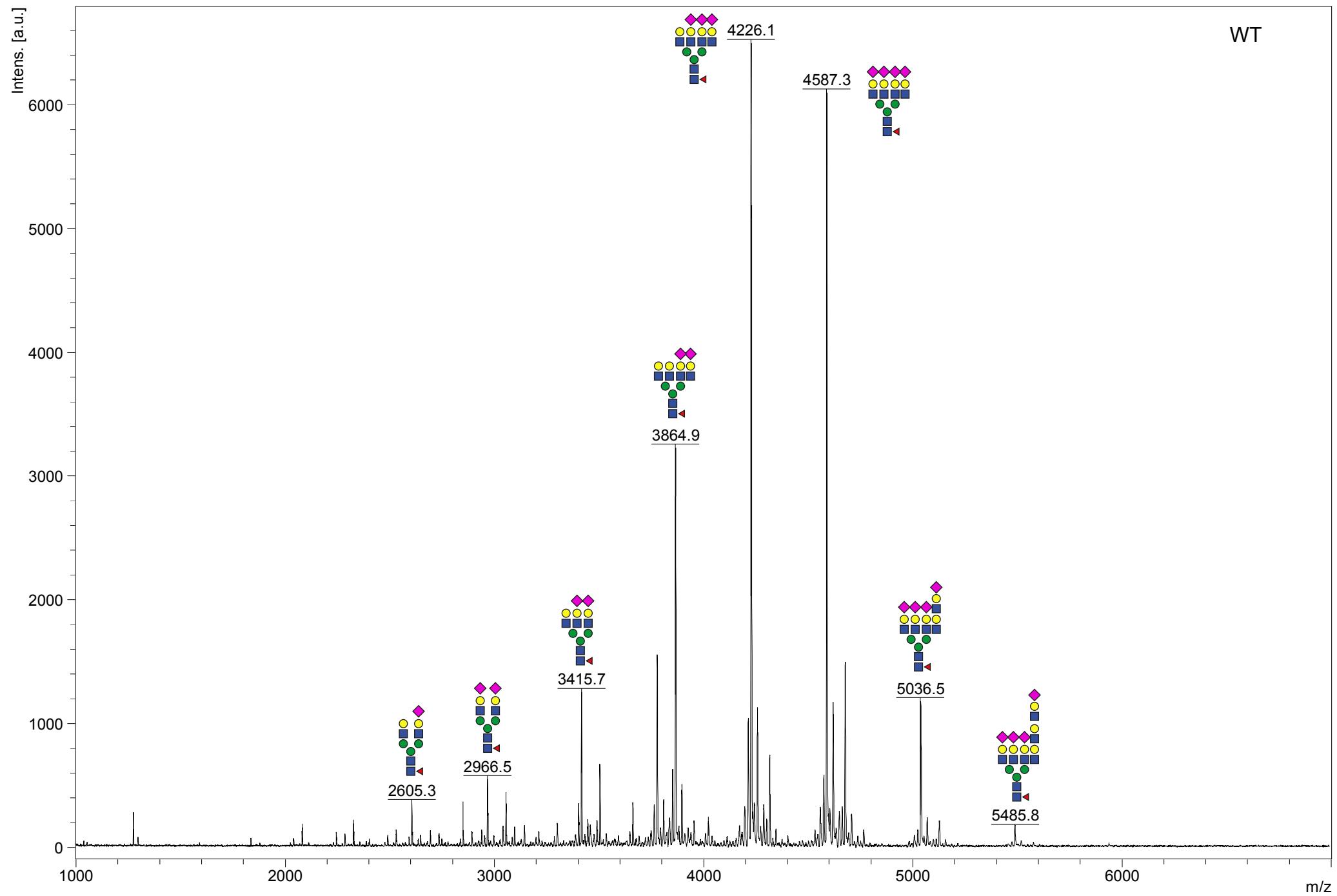
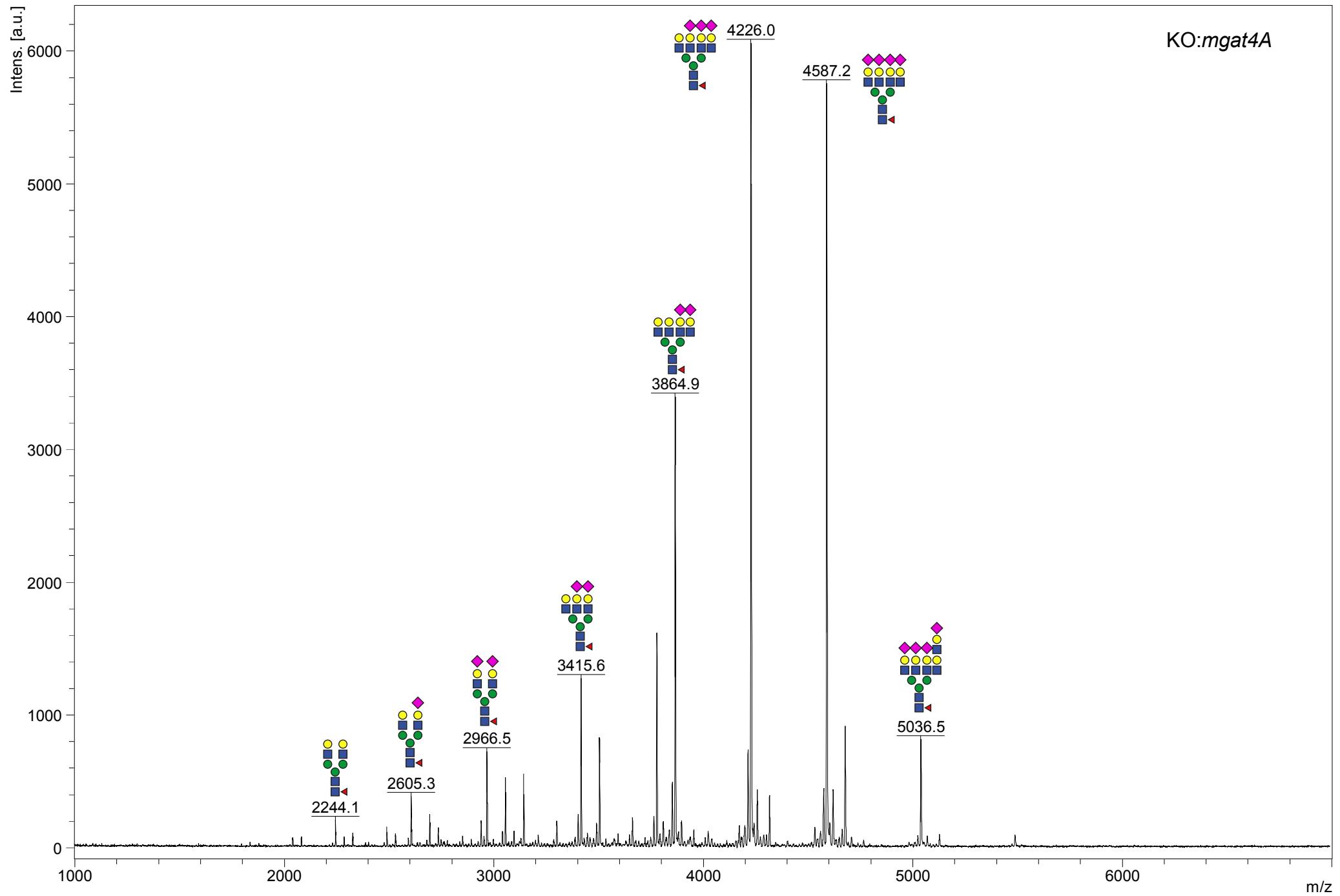


Supplementary Spectra

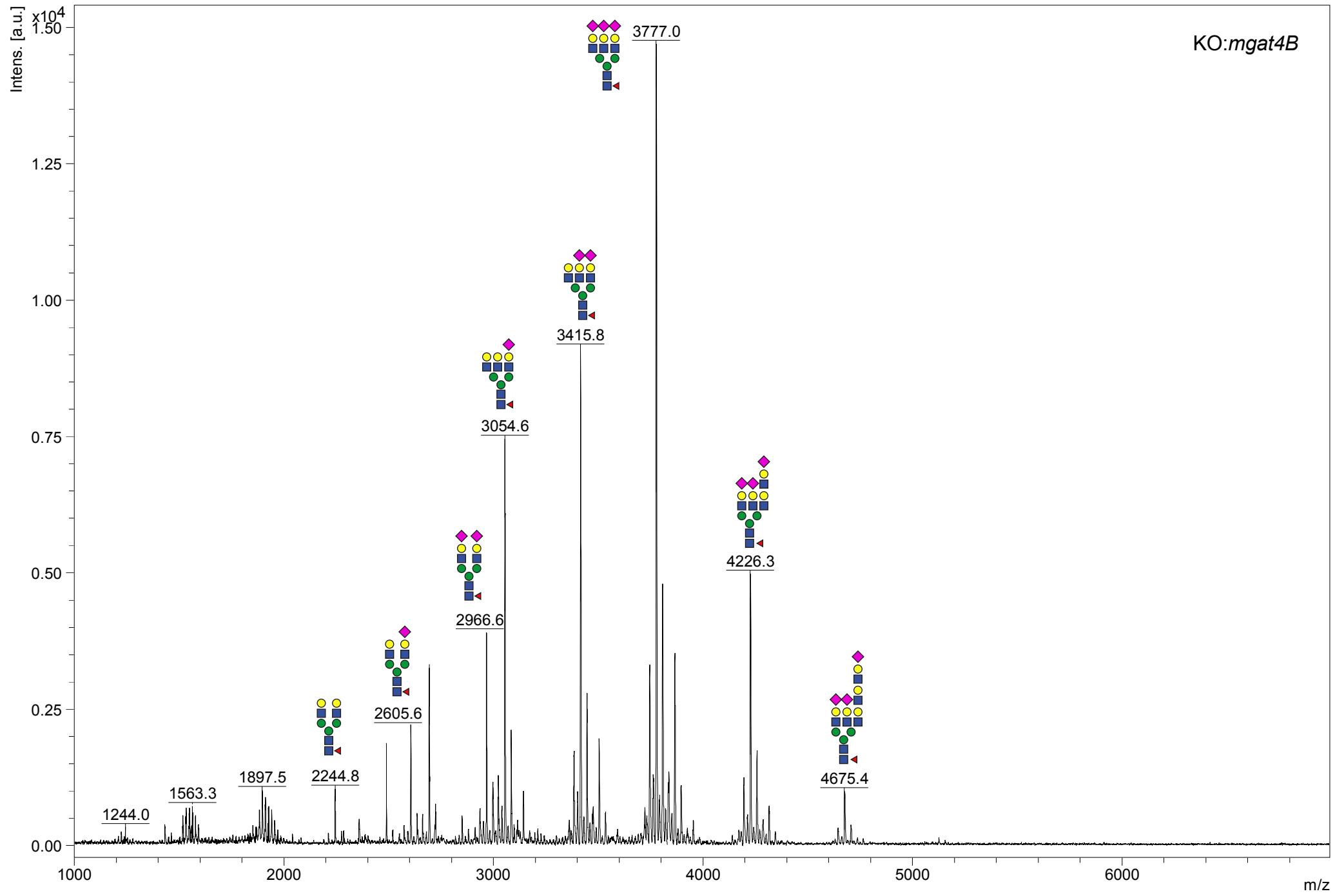
MALDI-TOF spectra of PNGase F released permethylated N-glycans with predicted structures for EPO (spectra #1-34) and IgG (spectra #35-38) samples.
#1



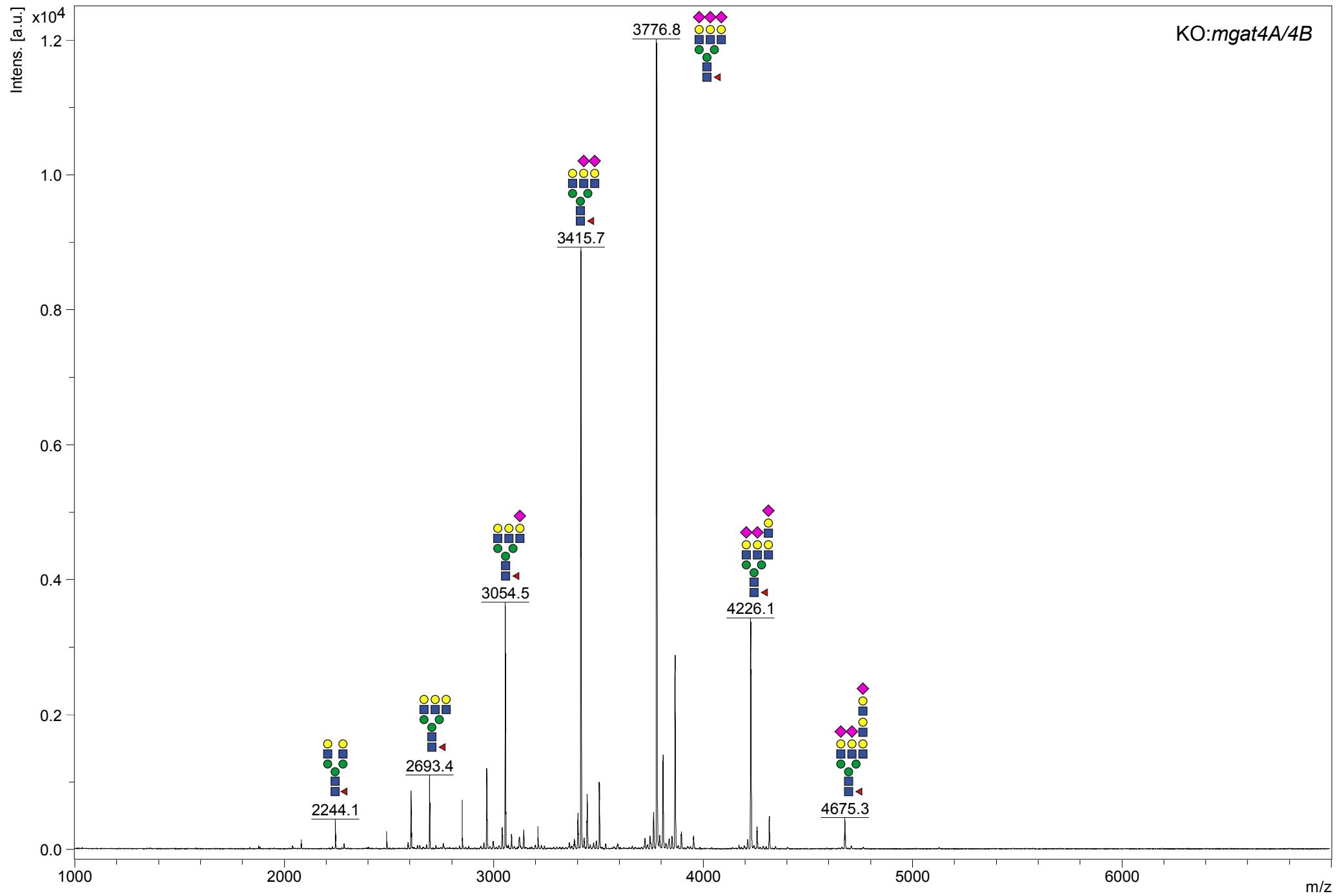
Supplementary Spectra
#2



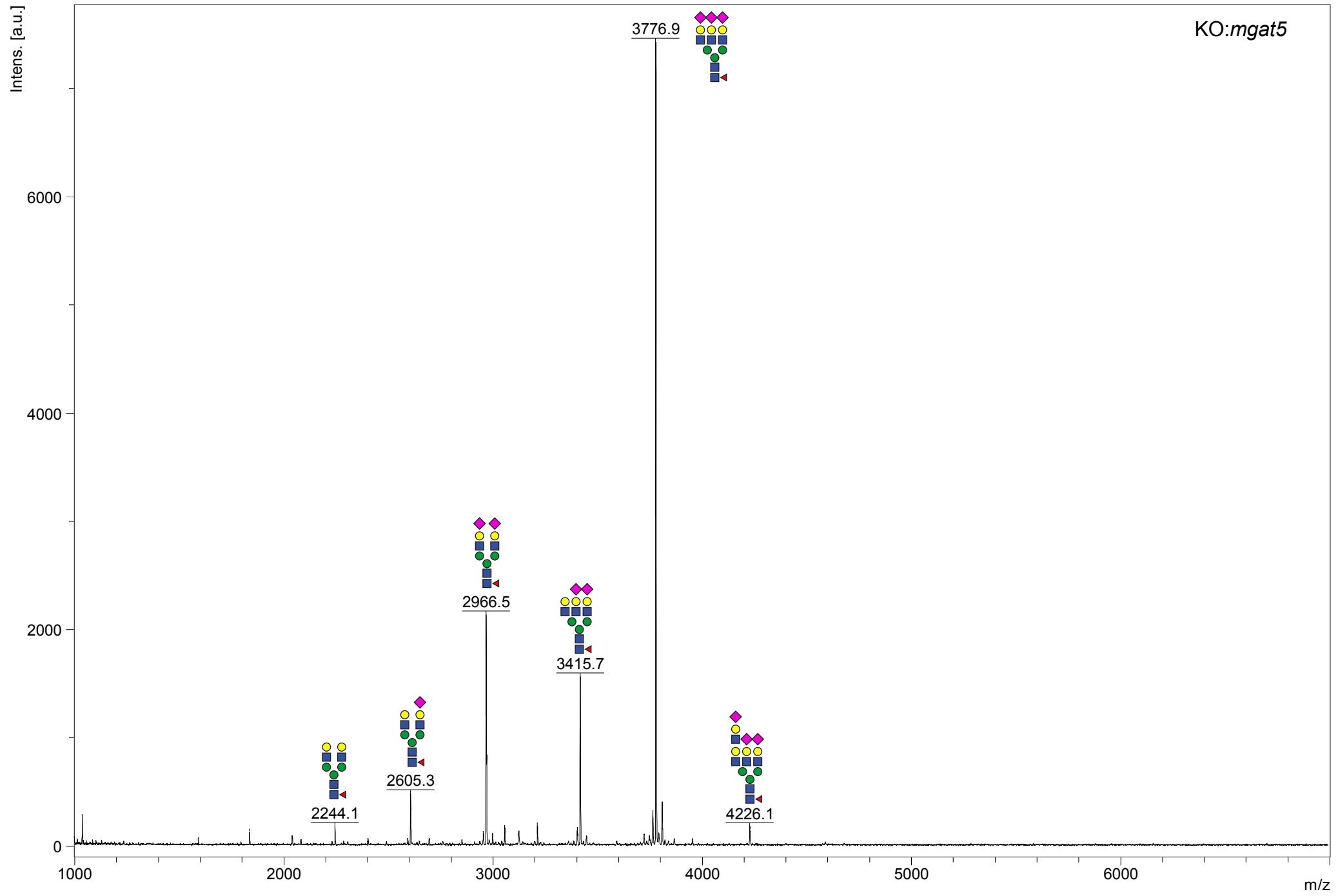
Supplementary Spectra
#3



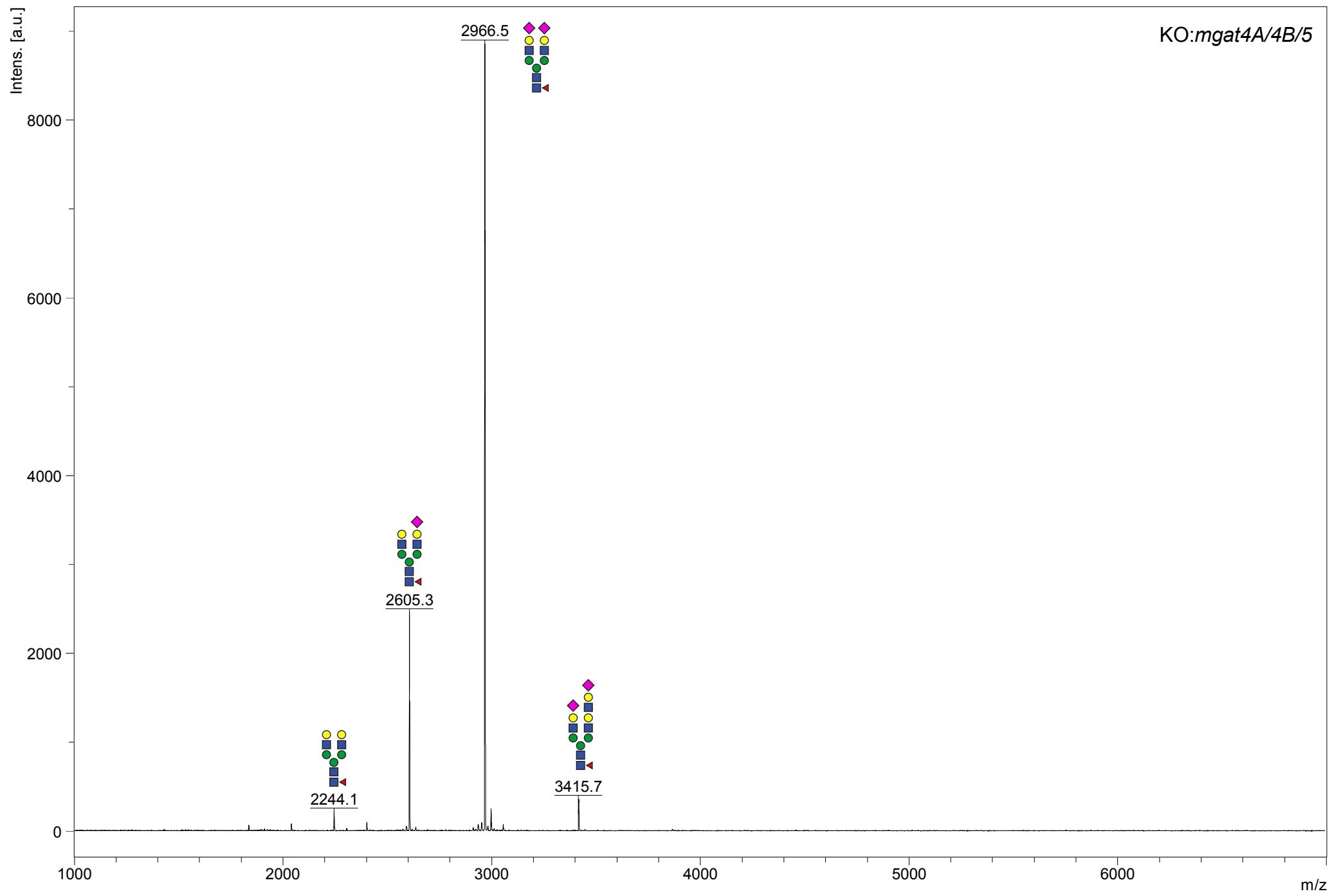
Supplementary Spectra
#4

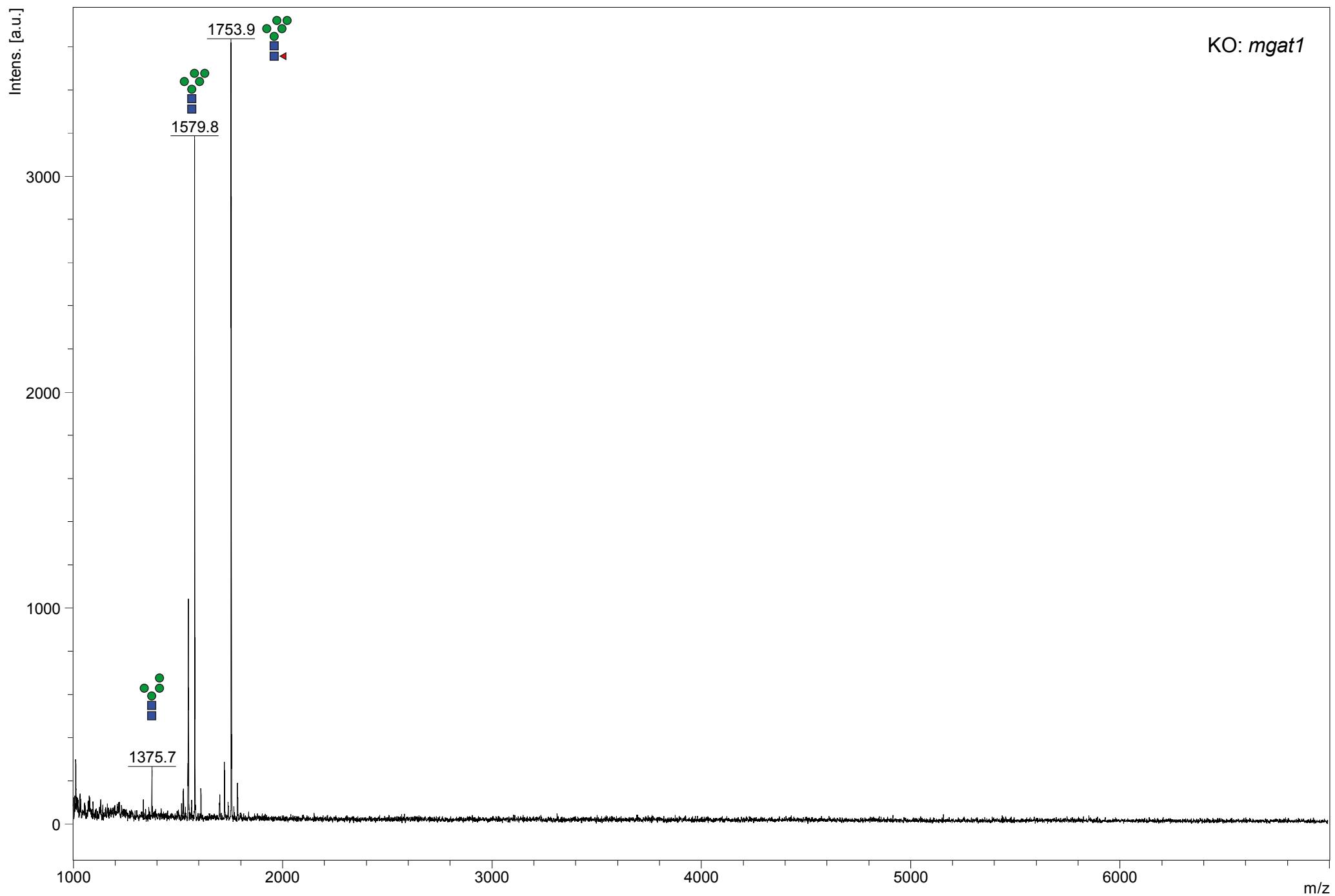


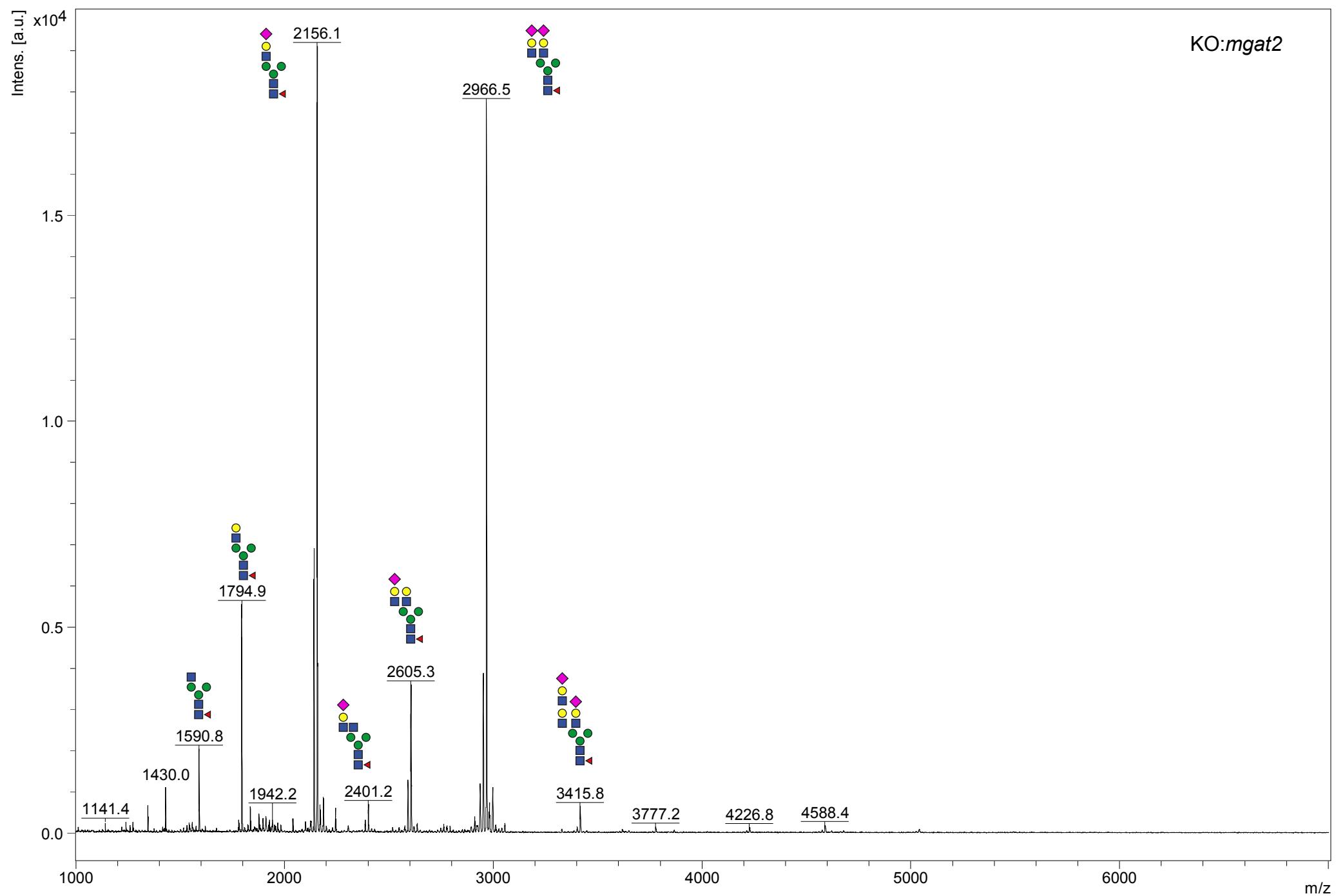
Supplementary Spectra
#5



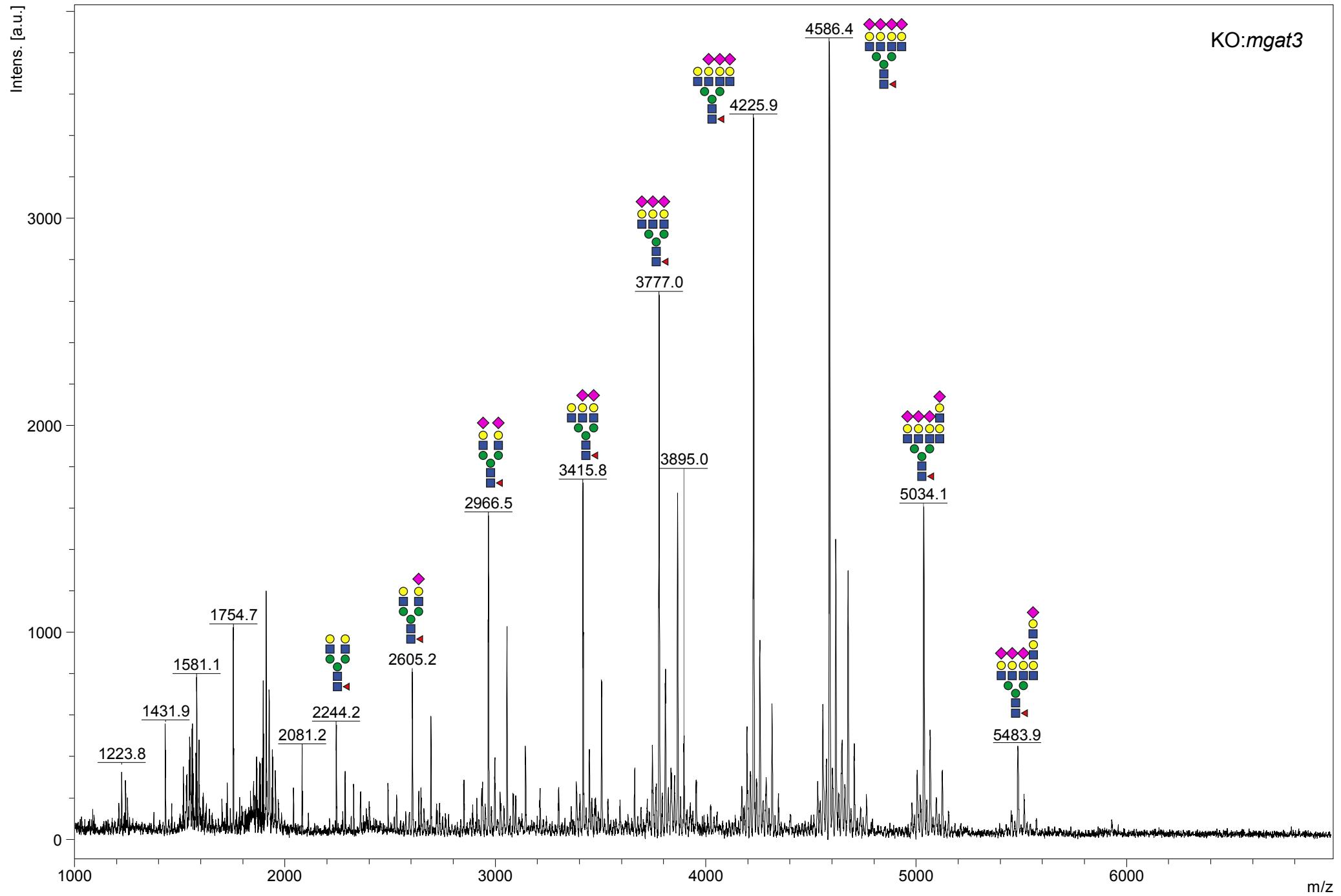
Supplementary Spectra
#6



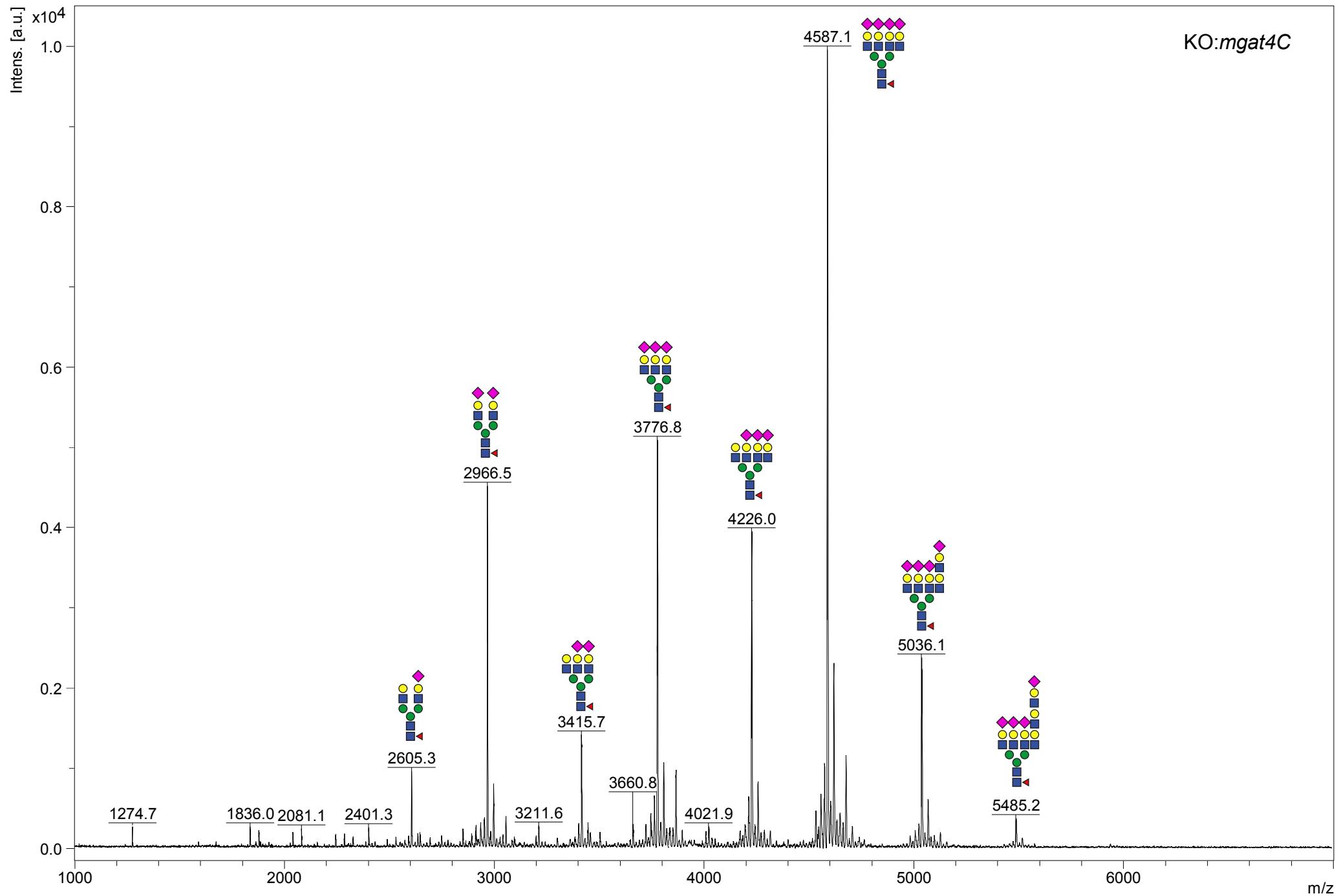




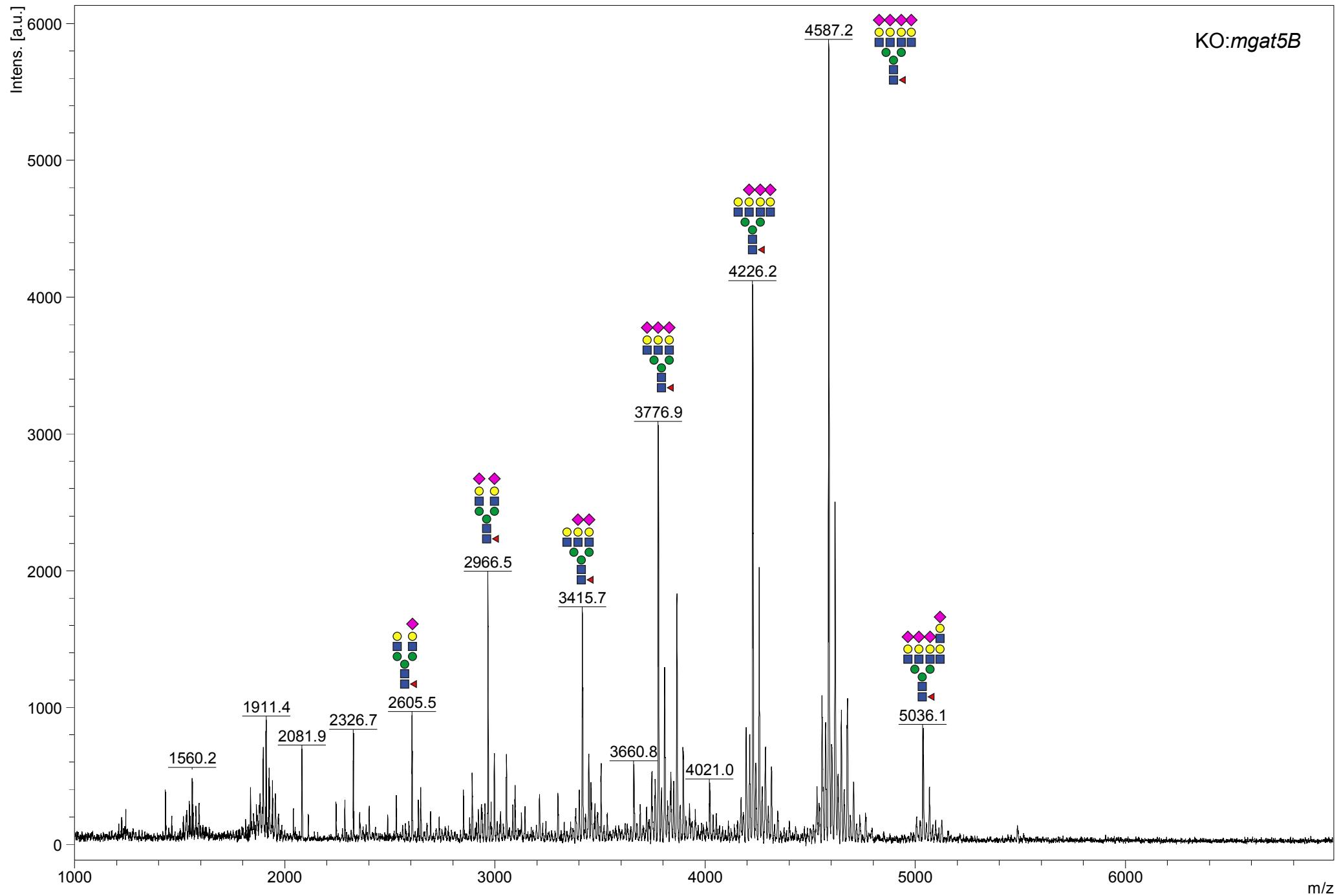
Supplementary Spectra
#9



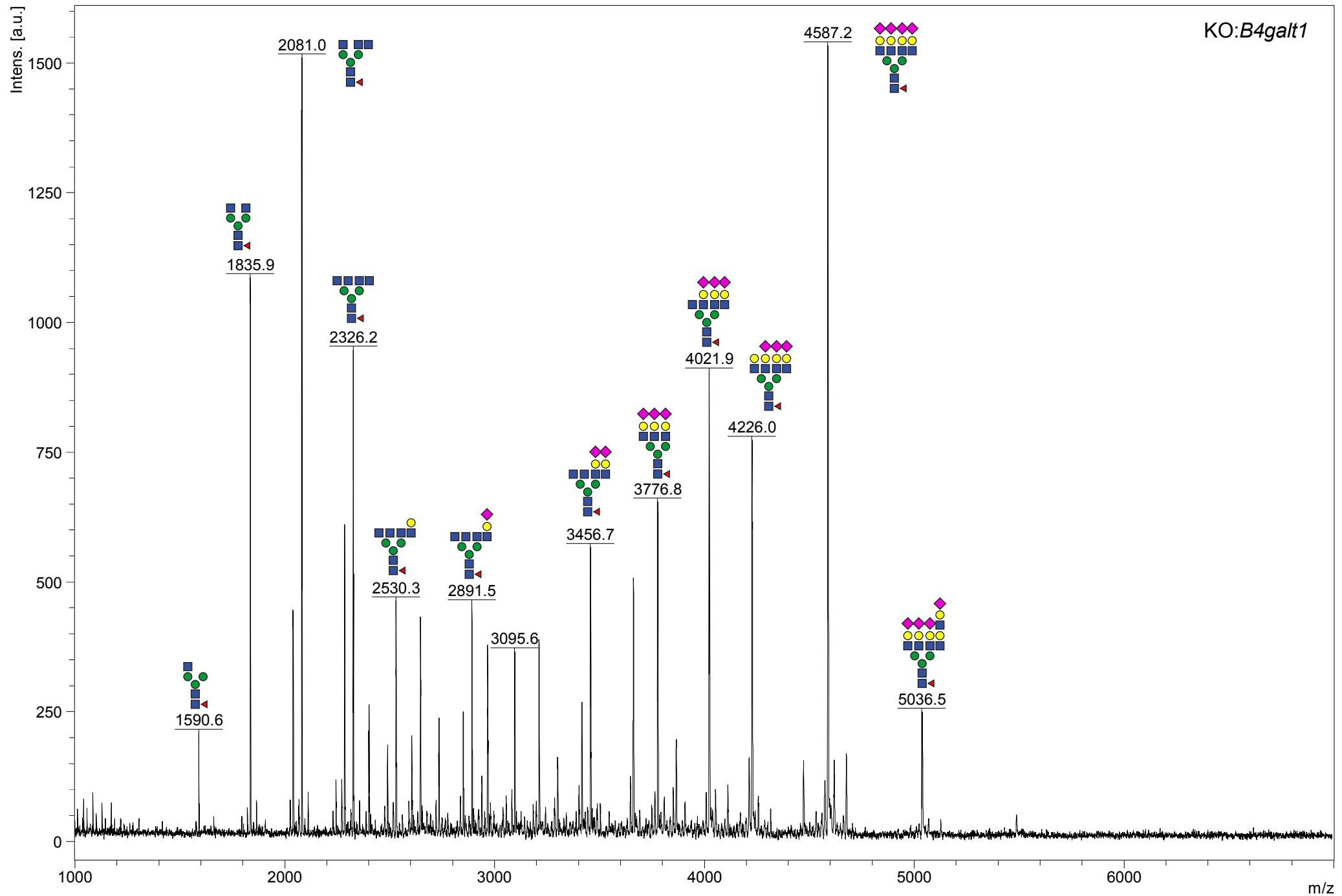
Supplementary Spectra
#10



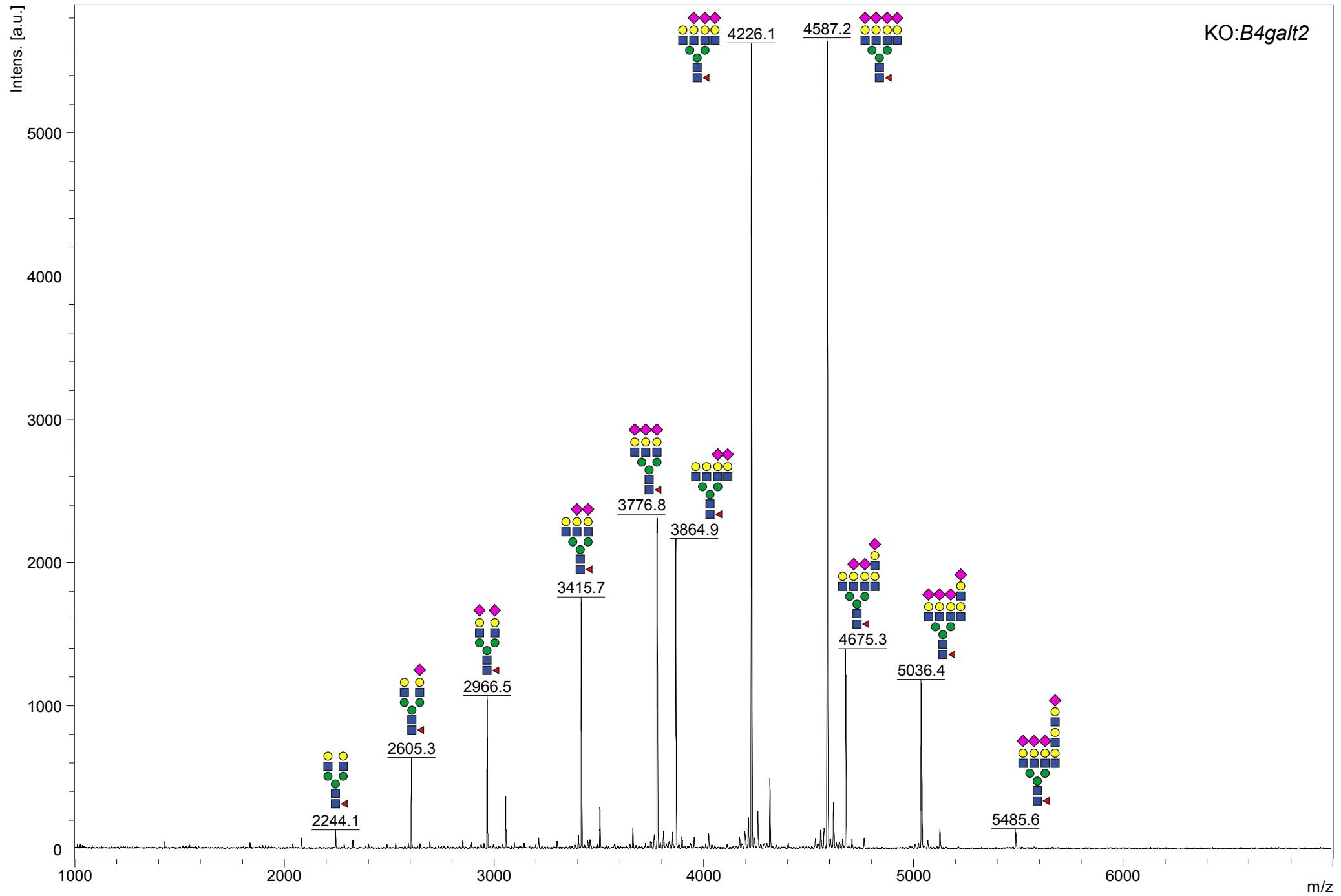
Supplementary Spectra
#11



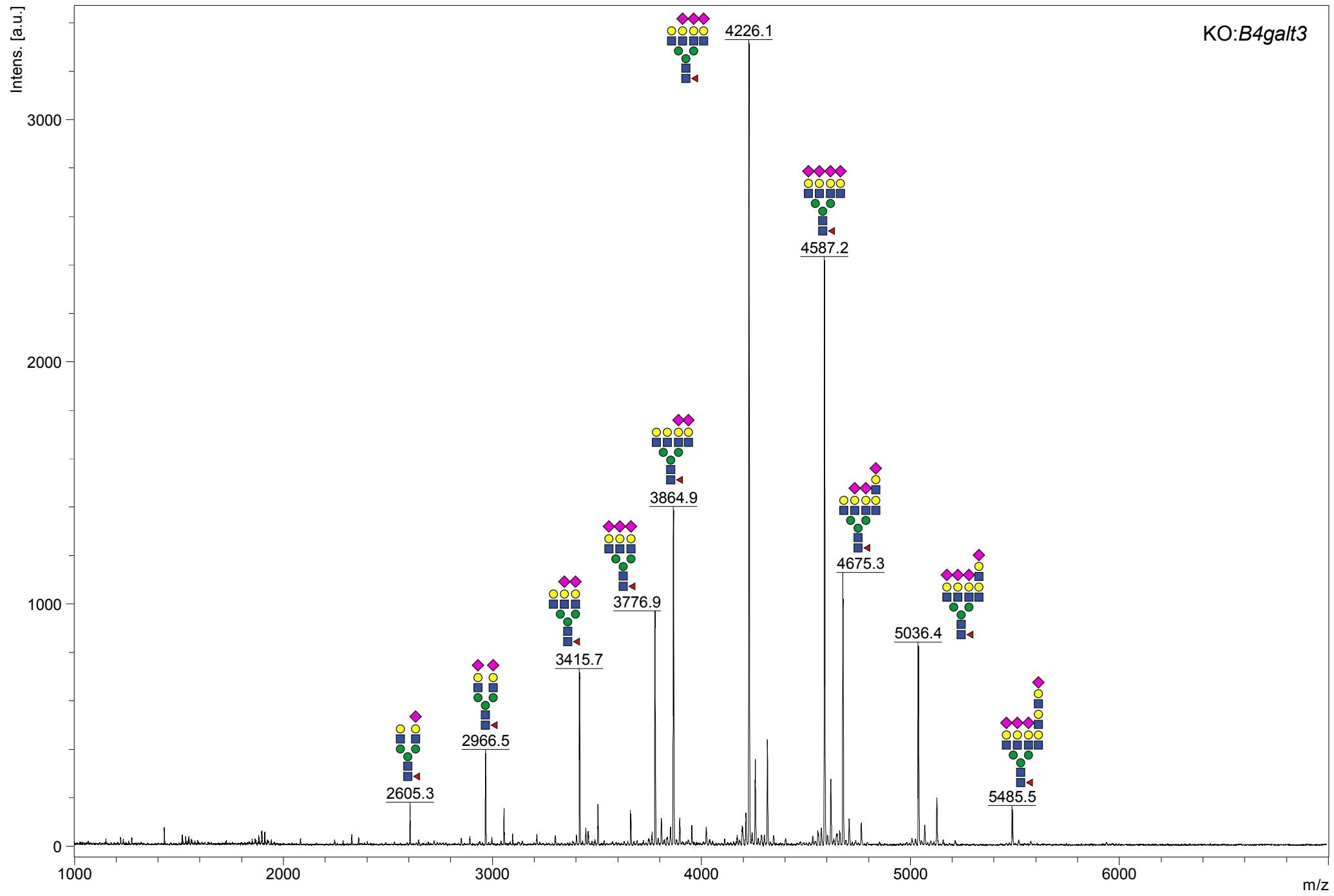
Supplementary Spectra
#12



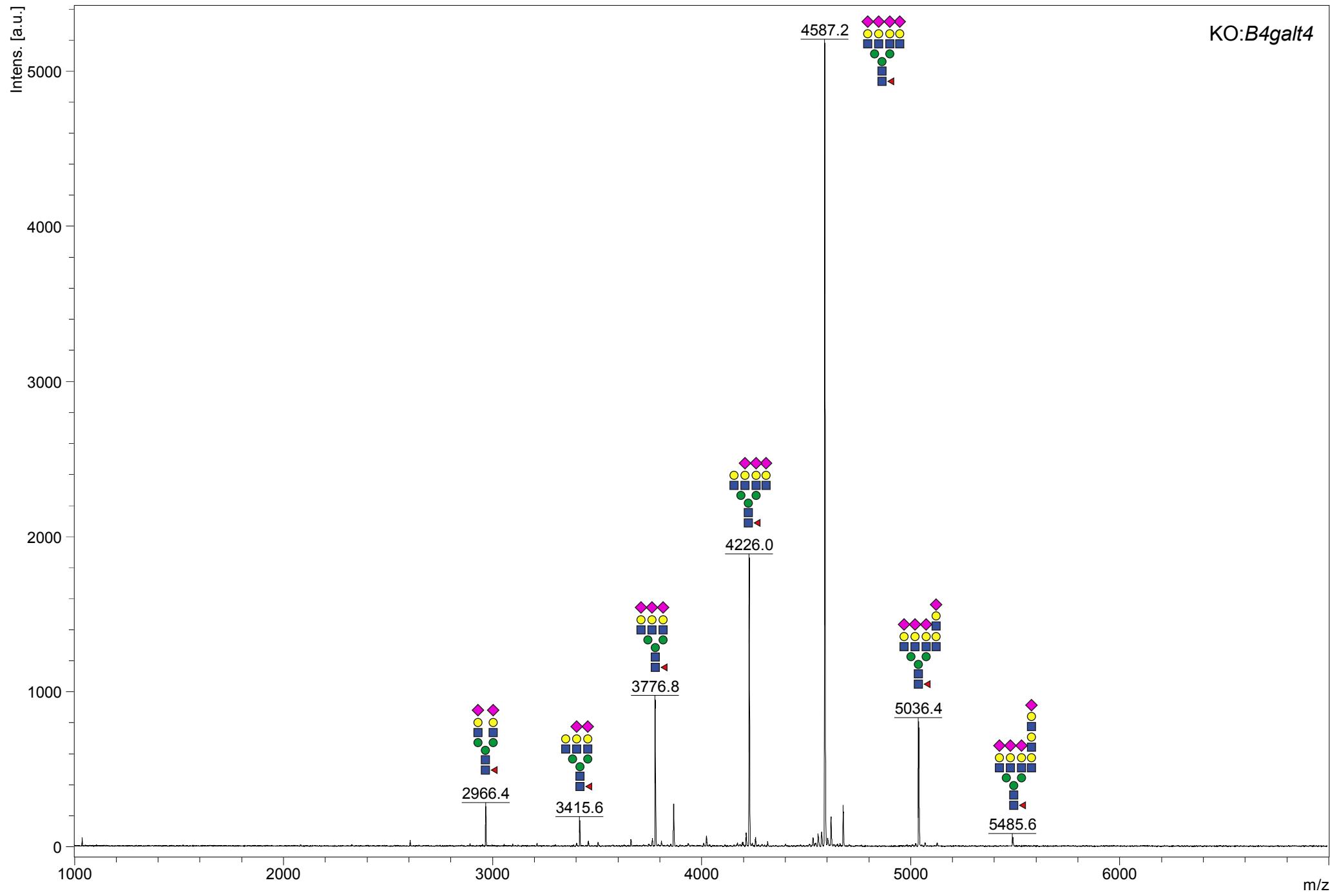
Supplementary Spectra
#13



Supplementary Spectra
#14

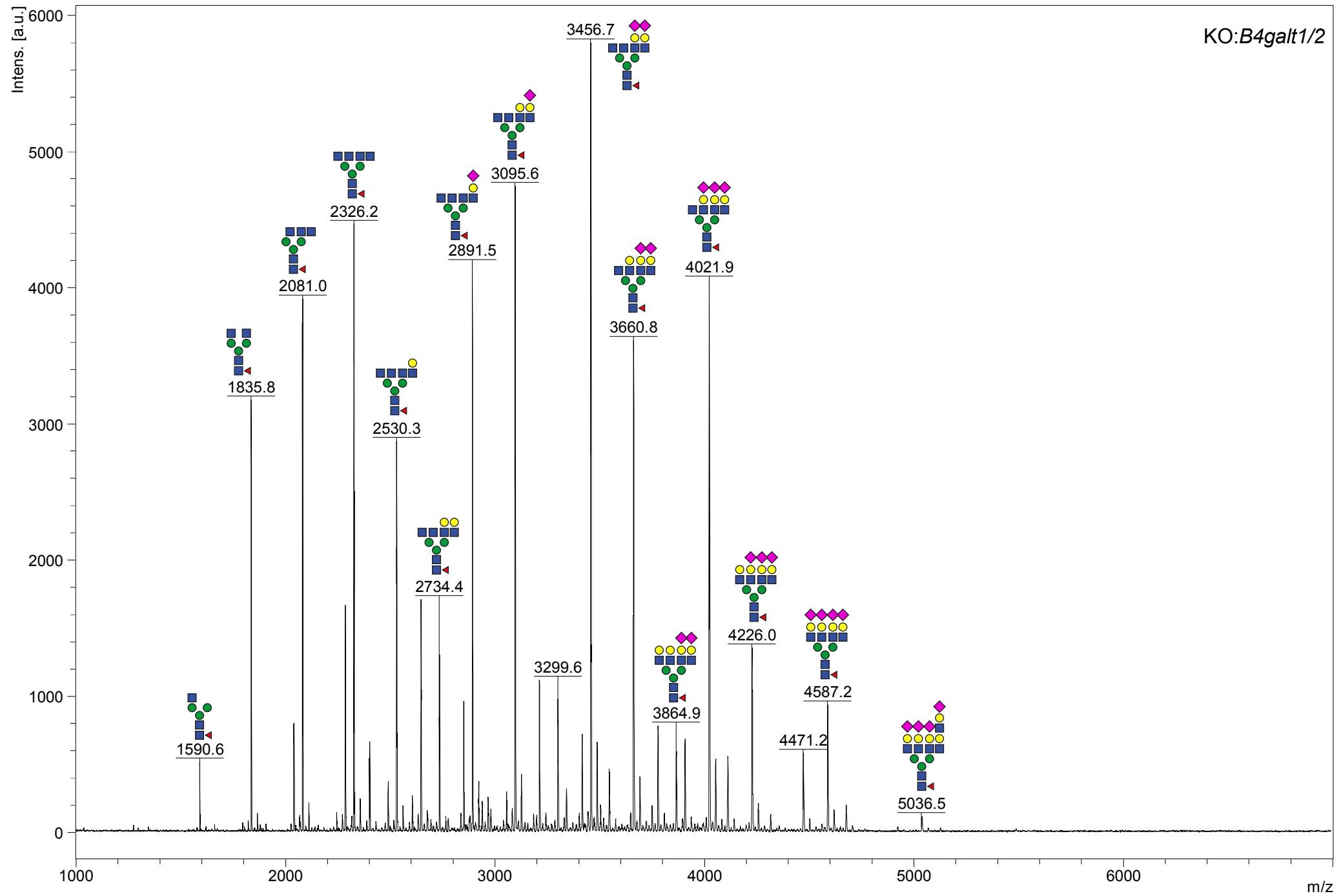


Supplementary Spectra
#15

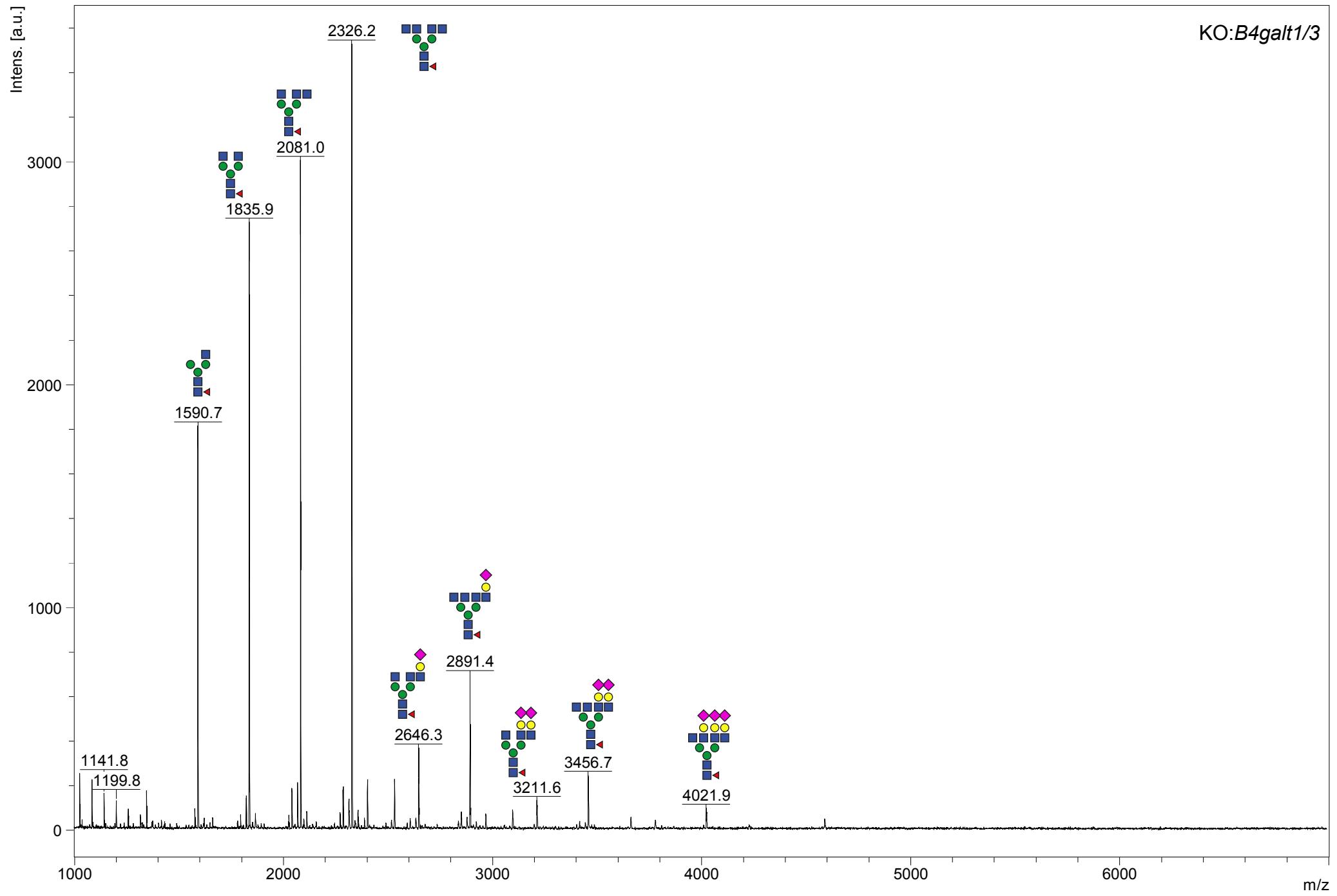


Supplementary Spectra
#16

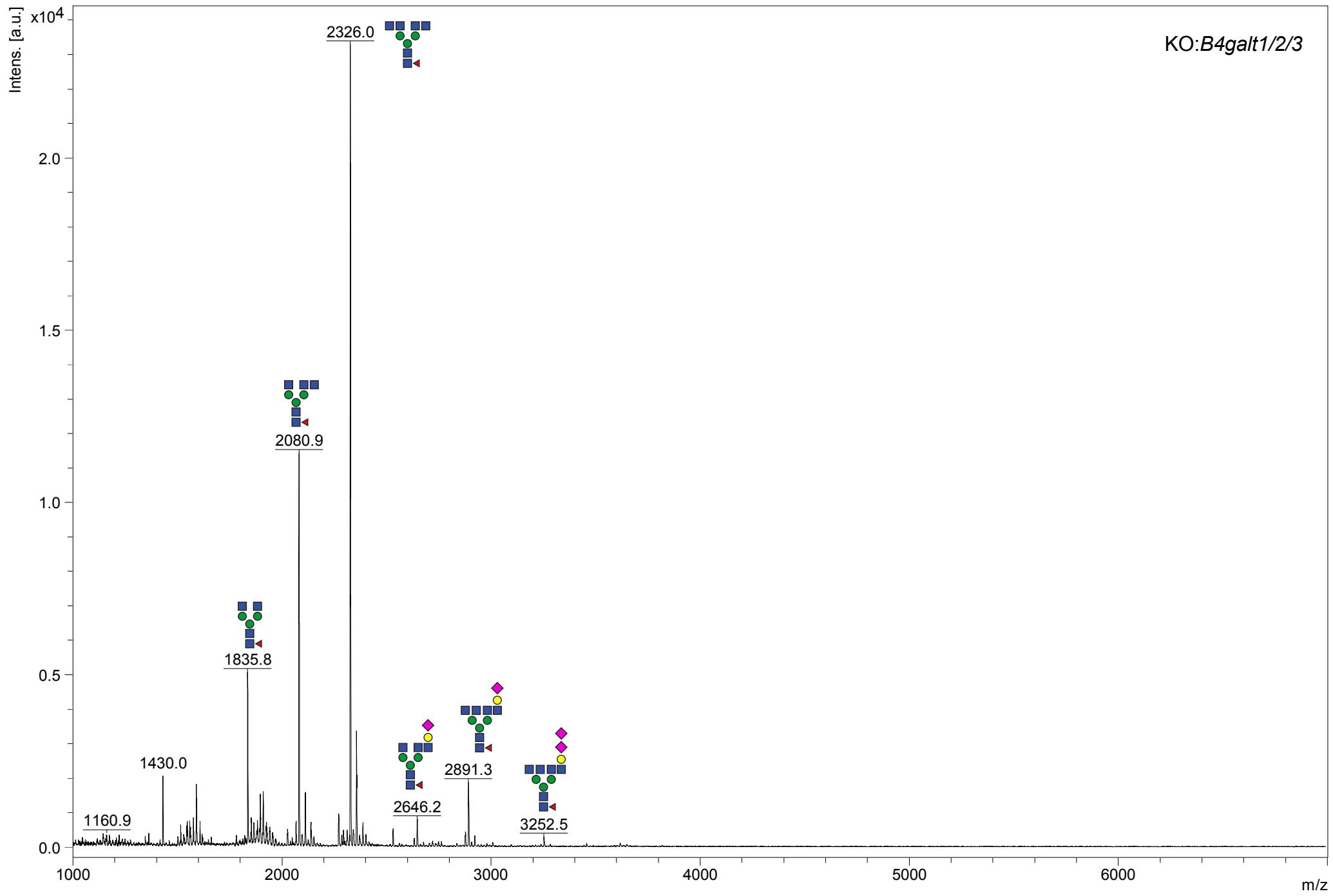
KO:*B4galt1/2*



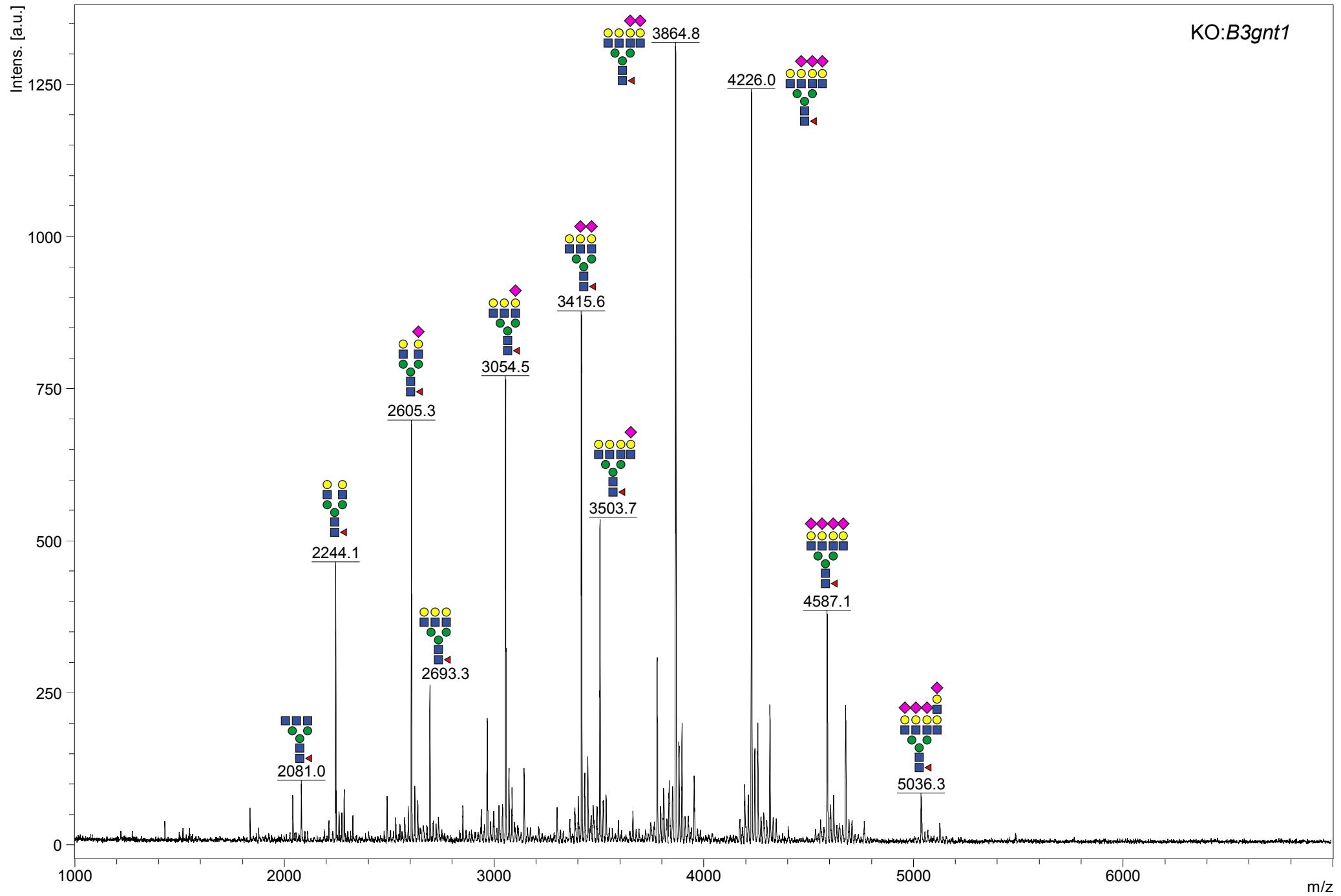
Supplementary Spectra
#17



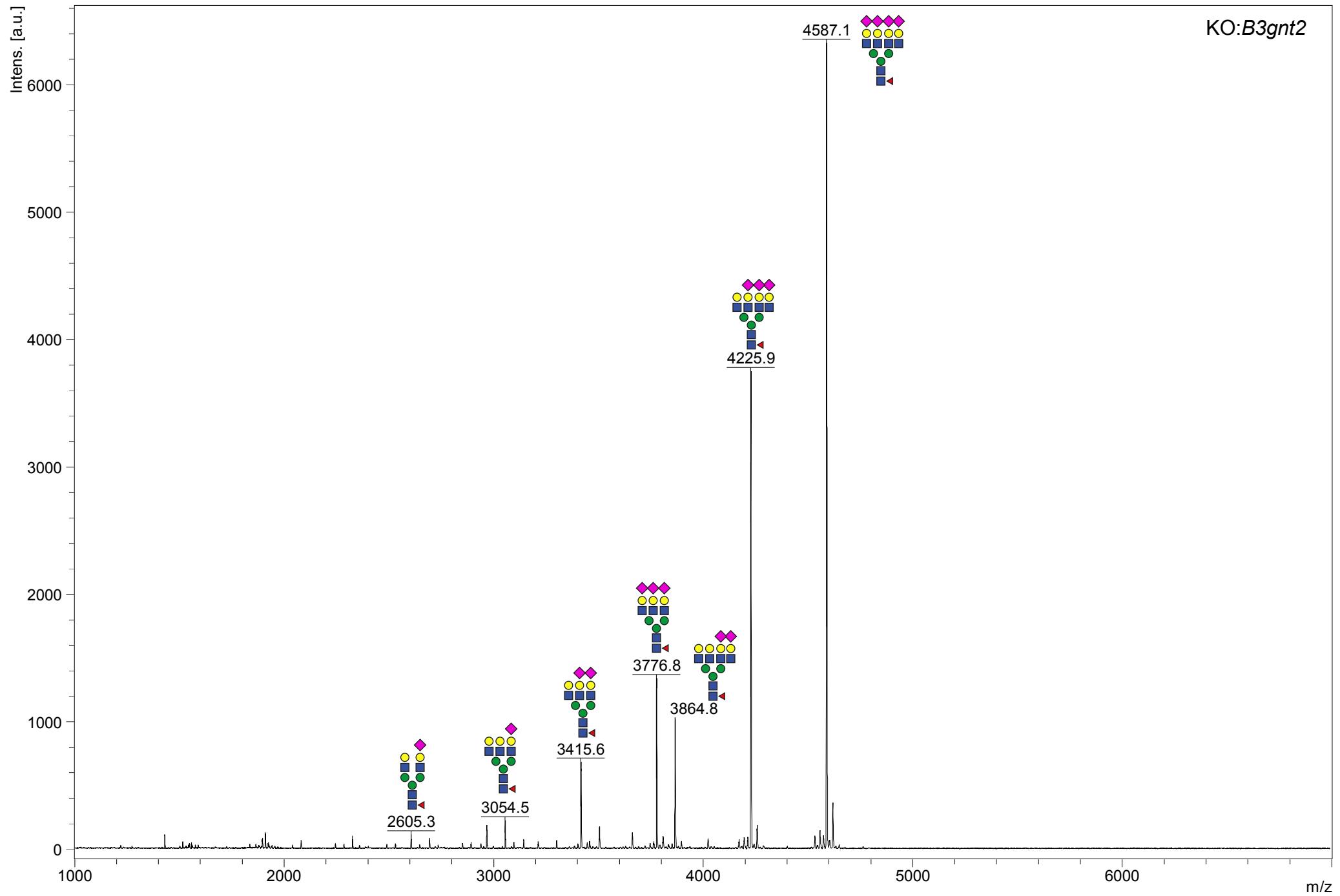
Supplementary Spectra
#18



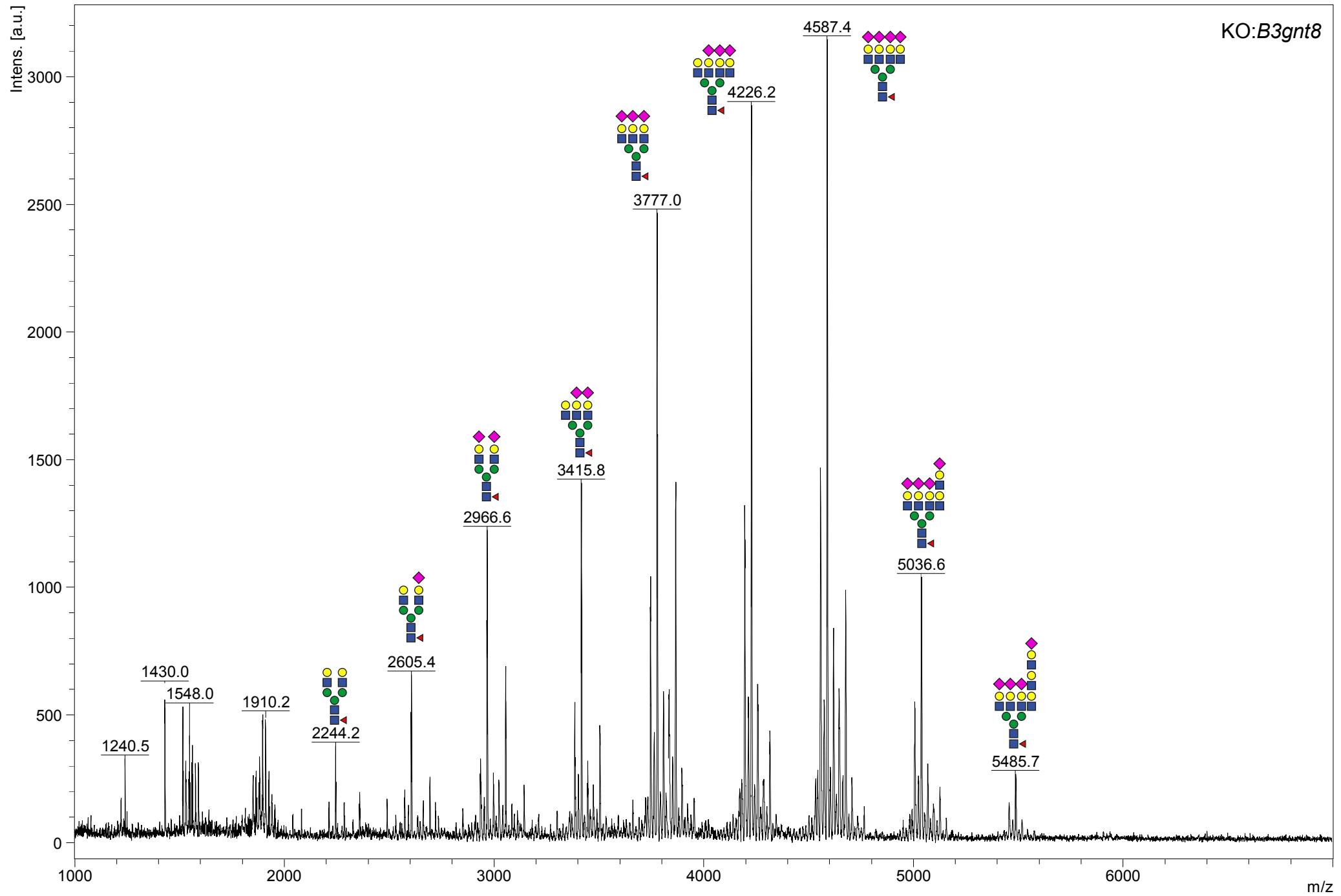
Supplementary Spectra
#19



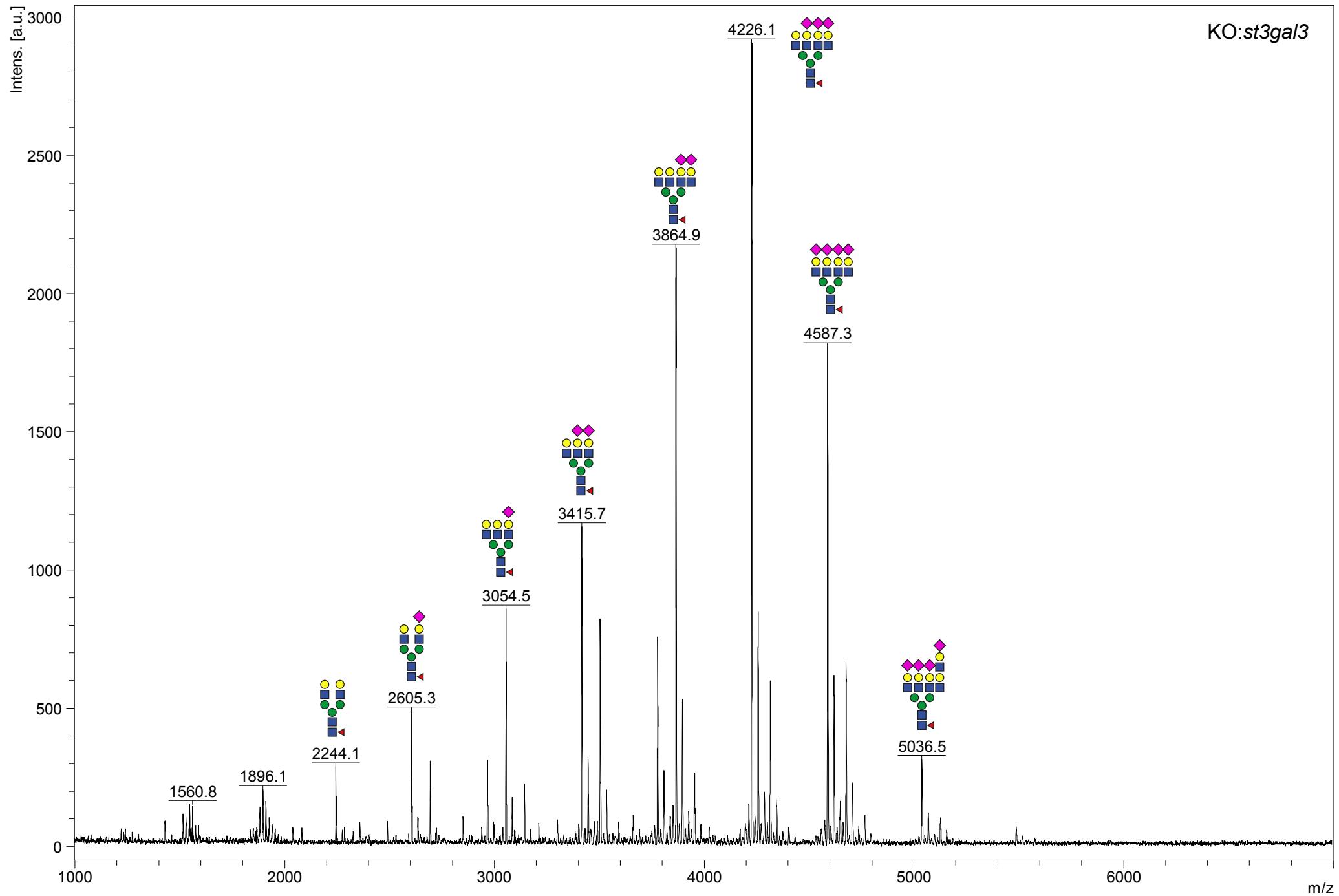
Supplementary Spectra
#20



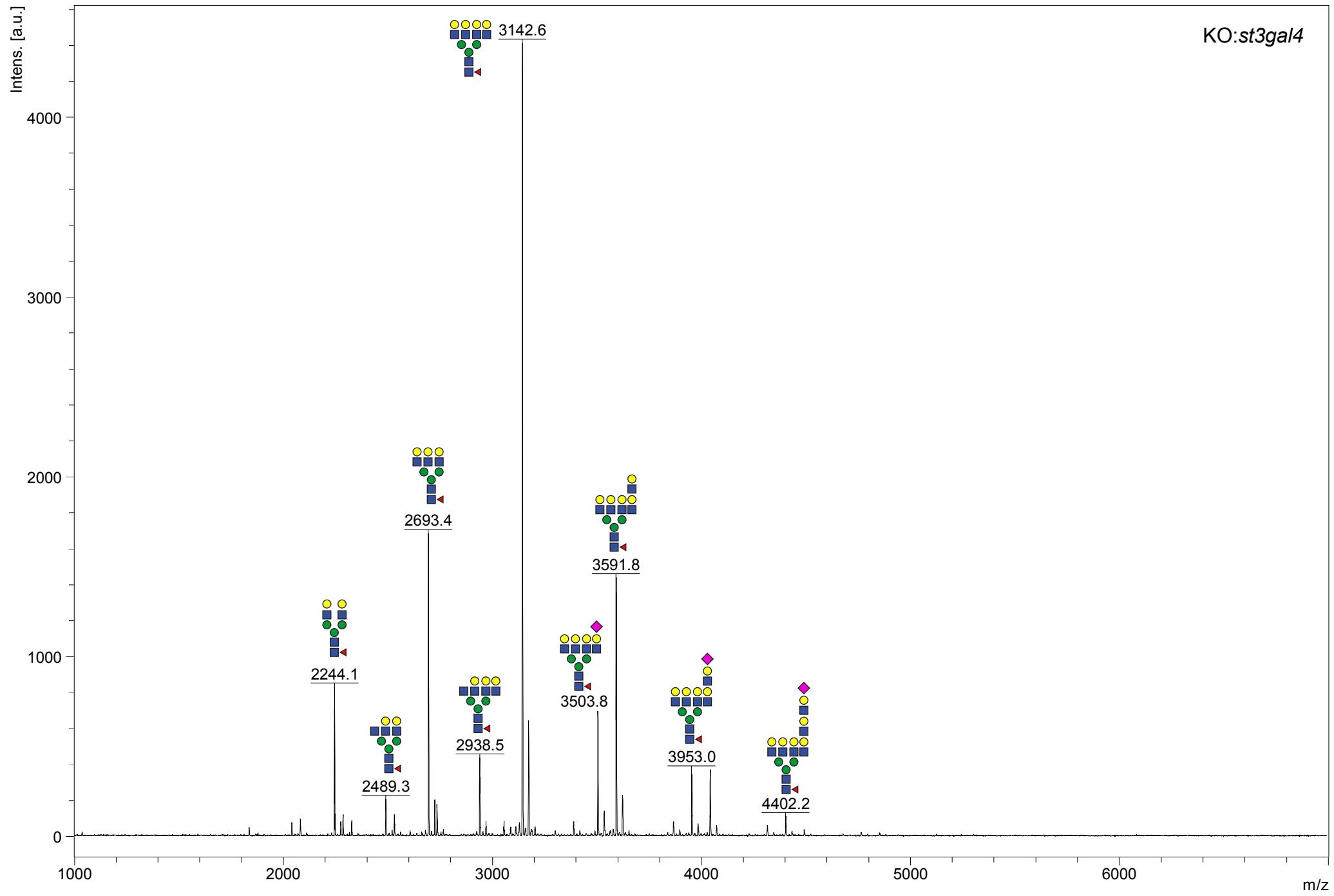
Supplementary Spectra
#21



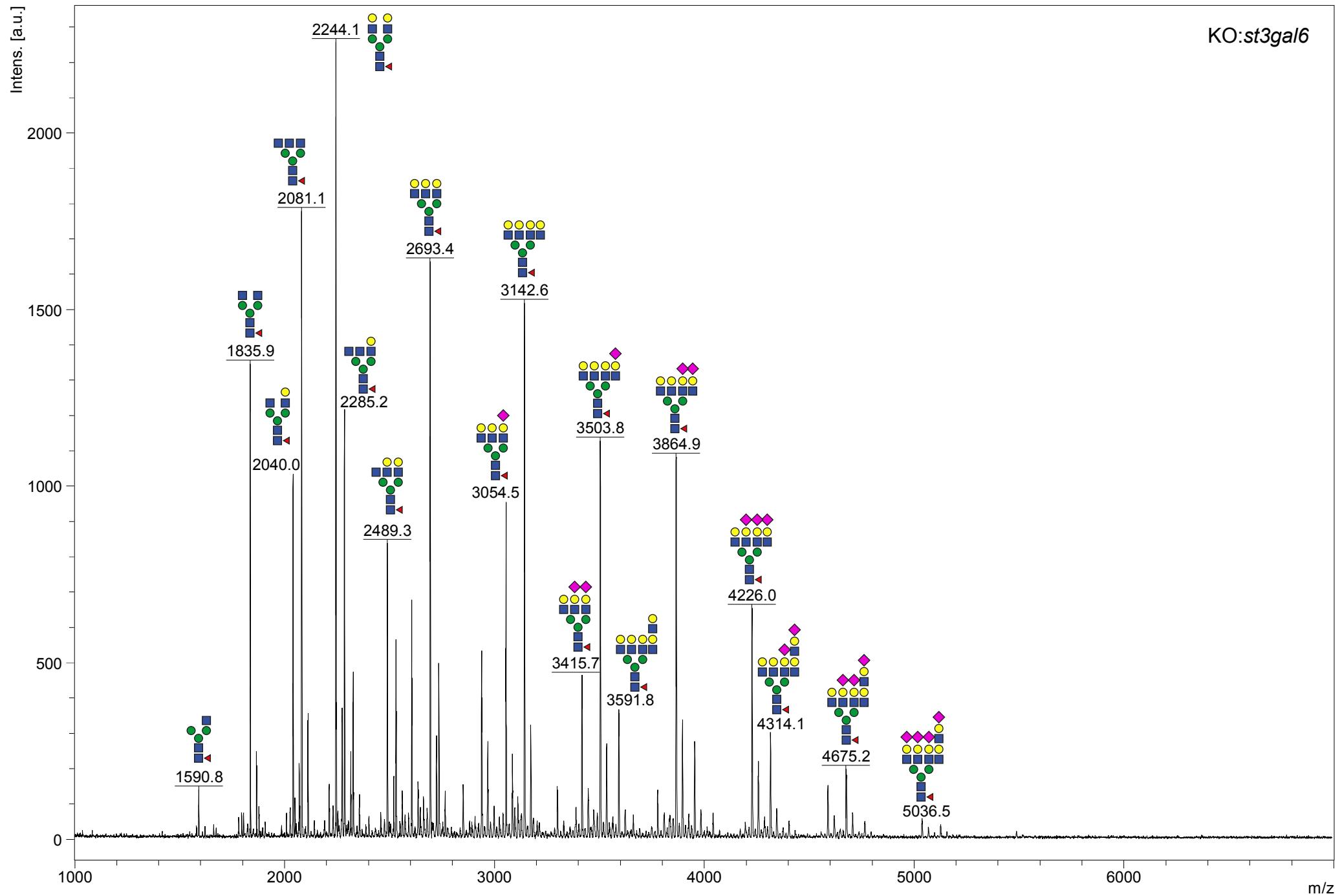
Supplementary Spectra
#22



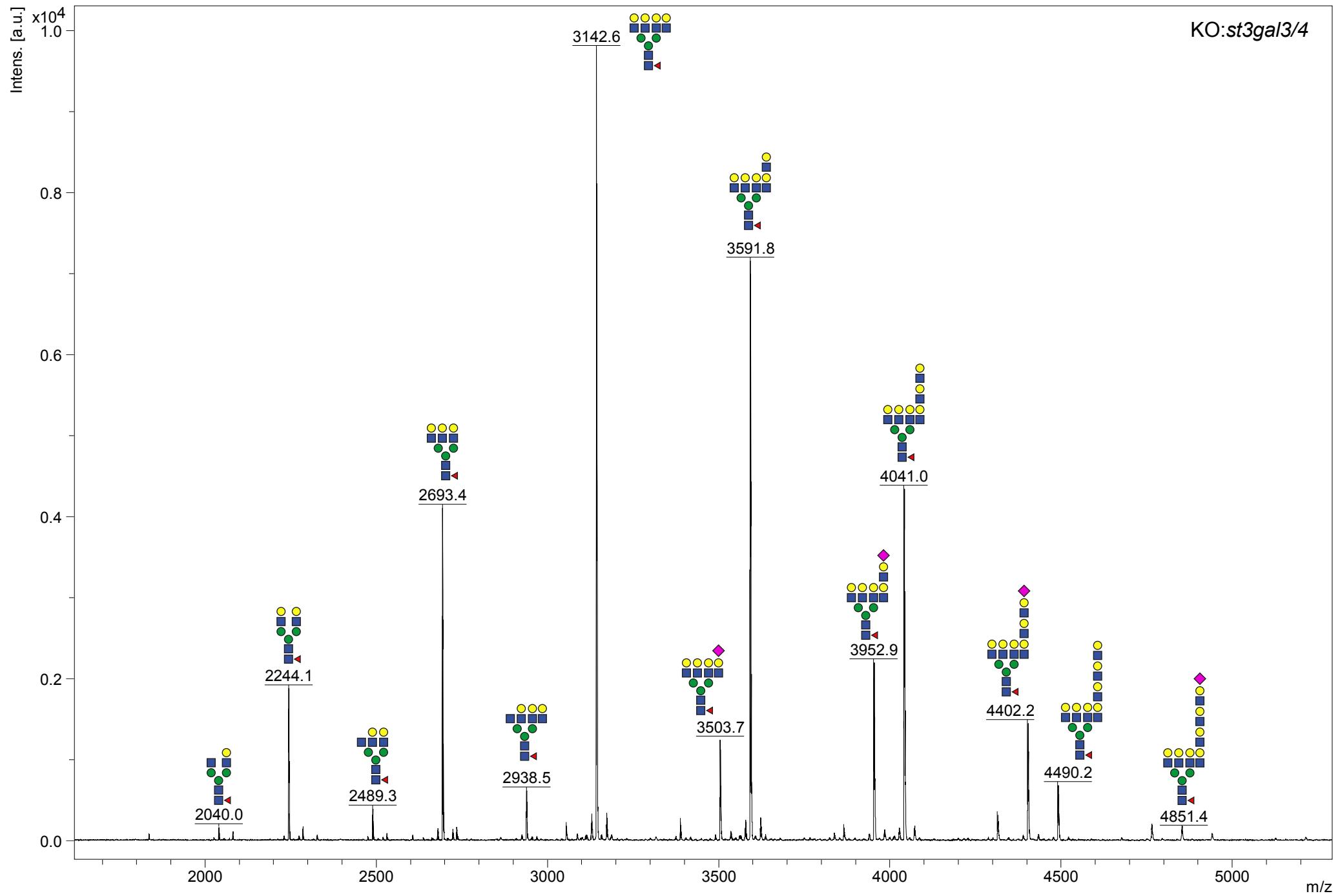
Supplementary Spectra
#23



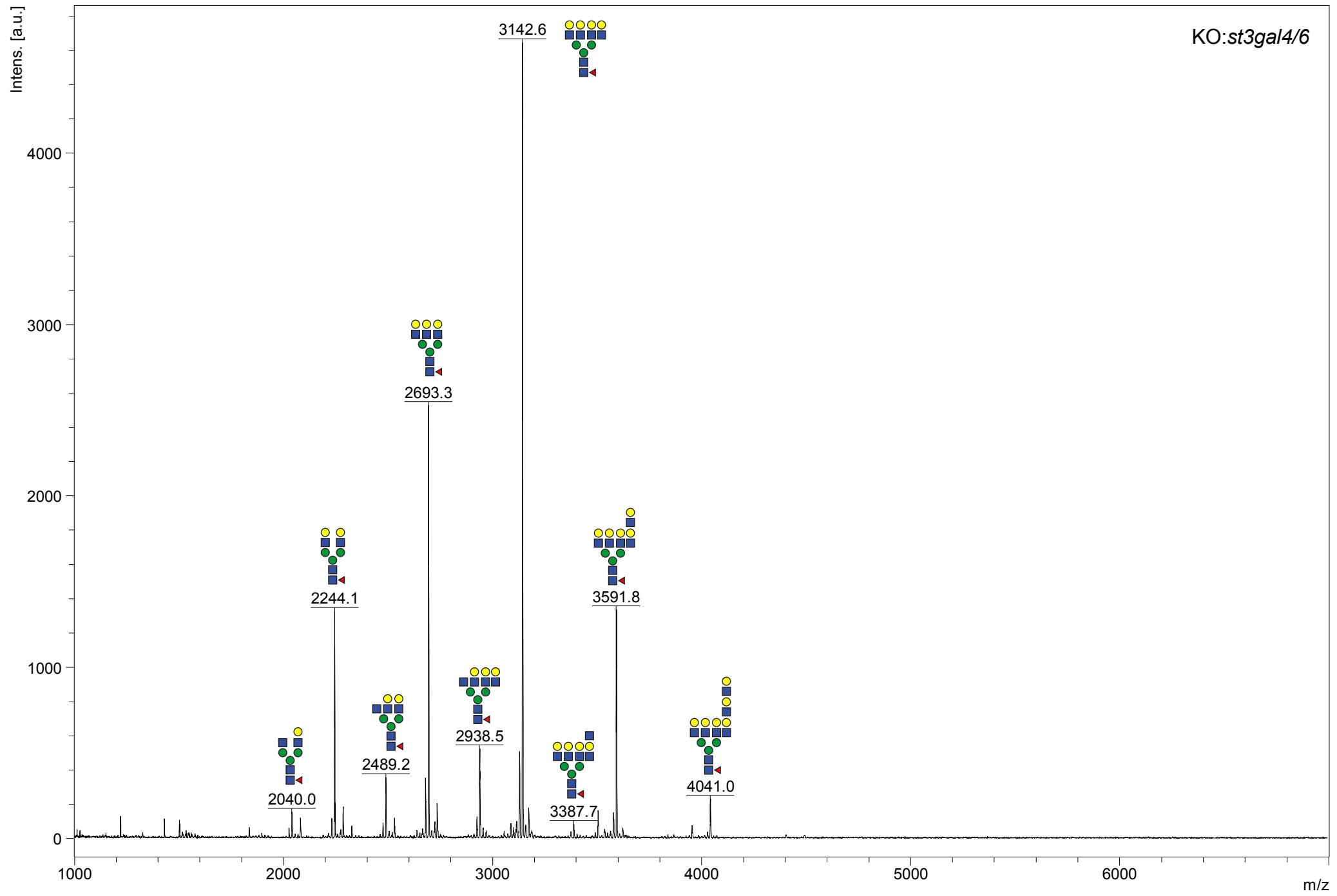
Supplementary Spectra
#24



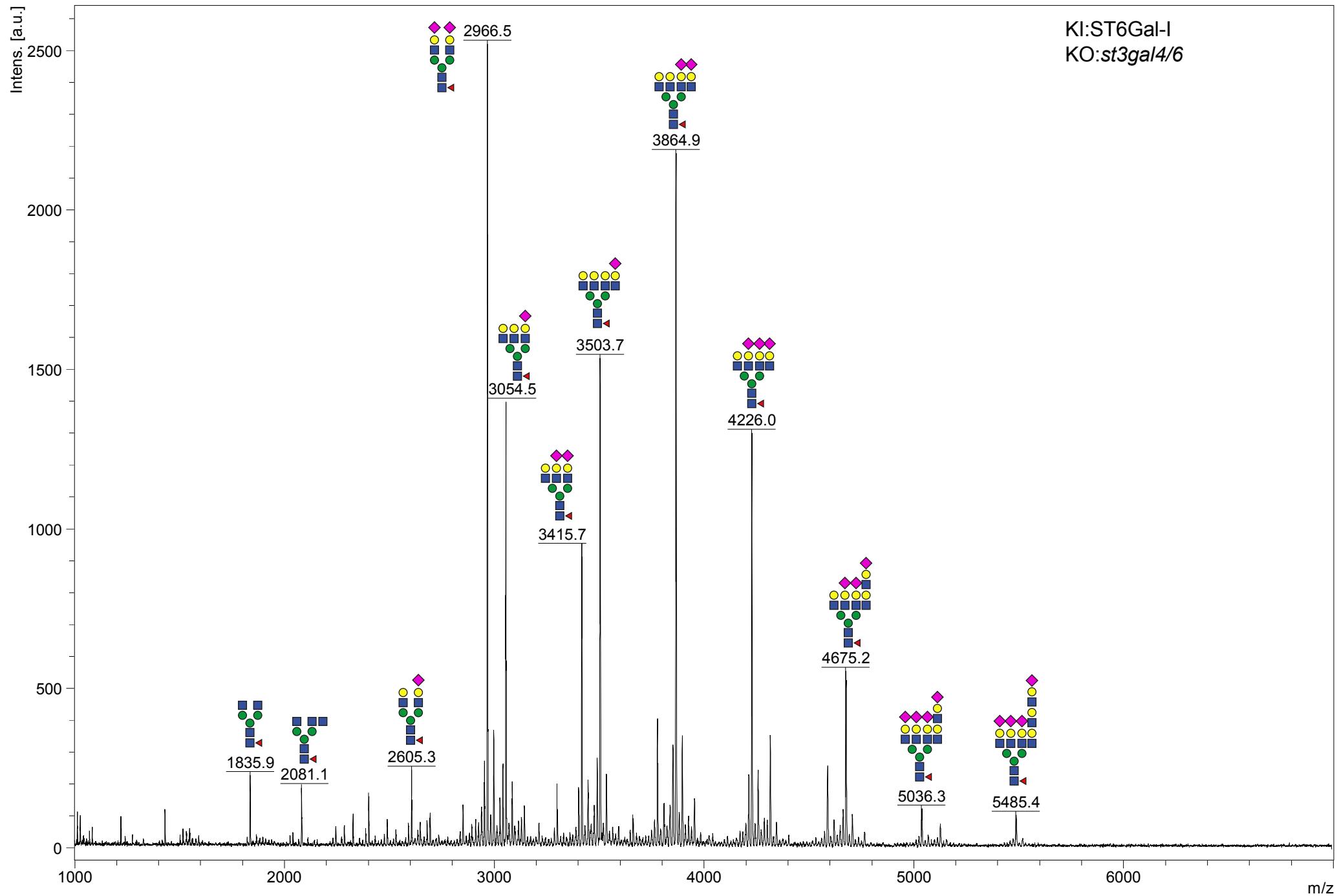
Supplementary Spectra
#25



Supplementary Spectra
#26

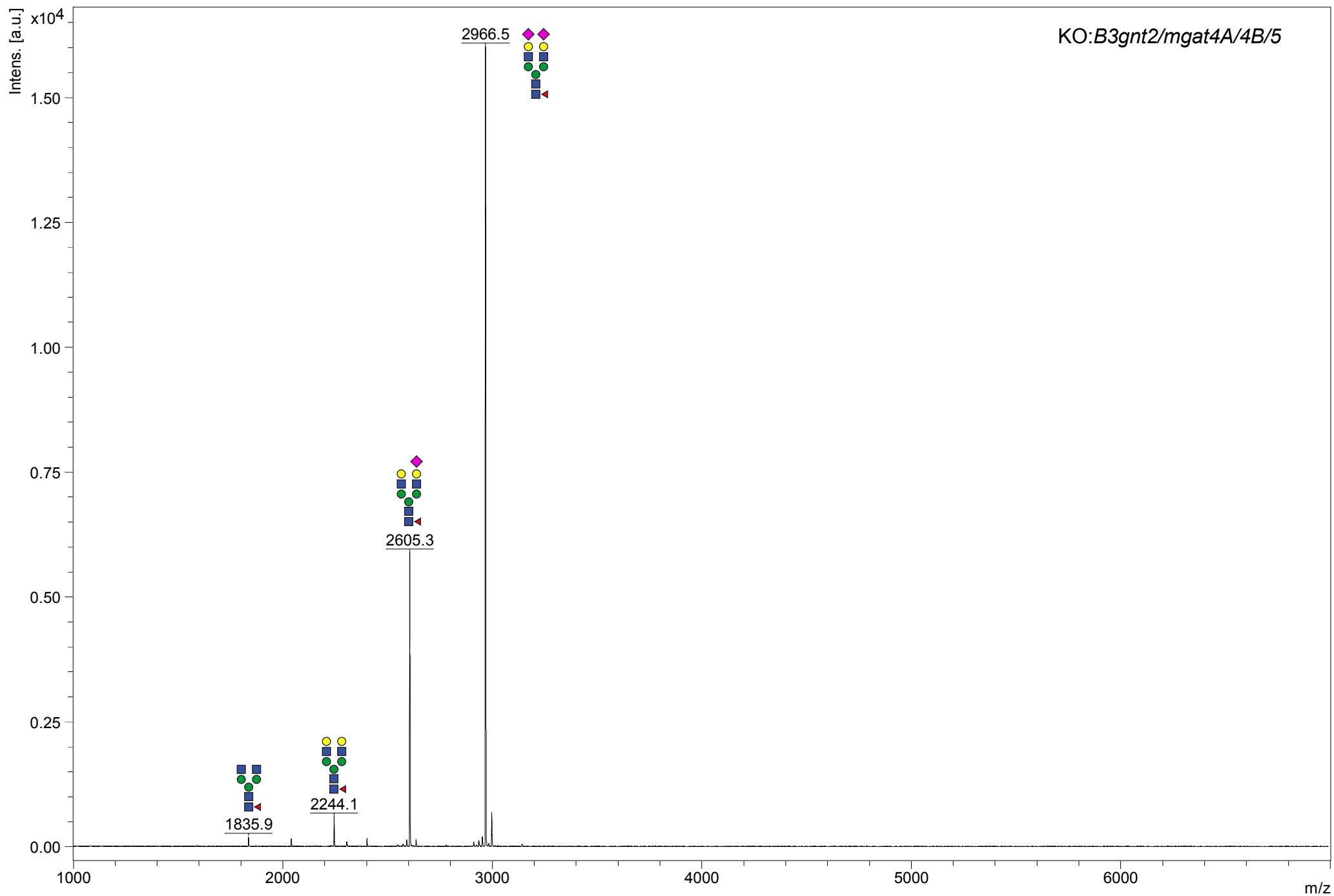


Supplementary Spectra
#27

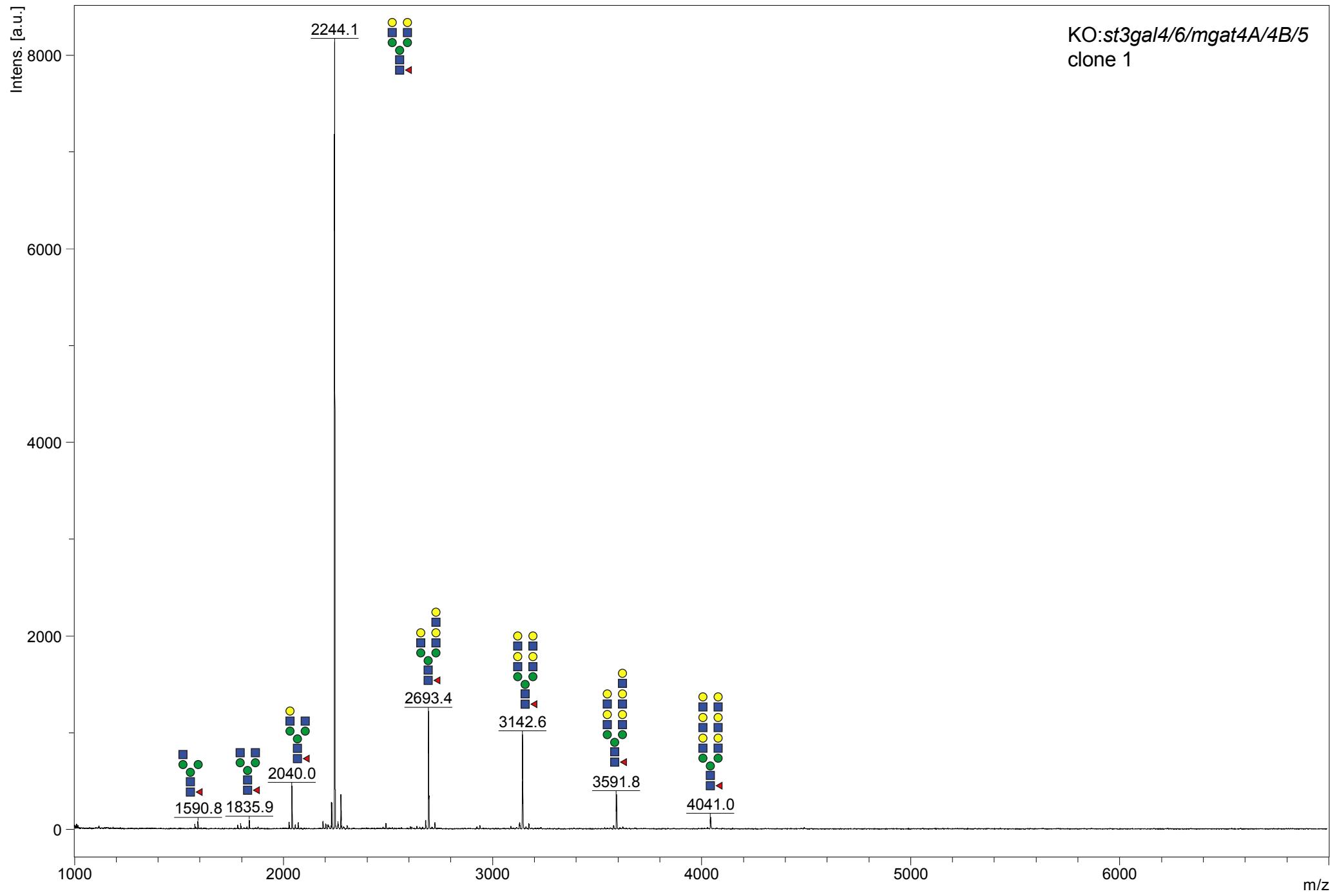


KI:ST6Gal-I
KO:st3gal4/6

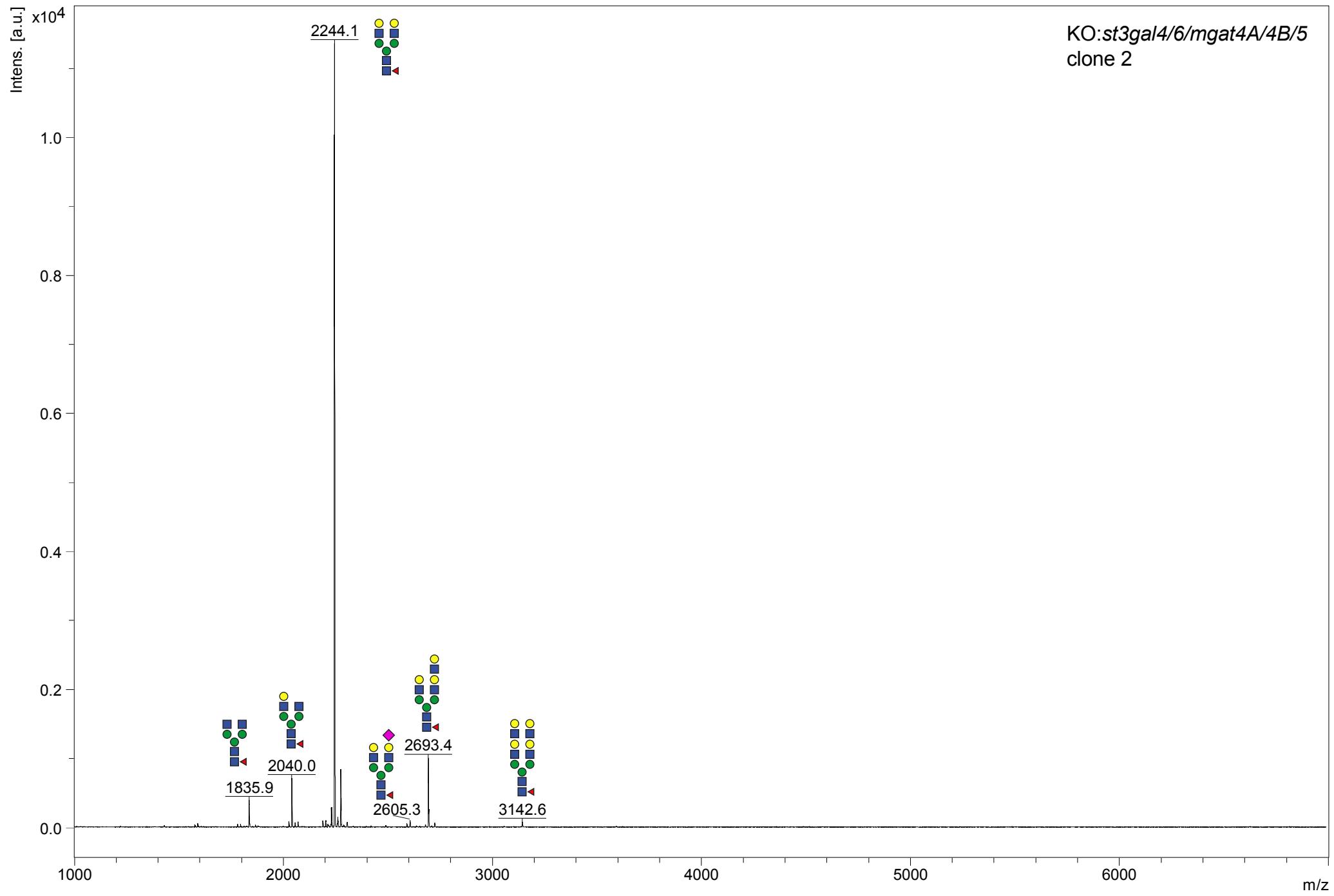
Supplementary Spectra
#28



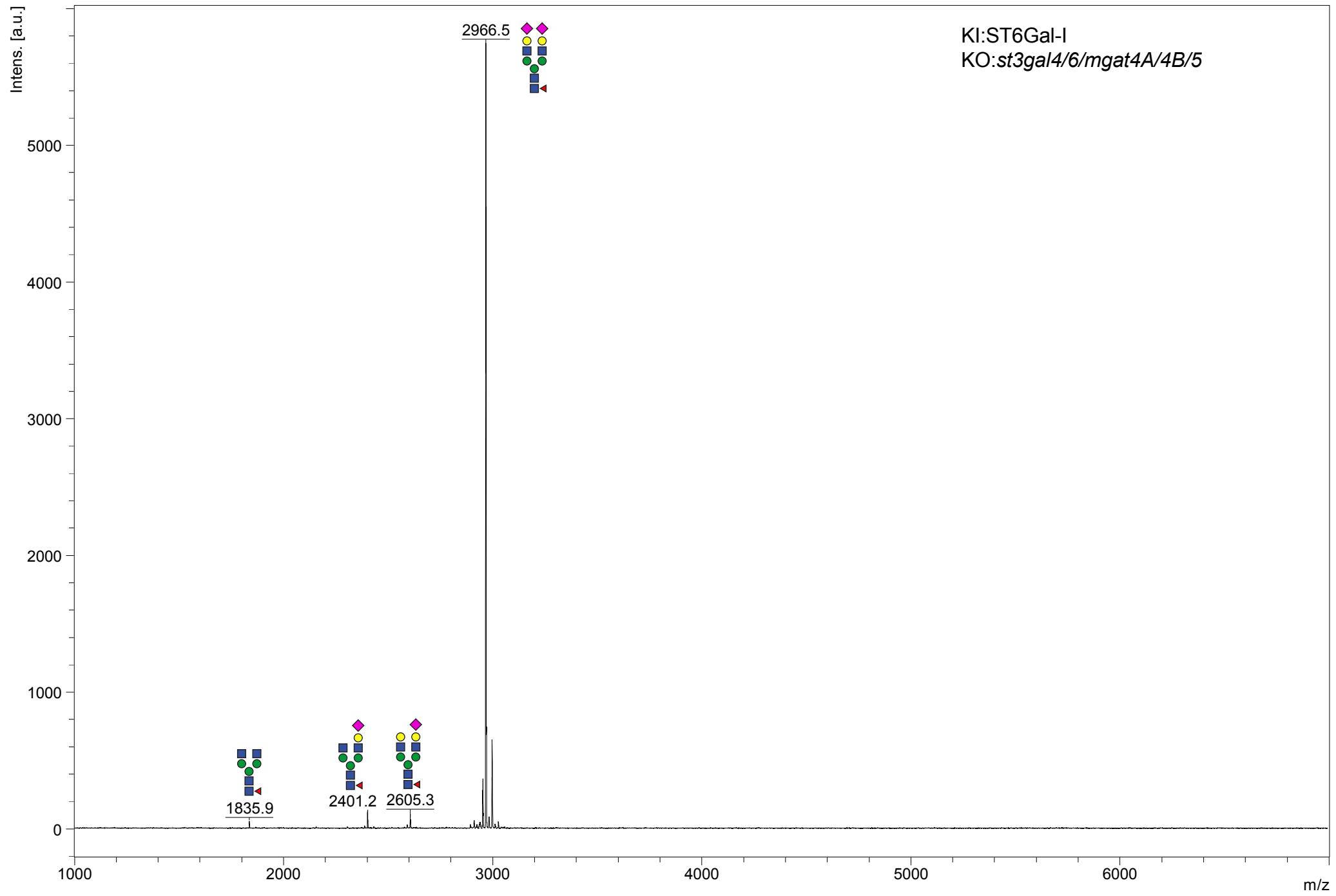
Supplementary Spectra
#29a



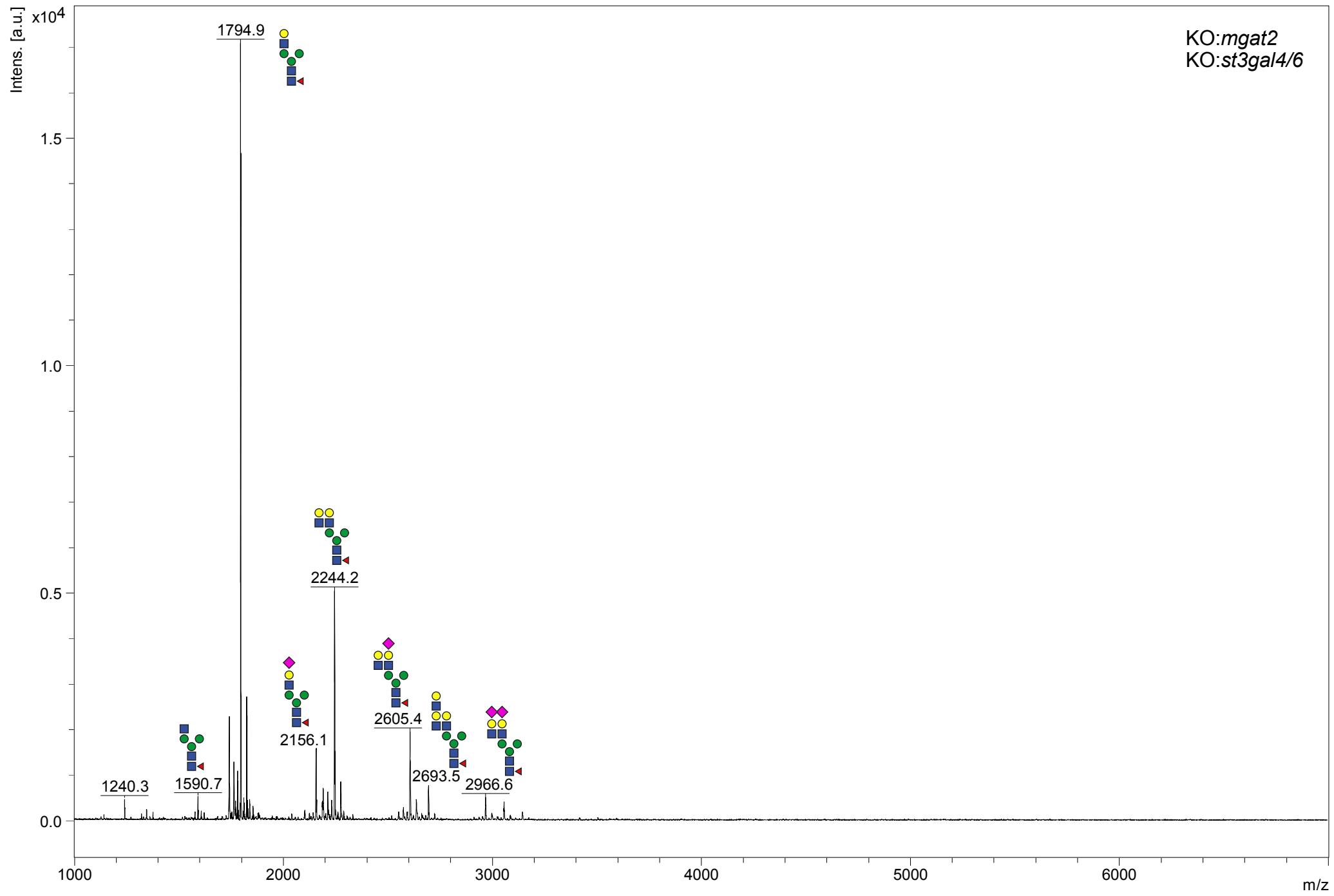
Supplementary Spectra
#29b



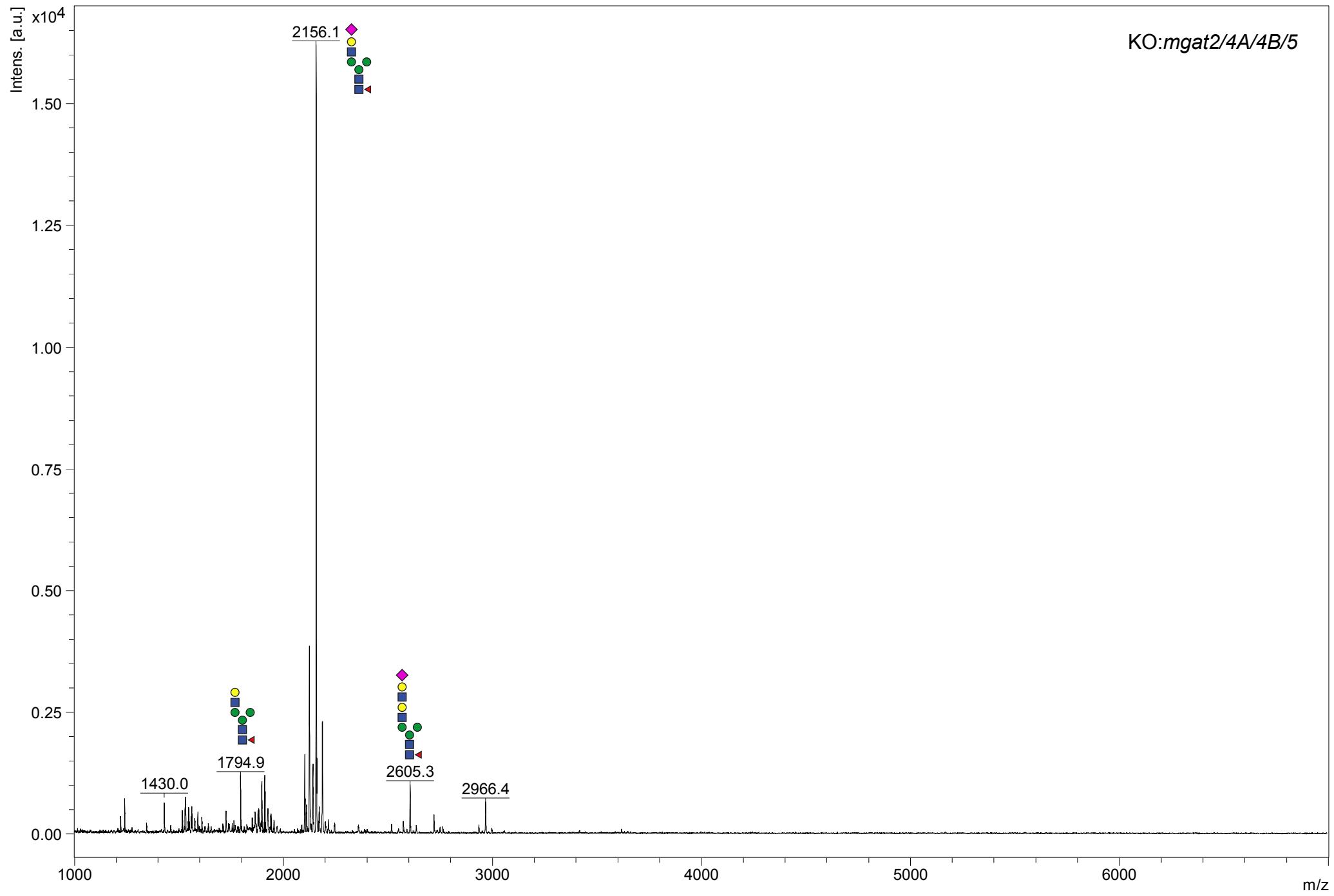
Supplementary Spectra
#30



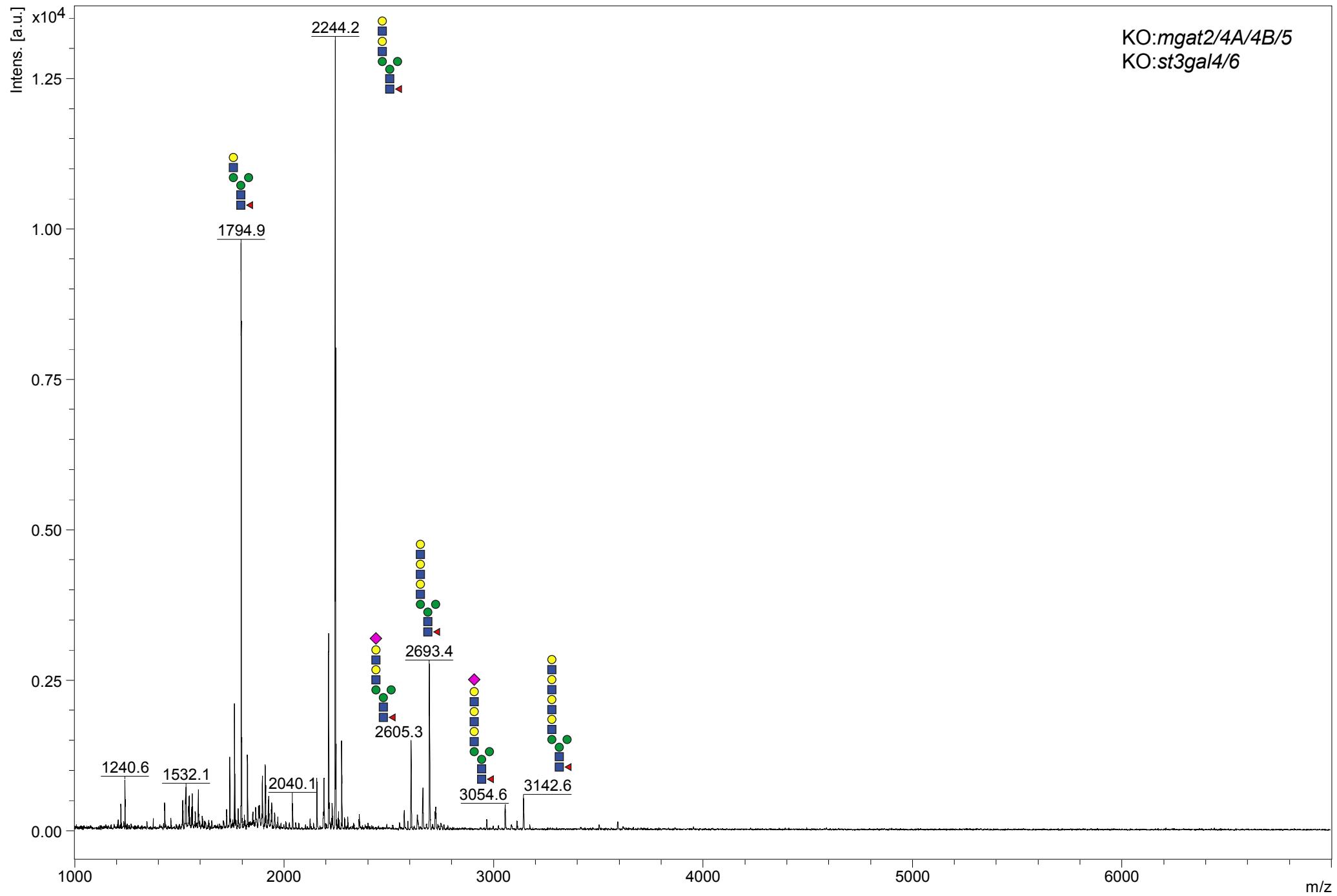
Supplementary Spectra
#31



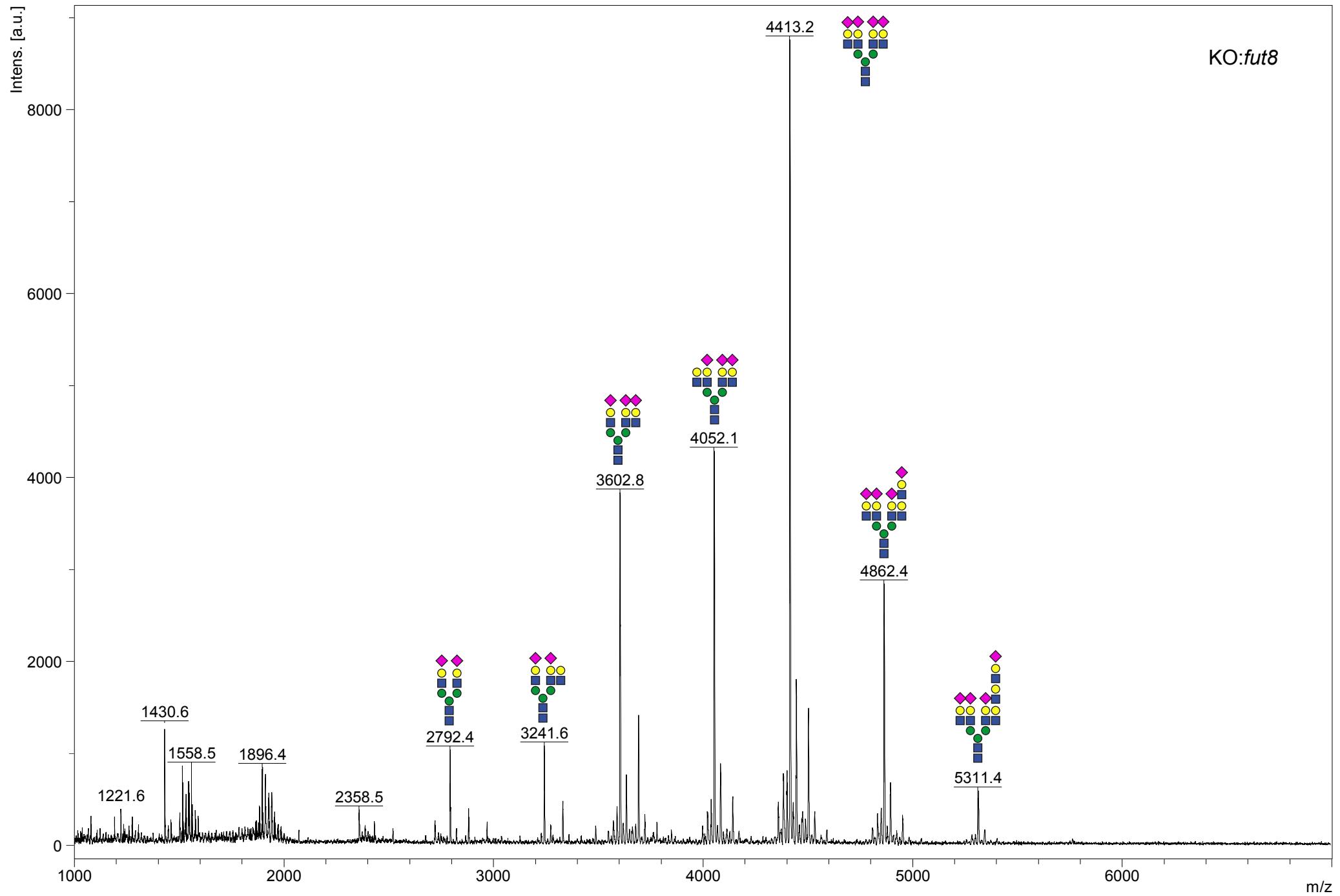
Supplementary Spectra
#32



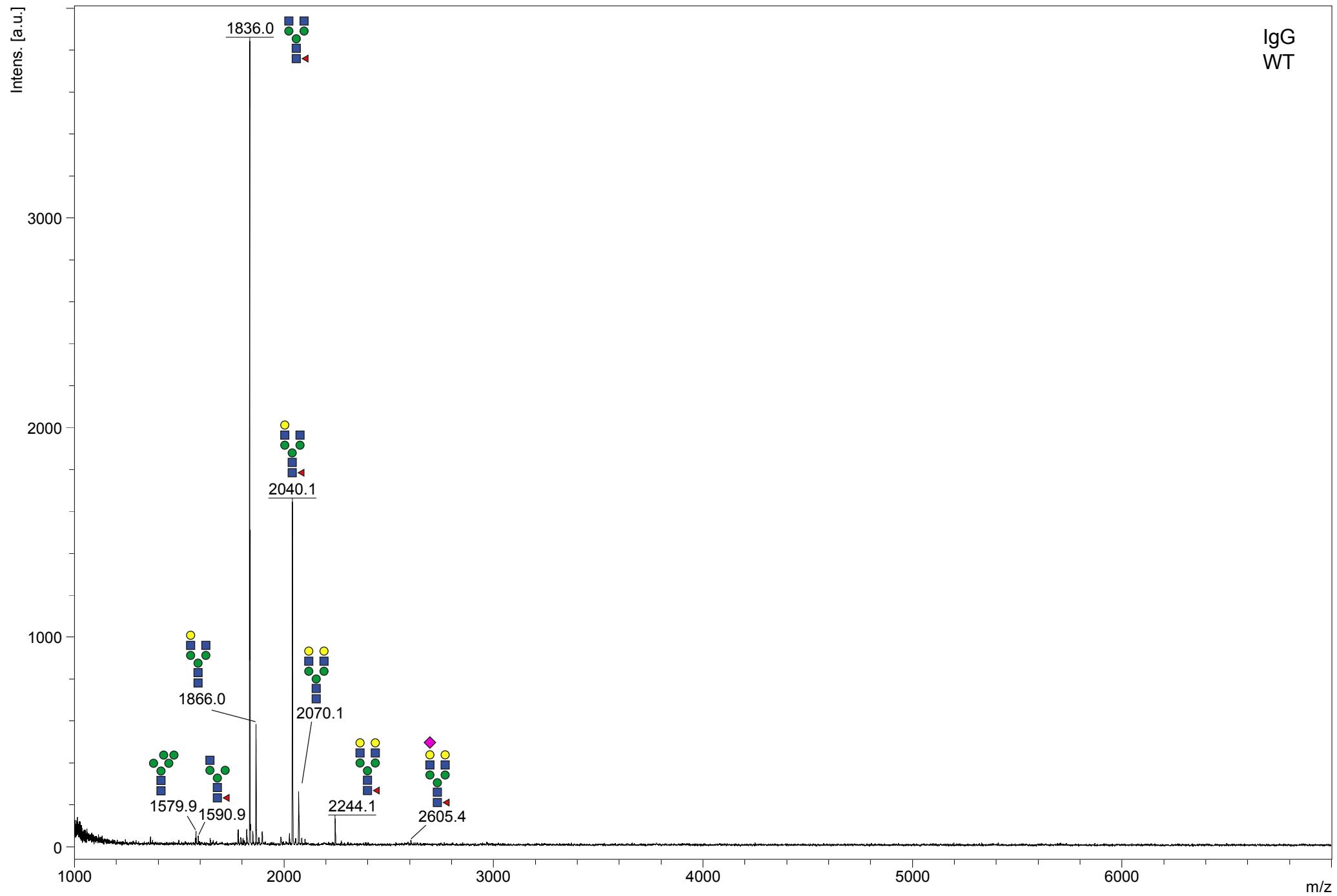
Supplementary Spectra
#33



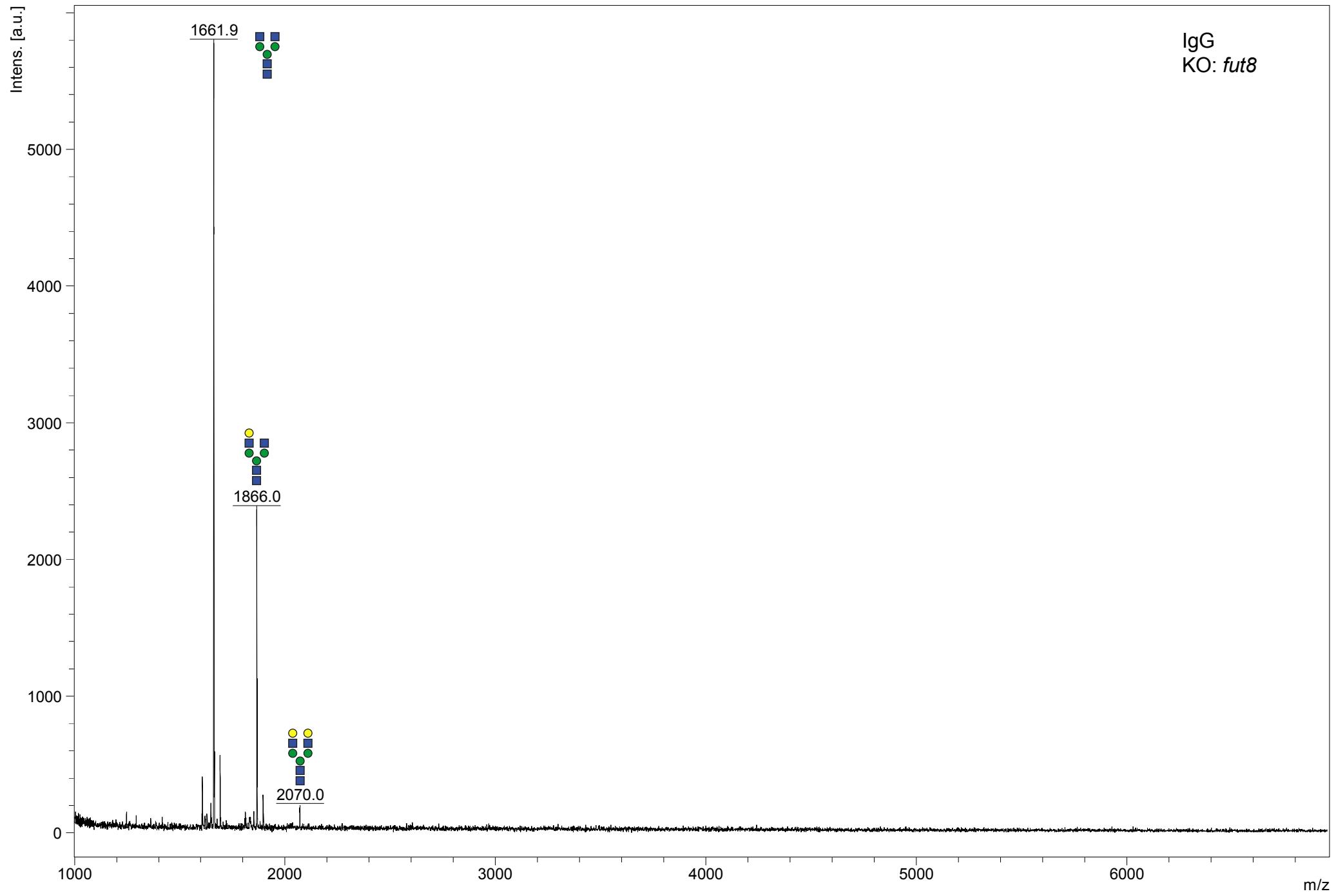
Supplementary Spectra
#34



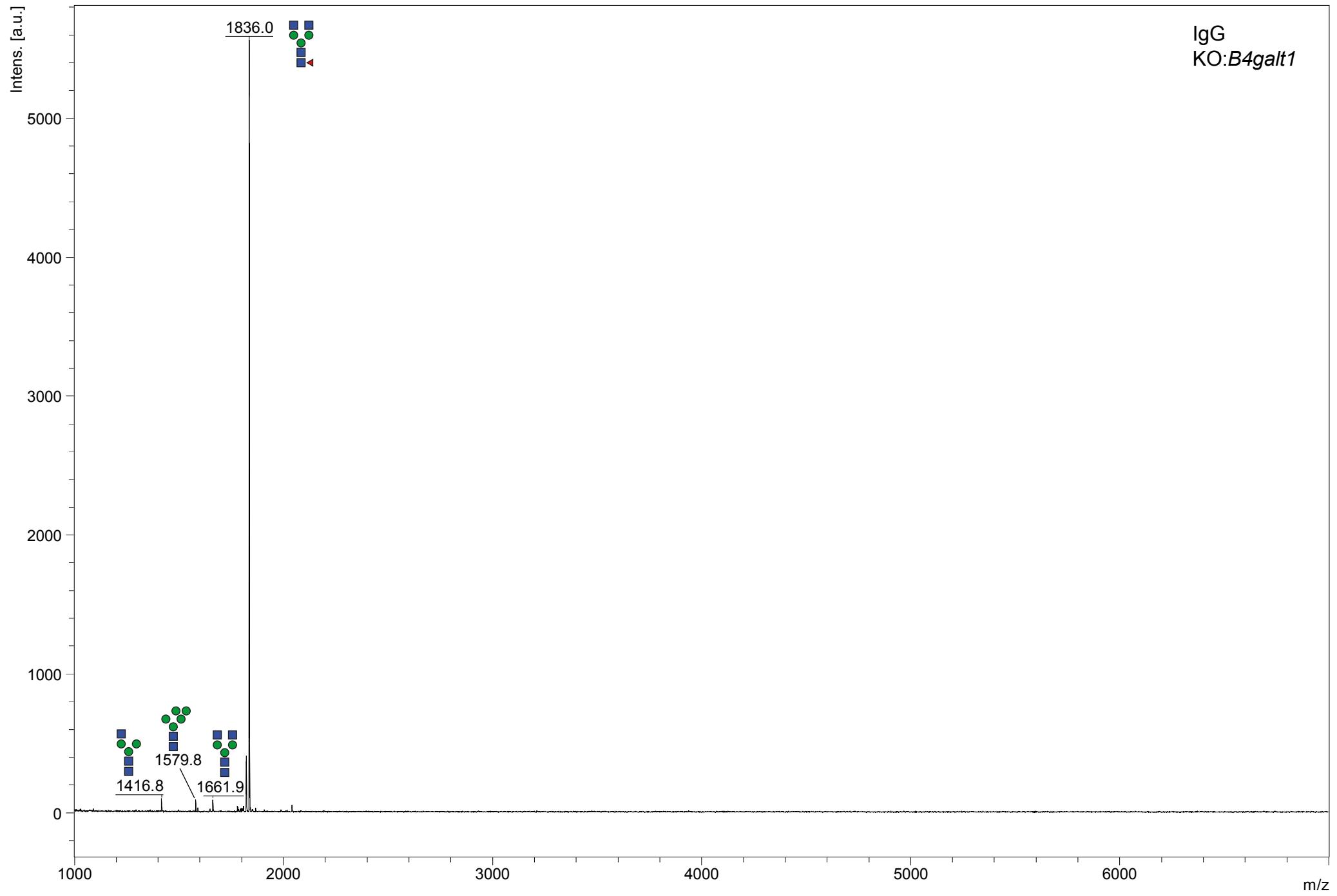
Supplementary Spectra
#35



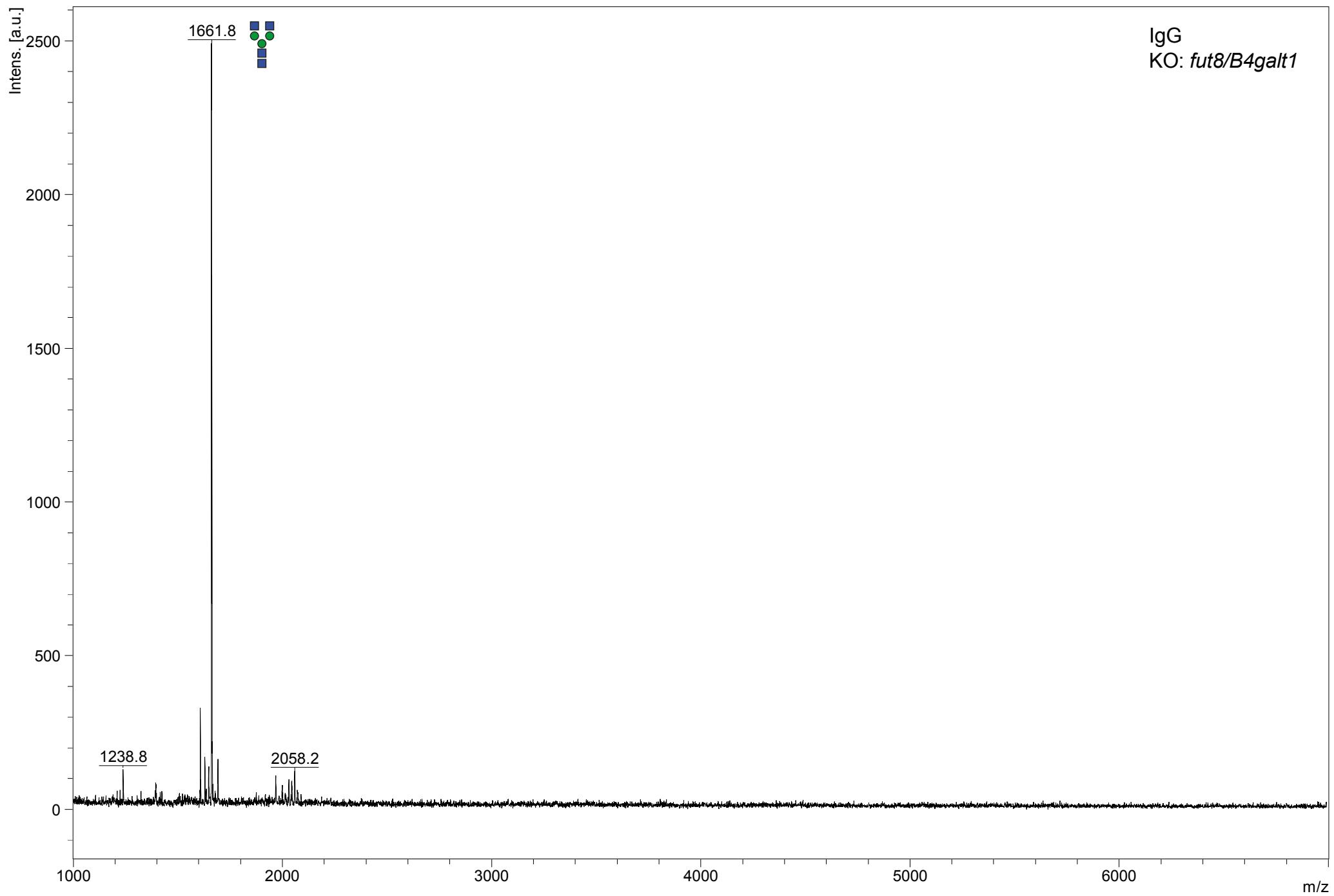
Supplementary Spectra
#36



Supplementary Spectra
#37



Supplementary Spectra
#38



Supplementary Table 1. Glycoengineered CHO clones and stacking ancestry.

Project number	Targeted genes	Parental CHO line	Glycoprofiled
ZFN127	KO: <i>mgat1</i>	WT	x
ZFN174	KO: <i>mgat2</i>	WT	x
ZFN048	KO: <i>mgat3</i>	WT	x
ZFN001	KO: <i>mgat4A</i>	WT	x
ZFN078	KO: <i>mgat4B</i>	WT	x
ZFN143	KO: <i>mgat4C</i>	WT	x
ZFN004	KO: <i>mgat5</i>	WT	x
ZFN104	KO: <i>mgat5B</i>	WT	x
ZFN056	KO: <i>mgat3/5</i>	ZFN048	
ZFN067	KO: <i>mgat4A/5</i>	ZFN001	x
ZFN229	KO: <i>mgat4B/5</i>	ZFN078	x
ZFN150	KO: <i>mgat5/5B</i>	ZFN104	
ZFN077	KO: <i>mgat4A/4B</i>	ZFN001	x
ZFN091	KO: <i>mgat4A/4B/5</i>	ZFN077	x
ZFN115	KO: <i>mgat3/4A/4B/5</i>	ZFN091	x
ZFN144	KO: <i>mgat4A/4B/4C</i>	ZFN077	x
ZFN173	KO: <i>mgat4A/4B/4C/5</i>	ZFN144	
ZFN110	KO: <i>B4galt1</i>	WT	x
ZFN111	KO: <i>B4galt2</i>	WT	x
ZFN112	KO: <i>B4galt3</i>	WT	x
ZFN113	KO: <i>B4galt4</i>	WT	x
ZFN120	KO: <i>B4galt1/2</i>	ZFN110	x
ZFN121	KO: <i>B4galt1/3</i>	ZFN110	x
ZFN200	KO: <i>B4galt1/2/3</i>	ZFN121	x
ZFN141	KO: <i>B4galt1/2/4</i>	ZFN120	
ZFN164	KO: <i>B4galt1/2/3/4</i>	ZFN141	
ZFN224	KO: <i>B4galt1/2/3/4</i>	ZFN200	x
ZFN084	KO: <i>B3gnt2</i>	WT	x
ZFN085	KO: <i>B3gnt1</i>	WT	x
ZFN275	KO: <i>B3gnt8</i>	WT	x
ZFN096	KO: <i>B3gnt1/2</i>	ZFN084	x
ZFN072	KO: <i>st3gal3</i>	WT	x
ZFN082	KO: <i>st3gal6</i>	WT	x

ZFN083	KO: <i>st3gal4</i>	WT	x
ZFN090	KO: <i>st3gal3/4</i>	ZFN072	x
ZFN095	KO: <i>st3gal3/6</i>	ZFN082	
ZFN098	KO: <i>st3gal4/6</i>	ZFN082	x
ZFN099	KO: <i>st3gal3/4/6</i>	ZFN090	x
 ZFN136	KO: <i>B4galt1/mgat4A/4B/5</i>	ZFN091	x
ZFN137	KO: <i>B4galt2/mgat4A/4B/5</i>	ZFN091	
ZFN161	KO: <i>B4galt3/mgat4A/4B/5</i>	ZFN091	x
ZFN162	KO: <i>B4galt4/mgat4A/4B/5</i>	ZFN091	
 ZFN119	KO: <i>B4galt1/st3gal4/6</i>	ZFN098	
ZFN138	KO: <i>B4galt2/stgal4/6</i>	ZFN098	
ZFN118	KO: <i>B3gnt2/st3gal4/6</i>	ZFN098	
ZFN167	KO: <i>B3gnt2/mgat4A/4B/5</i>	ZFN091	x
ZFN117	KO: <i>st3gal4/mgat4A/4B/5</i>	ZFN091	
ZFN163	KO: <i>st3gal4/6/mgat4A/4B/5</i>	ZFN117	x
ZFN225	KO: <i>st3gal3/4/6/mgat4A/4B/5</i>	ZFN163	
ZFN170	KO: <i>B3gnt2/st3gal4/6/mgat4A/4B/5</i>	ZFN163	x
 ZFN134	KI:ST6GAL-I/KO: <i>st3gal4/6</i>	ZFN098	x
ZFN168	KI:ST6GAL-I/KO: <i>B3gnt2/st3gal4/6</i>	ZFN118	x
ZFN169	KI:ST6GAL-I/KO: <i>st3gal4/6/mgat4A/4B/5</i>	ZFN163	x
ZFN184	KI:ST6GAL-I/KO: <i>B3gnt2/st3gal4/6/mgat4A/4B/5</i>	ZFN170	x
ZFN226	KI:ST6GAL-I/KO: <i>st3gal3/4/6/mgat4A/4B/5</i>	ZFN169	x
 ZFN174	KO: <i>mgat2</i>	WT	x
ZFN259	KO: <i>mgat2/stgal4/6</i>	ZFN98	x
ZFN261	KO: <i>mgat2/mgat4A/4B/5</i>	ZFN91	x
ZFN263	KO: <i>mgat2/st3gal4/6/mgat4A/4B/5</i>	ZFN163	x
ZFN281	KO: <i>B3gnt2/mgat2/st3gal4/6/mgat4A/4B/5</i>	ZFN263	x
ZFN266	KO: <i>fut8</i>	WT	x
ZFN268	KO: <i>fut8/B4galt1</i>	ZFN110	x

Supplementary Table 2. ZFN design and sequence analysis of CHO glycoengineered clones.

Clone	Inser & Del	Alignment
<i>KO:mgat4A</i>		
<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAATGCCATTGTCCAACAgtt-----tc <u>GCCGTGCAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-allele1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----CCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4A-allele2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAgttGAATTCTAGATGAt <u>cGCCGTGCAGGAGCAGCAACTAACGGAAG</u>
<i>KO:mgat4A/4B</i>		
<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAATGCCATTGTCCAACAgtt-----tc <u>GCCGTGCAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-allele1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----CCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4A-allele2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAgttGAATTCTAGATGAt <u>cGCCGTGCAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCTCTgaggactGGATGATCCTGGAGTT</u>
<i>mgat4B-allele1</i>	-7bp	<u>GCCCTCCAGCAGCCCTCTg-----GATGATCCTGGAGTT</u>
<i>KO:mgat5</i>		
<i>mgat5-WT</i>	ZFN	GGGGGATGATGCTTCTGCACTTCACCATCCAg-----cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele1</i>	+14bp	GGGGGATGATGCTTCTGCACTTCACCATCCAg <u>ca</u> GAATTCTAGATGAgcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele2</i>	-1bp	GGGGGATGATGCTTCTGCACTTCACCATCCAg-----agcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele3</i>	+61bp	GGGGGATGATGCTTCTGCACTTCACCATCCAg <u>cag</u> (+61bp) --cg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>KO:mgat4A/4B/5</i>		
<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAATGCCATTGTCCAACAgtt-----tc <u>GCCGTGCAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-allele1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----CCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4A-allele2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAgttGAATTCTAGATGAt <u>cGCCGTGCAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCTCTgaggactGGATGATCCTGGAGTT</u>
<i>mgat4B-allele1</i>	-7bp	<u>GCCCTCCAGCAGCCCTCTg-----GATGATCCTGGAGTT</u>
<i>mgat5-WT</i>	ZFN	GGGGGATGATGCTTCTGCACTTCACCATCCAg---cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele1</i>	+4bp	GGGGGATGATGCTTCTGCACTTCACCATCCAg <u>cagc</u> CAGCg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele2</i>	-4bp	GGGGGATGATGCTTCTGCACTTCACCATCCAg-----g <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele3</i>	-16bp	GGGGGATGATGCTTCTGCACTT-----CTCAGCCTGAGAGCAGCTCCATGTT
<i>KO:B3gnt1</i>		
<i>B3gnt1-WT</i>	ZFN	<u>TGCAGCTGCTCACCTGTC-----gctgc</u> TCTCCGGACTGCACG
<i>B3gnt1-allele1</i>	+11bp	TGCAGCTGCTCACCTGTC <u>cgac</u> TGCTCTACCTGTCTCCGGACTGCACG
<i>B3gnt1-allele2</i>	-5bp	TGCAGCTGCTCACCTGTC-----TCTCCGGACTGCACG
<i>KO:B3gnt2</i>		
<i>B3gnt2-WT</i>	ZFN	<u>TTCAGCCCTTCCcgggcGTACTGGAACAGAGAGCA</u>
<i>B3gnt2-allele1</i>	-1bp	TTCAGCCCTTCC-gggcGTACTGGAACAGAGAGCA
<i>B3gnt2-allele2</i>	-4bp	TTCAGCCCTTCC <u>cg</u> -----TACTGGAACAGAGAGCA
<i>KO:B4galt1</i>		
<i>B4galt1-WT</i>	ZFN	<u>TGCATCCGGTCCTACAGCgccagc</u> -----AACTGGACTATGGTA

<i>B4galt1</i> -alle1	+4bp	TGCATCCGGTCCTACAGCgccaggCAGCAACTGGACTATGGTA
<i>KO:B4galt2</i>		
<i>B4galt2</i> -WT	ZFN	CAGCCCCGCCACTTTgcc-----atcGCCATGGACAAGTTGGCT
<i>B4galt2</i> -alle1	+73bp	CAGCCCCGCCACTTTgcc-- (+73bp) --atcGCCATGGACAAGTTGGCT
<i>KO:B4galt3</i>		
<i>B4galt3</i> -WT	ZFN	CTAGCCCTCAAGTCAGGAtgt----tgCGGAGGCTGCTGGAGAGG
<i>B4galt3</i> -alle1	+5bp	CTAGCCCTCAAGTCAGGAtgtCGTGTtgCGGAGGCTGCTGGAGAGG
<i>B4galt3</i> -alle2	+2bp	CTAGCCCTCAAGTCAGGAtgt---tgCCCGGAGGCTGCTGGAGAGG
<i>KO:B4galt4</i>		
<i>B4galt4</i> -WT	ZFN	AACTGGGACTGCTTat----attcCACGATGTGGACCTGGTG
<i>B4galt4</i> -alle1	+1bp	AACTGGGACTGCTTat---TattcCACGATGTGGACCTGGTG
<i>B4galt4</i> -alle2	+4bp	AACTGGGACTGCTTatattTATTcCACGATGTGGACCTGGTG
<i>KO:B4galt1/2</i>		
<i>B4galt1</i> -WT	ZFN	TGCATCCGGTCCTACAGCgccagc---AACTGGACTATGGTA
<i>B4galt1</i> -alle1	+4bp	TGCATCCGGTCCTACAGCgccaggCAGCAACTGGACTATGGTA
<i>B4galt2</i> -WT	ZFN	CAGCCCCGCCACTTTgccatcGCCATGGACAAGTTGGCT
<i>B4galt2</i> -alle1	-8bp	CAGCCCCGCCAC-----cGCCATGGACAAGTTGGCT
<i>B4galt2</i> -alle2	-14bp	CAGCCC-----cGCCATGGACAAGTTGGCT
<i>KO:st3Gal3</i>		
<i>st3Gal3</i> -WT	ZFN	CTCTCTCTTGTCCTTGCTggcttCAAATGGCAGGACTTCAAG
<i>st3Gal3</i> -alle1	-5bp	CTCTCTCTTGTCCTTGCT----CAAATGGCAGGACTTCAAG
<i>st3Gal3</i> -alle2	-1bp	CTCTCTCTTGTCCTTGCTggct -CAAATGGCAGGACTTCAAG
<i>KO:st3Gal4</i>		
<i>st3Gal4</i> -WT	ZFN	GGCAGCCTCCAGTGTGTCgttgt----gTTGTGGTGGGAATGGC
<i>st3Gal4</i> -alle1	+4bp	GGCAGCCTCCAGTGTGTCgttgtTTGTgTTGTGGTGGGAATGGC
<i>st3Gal4</i> -alle2	-4bp	GGCAGCCTCCAGTGTGTCg----gTTGTGGTGGGAATGGC
<i>KO:st3Gal6</i>		
<i>st3Gal6</i> -WT	ZFN	CGGTACCTCTGATTTGCTtgccCTATGGGACAAGGCC
<i>st3Gal6</i> -alle1	-4bp	CGGTACCTCTGATTTGCTt---CTATGGGACAAGGCC
<i>st3Gal6</i> -alle2	-22bp	CGGTACCTCTGA-----AGGCC
<i>KO:st3Gal3/4</i>		
<i>st3Gal3</i> -WT	ZFN	CTCTCTCTTGTCCTTGCTggcttCAAATGGCAGGACTTCAAG
<i>st3Gal3</i> -alle1	-5bp	CTCTCTCTTGTCCTTGCT----CAAATGGCAGGACTTCAAG
<i>st3Gal3</i> -alle2	-1bp	CTCTCTCTTGTCCTTGCTggct -CAAATGGCAGGACTTCAAG
<i>st3Gal4</i> -WT	ZFN	GGCAGCCTCCAGTGTGTCgttgt----gTTGTGGTGGGAATGGC
<i>st3Gal4</i> -alle1	-4bp	GGCAGCCTCCAGTGTGTCg----gTTGTGGTGGGAATGGC

st3Gal4-alle2 +5bp GGCAGCCTCCAGTGTGTCgACACGttgtgTTGTGGTGGGAATGGC

KO:*st3Gal4/6*

<i>st3Gal4-WT</i>	ZFN	<u>GGCAGCCTCCAGTGTGTCgttgt</u> -----g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle1</i>	+4bp	GGCAGCCTCCAGTGTGTCgttgt <u>TTGTgTTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle2</i>	-4bp	GGCAGCCTCCAGTGTGTCg-----g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal6-WT</i>	ZFN	<u>CGGTACCTCTGATTGCTtgc</u> <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-alle1</i>	+1bp	CGGTACCTCTGATTGCTtgc <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-alle2</i>	-7bp	CGGTACCTCTGATTG----- <u>CTATGGGACAAGGCC</u>

KO:*B3gnt2/mgat4A/4B/5*

<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAAT <u>GCCATTGTCCAACAg</u> t-----tc <u>CCCGTGCAAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-alle1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----ccgtgcaggagcagcaactaacggaag
<i>mgat4A-alle2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAg <u>tGAATTCTAGATGAtcGCCGTGCAAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCCCT</u> <u>TCTgaggactGGATGATCCTGGAGTT</u>
<i>mgat4B-alle1</i>	-7bp	<u>GCCCTCCAGCAGCCCCCT</u> <u>Tg-----GATGATCCTGGAGTT</u>
<i>mgat5-WT</i>	ZFN	GGGGGATGATGCT <u>TCTGCACCCATCCAg</u> ---cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle1</i>	+4bp	GGGGGATGATGCTTCTGCACCCATCCAg <u>cagcgCAGCgGACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle2</i>	-4bp	GGGGGATGATGCTTCTGCACCCATCCAg <u>-----gACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle3</i>	-16bp	GGGGGATGATGCTTCTGCACCCATCC----- <u>CTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>B3gnt2-WT</i>	ZFN	<u>TTCAGCCCTTC</u> --cgggc <u>GTACTGGAACAGAGAGCA</u>
<i>B3gnt2-alle1</i>	+2bp	TTCAGCCCTTCCC <u>ccccccgggcGTACTGGAACAGAGAGCA</u>
<i>B3gnt2-alle2</i>	+1bp	TTCAGCCCTTCCC <u>-cgggcGTACTGGAACAGAGAGCA</u>

KO:*st3Gal4/6/mgat4A/4B/5*

<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAAT <u>GCCATTGTCCAACAg</u> t-----tc <u>CCCGTGCAAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-alle1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----ccgtgcaggagcagcaactaacggaag
<i>mgat4A-alle2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAg <u>tGAATTCTAGATGAtcGCCGTGCAAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCCCT</u> <u>TCTgaggactGGATGATCCTGGAGTT</u>
<i>mgat4B-alle1</i>	-7bp	<u>GCCCTCCAGCAGCCCCCT</u> <u>Tg-----GATGATCCTGGAGTT</u>
<i>mgat5-WT</i>	ZFN	GGGGGATGATGCT <u>TCTGCACCCATCCAg</u> ---cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle1</i>	+4bp	GGGGGATGATGCTTCTGCACCCATCCAg <u>cagcgCAGCgGACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle2</i>	-4bp	GGGGGATGATGCTTCTGCACCCATCCAg <u>-----gACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle3</i>	-16bp	GGGGGATGATGCTTCTGCACCCATCC----- <u>CTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>st3Gal4-WT</i>	ZFN	<u>GGCAGCCTCCAGTGTGTCgttgt</u> -----g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle1</i>	-5bp	GGCAGCCTCCAGTGTGTCgttgt-----g <u>GTGGGAAATGGC</u>
<i>st3Gal4-alle2</i>	+4bp	GGCAGCCTCCAGTGTGTCgttgt <u>TTGTgtgTTGTGGTGGGAATGGC</u>
<i>st3Gal6-WT</i>	ZFN	<u>CGGTACCTCTGATTGCTtgc</u> <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-alle1</i>	-4bp	CGGTACCTCTGATTGCTtgc <u>-----CTATGGGACAAGGCC</u>
<i>st3Gal6-alle2</i>	-2bp	CGGTACCTCTGATTGCTtgc <u>-CTATGGGACAAGGCC</u>

KI:*ST6GAL1*/KO:*st3gal4/6*

<i>st3Gal4-WT</i>	ZFN	<u>GGCAGCCTCCAGTGTGTCgttgt</u> -----g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle1</i>	+4bp	GGCAGCCTCCAGTGTGTCgttgt <u>TTGTgtgTTGTGGTGGGAATGGC</u>

<i>st3Gal4-alle2</i>	-4bp	GGCAGCCTCCAGTGTGTCg-----gTTGTGGTGGGAATGGC
<i>st3Gal6-WT</i>	ZFN	<u>CGGTACCTCTGATTGCT-ttgc</u> <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-alle1</i>	+1bp	CGGTACCTCTGATTGCTttgccCTATGGGACAAGGCC
<i>st3Gal6-alle2</i>	-7bp	CGGTACCTCTGATTG-----CTATGGGACAAGGCC

KI:ST6GAL1/KO:st3gal4/6/mgat4A/4B/5

<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAATGCCATTGTCCAACAgtt-----tc <u>CCCGTGAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-alle1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----CCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4A-alle2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAgttGAATTCTAGATGAtcGCCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCTCTgaggactGGATGATCCTGGAGTT</u>
<i>mgat4B-alle1</i>	-7bp	GCCCTCCAGCAGCCCTCTg-----GATGATCCTGGAGTT
<i>mgat5-WT</i>	ZFN	GGGGGATGATGCT <u>TCTGCACCTCACCATCCAg</u> ---cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle1</i>	+4bp	GGGGGATGATGCTTCTGCACCTCACCATCCAgcagcCAGCg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle2</i>	-4bp	GGGGGATGATGCTTCTGCACCTCACCATCCAg-----g <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-alle3</i>	-16bp	GGGGGATGATGCTTCTGCACCTTC-----CTCAGCCTGAGAGCAGCTCCATGTT
<i>st3Gal4-WT</i>	ZFN	<u>GGCAGCCTCCAGTGTGTCgttgt</u> ---g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle1</i>	-5bp	GGCAGCCTCCAGTGTGTCgttgt-----g <u>GTGGGAATGGC</u>
<i>st3Gal4-alle2</i>	+4bp	GGCAGCCTCCAGTGTGTCgttgtTTGTtgt <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal6-WT</i>	ZFN	<u>CGGTACCTCTGATTGCT-ttgc</u> <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-alle1</i>	-4bp	CGGTACCTCTGATTGCTttgc---CTATGGGACAAGGCC
<i>st3Gal6-alle2</i>	-2bp	CGGTACCTCTGATTGCTtgt---CTATGGGACAAGGCC

KO:mgat2

<i>mgat2-WT</i>	TALEN	<u>TCCTTGTCGCCATTGCTgtccagaggacgaagCCGCAGGCCACCACGA</u>
<i>mgat2-alle1</i>	-4bp	TCCTTGTCGCCATTGCTgtcc---gacgaagccgcaggcgccaccacga

KO:mgat2/stgal4/6

<i>mgat2-WT</i>	TALEN	<u>TCCTTGTCGCCATTGCTgtccagaggacgaagCCGCAGGCCACCACGA</u>
<i>mgat2-alle1</i>	-4bp	TCCTTGTCGCCATTGCTgtccag---cgaagCCGCAGGCCACCACGA
<i>st3Gal4-WT</i>	ZFN	<u>GGCAGCCTCCAGTGTGTCgttgt</u> ---g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle1</i>	+4bp	GGCAGCCTCCAGTGTGTCgttgtTTGTgt <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal4-alle2</i>	-4bp	GGCAGCCTCCAGTGTGTCgttgt---g <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal6-WT</i>	ZFN	<u>CGGTACCTCTGATTGCT-ttgc</u> <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-alle1</i>	+1bp	CGGTACCTCTGATTGCTttgccCTATGGGACAAGGCC
<i>st3Gal6-alle2</i>	-7bp	CGGTACCTCTGATTG-----CTATGGGACAAGGCC

KO:mgat2/mgat4A/4B/5

<i>mgat2-WT</i>	TALEN	<u>TCCTTGTCGCCATTGCTgtccagaggacgaagCCGCAGGCCACCACGA</u>
<i>mgat2-alle1</i>	-4bp	TCCTTGTCGCCATTGCTgtccag---cgaagCCGCAGGCCACCACGA
<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAAT <u>GCCATTGTCCAACAg</u> tt-----tc <u>CCCGTGAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-alle1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg-----CCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4A-alle2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAgttGAATTCTAGATGAtcGCCGTGCAGGAGCAGCAACTAACGGAAG
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCTCTgaggactGGATGATCCTGGAGTT</u>
<i>mgat4B-alle1</i>	-7bp	GCCCTCCAGCAGCCCTCTg-----GATGATCCTGGAGTT

<i>mgat5-WT</i>	ZFN	GGGGGATGATGCTTCTGCACCCATCCAg-----cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele1</i>	+4bp	GGGGGATGATGCTTCTGCACCCATCCAg <u>cagcgCAGCgGACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele2</i>	-4bp	GGGGGATGATGCTTCTGCACCCATCCAg----- <u>gACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele3</i>	-16bp	GGGGGATGATGCTTCTGCACCC----- <u>CTCAGCCTGAGAGCAGCTCCATGTT</u>

KO:mgat2/st3gal4/6/mgat4A/4B/5

<i>mgat2-WT</i>	TALEN	<u>TCCTTTGTCGCCATTGCT</u> gctccagaggacgaag <u>CCGCAGGCGGCCACCACGA</u>
<i>mgat2-allele1</i>	-5bp	TCCTTTGTCGCCATTGCTgctcca-----cgaa <u>CCGCAGGCGGCCACCACGA</u>
<i>mgat4A-WT</i>	ZFN	TTCTGAGTTGAATGCCATTGTCCAACAgtt-----tc <u>CCCGTGAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-allele1</i>	-5bp	TTCTGAGTTGAATGCCATTGTCCAACAg----- <u>CCGTGAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4A-allele2</i>	+13bp	TTCTGAGTTGAATGCCATTGTCCAACAgttGAATTCTAGATGA <u>tGCCGTGAGGAGCAGCAACTAACGGAAG</u>
<i>mgat4B-WT</i>	ZFN	<u>GCCCTCCAGCAGCCCTCT</u> gaggact <u>TGGATGATCCTGGAGTT</u>
<i>mgat4B-allele1</i>	-7bp	GCCCTCCAGCAGCCCTCTg----- <u>GATGATCCTGGAGTT</u>
<i>mgat5-WT</i>	ZFN	GGGGGATGATGCTTCTGCACCCATCCAg-----cagcg <u>GACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele1</i>	+4bp	GGGGGATGATGCTTCTGCACCCATCCAg <u>cagcgCAGCgGACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele2</i>	-4bp	GGGGGATGATGCTTCTGCACCCATCCAg----- <u>gACTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>mgat5-allele3</i>	-16bp	GGGGGATGATGCTTCTGCACCC----- <u>CTCAGCCTGAGAGCAGCTCCATGTT</u>
<i>st3Gal4-WT</i>	ZFN	<u>GGCAGCCTCCAGTGTGTCgttg</u> ---- <u>gTTGTGGTGGGAATGGC</u>
<i>st3Gal4-allele1</i>	-5bp	GGCAGCCTCCAGTGTGTCgttg----- <u>gGTGGGAATGGC</u>
<i>st3Gal4-allele2</i>	+4bp	GGCAGCCTCCAGTGTGTCgttgTTGTtgg <u>TTGTGGTGGGAATGGC</u>
<i>st3Gal6-WT</i>	ZFN	<u>CGGTACCTCTGATTGCTtgc</u> ct <u>ATGGGACAAGGCC</u>
<i>st3Gal6-allele1</i>	-4bp	CGGTACCTCTGATTGCTtgc----- <u>CTATGGGACAAGGCC</u>
<i>st3Gal6-allele2</i>	-2bp	CGGTACCTCTGATTGCTtgc----- <u>CTATGGGACAAGGCC</u>

KO:fut8

<i>fut8-WT</i>	CRISPR	CAAATA <u>CTTGATCCGTCCACAACCT</u> -TGGCTGGAAAGGGAA
<i>fut8-allele1</i>	-1bp	CAAATA <u>CTTGATCCGTCCACAACC</u> --TGGCTGGAAAGGGAA
<i>fut8-allele2</i>	+1bp	CAAATA <u>CTTGATCCGTCCACAACCTTGCTGGAAAGGGAA</u>

KO:fut8/B4galt1

<i>fut8-WT</i>	CRISPR	CAAATA <u>CTTGATCCGTCCACAACCT</u> -TGGCTGGAAAGGGAA
<i>fut8-allele1</i>	-4bp	CAAATA <u>CTTGATCCGTCCACAACC</u> --GGCTGGAAAGGGAA
<i>fut8-allele2</i>	+1bp	CAAATA <u>CTTGATCCGTCCACAACCTTGCTGGAAAGGGAA</u>
<i>B4galt1-WT</i>	ZFN	<u>TGCATCCGGTCTCACAGCgttgc</u> ----- <u>AACTGGACTATGGTA</u>
<i>B4galt1-allele1</i>	+4bp	TGCATCCGGTCTCACAGCgttgc----- <u>AACTGGACTATGGTA</u>

KO:mgat3

<i>mgat3-WT</i>	ZFN	<u>TTCCTGGACCACCTCCC</u> Acccggt----- <u>GGCCGGCAGGATGGC</u>
<i>mgat3-allele1</i>	-21bp	TTCCTGGACCACCT----- <u>ATGGC</u>
<i>mgat3-allele2</i>	+293bp	TTCCTGGACCAC <u>TGATA</u> Ccc--+293bp--cggt <u>GGCCGGCAGGATGGC</u>

KO:mgat4C

<i>mgat4C-WT</i>	ZFN	<u>ATACTTCAGACTATTatgt</u> AATGCTCGAAGATGATGTT
<i>mgat4C-allele1</i>	-13bp	ATACTTCAGACT----- <u>CGAAGATGATGTT</u>

mgat4C-allele2 - 8bp ATACTTCAGACTAT-----GCTCGAAGATGATGTT

KO:mgat5B

mgat5B-WT ZFN CGTGGCGCCCTCCGCAAGatgagtGACCTGCTGGAGCTG

mgat5B-allele1 - 7bp CAGCTCCAGCAGGT-----CTTGCGGAGGGCGCCACG

mgat5B-allele2 - 5bp CAGCTCCAGCAGGT-----ATCTGCGGAGGGCGCCACG

KO:B3gnt8

B3gnt8-WT TALEN TGGTCCAGAGATAGCTAATgaagcttctagggtgGAGAAGCTGGGCTGCTGA

B3gnt8-allele1 - 17bp TGGTCCAGAGG-----AGGGTGGAGAAGCTGGGCTGCTGA

KO:mgat1

mgat1-WT ZFN AACAAAGTTCAAGTTCccagcaGCTGTGGTAGTGGAGGAC

mgat1-allele1 - 2bp AACAAAGTTCAAGTTCc--gcaGCTGTGGTAGTGGAGGAC

☒ Gene targeting region underlined.

Supplementary Table 3. Summary of glycan composition list of permethylated N-glycans.

Supplementary Table 4. Production of recombinant human EPO and IgG in CHO clones.

EPO clones	Final yield (mg/L) ^a
WT	3.2
KO: <i>B4galt1</i>	2.2
KO: <i>B4galt2</i>	1.7
KO: <i>B4galt3</i>	0.7
KO: <i>B4galt4</i>	1.3
KO: <i>B4galt1/2</i>	0.6
KO: <i>B4galt1/3</i>	0.2
KO: <i>B4galt1/2/3</i>	0.4
KO: <i>st3gal3</i>	1.5
KO: <i>st3gal4</i>	2.9
KO: <i>st3gal6</i>	0.4
KO: <i>st3gal3/4</i>	3.5
KO: <i>st3gal4/6</i>	6.2
KO: <i>B3gnt1</i>	2.3
KO: <i>B3gnt2</i>	0.5
KO: <i>B3gnt8</i>	2.4
KO: <i>mgat4A</i>	1.1
KO: <i>mgat4A/4B</i>	2.5
KO: <i>mgat5</i>	1.0
KO: <i>mgat4A/4B/5</i>	4.3
KO: <i>mgat4B</i>	6.8
KO: <i>mgat1</i>	6.3
KO: <i>mgat2</i>	5.0
KO: <i>mgat3</i>	2.1
KO: <i>mgat4C</i>	10.4
KO: <i>mgat5B</i>	3.2
KO: <i>B3gnt2/mgat4A/4B/5</i>	3.4
KO: <i>st3gal4/6/mgat4A/4B/5</i>	1.3
KO: <i>B3gnt2/st3gal4/6/mgat4A/4B/5</i>	3.7
KI: ST6GAL-I/KO: <i>st3gal4/6/mgat4A/4B/5</i>	0.2
KI: ST6GAL-I/KO: <i>st3gal4/6</i>	0.8
KO: <i>st3gal4/6/mgat2</i>	10.3
KO: <i>mgat2/4A/4B/5</i>	1.5
KO: <i>st3gal4/6/mgat2/4A/4B/5</i>	0.4
KO: <i>fut8</i>	2.3
IgG clones	Final yield (mg/L) ^a
WT (Rituximab)	2.5
KO: <i>B4galt1</i> (Rituximab)	0.9
KO: <i>fut8</i> (Rituximab)	1.0
KO: <i>fut8/B4galt1</i> (Rituximab)	3.4
WT (SO57)	857
KO: <i>fut8/B4galt1</i> (SO57)	1,187

^aFinal yields represent purified proteins after NiNTA and HPLC purification for EPO after Protein G and ultrafiltration for IgG Rituximab. The SO57 cell line (clone 27) is a CHO-GS cell line producing anti-rabies IgG1 developed by Sigma¹². The production yield of IgG SO57 was determined by analysis of culture medium by ELISA using an internal IgG standard.

Supplementary reference

1. Xu, X. *et al.* The genomic sequence of the Chinese hamster ovary (CHO)-K1 cell line. *Nat. Biotechnol.* **29**, 735-741 (2011).
2. Sasaki, K. *et al.* Expression cloning of cDNA encoding a human beta-1,3-N-acetylglucosaminyltransferase that is essential for poly-N-acetyllactosamine synthesis. *Proc. Natl. Acad. Sci. U. S. A.* **94**, 14294-14299 (1997).
3. Bao, X. *et al.* Tumor suppressor function of laminin-binding alpha-dystroglycan requires a distinct beta3-N-acetylglucosaminyltransferase. *Proc. Natl. Acad. Sci. U. S. A.* **106**, 12109-12114 (2009).
4. Boersema, P.J. *et al.* Multiplex peptide stable isotope dimethyl labeling for quantitative proteomics. *Nat. Proto.* **4**, 484-494 (2009).
5. Walsh, G. & Jefferis, R. Post-translational modifications in the context of therapeutic proteins. *Nat. Biotechnol.* **24**, 1241-1252 (2006).
6. Maresca, M. *et al.* Obligate ligation-gated recombination (ObLiGaRe): custom-designed nuclease-mediated targeted integration through nonhomologous end joining. *Genome Res.* **23**, 539-546 (2013).
7. Wheeler, S.F. & Harvey, D.J. Negative ion mass spectrometry of sialylated carbohydrates: discrimination of N-acetylneurameric acid linkages by MALDI-TOF and ESI-TOF mass spectrometry. *Anal. Chem.* **72**, 5027-5039 (2000).
8. Harvey, D.J. Analysis of carbohydrates and glycoconjugates by matrix-assisted laser desorption/ionization mass spectrometry: an update for the period 2005-2006. *Mass Spec. Reviews* **30**, 1-100 (2011).
9. Yamane-Ohnuki, N. *et al.* Establishment of FUT8 knockout Chinese hamster ovary cells: an ideal host cell line for producing completely defucosylated antibodies with enhanced antibody-dependent cellular cytotoxicity. *Biotechnol. Bioeng.* **87**, 614-622 (2004).
10. Malphettes, L. *et al.* Highly efficient deletion of FUT8 in CHO cell lines using zinc-finger nucleases yields cells that produce completely nonfucosylated antibodies. *Biotechnol. Bioeng.* **106**, 774-783 (2010).
11. Sealover, N.R. *et al.* Engineering Chinese hamster ovary (CHO) cells for producing recombinant proteins with simple glycoforms by zinc-finger nuclease (ZFN)-mediated gene knockout of mannosyl (alpha-1,3)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase (Mgat1). *J. Biotechnol.* **167**, 24-32 (2013).
12. Lin, N. *et al.* Overexpression of Serpinb1 in Chinese hamster ovary cells increases recombinant IgG productivity. *J. Biotechnol.* **193**, 91-99 (2015).