.9	36.9	15.9	326.5	119.6	54.4	15.9	131.1	37.9	686.2	76.6	33.6	16.8	37.0	37.3
263	Neu5Aca2-3Galb1-4Glcb	303.3	67.6	310.1	86.6	193.3	62.0	220.5	73.3	312.0	113.0	317.5	95.4	1184.2	64.5	112.5	15.4	205.3	60.9
264	Neu5Aca2-3Galb1-4Glcb	661.4	30.1	617.9	33.8	613.4	154.0	783.9	136.9	576.3	36.4	332.1	52.8	2667.3	122.0	148.5	32.0	26.1	26.0
265	Neu5Aca2-6GalNAca	69.1	20.1	84.4	22.0	50.1	9.2	45.6	12.4	73.0	11.2	182.0	22.6	32.1	27.1	59.0	9.7	1.2	38.4
266	Neu5Aca2-6GalNAcb1-4GlcNAcb	0.0	21.1	0.0	10.6	5.7	17.4	20.6	26.5	1.8	5.5	7.3	22.8	0.0	19.8	7.4	12.7	0.1	19.5
267	Neu5Aca2-6Galb1-4(6S)GlcNAcb	196.1	52.7	89.3	8.6	81.1	17.6	67.0	7.4	65.3	21.9	163.3	17.3	0.0	24.2	92.9	26.7	3.0	11.7
268	Neu5Aca2-6Galb1-4GlcNAcb	78.7	16.7	1.2	15.3	36.1	22.4	42.5	17.7	42.0	40.0	38.9	19.9	81.3	14.5	27.4	19.8	0.1	16.7
269	Neu5Aca2-6Galb1-4GlcNAcb	94.2	8.9	76.4	40.9	157.3	20.3	264.9	311.6	80.2	19.1	27.0	13.4	106.8	73.4	0.0	19.5	40.8	9.3
270	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	71.5	10.0	146.0	8.8	169.9	53.2	142.8	42.9	79.9	35.0	234.0	116.8	597.5	63.0	49.5	24.6	32.2	20.0
271	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	47.3	21.1	60.8	18.6	25.9	24.6	87.1	124.6	80.4	21.1	8.4	19.7	56.1	4.4	11.8	38.1	50.4	23.2
272	Neu5Aca2-6Galb1-4Glcb	17.1	21.0	43.3	17.0	232.0	91.1	23.8	12.6	172.5	97.6	23.8	8.1	415.1	141.8	20.6	10.5	26.9	22.4
273	Neu5Aca2-6Galb1-4Glcb	76.8	11.6	96.1	21.5	92.6	12.3	83.5	36.0	121.9	18.8	406.3	42.2	1098.1	115.2	62.9	39.2	46.0	16.0

(Continued)

274	Neu5Aca2-6Galb	67.0	49.2	39.6	14.7	91.0	55.1	72.2	25.4	61.7	16.6	516.0	356.8	808.3	59.5	58.9	54.2	14.6	15.3
275	Neu5Aca2-8Neu5Aca	119.2	7.5	89.0	36.8	83.9	49.6	51.3	30.6	87.0	18.1	35.2	21.1	510.7	162.5	62.8	92.9	163.4	139.2
276	Neu5Aca2-8Neu5Aca2-3Galb1-4Glcb	11.9	14.8	22.8	16.2	37.1	33.9	54.9	10.4	76.9	35.8	102.2	38.7	44.2	11.9	40.1	7.2	40.0	23.4
277	Galb1-3(Fuca1-4)GlcNAcb1-3Galb1-3(Fuca1-4)GlcNAcb	131.1	13.1	48.5	4.7	0.0	38.5	37.9	14.7	43.7	16.5	114.6	19.5	40.4	12.2	116.7	4.9	65.9	12.0
278	Neu5Acb2-6GalNAca	0.0	17.4	12.0	48.5	53.8	62.5	27.7	14.9	0.0	16.5	0.0	11.2	41.2	34.7	23.3	39.4	0.1	21.3
279	Neu5Acb2-6Galb1-4GlcNAcb	26.1	15.1	35.5	25.7	25.0	3.9	71.0	18.6	24.1	10.6	54.0	55.4	110.5	39.9	0.0	102.6	11.3	20.4
280	Neu5Gca2-3Galb1-3(Fuca1-4)GlcNAcb	372.2	50.3	417.0	147.8	278.6	62.2	184.2	23.0	337.8	51.0	216.7	93.3	798.3	289.8	153.9	49.7	47.3	7.0
281	Neu5Gca2-3Galb1-3GlcNAcb	103.5	27.5	60.6	58.2	168.8	49.7	88.9	43.5	51.6	30.1	446.0	127.3	3762.7	260.8	46.0	38.9	25.9	22.0
282	Neu5Gca2-3Galb1-4(Fuca1-3)GlcNAcb	106.6	16.4	98.9	21.6	58.9	27.4	147.6	106.2	60.6	27.9	74.5	25.8	287.7	112.1	31.8	4.4	22.1	6.6
283	Neu5Gca2-3Galb1-4GlcNAcb	98.9	64.8	102.2	14.3	129.5	21.2	130.3	46.7	128.6	62.8	105.6	24.0	198.1	107.3	38.9	16.4	62.4	13.3
284	Neu5Gca2-3Galb1-4Glcb	95.0	38.9	201.7	43.2	347.8	169.0	117.8	313.2	167.6	69.7	234.2	36.4	797.9	38.9	145.0	58.5	13.6	32.6
285	Neu5Gca2-6GalNAca	153.0	51.1	67.7	49.0	431.5	352.8	174.2	90.0	89.0	33.1	142.4	43.6	137.4	140.9	95.1	70.8	60.8	12.1
286	Neu5Gca2-6Galb1-4GlcNAcb	168.9	44.9	118.7	81.4	139.2	28.4	206.8	67.3	178.2	62.5	334.8	85.3	581.6	173.4	67.1	34.3	32.0	10.9
287	Neu5Gca	167.2	32.6	220.6	20.2	190.5	55.3	168.7	80.3	49.7	30.6	133.3	105.7	830.1	55.2	0.0	24.5	0.1	40.5
288	Neu5Aca2-3Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	208.8	35.9	276.7	84.5	145.0	30.3	202.1	61.0	231.8	52.0	245.3	72.0	460.9	169.8	67.7	20.7	70.3	43.5
289	Galb1-3GlcNAcb1-3Galb1-3GlcNAcb	189.9	76.6	78.3	81.1	282.4	26.7	309.4	56.3	438.5	57.8	229.5	36.3	6567.7	376.2	93.4	34.1	0.1	8.2
290	Galb1-4(Fuca1-3)(6S)GlcNAcb	324.8	57.3	440.8	141.8	542.6	177.0	382.7	102.0	373.7	76.4	320.4	47.1	786.8	185.2	243.9	17.5	51.9	11.2
291	Galb1-4(Fuca1-3)(6S)Glcb	1027.1	50.2	1053.5	193.8	1084.3	121.4	1262.9	94.2	1152.6	113.8	1053.0	103.9	1608.0	202.6	431.5	50.9	67.3	23.3
292	Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-3(Fuca1-4)GlcNAcb	32.1	28.8	35.6	31.8	30.4	25.2	74.4	74.5	68.4	40.9	52.8	26.7	0.0	323.9	58.1	20.8	0.1	20.9
293	Galb1-4GlcNAcb1-3Galb1-3GlcNAcb	34.1	36.9	40.0	15.7	2.7	12.8	20.3	9.9	26.8	7.3	55.9	42.4	2412.4	112.3	58.6	34.2	53.6	28.1
294	Neu5Aca2-3Galb1-3GlcNAcb1-3Galb1-3GlcNAcb	162.9	61.0	116.5	14.1	150.4	72.0	85.9	53.2	164.8	11.2	110.5	31.5	4626.9	758.2	33.0	20.7	46.7	11.1
295	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-3GlcNAcb	0.0	16.3	21.0	18.4	39.8	14.0	52.3	56.0	0.0	20.8	59.8	28.8	2994.8	404.3	30.8	9.0	5.5	25.0
296	4S(3S)Galb1-4GlcNAcb	128.1	38.6	108.4	32.7	92.9	35.4	112.1	27.1	164.1	36.9	408.1	88.9	429.8	46.0	29.8	8.4	67.7	14.8
297	(6S)Galb1-4(6S)GlcNAcb	57.8	42.0	51.9	31.3	90.5	48.6	131.6	52.2	36.5	12.6	367.1	134.4	193.5	170.5	22.9	20.5	0.1	39.8
298	(6P)Glcb	326.6	16.8	343.3	34.8	220.4	77.6	180.3	61.2	427.4	96.4	523.2	51.3	1755.2	46.5	119.7	43.3	100.1	35.1
299	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-6(Galb1-3)GalNAca	129.5	53.6	124.2	33.5	103.8	21.6	126.2	8.4	109.7	38.4	205.3	32.7	684.9	258.1	70.5	26.5	43.0	25.5
300	Galb1-3Galb1-4GlcNAcb	93.3	36.7	162.1	61.7	247.9	52.0	124.2	18.9	125.3	5.2	116.6	43.8	1025.4	61.6	0.0	124.6	26.6	16.2
301	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	22.5	18.0	86.5	22.6	58.9	29.1	50.0	49.4	93.8	19.9	143.4	66.6	392.6	60.2	63.2	50.8	41.1	7.8
302	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3)Galb1-4GlcNAc	159.9	152.4	51.9	28.4	50.4	27.6	103.7	9.8	70.0	44.0	37.4	51.6	289.6	206.3	0.0	74.2	15.2	34.5
303	GlcNAcb1-6(Galb1-4GlcNAcb1-3)Galb1-4GlcNAc	271.4	46.6	151.2	18.9	90.1	16.8	145.7	67.8	137.0	50.3	147.1	30.7	303.3	198.1	77.2	20.7	34.2	16.1
304	Galb1-4GlcNAca1-6Galb1-4GlcNAcb	500.2	78.3	150.9	80.7	345.2	25.4	247.0	81.4	51.3	46.2	70.1	9.0	1436.7	155.9	38.9	28.6	0.1	17.3
305	Galb1-4GlcNAcb1-6Galb1-4GlcNAcb	0.0	54.0	0.0	67.6	10.5	25.4	0.0	50.7	7.0	157.9	0.0	62.7	0.0	75.8	0.0	50.7	0.1	110.4
306	GalNAcb1-3Galb	6.5	49.9	1.0	12.5	143.3	120.4	45.9	18.0	26.0	2.6	0.0	28.0	669.0	16.5	52.7	20.4	16.5	50.6
307	GlcAb1-3GlcNAcb	270.7	25.1	106.0	38.6	130.2	33.7	161.0	159.2	127.1	22.4	725.2	143.8	2831.7	183.5	75.6	12.2	0.1	6.5
308	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	53.6	43.1	63.9	43.9	209.3	58.0	106.7	33.3	41.3	48.6	88.0	62.9	155.5	128.3	43.1	74.0	49.9	19.8
309	GlcNAcb1-3Man	371.5	127.9	248.7	24.2	157.0	40.0	152.5	54.1	174.2	42.1	495.9	30.3	1443.3	157.2	48.3	30.2	0.1	28.3
310	GlcNAcb1-4GlcNAcb	252.8	71.5	153.2	33.2	152.0	65.2	89.9	27.7	169.1	32.6	204.7	50.2	1429.7	199.0	80.5	26.0	3.5	19.2
311	GlcNAcb1-4GlcNAcb	128.9	22.1	73.1	15.5	45.7	54.2	0.0	106.1	41.5	17.3	311.6	50.1	1713.6	218.8	31.5	25.6	20.3	12.1
312	MurNAcb1-4GlcNAcb	2281.6	291.4	1445.2	65.4	1166.4	95.5	1252.5	132.4	1433.8	128.5	2037.8	371.3	5927.1	277.3	318.2	95.3	29.2	31.5
313	Mana1-6Manb	19.7	20.3	59.4	46.0	161.4	84.9	19.5	31.9	112.2	44.7	106.4	28.3	1004.4	90.8	52.7	20.7	29.8	31.4
314	Mana1-6(Mana1-3)Mana1-6(Mana1-3)Manb	91.5	53.0	52.9	53.3	157.2	118.0	189.0	35.4	242.0	103.4	113.1	45.4	5057.1	2460.9	38.6	11.0	17.9	23.2
315	Mana1-2Mana1-6(Mana1-3)Mana1-6(Mana1-2Mana1-2Mana1-3)Mana	28.8	49.9	0.0	34.4	53.5	77.2	50.7	30.1	125.7	35.1	44.3	12.3	1547.5	180.6	18.9	24.9	0.1	29.7
316	Mana1-2Mana1-6(Mana1-2Mana1-3)Mana1-6(Mana1-2Mana1-2Mana1-3)Mana	254.7	104.5	32.8	38.1	70.0	28.4	109.9	26.1	105.2	62.8	73.0	29.4	593.6	105.2	82.6	19.4	34.6	11.2
317	Neu5Aca2-3Galb1-4GlcNAcb1-6(Neu5Aca2-3Galb1-3)GalNAca	0.0	14.6	0.0	70.4	25.3	10.0	23.5	14.2	26.0	24.7	42.8	12.8	189.1	66.6	25.6	7.7	21.8	13.7
318	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	0.0	13.4	19.2	24.8	34.0	12.6	118.4	211.6	2.9	45.9	163.8	176.0	165.6	117.5	52.8	29.9	13.1	4.1
319	Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	175.4	38.5	142.9	30.7	122.4	6.0	121.2	85.2	179.5	52.4	223.7	103.5	233.2	31.7	75.7	9.3	56.2	28.3
320	GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	166.3	40.9	153.3	40.8	188.4	18.3	207.2	35.0	32.6	33.6	173.0	125.8	263.3	83.9	52.4	31.5	0.1	48.0
321	Neu5Aca2-8Neu5Acb	90.8	29.0	60.3	7.4	87.8	17.3	66.0	15.3	182.9	50.5	113.0	64.6	350.6	38.1	27.1	43.9	39.7	15.2
322	Neu5Aca2-8Neu5Aca2-8Neu5Acb	17.8	31.7	45.1	39.2	94.3	15.1	72.4	22.0	24.6	32.5	159.3	106.7	230.8	219.6	14.6	12.8	19.6	18.7
323	Neu5Gcb2-6Galb1-4GlcNAc	102.3	45.9	110.7	14.2	90.8	25.0	50.3	31.9	18.9	15.9	84.8	39.9	1169.4	79.8	0.0	3.8	0.1	32.8
324	Galb1-3GlcNAcb1-2Mana1-6(Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	6.2	46.4	430.4	74.2	396.6	306.3	278.5	98.5	369.0	69.9	396.4	103.9	627.4	16.6	67.0	21.1	101.8	48.3
325	Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	2.3	25.0	13.7	16.0	22.6	19.8	0.0	23.1	9.7	28.4	0.0	44.1	518.7	72.3	0.0	16.5	0.1	45.7
326	Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	445.1	42.7	330.4	119.0	460.8	58.6	338.7	67.0	436.0	110.7	374.8	107.3	650.3	67.9	163.8	97.6	33.2	14.5
327	Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-6(Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	95.8	39.1	61.2	28.8	35.0	9.1	99.2	33.9	20.4	15.2	46.6	24.3	264.2	70.8	2.2	18.0	0.1	10.7
328	Neu5,9Ac2a2-3Galb1-4GlcNAcb	402.8	85.6	312.4	102.7	281.1	79.4	362.3	231.4	228.9	49.1	354.6	17.0	1149.1	51.4	327.8	101.9	323.9	33.8
329	Neu5,9Ac2a2-3Galb1-3GlcNAcb	120.8	37.1	90.9	44.5	151.3	106.7	187.2	62.3	123.9	13.2	565.9	172.2	3580.3	468.2	34.2	14.3	0.1	16.4
330	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-3GlcNAcb	264.6	59.3	275.1	57.9	179.1	39.0	548.3	374.9	211.5	19.5	244.2	118.7	4941.7	876.2	128.5	42.2	55.9	43.4
331	Neu5Aca2-3Galb1-3(Fuca1-4)GlcNAcb1-3Galb1-3(Fuca1-4)GlcNAcb	971.2	61.4	893.1	68.3	625.4	51.3	838.0	46.5	903.0	61.8	1147.2	54.9	1195.6	497.7	321.2	36.2	0.1	19.3
332	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	141.9	42.7	286.8	37.6	124.3	81.3	0.0	496.5	125.1	32.0	233.1	83.7	474.6	80.5	109.3	53.1	13.4	8.6
333	Gala1-4Galb1-4GlcNAcb1-3Galb1-4Glcb	0.0	132.8	0.0	82.0	86.2	92.2	0.0	87.8	0.0	91.9	80.0	184.7	4995.4	957.0	0.0	43.2	0.1	109.3
334	GalNAcb1-3Gala1-4Galb1-4GlcNAcb1-3Galb1-4Glcb	107.3	85.0	193.5	93.7	143.8	12.4	83.7	66.8	69.0	13.9	602.5	32.8	4110.8	991.3	130.2	48.5	34.5	10.7
335	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	0.0	28.5	392.1	59.5	308.1	57.4	289.8	21.3	36.7	32.4	0.0	5.8	941.7	131.1	0.0	22.9	14.9	39.8
336	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	221.6	53.0	668.0	31.0	345.2	15.2	465.1	31.3	218.0	72.9	158.6	100.6	1224.1	73.0	118.1	30.6	92.4	37.1
337	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-6(Neu5Aca2-3Galb1-3)GalNAc	376.1	69.1	356.7	54.2	580.3	43.1	237.7	83.6	769.2	58.0	548.3	52.4	446.0	167.3	101.6	71.1	32.3	25.2
338	GlcNAca1-4Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	0.0	7.9	0.4	30.4	0.0	69.8	7.4	11.4	41.1	32.9	0.4	39.6	243.8	47.5	29.6	14.2	27.2	27.0
339	GlcNAca1-4Galb1-4GlcNAcb	0.0	15.3	0.0	12.0	4.1	21.3	0.0	31.6	0.0	34.2	5.5	15.7	168.8	41.3	14.7	13.5	0.1	30.5
340	GlcNAca1-4Galb1-3GlcNAcb	4.6	51.5	34.0	59.0	66.2	13.1	50.7	64.0	136.6	43.5	75.5	63.0	979.5	151.3	23.4	55.3	23.6	23.2
341	GlcNAca1-4Galb1-4GlcNAcb1-3Galb1-4Glcb	0.0	24.7	0.0	16.4	25.4	39.1	69.6	15.9	0.0	22.6	72.7	44.0	661.9	94.5	27.3	31.2	0.1	26.5
342	GlcNAca1-4Galb1-4GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb	216.7	21.1	231.3	39.3	255.4	47.4	388.5	81.3	407.8	90.7	460.6	36.2	1011.8	131.6	183.5	59.3	67.7	19.0
343	GlcNAca1-4Galb1-4GlcNAcb1-3Galb1-4GlcNAcb	64.0	37.0	39.5	11.6	40.6	16.1	93.8	31.1	7.0	17.5	59.0	42.8	376.7	77.4	44.3	29.4	0.1	27.3
344	GlcNAca1-4Galb1-3GalNAc	68.2	40.6	162.9	17.4	263.2	61.8	229.9	25.8	723.2	123.5	2465.4	341.7	832.2	74.5	92.7	44.0	46.6	33.8
345	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	0.0	3.7	13.6	27.0	0.0	164.3	0.0	69.9	10.3	33.8	186.7	64.1	243.4	65.2	18.4	15.5	2.7	12.4
346	Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	368.4	180.0	557.9	36.8	451.7	127.6	364.9	128.1	529.9	112.5	195.2	69.4	1256.5	153.7	90.6	31.2	65.7	27.1
347	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6Manb1-4GlcNAcb1-4GlcNAc	0.0	23.0	46.3	32.1	528.2	961.7	87.1	26.7	7.0	33.0	0.0	23.3	614.2	36.5	0.0	17.6	15.2	41.1
348	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3Manb1-4GlcNAcb1-4GlcNAc	69.8	19.6	105.6	37.9	97.6	52.4	121.2	72.2	134.3	33.9	84.2	19.1	772.9	157.7	188.6	158.3	70.7	49.0
349	Galb1-4GlcNAcb1-2Mana1-3Manb1-4GlcNAcb1-4GlcNAc	167.6	33.4	87.6	43.2	78.8	13.6	91.5	47.8	26.7	16.1	0.0	60.1	265.6	79.4	38.7	15.1	0.1	10.7
350	Galb1-4GlcNAcb1-2Mana1-6Manb1-4GlcNAcb1-4GlcNAc	47.9	28.4	31.8	31.0	31.5	3.7	38.8	29.6	73.1	33.5	32.9	26.8	118.9	14.8	0.0	14.9	29.5	9.3
351	Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	354.6	100.3	249.0	25.5	202.6	61.7	283.0	241.9	106.6	52.0	146.1	31.6	409.6	118.3	95.0	48.2	0.1	31.7
352	GlcNAcb1-2Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	75.1	34.4	39.4	26.8	27.7	69.2	35.1	55.2	57.1	37.4	123.4	26.0	192.8	19.3	33.9	13.4	17.4	29.8
353	Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	190.5	63.3	341.8	75.8	224.4	84.0	199.2	58.2	241.2	45.0	310.4	95.0	572.8	159.3	225.8	83.3	87.6	35.2
354	Galb1-3GlcNAcb1-2Mana1-6(Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	134.8	20.3	85.3	246.1	278.0	81.0	371.2	140.5	348.3	105.2	137.8	71.9	642.1	96.4	66.3	22.5	80.9	70.0
355	(6S)GlcNAcb1-3Galb1-4GlcNAcb	191.6	46.2	358.8	146.8	56.3	118.7	0.0	365.3	110.9	20.1	229.0	74.1	361.1	112.8	30.5	25.7	29.3	34.7
356	KDNa2-3Galb1-4(Fuca1-3)GlcNAc	144.9	20.8	106.1	32.1	190.1	98.2	172.1	20.0	231.5	87.6	209.6	65.3	797.2	83.6	48.9	22.2	35.3	38.9
357	KDNa2-6Galb1-4GlcNAc	63.2	42.4	84.3	6.0	63.1	2.3	42.2	78.3	78.4	28.1	110.5	25.8	514.7	80.6	46.0	41.2	5.8	22.4
358	KDNa2-3Galb1-4Glc	148.7	61.7	0.0	30.3	170.1	45.1	159.2	25.8	229.0	57.0	164.4	25.0	1493.0	148.2	52.1	39.0	18.6	22.7
359	KDNa2-3Galb1-3GalNAca	233.9	81.3	115.4	141.7	375.1	187.6	173.5	54.7	612.0	231.3	411.0	117.3	1715.5	1165.6	105.6	51.8	0.1	26.5
360	Fuca1-2Galb1-3GlcNAcb1-2Mana1-6(Fuca1-2Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	873.4	52.1	1051.9	58.9	1004.5	164.0	1137.1	78.3	1396.8	42.9	1445.1	150.0	2999.6	785.8	448.0	219.0	238.0	28.4
361	Fuca1-2Galb1-4GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	588.7	51.4	485.0	27.5	536.5	204.3	649.1	77.3	772.4	47.0	606.4	115.8	914.5	90.7	302.7	77.5	49.7	26.4
362	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAb	968.8	172.8	620.9	41.9	321.9	121.3	1044.6	38.8	150.4	40.8	1075.8	207.0	572.8	104.9	448.9	59.3	74.9	36.9
363	Gala1-3Galb1-4GlcNAcb1-2Mana1-6(Gala1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	270.1	127.4	219.7	99.5	255.6	121.3	35.4	265.3	477.2	106.4	179.0	40.5	5926.8	294.0	114.5	13.8	64.8	18.0
364	Galb1-4GlcNAcb1-2Mana1-6(Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	0.0	13.0	0.0	15.3	0.0	101.9	69.8	11.3	0.0	5.8	17.0	17.4	2181.1	311.2	12.8	35.1	0.1	13.6
365	Fuca1-4(Galb1-3)GlcNAcb1-2Mana1-6(Fuca1-4(Galb1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1306.4	172.4	1116.8	272.5	1342.1	290.0	1383.3	160.6	1987.8	187.1	1819.0	112.4	3917.9	740.6	729.0	64.9	229.0	80.5
366	Neu5Aca2-6GlcNAcb1-4GlcNAc	1.0	42.1	0.0	11.4	0.0	130.1	36.8	34.4	0.0	13.3	18.3	2.4	144.0	25.8	26.5	11.5	0.1	9.5
367	Neu5Aca2-6GlcNAcb1-4GlcNAcb1-4GlcNAc	0.0	7.3	0.0	40.4	37.8	31.4	0.0	20.7	66.6	54.3	233.1	134.7	401.0	109.2	104.1	78.0	0.1	43.6
368	Galb1-4(Fuca1-3)GlcNAcb1-6(Fuca1-2Galb1-4GlcNAcb1-3)Galb1-4Glc	135.3	43.1	129.5	17.8	62.1	67.1	79.9	94.2	76.7	23.7	150.8	73.7	576.8	65.4	38.7	18.8	28.7	39.2
369	Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-4(Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	122.1	93.9	50.5	68.5	191.0	29.0	91.2	60.2	286.4	69.4	138.2	110.1	1500.8	125.7	149.7	198.5	0.1	31.3
370	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-6(GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	246.8	85.1	560.6	22.1	424.9	186.3	602.0	134.6	485.5	27.1	704.9	110.2	622.6	71.3	23.6	9.7	62.2	32.9
371	Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-6(Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	305.5	27.9	336.3	65.6	384.8	58.8	211.6	35.9	391.6	42.9	209.4	72.9	1076.6	150.6	166.2	65.6	85.2	5.0
372	Gala1-3Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-6(Gala1-3Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	3197.8	876.2	1754.3	735.6	1840.3	803.2	1918.6	527.9	2512.4	969.0	2819.6	1013.0	4714.3	696.1	1462.2	514.1	541.6	219.2
373	GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-6(GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	3014.4	863.5	1494.0	515.6	505.0	180.2	1654.6	573.8	262.6	137.5	595.3	244.2	51.2	65.1	93.5	64.8	20.1	44.1
374	Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-6(Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	2386.5	422.1	1895.3	387.8	1419.2	152.0	2202.6	534.7	1052.2	293.1	665.6	199.5	1059.3	180.0	626.1	159.8	63.3	22.3
375	Fuca1-4(Fuca1-2Galb1-3)GlcNAcb1-2Mana1-3(Fuca1-4(Fuca1-2Galb1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	1049.2	399.6	673.0	310.1	1266.3	424.1	807.7	219.8	1171.0	655.3	1129.1	508.7	2500.9	27.8	649.3	269.8	13.9	73.4
376	Neu5Aca2-3Galb1-4GlcNAcb1-3GalNAc	588.5	183.5	510.4	182.2	320.2	62.9	603.6	446.1	463.5	211.4	423.1	204.4	412.0	91.8	213.1	52.0	78.5	51.5
377	Neu5Aca2-6Galb1-4GlcNAcb1-3GalNAc	2744.9	1548.9	1845.7	1086.3	1917.7	1166.7	1468.9	901.3	1487.2	1207.4	3625.0	2195.6	1178.9	99.6	841.5	493.8	0.1	10.9
378	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-3GalNAca	544.1	211.3	533.4	209.4	446.2	255.0	454.9	221.9	635.4	303.0	470.2	258.9	410.4	71.1	354.2	172.6	91.6	41.0
379	GalNAcb1-4GlcNAcb1-2Mana1-6(GalNAcb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	998.1	457.6	940.8	355.1	669.0	159.1	807.6	291.8	873.7	421.9	777.7	425.8	648.8	49.8	381.5	187.9	346.2	75.6
380	Galb1-3GalNAca1-3(Fuca1-2)Galb1-4Glc	1473.1	575.1	1152.5	240.0	1091.3	221.4	1374.9	310.2	598.4	183.0	443.6	202.9	1300.3	120.8	249.4	146.2	50.4	26.2
381	Galb1-3GalNAca1-3(Fuca1-2)Galb1-4GlcNAc	2788.4	628.2	1850.8	577.9	1754.5	275.3	2416.7	649.4	598.6	153.6	639.0	271.2	807.0	138.6	72.3	56.9	26.7	8.5
382	Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3GlcNAcb1-3)Galb1-4Glcb	270.1	88.4	352.7	44.9	107.6	16.9	120.5	21.7	148.1	26.1	152.1	48.3	269.4	40.9	102.9	37.1	2.6	12.8
383	Galb1-4(Fuca1-3)GlcNAcb1-6(Galb1-3GlcNAcb1-3)Galb1-4Glc	894.8	37.9	739.9	101.8	481.2	141.2	675.3	211.1	926.2	128.0	565.3	141.9	340.4	86.5	147.6	62.5	45.7	37.1
384	Galb1-4GlcNAcb1-6(Fuca1-4(Fuca1-2Galb1-3)GlcNAcb1-3)Galb1-4Glc	211.3	169.8	0.0	24.1	0.0	12.4	14.5	58.9	64.9	19.0	18.7	15.6	44.8	15.8	13.1	19.3	40.4	10.9

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385	Galb1-4(Fuca1-3)GlcNAcb1-6(Fuca1-4(Fuca1-2Galb1-3)GlcNAcb1-3)Galb1-4Glc	148.2	77.1	0.0	24.0	0.0	7.7	5.1	32.0	83.1	32.2	102.1	68.8	0.0	34.2	6.1	19.8	62.5	43.2
386	Galb1-3GlcNAcb1-3Galb1-4(Fuca1-3)GlcNAcb1-6(Galb1-3GlcNAcb1-3)Galb1-4Glc	156.0	52.3	73.4	23.8	65.7	14.3	21.1	18.9	80.2	53.9	46.0	30.8	49.1	34.6	37.0	50.2	39.1	45.9
387	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(Galb1-4GlcNAcb1-4(Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	5.4	51.6	0.0	25.4	0.0	10.1	0.0	194.7	69.8	23.8	101.0	20.7	8.6	64.5	24.5	4.9	123.0	33.6
388	GlcNAcb1-2Mana1-6(GlcNAcb1-4(GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	516.7	189.3	386.5	93.0	248.5	105.2	181.9	148.2	312.5	168.6	113.6	33.4	246.4	41.7	113.2	28.1	87.6	7.5
389	Fuca1-2Galb1-3GalNAca1-3(Fuca1-2)Galb1-4Glcb	306.9	227.6	317.1	157.8	504.0	223.1	33.1	125.6	438.2	207.8	320.9	163.4	596.0	43.8	82.2	46.8	237.0	43.2
390	Fuca1-2Galb1-3GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb	629.6	273.1	680.0	200.8	892.5	215.7	727.9	404.9	412.1	260.3	276.2	205.4	252.4	104.9	109.8	63.5	68.0	40.8
391	Galb1-3GlcNAcb1-3GalNAca	269.3	178.4	173.4	236.9	313.2	176.9	49.0	114.6	310.6	225.8	296.6	174.5	746.2	105.6	55.6	12.0	60.5	10.7
392	GalNAcb1-4(Neu5Aca2-3)Galb1-4GlcNAcb1-3GalNAca	1766.5	66.7	1668.9	96.9	1084.3	60.3	1511.5	315.9	1152.4	125.0	970.9	312.3	666.1	52.2	1087.9	107.8	1225.4	271.1
393	GalNAca1-3(Fuca1-2)Galb1-3GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb	939.4	361.5	1626.6	547.6	1898.0	188.7	1051.8	369.7	639.4	220.1	231.6	137.9	1499.9	288.2	45.5	21.1	25.2	14.9
394	Gala1-3Galb1-3GlcNAcb1-2Mana1-6(Gala1-3Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	877.0	287.2	684.1	251.4	771.0	244.3	941.7	81.1	902.9	356.5	2176.5	1137.4	1143.5	182.1	345.8	155.5	379.9	189.7
395	Gala1-3Galb1-3(Fuca1-4)GlcNAcb1-2Mana1-6(Gala1-3Galb1-3(Fuca1-4)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	3508.4	168.3	3981.3	361.0	3621.6	245.0	3404.1	139.0	4332.4	618.3	4604.1	837.1	4899.8	646.3	1238.8	73.8	499.6	72.8
396	Neu5Aca2-3Galb1-3GlcNAcb1-2Mana1-6(Neu5Aca2-3Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	671.3	153.6	571.7	170.1	242.7	187.1	641.0	114.7	510.3	182.7	645.1	136.8	3015.1	804.0	299.6	68.4	385.1	99.6
397	GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	229.5	123.6	0.0	48.4	62.5	36.7	120.2	54.0	87.2	54.6	311.7	34.7	80.9	101.7	21.6	33.7	0.1	16.1
398	Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	523.9	80.6	293.0	41.1	266.1	71.6	338.6	50.4	477.3	74.3	422.9	39.7	1460.2	426.4	85.2	23.7	0.1	28.8
399	Neu5Aca2-3Galb1-3GlcNAcb1-3GalNAca	293.9	103.8	185.9	93.0	276.1	30.6	190.7	26.6	175.3	49.2	161.7	21.7	171.1	52.4	134.0	31.7	65.8	38.4
400	Fuca1-2Galb1-4GlcNAcb1-3GalNAca	522.8	120.5	367.9	110.3	249.0	13.9	651.5	320.2	369.7	80.2	303.4	102.7	260.1	26.3	103.5	55.6	0.1	27.7
401	Galb1-4(Fuca1-3)GlcNAcb1-3GalNAca	366.8	83.6	332.4	128.6	498.1	111.4	322.3	78.5	428.8	120.3	413.4	115.3	283.4	176.8	160.6	9.1	91.8	21.8
402	GalNAca1-3GalNAcb1-3Gala1-4Galb1-4GlcNAcb	4337.0	1435.3	2055.9	724.0	1982.7	470.7	2705.0	952.9	1863.9	580.9	1600.1	541.8	4633.4	268.2	1418.3	424.6	24.2	14.9
403	Gala1-4Galb1-3GlcNAcb1-2Mana1-6(Gala1-4Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	4774.4	1605.3	2692.0	856.9	2926.8	367.2	2875.3	785.6	2195.3	547.3	2407.9	388.8	5448.2	588.8	971.8	584.7	81.7	33.3
404	Gala1-4Galb1-4GlcNAcb1-2Mana1-6(Gala1-4Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	4582.6	803.7	2004.4	235.9	1838.7	211.0	1917.6	289.8	1629.7	241.9	1757.0	208.2	9304.6	826.0	1344.0	161.7	65.9	11.5
405	Gala1-3Galb1-4GlcNAcb1-3GalNAca	959.3	356.1	504.7	219.0	908.9	424.5	496.6	233.9	596.4	123.9	380.7	134.2	1875.1	173.2	102.7	43.7	37.3	8.4
406	Galb1-3GlcNAcb1-6Galb1-4GlcNAcb	281.7	75.7	324.8	29.3	142.5	37.0	129.6	21.6	240.4	39.6	436.5	119.0	604.3	47.8	48.1	39.9	11.1	27.5
407	Galb1-3GlcNAca1-6Galb1-4GlcNAcb	7764.9	733.5	3594.8	940.0	2696.7	473.6	3803.6	742.7	2271.9	139.2	829.7	59.2	6710.0	706.4	885.6	73.3	22.6	19.8
408	GalNAcb1-3Gala1-6Galb1-4Glcb	2126.3	836.9	49.6	28.4	0.0	11.1	731.8	230.9	128.8	47.6	398.9	138.5	165.8	55.9	24.7	22.2	33.5	12.2
409	Gala1-3(Fuca1-2)Galb1-4(Fuca1-3)Glcb	194.4	105.8	195.7	142.7	18.1	11.7	70.3	21.8	139.4	35.7	231.6	31.6	289.5	164.3	22.9	24.5	122.1	36.8
410	Galb1-4GlcNAcb1-6(Neu5Aca2-6Galb1-3GlcNAcb1-3)Galb1-4Glc	190.9	43.9	192.3	150.0	68.2	103.3	34.4	97.6	118.4	18.6	58.3	20.6	178.3	33.0	31.9	45.3	45.0	33.4
411	Galb1-3GalNAcb1-4(Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	2102.1	229.9	1017.1	270.8	94.4	53.5	566.7	101.7	921.4	116.3	728.1	145.4	1390.9	93.7	102.5	45.3	175.4	31.8
412	Neu5Aca2-3Galb1-3GalNAcb1-4(Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	1070.8	350.6	523.9	292.1	567.2	180.9	378.6	149.1	652.8	69.2	329.1	106.1	736.0	270.4	278.2	61.6	52.7	20.1
413	Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-3GalNAca	361.7	137.0	320.9	96.4	319.4	55.3	252.8	85.5	291.0	66.0	205.9	52.3	472.0	86.5	100.6	34.7	187.0	36.3
414	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-3GalNAca	525.2	53.7	673.5	28.0	691.2	88.3	594.1	120.0	563.9	48.7	508.4	34.8	265.0	171.5	148.4	34.9	0.1	8.1
415	GalNAca1-3GalNAcb1-3Gala1-4Galb1-4Glcb	4217.7	1547.9	3175.7	920.3	3077.4	603.6	1956.0	828.4	3063.6	862.6	1915.6	451.4	5110.7	1015.3	91.7	13.8	103.7	16.1
416	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-3GalNAca	430.7	115.0	271.4	65.8	910.2	422.4	1072.3	569.5	469.2	86.5	577.4	195.5	369.8	121.2	137.7	60.7	104.0	75.8
417	Gala1-3(Fuca1-2)Galb1-4(Fuca1-3)GlcNAcb1-3GalNAc	1107.5	220.7	1185.2	119.2	1023.8	82.4	825.4	220.0	1410.7	147.1	1126.4	134.3	422.0	47.1	182.7	59.9	47.0	4.1
418	GalNAca1-3(Fuca1-2)Galb1-4(Fuca1-3)GlcNAcb1-3GalNAc	353.0	205.7	133.7	51.5	329.7	84.1	330.0	49.2	283.3	64.6	420.9	93.8	515.6	21.5	73.5	83.8	123.6	53.0
419	Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-6(Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	372.9	66.3	384.3	106.0	314.5	110.7	266.9	54.5	185.9	52.7	456.0	147.3	463.7	176.0	177.8	46.4	118.2	22.7
420	Fuca1-2Galb1-4GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1972.5	498.2	1718.9	192.2	1443.0	132.9	1495.1	193.1	1606.7	130.3	1917.4	97.9	2433.6	92.5	832.7	134.5	187.9	45.4
421	GlcNAcb1-2(GlcNAcb1-6)Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	85.6	9.3	38.0	29.8	14.5	14.2	39.1	23.2	81.7	63.8	95.8	34.0	56.7	15.2	35.7	11.7	2.0	31.9
422	Fuca1-2Galb1-3GlcNAcb1-3GalNAc	204.7	106.3	190.2	39.2	198.8	54.1	0.0	361.1	180.0	45.6	471.7	89.9	100.7	95.9	51.4	28.4	60.5	19.4
423	Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-3GalNAc	630.4	156.2	344.4	157.1	619.4	68.8	433.3	25.2	347.7	107.7	546.4	72.1	1244.2	170.0	111.4	23.1	25.7	30.6
424	GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb1-3GalNAc	794.3	175.0	536.9	91.6	370.6	39.9	650.3	36.7	426.2	46.1	474.1	116.9	453.5	25.2	242.4	30.9	83.1	7.6
425	Gala1-3Galb1-3GlcNAcb1-3GalNAc	1059.8	150.4	694.8	143.9	701.1	178.9	697.2	43.1	999.6	178.4	1241.3	70.4	1360.6	126.8	481.2	122.1	43.8	4.6
426	Fuca1-2Galb1-3GlcNAcb1-2Mana1-6(Fuca1-2Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1755.5	310.3	1745.7	328.4	1222.5	146.3	1367.9	145.8	1383.3	91.2	1477.5	79.2	2485.5	102.0	825.1	65.5	96.1	19.8
427	Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-6(Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1814.5	487.8	1546.7	387.9	1880.7	339.6	1859.6	552.8	1592.8	175.8	913.3	158.3	1545.7	244.5	135.8	75.7	31.8	7.3
428	Galb1-3GlcNAcb1-6(Galb1-3GlcNAcb1-2)Mana1-6(Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	1229.5	248.9	1758.9	361.0	1066.7	155.5	1209.5	203.9	1125.5	184.9	784.2	126.5	1376.1	150.5	671.7	25.7	82.6	13.2
429	Galb1-4GlcNAcb1-6(Fuca1-2Galb1-3GlcNAcb1-3)Galb1-4Glc	153.4	59.1	166.5	67.1	395.2	170.4	210.6	143.2	104.1	25.2	123.5	20.6	349.4	42.1	61.2	24.4	0.7	22.3
430	Fuca1-3GlcNAcb1-6(Galb1-4GlcNAcb1-3)Galb1-4Glc	598.3	213.3	539.2	168.6	288.5	125.1	383.2	174.2	407.2	65.1	428.2	167.5	709.6	33.7	250.0	73.7	190.6	63.8
431	GlcNAcb1-2Mana1-6(GlcNAcb1-4)(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	339.1	84.5	408.1	54.1	1235.6	138.7	309.8	100.2	854.9	145.7	247.0	32.8	142.2	33.3	141.1	93.3	22.2	16.4
432	GlcNAcb1-2Mana1-6(GlcNAcb1-4)(GlcNAcb1-4(GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	169.2	34.4	52.2	28.2	20.4	15.5	53.4	14.3	21.4	11.8	62.2	38.4	52.3	12.1	20.3	14.9	5.6	16.7
433	GlcNAcb1-6(GlcNAcb1-2)Mana1-6(GlcNAcb1-4)(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	666.8	108.0	230.0	111.7	10.3	7.9	131.1	106.6	272.8	105.2	219.7	48.7	58.1	22.9	57.8	15.8	132.9	45.8
434	GlcNAcb1-6(GlcNAcb1-2)Mana1-6(GlcNAcb1-4)(GlcNAcb1-4(GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	153.4	26.6	95.5	26.9	35.7	17.8	75.2	89.3	63.1	23.3	50.9	45.9	60.5	50.6	16.5	12.2	43.2	38.9
435	Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-4)(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	681.5	161.5	608.5	51.0	30.0	6.1	390.2	109.9	570.5	91.7	270.9	99.4	311.6	70.3	64.9	17.4	113.2	15.6
436	Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-4)(Galb1-4GlcNAcb1-4(Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	10135.2	972.6	5181.2	784.9	7124.1	1050.1	5479.8	1932.7	5223.8	542.8	1872.1	130.4	4858.1	457.7	2499.5	479.4	111.4	36.4
437	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(GlcNAcb1-4)(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	124.2	53.0	59.2	56.8	85.6	54.3	150.0	63.3	76.1	33.1	172.4	20.2	65.0	11.7	83.1	3.9	77.1	28.5
438	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(GlcNAcb1-4)(Galb1-4GlcNAcb1-4(Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	249.4	99.2	245.0	62.9	192.0	80.1	159.5	78.4	254.0	83.3	280.7	42.0	538.0	368.5	41.2	25.9	74.5	29.9
439	Galb1-4Galb	220.5	76.3	188.4	37.9	171.7	16.1	198.2	164.2	71.2	19.7	353.6	37.5	405.4	26.9	69.2	28.1	3.2	23.0
440	Galb1-6Galb	1346.4	178.9	1188.0	238.2	1098.4	243.3	909.4	175.8	2286.8	375.3	12018.5	3062.6	3987.6	114.0	170.0	50.6	90.3	22.5
441	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb	0.0	44.3	12.1	31.4	110.8	44.2	188.5	63.6	13.2	32.0	123.7	51.7	352.8	136.9	79.0	30.8	8.4	16.0
442	GalNAcb1-6GalNAcb	783.9	190.3	700.5	315.1	709.4	170.7	468.8	158.9	849.8	314.1	488.6	21.6	860.8	101.1	215.5	50.6	154.9	39.2
443	(6S)Galb1-3GlcNAcb	914.0	300.7	853.8	245.3	1139.0	303.1	938.3	267.6	932.0	254.7	2222.3	494.3	4384.4	2452.6	187.1	75.2	273.0	94.3
444	(6S)Galb1-3(6S)GlcNAc	753.9	260.6	211.4	43.1	349.0	33.1	0.0	78.0	157.5	19.4	333.9	42.3	151.7	67.8	156.3	28.3	38.3	4.7
445	Fuca1-2Galb1-4 GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4GlcNAcb1-2(Fuca1-2Galb1-4GlcNAcb1-4)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	598.6	116.3	0.0	35.6	21.5	6.5	219.7	35.0	173.8	22.6	357.3	127.4	465.6	33.5	21.0	22.3	19.7	36.1
446	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-4(Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	317.9	62.9	376.5	119.9	301.0	86.5	563.6	224.6	199.9	25.0	248.9	40.9	290.6	103.1	255.9	146.8	106.5	41.1
447	Galb1-4(Fuca1-3)GlcNAcb1-6GalNAc	452.6	197.4	257.5	129.3	453.2	21.9	336.4	131.7	403.6	104.4	436.6	174.1	208.4	129.2	250.7	117.9	14.8	20.9
448	Galb1-4GlcNAcb1-2Mana	3015.8	1021.9	1602.2	519.7	1268.3	206.9	1646.6	256.0	1114.9	320.9	922.3	206.6	1301.9	177.3	1071.1	304.2	69.1	18.1
449	Fuca1-2Galb1-4GlcNAcb1-6(Fuca1-2Galb1-4GlcNAcb1-3)GalNAc	1728.8	41.8	1376.9	12.8	1195.2	58.8	1160.3	54.7	1469.4	121.2	2466.0	186.3	1044.8	91.6	706.1	148.4	25.7	22.6
450	Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-6(Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-3)GalNAc	1525.3	393.1	1374.4	205.1	1023.2	158.7	843.9	767.6	1239.1	225.2	1206.8	282.9	763.8	89.6	510.9	203.8	340.3	34.0
451	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-6(GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-3)GalNAc	1241.9	412.4	1936.9	762.7	1784.7	372.0	2011.7	899.4	513.7	99.4	544.4	92.7	639.2	59.6	57.1	30.3	45.8	17.4
452	Neu5Aca2-8Neu5Aca2-3Galb1-3GalNAcb1-4(Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	474.2	83.3	455.5	106.5	381.5	31.7	243.1	93.0	438.4	60.2	357.6	104.8	373.6	114.5	129.0	66.5	39.6	8.8
453	GalNAcb1-4Galb1-4Glcb	632.7	218.4	471.2	62.1	715.2	110.6	515.3	144.0	496.5	75.1	786.7	90.7	2002.1	63.9	150.8	26.4	108.1	48.3
454	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-6(GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1552.2	409.3	1960.8	688.7	1393.0	311.4	1368.7	422.5	1098.9	261.6	1066.5	67.5	1776.6	78.0	367.3	149.2	126.9	20.9
455	Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-6(Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	3385.3	952.2	3369.9	879.0	3326.9	697.0	3775.6	977.3	2927.3	574.9	1621.5	242.1	2984.1	202.0	1207.2	263.8	464.6	62.7
456	Neu5Aca2-6Galb1-4GlcNAcb1-6(Fuca1-2Galb1-3GlcNAcb1-3)Galb1-4Glc	186.7	22.2	17.2	36.1	0.0	28.7	116.8	34.2	80.2	17.5	83.8	22.8	71.2	13.2	28.9	75.2	20.8	17.7
457	GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-6(GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1149.0	405.6	749.6	330.1	26.9	18.3	778.1	365.5	175.2	78.8	219.4	52.2	303.8	65.5	110.8	15.2	115.9	47.8
458	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	503.7	170.5	147.6	95.8	80.1	38.3	67.9	170.3	293.7	55.9	51.3	25.0	641.5	178.6	53.0	21.5	0.1	30.9
459	Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-4)(Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	119.7	8.4	58.5	10.6	59.1	25.5	91.9	20.9	70.3	51.9	111.9	9.9	50.8	50.7	93.7	13.5	88.4	16.5
460	Neu5Aca2-3Galb1-4GlcNAcb1-4Mana1-6(GlcNAcb1-4)(Neu5Aca2-3Galb1-4GlcNAcb1-4(Neu5Aca2-3Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	93.1	22.2	88.4	38.1	93.0	10.9	121.0	13.7	67.9	22.4	60.9	46.6	195.0	103.6	62.9	44.6	19.5	15.7
461	Neu5Aca2-3Galb1-4GlcNAcb1-6(Neu5Aca2-3Galb1-4GlcNAcb1-2)Mana1 6(GlcNAcb1-4)(Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	0.0	25.6	30.3	44.6	73.7	25.1	0.0	11.9	45.1	25.0	57.4	29.0	8.5	17.0	52.2	14.0	93.8	11.1
462	Neu5Aca2-3Galb1-4GlcNAcb1-6(Neu5Aca2-3Galb1-4GlcNAcb1-2)Mana1 6(GlcNAcb1-4)(Neu5Aca2-3Galb1-4GlcNAcb1-4(Neu5Aca2-3Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	125.8	56.6	26.1	11.1	60.9	26.1	103.8	17.1	68.9	31.4	114.4	50.8	86.7	39.9	27.6	68.5	46.7	35.5
463	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-4)(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	133.6	29.6	140.7	20.4	168.4	7.4	126.5	14.6	90.6	33.2	147.8	13.3	198.7	75.7	56.5	11.3	61.8	31.9
464	Neu5Aca2-6Galb1-4GlcNAcb1-4Mana1-6(GlcNAcb1-4)(Neu5Aca2-6Galb1-4GlcNAcb1-4(Neu5Aca2-6Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	139.2	64.4	164.5	42.4	148.3	65.2	139.4	64.2	223.3	67.5	190.0	51.7	801.0	390.7	190.0	89.4	57.3	25.6
465	Neu5Aca2-6Galb1-4GlcNAcb1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2)Mana1 6(GlcNAcb1-4)(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	53.9	48.3	163.6	35.1	169.0	113.5	115.4	76.1	126.5	39.5	181.6	23.5	0.0	48.3	47.4	10.1	110.8	20.1
466	Neu5Aca2-6Galb1-4GlcNAcb1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2)Mana1 6(GlcNAcb1-4)(Neu5Aca2-6Galb1-4GlcNAcb1-4(Neu5Aca2-6Galb1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	71.7	101.9	52.5	12.6	125.9	23.4	79.4	47.3	222.1	61.6	390.2	110.5	133.6	79.3	30.8	12.8	90.4	9.5
467	Gala1-3(Fuca1-2)Galb1-3GalNAca	980.4	261.7	1447.3	425.5	1435.7	230.7	1307.7	327.2	995.5	248.7	1199.1	84.8	2745.7	280.4	96.8	49.4	99.4	37.0
468	Gala1-3(Fuca1-2)Galb1-3GalNAcb	1828.2	435.1	1360.8	241.5	1139.1	196.0	1511.4	381.6	575.9	142.7	376.3	95.6	3115.5	629.2	576.6	79.3	61.7	8.6
469	Glca1-6Glca1-6Glca1-6Glcb	403.3	194.6	390.8	168.9	388.1	142.9	465.9	358.7	532.9	225.1	470.1	159.3	2389.7	110.4	114.8	49.7	32.6	19.3
470	Glca1-4Glca1-4Glca1-4Glcb	414.2	118.2	132.2	128.8	513.6	168.4	39.1	150.7	286.0	303.7	11.6	79.0	2862.9	862.1	200.6	69.9	0.1	37.6
471	Neu5Aca2-3Galb1-4GlcNAcb1-6(Neu5Aca2-3Galb1-4GlcNAcb1-3)GalNAca	34.2	38.2	134.9	45.3	121.2	4.2	245.5	28.6	90.6	56.0	41.7	59.2	457.2	107.9	30.1	12.1	14.0	27.0
472	Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1170.6	82.2	1113.9	35.9	964.4	99.7	796.3	22.7	992.8	105.6	3104.3	290.3	518.4	42.8	537.0	124.5	234.6	21.7
473	Fuca1-2Galb1-3(Fuca1-4)GlcNAcb1-2Mana1-6(Fuca1-2Galb1-3(Fuca1-4)GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb1-4(Fuca1 6)GlcNAcb	1532.0	290.0	170.2	62.6	54.2	32.3	810.6	221.2	630.4	129.1	1365.4	141.7	90.3	12.8	128.7	42.0	25.8	6.7

(Continued)

474	Neu5Aca2-3Galb1-3GlcNAcb1-6(Neu5Aca2-3Galb1-3GlcNAcb1-2)Mana1-6(Neu5Aca2-3Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	849.5	105.7	685.2	39.0	593.7	43.7	744.0	360.4	1035.8	65.5	2856.4	468.5	762.0	76.7	437.4	92.1	102.2	24.6
475	GlcNAcb1-6(GlcNAcb1-2)Mana1-6(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	462.3	80.3	211.5	54.7	342.5	94.8	197.5	52.8	360.6	90.4	512.2	67.7	105.0	58.7	125.1	45.4	84.3	28.9
476	Galb1-3GlcNAcb1-2Mana1-6(GlcNAcb1-4)(Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	4573.0	823.6	5245.3	1681.0	3990.0	338.0	5453.1	904.6	3084.2	284.5	3488.9	493.8	1518.8	118.9	2482.4	291.6	196.8	61.1
477	Neu5Aca2-6Galb1-4GlcNAcb1-6(Galb1-3GlcNAcb1-3)Galb1-4Glcb	414.2	100.5	311.7	67.2	300.4	47.5	229.7	22.0	349.0	72.3	304.7	50.5	146.9	18.7	96.5	54.7	102.3	71.3
478	Neu5Aca2-3Galb1-4GlcNAcb1-2Mana	1093.6	46.4	933.4	107.6	668.6	83.8	946.5	220.2	726.3	162.0	3165.9	363.8	563.4	130.4	545.7	128.2	116.9	23.3
479	Neu5Aca2-3Galb1-4GlcNAcb1-6GalNAca	3472.3	325.8	1756.2	488.9	2002.4	92.8	2328.3	129.7	2263.7	374.1	1455.4	127.5	2906.4	511.7	1023.5	139.5	61.5	43.8
480	Neu5Aca2-6Galb1-4GlcNAcb1-6GalNAca	436.0	189.6	471.9	131.5	743.9	306.9	454.9	27.4	1007.7	590.8	666.5	367.7	906.2	44.0	101.0	55.4	78.2	21.4
481	Neu5Aca2-6Galb1-4 GlcNAcb1-6(Neu5Aca2-6Galb1-4GlcNAcb1-3)GalNAca	223.0	38.8	317.2	97.5	232.9	65.3	240.9	64.9	359.0	143.1	362.7	153.1	554.7	23.3	27.4	33.5	0.1	34.5
482	Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	100.1	91.2	66.9	67.7	109.5	155.1	36.6	69.2	45.8	99.0	68.7	140.6	185.5	72.5	259.3	446.7	34.4	9.8
483	Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	86.5	57.2	58.8	17.3	51.1	41.3	63.6	10.9	112.0	45.3	113.7	48.8	0.0	30.9	72.3	49.9	70.8	22.9
484	Mana1-6(Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	948.7	54.6	920.2	156.2	45.9	16.9	1187.4	249.6	121.2	49.4	307.2	103.1	101.5	42.3	127.5	11.1	74.1	15.5
485	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1286.1	96.0	727.1	145.1	212.5	106.3	723.9	46.8	710.9	102.4	288.4	86.9	413.4	58.7	102.2	77.0	0.1	34.5
486	Neu5Aca2-3Galb1-3GlcNAcb1-2Mana1-6(GlcNAcb1-4)(Neu5Aca2-3Galb1-3GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	316.4	52.5	286.2	81.4	40.8	33.6	378.9	45.7	62.9	24.8	326.7	44.7	310.6	73.9	106.7	37.0	25.8	11.4
487	Neu5Aca2-6Galb1-4GlcNAcb1-6(Fuca1-2Galb1-4(Fuca1-3)GlcNAcb1-3)Galb1-4Glc	503.3	111.8	436.3	43.2	329.9	31.5	273.0	58.5	365.6	120.8	196.5	104.1	279.4	30.4	73.3	63.0	0.1	51.3
488	Galb1-3GlcNAcb1-6GalNAca	752.4	224.9	331.9	148.1	161.2	37.6	188.0	121.9	44.1	7.7	416.7	98.7	1284.6	107.6	83.4	16.0	7.0	37.4
489	Gala1-3Galb1-3GlcNAcb1-6GalNAca	1692.9	543.4	610.9	284.1	793.4	155.4	483.6	156.7	914.2	229.2	493.1	111.0	1947.3	126.0	92.2	29.4	27.0	17.1
490	Galb1-3(Fuca1-4)GlcNAcb1-6GalNAca	100.0	72.7	34.8	55.7	124.5	59.0	244.1	121.4	0.0	16.3	476.1	101.5	154.3	28.8	31.9	4.6	0.1	27.9
491	Neu5Aca2-3Galb1-3GlcNAcb1-6GalNAca	1630.9	468.3	787.7	269.1	387.7	134.8	498.3	174.0	535.3	124.5	605.2	188.9	2484.0	192.8	434.8	173.3	0.1	33.8
492	(3S)Galb1-3(Fuca1-4)GlcNAcb	300.8	74.7	257.4	58.2	190.1	13.4	484.0	224.0	277.9	67.6	356.6	88.2	509.1	152.5	118.6	31.9	42.7	14.7
493	Galb1-4(Fuca1-3)GlcNAcb1-6(Neu5Aca2-6(Neu5Aca2-3Galb1-3)GlcNAcb1-3)Galb1-4Glc	449.8	109.2	267.1	39.1	198.8	33.3	146.4	43.0	250.0	60.5	126.8	25.9	408.6	42.2	192.5	12.9	53.9	10.6
494	Fuca1-2Galb1-4GlcNAcb1-6GalNAca	571.1	36.0	629.0	17.0	563.4	56.6	1181.9	185.7	869.3	21.7	1008.4	49.7	1236.0	171.8	212.5	8.4	0.1	20.7
495	Gala1-3Galb1-4GlcNAcb1-6GalNAca	1664.8	340.0	394.8	169.3	595.0	260.2	368.2	106.6	389.7	182.3	389.6	67.2	2026.7	534.1	165.4	68.9	0.1	49.6
496	Galb1-4(Fuca1-3)GlcNAcb1-2Mana	755.2	78.9	3.4	8.4	0.0	60.4	296.6	84.4	166.9	19.2	584.8	33.8	73.4	30.0	34.8	21.4	24.6	40.8
497	Fuca1-2(6S)Galb1-3GlcNAcb	2091.2	235.0	1640.1	189.2	1437.4	135.6	1241.1	187.7	1892.5	144.4	2118.4	140.7	656.5	65.4	815.4	146.7	11.8	29.4
498	Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-6GalNAca	1207.6	256.3	552.7	214.5	713.3	111.8	900.1	116.9	1029.3	187.1	899.7	202.5	648.6	77.4	424.9	141.8	104.3	31.7
499	Fuca1-2Galb1-4GlcNAcb1-2Mana	786.0	79.1	621.3	92.6	387.5	54.0	2194.6	2535.0	425.8	181.2	629.8	168.8	397.6	204.0	178.3	16.7	52.7	25.1
500	Fuca1-2Galb1-3(6S)GlcNAcb	1934.5	160.1	1374.9	253.8	1503.2	131.5	983.3	166.2	1641.5	52.7	1455.3	114.5	803.0	30.5	464.0	68.9	74.3	18.6
501	Fuca1-2(6S)Galb1-3(6S)GlcNAcb	7697.0	1565.8	5649.7	129.2	4076.2	366.0	4738.5	458.2	4624.3	696.5	9232.3	473.4	3608.9	895.2	1968.1	189.1	150.1	16.6
502	Neu5Aca2-6GalNAcb1-4(6S)GlcNAcb	614.2	195.3	534.0	189.5	495.5	106.3	567.1	105.9	713.7	81.7	459.5	135.9	488.9	33.6	37.2	11.5	23.5	9.0
503	GalNAcb1-4(Fuca1-3)(6S)GlcNAcb	640.2	105.2	651.8	99.8	456.2	63.3	567.9	122.2	1110.0	178.5	1472.6	187.3	835.1	336.7	119.4	14.4	22.2	32.5
504	(3S)GalNAcb1-4(Fuca1-3)GlcNAcb	344.5	139.4	367.0	191.4	515.2	93.0	428.2	144.7	447.9	169.3	463.3	13.2	2308.2	136.4	60.5	39.9	89.3	7.9
505	Fuca1-2Galb1-3GlcNAcb1-6(Fuca1-2Galb1-3GlcNAcb1-3)GalNAca	733.6	18.8	1030.4	63.3	679.0	68.6	802.1	72.0	1567.4	84.8	1348.5	50.9	869.5	100.5	234.0	12.3	100.9	33.5
506	GalNAca1-3(Fuca1-2)Galb1-3GlcNAcb1-6GalNAca	1316.5	452.4	95.4	48.0	0.0	15.7	840.4	138.0	179.6	36.1	114.6	28.1	61.6	9.3	41.0	11.2	12.2	37.7
507	GlcNAcb1-6(GlcNAcb1-2)Mana1-6(GlcNAcb1-4)(GlcNAcb1-4(GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAc	75.6	28.0	10.6	7.9	0.0	6.0	325.3	459.1	8.3	12.8	156.8	38.3	0.0	24.3	83.3	15.1	76.5	29.0
508	Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-2)Mana1-6(GlcNAcb1-4)Galb1-4GlcNAcb1-4(Gal b1-4GlcNAcb1-2)Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAc	215.3	79.8	98.2	43.1	22.4	12.9	70.2	11.6	148.0	17.9	40.8	15.5	13.7	39.9	26.1	36.5	0.1	26.3
509	Galb1-3GlcNAca1-3Galb1-4GlcNAcb	6718.5	1177.4	2515.4	600.6	340.7	94.5	3655.4	542.3	909.4	361.7	431.8	102.3	3351.0	707.0	97.8	13.7	1.4	20.6
510	Galb1-3(6S)GlcNAcb	472.1	99.8	150.5	72.0	248.0	139.4	201.1	82.4	230.2	31.6	94.0	33.1	412.7	47.3	81.5	17.8	0.1	39.9
511	(6S)(4S)GalNAcb1-4GlcNAc	63.7	7.2	103.1	50.5	89.1	17.1	140.3	45.7	14.7	30.7	323.1	59.5	74.5	28.3	65.8	9.8	0.1	6.1
512	(6S)GalNAcb1-4GlcNAc	198.2	54.0	189.0	65.5	153.5	20.1	139.4	22.5	191.4	89.7	43.5	38.1	255.8	64.9	47.5	42.5	0.1	16.5
513	(3S)GalNAcb1-4(3S)GlcNAc	444.5	260.0	441.0	32.2	554.4	36.3	286.1	69.9	80.8	85.8	707.5	124.9	1003.7	296.5	0.0	56.1	8.9	12.3
514	GalNAcb1-4(6S)GlcNAc	817.9	175.2	689.5	189.5	363.3	136.4	507.7	189.5	467.2	177.4	1443.5	343.7	1454.7	227.8	232.6	72.8	0.1	29.9
515	(3S)GalNAcb1-4GlcNAc	946.9	246.9	276.0	265.4	781.4	383.2	282.4	138.5	408.2	67.9	263.9	159.1	1299.3	424.6	120.2	15.4	5.7	21.4
516	(4S)GalNAcb	337.6	116.2	244.6	211.9	264.7	47.1	214.4	132.4	375.3	88.1	361.6	65.8	646.9	96.5	97.9	49.8	29.3	31.0
517	Galb1-4(6P)GlcNAcb	120.4	67.0	129.8	88.2	255.5	210.9	309.6	185.6	152.2	93.3	352.7	81.1	495.0	86.4	29.5	24.4	0.1	9.7
518	(6P)Galb1-4GlcNAcb	3895.4	163.1	802.5	284.2	1474.9	125.4	1225.8	170.8	1477.9	248.5	877.0	68.3	6144.1	293.9	1098.5	123.8	241.5	19.4
519	GalNAca1-3(Fuca1-2)Galb1-4GlcNAcb1-6GalNAc	2254.5	309.6	1178.3	284.2	628.4	152.7	1249.3	227.5	768.4	108.9	1289.4	55.0	62.5	48.4	292.9	150.9	42.8	14.7
520	Neu5Aca2-6Galb1-4GlcNAcb1-2Man	554.9	51.1	463.3	57.5	346.4	93.8	402.8	65.1	266.6	102.3	378.2	51.1	218.7	22.8	233.2	46.2	157.2	73.1
521	Gala1-3Galb1-4GlcNAcb1-2Mana	871.8	92.8	450.4	122.6	613.3	152.6	489.0	41.4	539.5	23.2	575.8	43.7	1289.7	112.5	258.9	66.4	71.2	46.6
522	Gala1-3(Fuca1-2)Galb1-4GlcNAcb1-2Mana	661.7	36.1	509.2	72.9	530.5	58.6	756.2	357.7	491.3	55.5	558.8	47.8	262.7	86.3	216.0	58.4	66.8	23.9
523	GalNAca1-3(Fuca1-2)Galb1-4 GlcNAcb1-2Mana	1097.1	83.2	1610.0	210.3	1547.6	37.1	1664.9	190.6	229.0	49.3	456.6	64.8	114.5	12.3	200.5	90.9	5.0	21.6
524	Galb1-3GlcNAcb1-2Mana	562.3	225.4	315.9	83.5	205.0	82.1	278.9	47.8	246.3	75.4	274.8	42.8	423.9	48.0	133.9	40.6	5.8	24.3
525	Gala1-3(Fuca1-2)Galb1-3GlcNAcb1-6GalNAc	1536.2	220.3	1108.0	62.2	914.1	67.3	1023.4	126.9	640.9	61.1	527.1	69.7	1032.2	132.6	186.3	88.0	0.1	28.5
526	Neu5Aca2-3Galb1-3GlcNAcb1-2Mana	464.5	58.7	483.1	122.8	379.8	74.6	377.6	96.3	502.1	77.2	503.5	124.6	435.2	123.3	118.7	19.1	202.0	60.4
527	Gala1-3Galb1-3GlcNAcb1-2Mana	666.1	216.9	743.8	151.6	634.1	171.9	636.8	93.9	1189.8	314.5	978.3	226.6	5823.7	426.3	76.1	46.1	38.2	23.9
528	GalNAcb1-4GlcNAcb1-2Mana	120.3	54.1	186.3	26.8	138.7	32.8	145.4	141.1	264.8	125.1	59.6	29.1	230.3	93.1	17.6	25.9	45.1	32.2
529	Neu5Aca2-3Galb1-3GalNAcb1-4Galb1-4Glcb	515.9	58.2	38.5	20.7	5.5	14.9	156.7	40.5	97.8	42.7	106.1	24.1	113.3	31.5	39.7	4.2	75.3	29.0
530	GlcNAcb1-2 Mana1-6(GlcNAcb1-4)(GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAc	183.8	16.9	99.2	19.0	88.0	27.9	71.5	45.0	99.5	30.4	131.6	47.3	27.6	30.6	58.4	13.7	31.7	14.6
531	Galb1-4GlcNAcb1-2 Mana1-6(GlcNAcb1-4)(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAc	86.5	21.7	61.5	13.7	18.9	4.3	26.3	12.5	103.9	40.0	103.9	15.1	0.0	16.3	47.9	5.0	145.4	48.3
532	Galb1-4GlcNAcb1-2 Mana1-6(Galb1-4GlcNAcb1-4)(Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAc	1201.9	78.5	830.1	86.3	651.0	182.7	548.1	200.4	1315.0	104.8	608.2	50.5	337.6	23.9	422.0	44.3	45.5	8.0
533	Fuca1-4(Galb1-3)GlcNAcb1-2 Mana	2202.8	833.0	1127.0	69.7	992.2	199.8	772.7	37.5	1501.6	141.8	1905.1	208.8	3005.9	241.6	0.0	184.9	273.3	41.2
534	Neu5Aca2-3Galb1-4(Fuca1-3)GlcNAcb1-2Mana	119.8	58.8	110.5	57.5	115.3	43.2	174.9	44.8	189.8	55.0	229.8	164.6	8.9	160.4	67.9	48.4	12.1	25.9
535	GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3)Galb1-4GlcNAc	58.2	35.9	77.4	49.6	104.8	33.9	67.0	19.5	115.1	26.5	118.9	47.1	107.5	20.2	74.3	7.4	69.0	26.8
536	GalNAca1-3(Fuca1-2)Galb1-3GalNAcb1-3Gala1-4Galb1-4Glc	1998.9	818.6	2733.3	885.4	1613.7	241.9	2284.7	636.0	590.0	74.8	546.3	88.4	2062.6	130.8	476.4	196.2	49.9	25.2
537	Gala1-3(Fuca1-2)Galb1-3GalNAcb1-3Gala1-4Galb1-4Glc	573.5	151.0	583.7	262.3	854.1	273.9	743.1	210.4	425.2	118.2	390.7	120.7	2533.9	171.2	82.5	40.9	78.2	63.9
538	Galb1-3GalNAcb1-3Gal	305.0	107.9	203.3	104.8	397.2	103.1	397.0	70.6	225.7	55.1	393.2	159.8	1337.6	61.4	148.3	34.4	75.8	36.0
539	GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	559.8	88.9	372.7	36.7	255.8	24.0	344.5	40.5	275.2	23.2	322.0	11.7	305.7	14.7	173.7	10.3	109.9	65.6
540	GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	170.5	46.6	218.6	143.7	401.5	227.7	405.3	179.5	236.9	40.6	267.6	109.7	314.4	52.7	77.5	35.9	127.0	122.1
541	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	5186.7	95.0	3923.7	234.3	2916.5	352.8	4168.4	801.2	3730.8	128.0	6150.9	805.4	3535.7	138.1	3284.7	365.2	3244.7	114.7
542	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	134.8	35.5	12.8	5.4	45.1	36.5	53.4	15.3	98.0	34.2	148.4	43.1	184.9	263.5	19.3	17.6	0.1	25.6
543	Neu5Gca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Gca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	250.3	38.6	218.4	48.5	279.1	33.7	145.3	36.7	183.4	59.1	283.1	53.8	164.5	81.8	165.4	27.1	0.1	44.0
544	Fuca1-2Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Fuca1-2Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	1145.3	99.4	1179.9	171.0	744.6	74.6	1108.0	154.7	1086.4	149.2	928.1	89.8	807.6	38.2	606.0	45.9	76.1	6.8
545	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	2686.0	245.6	2428.9	61.9	1748.9	66.5	2133.8	76.2	1963.0	224.4	2056.9	172.5	2289.3	196.8	1609.9	157.6	1756.8	87.4
546	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	232.8	41.0	186.7	53.5	461.7	260.9	81.5	56.4	229.3	112.4	276.5	49.7	125.7	35.1	160.1	49.5	41.0	17.6
547	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	3027.8	166.7	2571.6	267.2	2311.2	73.7	2387.5	168.0	2074.9	384.2	2723.8	39.4	2460.4	154.9	1836.2	158.5	2690.7	195.0
548	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	252.6	45.7	222.3	24.3	243.3	64.3	197.4	56.4	197.5	44.7	164.5	40.8	77.7	50.2	263.6	222.0	1.0	20.5
549	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	32.5	65.3	339.8	232.0	237.9	95.7	114.8	17.4	178.6	30.0	130.7	19.2	415.8	81.2	0.0	806.1	0.1	20.5
550	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	30.1	77.4	96.8	55.6	162.1	162.3	152.1	104.4	218.7	34.1	154.8	114.8	366.6	58.5	31.7	49.6	122.9	292.0
551	Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	1095.9	45.3	0.0	159.5	15.8	18.0	615.3	52.0	238.6	157.0	247.1	156.3	35.5	34.6	13.7	6.9	40.3	42.2
552	Neu5Gca2-8Neu5Gca2-3Galb1-4GlcNAc	195.9	67.4	104.7	7.9	22.1	4.7	83.8	8.4	108.5	5.4	260.0	75.8	0.0	29.4	44.6	16.7	91.0	64.8
553	Neu5Aca2-8Neu5Gca2-3Galb1-4GlcNAc	147.2	30.2	0.0	74.4	18.0	29.9	18.1	52.6	112.0	37.6	124.0	56.5	0.0	43.4	86.5	13.6	0.1	36.1
554	Neu5Gca2-8Neu5Aca2-3Galb1-4GlcNAc	228.5	82.6	42.7	17.1	6.9	28.8	0.0	59.6	138.8	9.4	162.6	26.2	0.0	24.5	101.3	37.3	189.7	35.1
555	Neu5Gca2-8Neu5Gca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAc	270.0	105.8	165.3	114.1	446.7	41.1	146.1	96.0	128.5	21.7	155.7	46.1	0.0	94.0	111.5	52.8	2.9	10.7
556	Neu5Gca2-8Neu5Gca2-6Galb1-4GlcNAc	585.9	71.1	384.8	82.6	443.3	129.5	194.4	26.4	615.0	47.9	380.1	40.6	1652.1	338.7	79.7	12.2	195.6	37.6
557	Neu5Aca2-8Neu5Aca2-3Galb1-4GlcNAc	0.0	104.6	0.0	12.0	39.1	24.5	170.5	221.7	19.2	17.7	77.4	26.5	5.3	36.5	13.9	10.4	0.1	31.9
558	GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-4GlcNAcb1-2)Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-2Man a1-3)Manb1-4GlcNAcb1-4GlcNAc	1066.3	85.9	1175.9	786.3	996.9	187.0	570.9	61.5	741.9	119.1	393.6	60.2	698.9	105.3	142.0	63.1	44.3	21.9
559	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2)Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Mana1-4GlcNAcb1-4GlcNAc	1058.5	49.8	890.8	40.0	636.2	160.0	758.5	251.0	534.9	266.9	2216.5	743.2	687.3	41.7	364.5	119.7	103.4	56.5
560	Gala1-3Galb1-4GlcNAcb1-2Mana1-6(Gala1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAc	5184.5	456.1	2104.8	355.4	2223.8	209.0	2842.8	286.0	2158.4	398.4	844.9	181.5	5885.0	338.2	577.7	64.0	3.5	27.2
561	GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-3)GalNAca	499.8	67.3	503.2	75.0	293.3	75.8	217.2	71.6	300.2	41.9	471.8	81.2	241.9	38.1	212.5	86.9	0.1	17.2
562	GalNAcb1-3GlcNAcb	8162.2	2061.9	4344.9	1197.7	3211.1	443.2	3937.5	955.7	2893.4	761.9	3256.0	612.7	8646.8	580.4	1652.5	378.2	0.1	30.6
563	GalNAcb1-4GlcNAcb1-3GalNAcb1-4GlcNAcb	200.1	30.5	249.8	85.0	242.8	55.7	109.9	81.6	132.2	32.0	155.1	31.3	224.2	92.0	76.8	32.3	0.1	7.9
564	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	139.6	47.2	252.3	73.5	190.3	74.4	161.2	35.7	200.8	33.6	136.6	55.9	523.6	34.1	0.0	115.9	0.1	17.8
565	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	64.2	169.2	10.4	14.5	138.7	94.1	50.6	16.0	85.4	26.4	7.7	6.8	227.4	263.2	22.1	22.0	28.8	104.5
566	GlcNAb1-3Galb1-3GalNAc	403.7	110.9	475.0	38.7	374.8	50.6	389.5	83.7	456.3	59.0	548.6	75.0	461.2	60.9	92.6	28.2	93.1	23.3
567	Galb1-3GlcNAcb1-6(Galb1-3)GalNAc	1325.0	186.9	936.5	227.0	1114.3	37.3	990.7	87.2	1655.1	67.7	1873.9	307.0	1148.9	28.2	325.7	35.7	33.2	17.9

(Continued)

Supplemental Table 2. (Continued)

568	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	245.4	43.7	323.4	31.6	254.4	82.0	175.2	55.9	90.3	18.2	635.0	203.2	223.3	101.0	73.5	44.4	31.8	24.4
569	(3S)GlcAb1-3Galb1-4GlcNAcb1-3Galb1-4Glc	472.0	212.3	513.4	118.9	696.2	80.8	324.3	173.3	865.8	200.3	720.3	66.2	1382.6	120.1	78.8	13.7	0.1	17.7
570	(3S)GlcAb1-3Galb1-4GlcNAcb1-2Mana	73.2	26.5	114.1	65.5	186.5	28.7	0.0	91.9	60.4	32.3	640.5	397.9	72.3	21.7	43.5	14.8	35.2	17.7
571	Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	3935.1	151.9	4924.8	144.1	3721.9	192.3	3385.6	90.0	6593.8	498.7	7864.6	1603.1	7203.5	450.6	1050.9	46.0	106.9	22.1
572	Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(Galb1-3GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	1217.0	318.5	2680.5	241.2	1959.4	96.9	1849.4	150.9	3212.7	232.6	4578.0	96.7	2769.4	317.1	124.1	30.8	149.0	26.5
573	Neu5Aca2-8Neu5Aca2-3Galb1-3GalNAcb1-4(Neu5Aca2-3)Galb1-4Glc	169.6	63.9	164.4	36.2	260.8	58.1	170.5	37.1	523.7	126.4	360.0	93.0	236.5	49.8	345.0	268.3	101.1	24.0
574	GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	461.0	33.5	428.0	108.5	273.6	92.0	109.3	172.7	276.0	8.0	503.5	40.9	159.3	126.6	201.8	33.6	38.1	18.6
575	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	300.4	82.3	280.8	49.3	280.3	55.1	266.6	27.2	259.0	76.7	251.8	58.6	245.3	92.8	126.7	26.1	125.7	26.7
576	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	225.9	110.6	276.8	192.5	206.2	51.4	0.0	101.1	193.1	39.8	207.1	91.2	396.1	71.5	147.3	42.6	59.5	26.0
577	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	147.4	26.9	246.1	55.4	146.6	16.3	226.4	121.6	193.3	28.8	74.0	16.8	238.2	63.8	97.5	27.5	50.0	32.9
578	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	225.4	34.4	296.6	48.2	125.3	44.9	110.4	46.2	188.5	54.8	198.1	54.3	828.2	165.1	92.1	16.7	83.1	44.5
579	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	151.7	27.5	187.8	101.6	129.8	40.6	119.5	47.6	260.4	131.4	146.9	35.8	501.1	66.3	132.5	43.6	58.0	9.9
580	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	0.0	24.5	71.4	44.9	95.7	101.7	0.0	50.7	113.9	23.0	73.9	19.6	309.8	55.8	39.0	37.6	42.1	30.3
581	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	0.0	41.6	16.9	113.7	98.0	35.7	0.0	14.6	72.7	28.3	5.3	39.4	274.4	28.8	73.0	116.2	82.2	32.6
582	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	322.2	43.3	302.8	20.3	196.3	29.4	261.1	182.9	451.3	53.4	423.6	97.1	450.3	62.9	162.2	34.4	138.7	43.7
583	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	65.3	39.9	146.2	41.8	144.3	43.1	81.2	99.6	15.8	14.5	629.3	107.2	413.1	107.4	34.4	22.9	37.8	9.2
584	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb 4	220.6	46.1	190.3	24.0	0.0	210.4	97.2	21.0	274.4	53.5	187.8	42.7	377.3	76.2	500.9	470.7	73.5	19.2
585	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	107.8	114.6	135.3	47.6	204.9	24.2	171.7	42.4	182.9	40.5	439.0	43.9	496.4	67.4	38.1	19.0	64.9	35.2
586	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	91.7	20.6	257.2	68.4	246.9	27.0	80.1	18.0	232.6	68.6	210.8	81.4	537.2	55.1	430.8	373.7	90.6	40.9
587	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	230.1	93.3	423.6	57.7	363.0	92.3	344.5	36.8	552.4	19.4	716.8	84.1	881.1	27.7	61.1	19.8	88.4	31.2
588	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAb1-2)Mana1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4(Fuca1-6)GlcNAcb	474.1	105.6	494.1	143.3	505.6	67.0	350.3	22.6	825.7	24.8	1043.3	343.9	890.4	67.1	892.0	865.1	11.6	68.3
589	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3GalNAca	1969.6	83.1	1850.1	75.5	1241.2	70.5	1702.2	54.3	1456.2	59.3	1356.3	127.3	1033.2	104.4	1277.8	64.4	1393.5	116.1
590	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	3344.0	133.9	2714.7	551.5	1954.7	108.8	2238.5	137.1	2676.0	158.5	2882.2	123.5	1414.9	101.4	1493.0	118.4	1240.7	58.9
591	Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3)GalNAca	1816.5	109.0	1518.5	103.0	962.0	67.4	1160.3	135.9	1364.2	63.4	1163.1	53.7	2157.8	75.7	926.5	26.3	955.1	19.6
592	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3GalNAca	412.3	45.9	364.7	75.1	225.2	24.2	436.1	95.8	318.0	34.6	234.1	32.6	269.7	115.5	116.3	22.8	75.0	18.5
593	GlcNAcb1-3Galb1-4GlcNAcb1-3GalNAca	438.0	67.5	324.9	41.0	135.5	51.4	146.0	42.4	272.0	57.9	190.9	28.4	311.8	92.7	98.9	50.5	51.8	31.6
594	GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	410.7	83.3	281.3	79.4	189.7	55.8	376.8	123.5	383.1	71.7	281.8	62.0	320.2	68.6	17.5	19.1	0.1	24.1
595	GlcNAcb1-3Galb1-4GlcNAcb1-6(GlcNAcb1-3Galb1-4GlcNAcb1-3)GalNAca	586.5	96.5	532.5	81.8	15.9	59.4	159.9	68.2	541.8	74.0	393.9	51.6	258.4	191.8	215.7	18.5	40.5	16.6
596	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3)GalNAca	884.1	75.6	826.0	95.3	574.9	27.4	1005.3	64.9	859.0	64.7	482.1	48.7	1720.4	252.3	357.8	30.3	301.7	61.8
597	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3GalNAca	468.6	49.2	306.9	17.5	327.1	69.6	214.4	62.8	287.4	83.4	267.3	87.2	280.8	91.1	131.3	63.6	78.7	3.9
598	GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3GalNAca	186.2	28.1	183.5	14.2	212.3	59.9	162.5	21.9	180.0	64.7	194.4	71.8	169.1	55.3	90.4	28.5	15.2	19.2
599	Galb1-4GlcNAcb1-3Galb1-3GalNAca	453.4	122.5	316.5	64.9	283.9	86.2	253.6	47.9	406.0	22.4	414.2	161.7	213.5	51.9	81.0	26.7	91.7	19.4
600	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	454.9	33.0	424.6	21.3	435.0	39.9	472.3	26.5	481.3	55.7	456.1	64.3	227.1	158.7	184.9	20.1	309.9	54.8
601	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	792.0	52.2	759.2	55.4	699.0	62.1	662.3	177.2	1050.0	119.3	1021.9	72.6	690.6	35.4	297.3	83.8	217.5	29.5
602	Neu5Aca2-6Galb1-4GlcNAcb1-6(Galb1-3)GalNAca	372.9	39.0	463.8	48.8	311.8	24.2	160.7	149.5	674.6	183.9	491.1	8.3	321.1	204.0	162.4	101.5	130.8	40.9
603	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	2592.2	131.2	2080.1	145.1	1551.7	114.4	1932.3	99.0	2369.8	106.3	2486.6	192.2	1723.8	835.9	1413.3	60.9	1320.0	106.7
604	GlcNAcb1-6(Neu5Aca2-3Galb1-3)GalNAca	653.8	122.4	478.7	50.0	432.0	39.2	276.2	280.5	674.2	47.6	1144.2	167.2	386.1	45.2	161.7	67.6	68.6	32.0
605	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-6(Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3)GalNAca	1949.4	114.3	1785.4	81.6	1047.7	53.3	1574.6	99.3	1527.4	124.8	1324.2	136.9	1877.9	83.7	1108.9	64.7	1303.4	30.4
606	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	1933.3	103.1	1669.8	191.2	1578.6	338.6	1713.3	158.7	1630.5	164.2	1435.5	77.2	2501.3	464.6	1073.9	37.0	1220.8	16.2
607	Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	4096.4	256.7	3727.5	749.4	2855.7	502.4	2825.6	99.7	3922.8	253.6	4294.9	134.0	7929.5	375.0	1939.9	137.5	2099.8	105.7
608	Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-6(Neu5Aca2-6Galb1-4GlcNAcb1-3Galb1-4GlcNAcb1-2Mana1-3)Manb1-4GlcNAcb1-4GlcNAcb	1814.8	163.2	1538.2	120.1	1063.5	79.5	1522.1	199.6	2016.6	157.0	1363.0	61.3	2417.3	192.6	508.8	1.2	245.1	68.4
609	GlcNAcb1-3Fuca	3225.2	162.9	3310.6	562.6	2894.3	180.7	2548.6	520.6	3042.2	374.1	5881.3	404.9	15112.5	507.9	1594.8	211.3	77.0	28.0
610	Galb1-3GalNAcb1-4(Neu5Aca2-8Neu5Aca2-8Neu5Aca2-3)Galb1-4Glcb	707.6	97.2	823.0	85.0	807.7	112.1	1678.6	310.5	778.3	55.2	652.1	76.9	974.5	192.5	207.7	58.3	78.2	13.0