

Table S1: Literature containing carbohydrate CCSs. CCSs from bold references are implemented in Table S2 if not noted otherwise. References in green contain large CCS collections.

reference label	instru- ment type†	CCS drift gas #	investigated carbohydrates	comment	reference
Both2014	TW	He	mono-, disaccharides, glycopeptide fragment CCSs available		P. Both et al., Nat. Chem. 2014, 6, 65-74.
Clowers2005	DT	N ₂	di-, trisaccharides		B. H. Clowers et al., J. Am. Soc. Mass Spectrom. 2005, 16, 660-669.
Dwivedi2007	DT	N ₂ , He, CO ₂ , SF ₆	methyl-monosaccharides	different adducts, different drift gases	P. Dwivedi et al., J. Am. Soc. Mass Spectrom. 2007, 18, 1163-1175.
Fenn2011	DT	He	milk oligosaccharides of different sizes	underivatized and Sp derivatized carbohydrates, CCSs of intact ions and their fragments, adducts	L. S. Fenn, J. A. McLean, Phys. Chem. Chem. Phys. 2011, 13, 2196-2205.
Gaye2015	DT	He	mono-, di- and trisaccharides	fragment CCS available	M. M. Gaye et al., Analyst 2015, 140, 6922-6932.
Gelb2014	DT	He	milk oligosaccharides		A. S. Gelb et al., Anal. Chem. 2014, 86, 11396-11402.
Guttman2016	TW	N ₂	glycopeptides, trisaccharides	sialic acid linkage identification, further fragment CCSs available	M. Guttman, K. K. Lee, Anal. Chem. 2016, 88, 5212-5217.
Harvey2014	TW	N ₂ , He	N-glycans		D. J. Harvey et al., Rapid Commun. Mass Spectrom. 2014, 28, 2008-2018.
Harvey2016	TW	N ₂	reduced N-glycans		D. J. Harvey, J. L. Abrahams, Rapid Commun. Mass Spectrom. 2016, 30, 627-634.
Harvey2016	TW	N ₂ , He	N-glycans	CCSs of intact and fragment ions	D. J. Harvey et al., J. Mass Spectrom. 2016, 51, 1064-1079.
Hinneburg2016	TW	N ₂	glycopeptides	sialic acid linkage identification	H. Hinneburg et al., Chem. Commun. 2016, 52, 4381-4384.
Hofmann2014	DT	N ₂ , He	large number of N-glycans, dextran	CCSs of intact and fragment ions available	J. Hofmann et al., Anal. Chem. 2014, 86, 10789-10795.
Hofmann2015	TW	N ₂	trisaccharides	data in pos+neg ion mode	J. Hofmann et al., Nature 2015, 526, 241-244.
Hofmann2017	TW	N ₂	Lewis and blood group tri- and tetrasaccharides, milk oligosaccharides, N-glycans	CCSs of intact and fragment ions available	J. Hofmann et al., Anal. Chem. 2017, 89, 2318-2325.
Huang2013	TW	He	di-, trisaccharides, LNFP, LNDFH	different adducts	Y. Huang, E. D. Dodds, Anal. Chem. 2013, 85, 9728-9735.
Huang2015	TW	He	trisaccharides and milk oligosaccharides	different adducts and additional CCSs, ETD product ions	Y. T. Huang, E. D. Dodds, Analyst 2015, 14, 6912-6921.
Lee1997	DT	He	tetra- and hexasaccharides	theoretical CCS calculations	S. Lee et al., Int. J. Mass Spectrom. Ion Process. 1997, 167, 605-614.
Li2012	DT	N ₂	methyl-monosaccharides		H. Li et al., Anal. Chem. 2012, 84, 3231-3239.
May2014	DT	N ₂	large number of different saccharides	different adducts	J. C. May et al., Anal. Chem. 2014, 86, 2107-2116.
Pagel2013	DT	N ₂ , He	large number of N-glycans	CCSs of intact and fragment ions available	K. Pagel, D. J. Harvey, Anal. Chem. 2013, 85, 5138-5145.
Paglia2014	TW	N ₂	mono-, di- and trisaccharides	data in pos+neg ion mode, validation of CCS on 3 instruments in different labs, theoretical calculations available	G. Paglia et al., Anal. Chem. 2014, 86, 3985-3993.

reference label	instru- ment type†	CCS drift gas #	investigated carbohydrates	comment	reference
Plasencia2008	DT	He	N-glycans		M. D. Plasencia et al., J. Am. Soc. Mass Spectrom. 2008, 19, 1706-1715.
Pu2016	TIMS, DT	N ₂	permethylated LNnT/LNT		Y. Pu et al., Anal. Chem. 2016, 88, 3440-3443.
Rashid2014	TW	He	dextran, pullulan, maltooligosaccharide	neg ion mode	A. M. Rashid et al., Rapid Commun. Mass Spectrom. 2014, 28, 191-199.
Seo2011	TW	N ₂	heparin octasaccharides	different adducts	Y. Seo et al., Int. J. Mass Spectrom. 2011, 303, 191-198.
Seo2012	TW	N ₂	heparin octasaccharides	different adducts	Y. Seo et al., Anal. Chem. 2012, 84, 2416-2423.
Struwe2015	DT	N ₂	high-mannose glycans	data in pos+neg ion mode	W. B. Struwe et al., Analyst 2015, 140, 6799-6803.
Struwe2016	TW	He	milk oligosaccharides (synthetic and biol. source)	data in pos+neg ion mode, theoretical calculations available	W. B. Struwe et al., Chem. Commun. 2016, 52, 12353-12356.
Williams2010	DT	He	N-glycans and milk oligosaccharides	further CCSs available	J. P. Williams et al., Int. J. Mass Spectrom. 2010, 298, 119-127.
Yang2016	TW	He	mono-, disaccharides	underivatized, and labeled with phenylhydrazine (PHN), and 1-phenyl-3- methyl-5-pyrazolone (PMP)	H. M. Yang et al., Sci. Rep. 2016, 6, 28079.
Zheng2017	DT	N ₂	mono- to hexasaccharides	data in pos+neg ion mode; several isomer separations, but no CCS reported	X. Zheng et al., Anal. Bioanal. Chem. 2017, 409, 467-476.

† TW and TIMS CCSs were obtained from calibrations with reference substances with known CCSs.

Drift gas for which the CCS was calculated or estimated. This is not necessarily the drift gas used during the measurement. Often He calibrant values are used to estimate He CCS from N₂ measurements.

The CCS values of Table S2 are collected from references listed in Table S1. Please cite the appropriate publication(s), if you use any of these values for your research.

Table S2: CCS values in Å² of carbohydrates reported in the references of Table S1. CCS errors correspond to the standard deviations of replicate measurements. Blue values result from measurements in the negative ion mode.

Monosaccharides

molecule	MW in Da	type	<i>m/z</i>	^{TW} CCS _{He} Both14	^{TW} CCS _{He} Yang16	^{DT} CCS _{N₂} May14	^{DT} CCS _{N₂} Li12	^{TW} CCS _{N₂} Paglia14
α-Me-All	178.18	[M + Na] ⁺	201.07				133.04	
α-Me-Alt	178.18	[M + Na] ⁺	201.07				138.74	
α-Me-Gal	178.18	[M + Na] ⁺	201.07				140.8	
α-Me-Glc	178.18	[M + Na] ⁺	201.07				141.77	
α-Me-Gul	178.18	[M + Na] ⁺	201.07				131.34	
α-Me-Ido	178.18	[M + Na] ⁺	201.07				133.04	
α-Me-Man	178.18	[M + Na] ⁺	201.07				136.8	
α-Me-Tal	178.18	[M + Na] ⁺	201.07				126.98	
β-Me-All	178.18	[M + Na] ⁺	201.07				139.83	
β-Me-Alt	178.18	[M + Na] ⁺	201.07				131.34	
β-Me-Gal	178.18	[M + Na] ⁺	201.07				135.83	
β-Me-Glc	178.18	[M + Na] ⁺	201.07				139.83	
β-Me-Gul	178.18	[M + Na] ⁺	201.07				131.34	
β-Me-Ido	178.18	[M + Na] ⁺	201.07				132.19	
β-Me-Man	178.18	[M + Na] ⁺	201.07				134.01	
β-Me-Tal	178.18	[M + Na] ⁺	201.07				131.34	
Glucosamine	179.17	[M + H] ⁺	180.09				135	
Fructose	180.16	[M - H] ⁻	179.06				135	
Fructose	180.16	[M + Na] ⁺	203.05		83.7 ± 0.4		135	
Galactose	180.16	[M - H] ⁻	179.06				130	
Galactose	180.16	[M + Na] ⁺	203.05	75	83.5 ± 0.3		136	
Glucose	180.16	[M - H] ⁻	179.06				135	
Glucose	180.16	[M + Na] ⁺	203.05	78			138	
Mannose	180.16	[M - H] ⁻	179.06				130	
Mannose	180.16	[M + Na] ⁺	203.05	76	83.4 ± 0.5		138	
Mannitol	182.17	[M - H] ⁻	181.07				131	
Mannitol	182.17	[M + Li] ⁺	189.10			144.5		
Mannitol	182.17	[M + Na] ⁺	205.07			140.6	137	
Sorbitol	182.17	[M + H] ⁺	183.09			147.2		

molecule	MW in Da	type	<i>m/z</i>	^{TW} CCS _{He} Both14	^{TW} CCS _{He} Yang16	^{DT} CCS _{N2} May14	^{DT} CCS _{N2} Li12	^{TW} CCS _{N2} Paglia14
Sorbitol	182.17	[M + Na] ⁺	205.07			139.4		
Glucuronic acid	194.14	[M - H] ⁻	193.03					130
Glucuronic acid	194.14	[M + Na] ⁺	217.03					140
Gluconic acid	196.16	[M - H] ⁻	195.05					132
Gluconic acid	196.16	[M + Na] ⁺	219.05					148
α -Me-GalNAc	219.24	[M + Na] ⁺	242.10			145.05		
α -Me-GlcNAc	219.24	[M + Na] ⁺	242.10				151.84	
β -Me-GalNAc	219.24	[M + Na] ⁺	242.10			146.14		
β -Me-GlcNAc	219.24	[M + Na] ⁺	242.10				151.84	
N-Acetylgalactosamine	221.21	[M + Na] ⁺	244.08	85				
N-Acetylglucosamine	221.21	[M - H] ⁻	220.08					143
N-Acetylglucosamine	221.21	[M + Na] ⁺	244.08	90				153
N-Acetylmannosamine	221.21	[M + Na] ⁺	244.08	84				
Ribose 5-phosphate	230.11	[M - H] ⁻	229.01					137
Ribose 5-phosphate	230.11	[M + Na] ⁺	253.01				146	
Glucosamine 1-phosphate	259.15	[M - H] ⁻	258.04					146
Glucosamine 1-phosphate	259.15	[M + H] ⁺	260.05				153	
Glucosamine 6-phosphate	259.15	[M - H] ⁻	258.04					146
Glucosamine 6-phosphate	259.15	[M + H] ⁺	260.05				150	
Fructose 6-phosphate	260.14	[M + Na] ⁺	259.02					150
Fructose 6-phosphate	260.14	[M - H] ⁻	259.02					143
Glucose 6-phosphate	260.14	[M - H] ⁻	259.02					143
Glucose 6-phosphate	260.14	[M + Na] ⁺	283.02					154
Fructose (PHN label)	270.29	[M + Na] ⁺	293.11		100.3 ± 0.5			
Galactose (PHN label)	270.29	[M + Na] ⁺	293.11		98.2 ± 0.4			
Mannose (PHN label)	270.29	[M + Na] ⁺	293.11		98.0 ± 0.5			
6-Phosphogluconate	276.14	[M - H] ⁻	275.02					142
6-Phosphogluconate	276.14	[M + Na] ⁺	299.01					159
Fructose 1,6-diphosphate	340.12	[M - H] ⁻	338.99					150
Fructose (PMP lable)	510.55	[M + H] ⁺	511.22			154.4 ± 0.2		
Galactose (PMP label)	510.55	[M + H] ⁺	511.22			152.8 ± 0.1		
Mannose (PMP label)	510.55	[M + H] ⁺	511.22			153.7 ± 0.2		

Milk oligosaccharides

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Fenn11	^{DT} CCS _{He} Williams10	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang15	^{DT} CCS _{N2} May14	TIMS ^{DT} CCS _{N2} Pu16 §	^{TW} CCS _{N2} Hofmann17	^{TW} CCS _{N2} Struwe16
blood group A trisaccharide	529.49	[M + Na] ⁺	552.19								225 ± 1	
blood group H type 1 trisaccharide	529.49	[M + Na] ⁺	552.19								224 ± 1	
blood group H type 2 trisaccharide	529.49	[M + Na] ⁺	552.19								229 ± 2	
Lewis a trisaccharide	529.49	[M + Na] ⁺	552.19								223 ± 2	
Lewis x trisaccharide	529.49	[M + Na] ⁺	552.19								215 ± 2	
Lewis b tetrasaccharide	675.63	[M + Na] ⁺	698.25								252 ± 1	
Lewis y tetrasaccharide	675.63	[M + Na] ⁺	698.25								248 ± 1	
Lacto-N-neotetraose (LNnT)	707.63	[M - H] ⁻	706.24								186.3 ± 0.8	
Lacto-N-neotetraose (LNnT)	707.63	[M + H] ⁺	708.26								189.1 ± 0.4	
Lacto-N-neotetraose (LNnT)	707.63	[M + Na] ⁺	730.24								169.6 ± 1.0	
Lacto-N-neotetraose (LNnT)	707.63	[M + Cl] ⁻	742.22								180.4 ± 0.8	
Lacto-N-neotetraose (LNnT) - permethylated	707.63	[M + Na] ⁺	926.46							300 299		
Lacto-N-tetraose (LNT)	707.63	[M - H] ⁻	706.24								172.2 ± 1.0	
Lacto-N-tetraose (LNT)	707.63	[M + H] ⁺	708.26								177.9 ± 0.8	
Lacto-N-tetraose (LNT)	707.63	[M + Na] ⁺	730.24								170.5 ± 1.0	
Lacto-N-tetraose (LNT)	707.63	[M + Cl] ⁻	742.22								182.1 ± 1.2	
Lacto-N-tetraose (LNT) - permethylated	707.63	[M + Na] ⁺	926.46							291 293		
Lacto-N-fucopentaose I	853.77	[M + Be] ²⁺	431.16					204.0 ± 0.3				
Lacto-N-fucopentaose I	853.77	[M + Mg] ²⁺	438.65					201.1 ± 0.4				
Lacto-N-fucopentaose I	853.77	[M + Ca] ²⁺	446.63					198.2 ± 0.8				
Lacto-N-fucopentaose I	853.77	[M + Sr] ²⁺	470.61					200.8 ± 1.0				
Lacto-N-fucopentaose I	853.77	[M + Ba] ²⁺	495.61					193.5 ± 0.9				
Lacto-N-fucopentaose I	853.77	[M + Li] ⁺	860.32	203.1 ± 1.2		198.1 ± 0.2	200.2 ± 0.3		269.6			
Lacto-N-fucopentaose I	853.77	[M + Na] ⁺	876.30	204.4 ± 1.4	196	201.5 ± 0.3	203.8 ± 0.3		276.1			
Lacto-N-fucopentaose I	853.77	[M + K] ⁺	892.27	205.0 ± 0.7		201.1 ± 0.3	203.4 ± 0.3		274.7			
Lacto-N-fucopentaose I	853.77	[M + Rb] ⁺	938.22	198.4 ± 1.5		200.8 ± 0.2	203.1 ± 0.2		275.2			
Lacto-N-fucopentaose I	853.77	[M + Cs] ⁺	986.21	204.0 ± 2.1		202.0 ± 0.6	204.3 ± 0.6		275.6			
Lacto-N-fucopentaose I (peak 1)	853.77	[M + H] ⁺	854.31		185							

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Fenn11	^{DT} CCS _{He} Williams10	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang15	^{DT} CCS _{N2} May14	TIMS ^{DT} CCS _{N2} Pu16 §	^{TW} CCS _{N2} Hofmann17	^{TW} CCS _{N2} Struwe16
Lacto-N-fucopentaose I (peak2)	853.77	[M + H] ⁺	854.31		201							
Lacto-N-fucopentaose II	853.77	[M + Be] ²⁺	431.16					201.2 ± 0.4				
Lacto-N-fucopentaose II	853.77	[M + Mg] ²⁺	438.65					203.2 ± 1.2				
Lacto-N-fucopentaose II	853.77	[M + Ca] ²⁺	446.63					201.7 ± 0.5				
Lacto-N-fucopentaose II	853.77	[M + Sr] ²⁺	470.61					201.5 ± 0.5				
Lacto-N-fucopentaose II	853.77	[M + Ba] ²⁺	495.61					205.5 ± 0.5				
Lacto-N-fucopentaose II	853.77	[M + Li] ⁺	860.32	198.7 ± 1.3				194.1 ± 0.5				
Lacto-N-fucopentaose II	853.77	[M + Na] ⁺	876.30	201.3 ± 1.2				195.6 ± 0.6	271.1			
Lacto-N-fucopentaose II	853.77	[M + K] ⁺	892.27	202.6 ± 1.6				198.5 ± 0.4	267.2			
Lacto-N-fucopentaose II	853.77	[M + Rb] ⁺	938.22	197.5 ± 1.7				200.0 ± 0.4	278.4			
Lacto-N-fucopentaose II	853.77	[M + Cs] ⁺	986.21					200.9 ± 0.4				
Lacto-N-fucopentaose III	853.77	[M + Be] ²⁺	431.16					201.3 ± 0.7				
Lacto-N-fucopentaose III	853.77	[M + Mg] ²⁺	438.65					199.8 ± 0.8				
Lacto-N-fucopentaose III	853.77	[M + Ca] ²⁺	446.63					195.9 ± 0.9				
Lacto-N-fucopentaose III	853.77	[M + Sr] ²⁺	470.61					196.2 ± 1.0				
Lacto-N-fucopentaose III	853.77	[M + Ba] ²⁺	495.61					197.4 ± 1.1				
Lacto-N-fucopentaose III	853.77	[M + Li] ⁺	860.32					192.9 ± 0.2				
Lacto-N-fucopentaose III	853.77	[M + Na] ⁺	876.30					196.6 ± 0.4				
Lacto-N-fucopentaose III	853.77	[M + Na] ⁺	876.30	199.2 ± 1.0								
Lacto-N-fucopentaose III	853.77	[M + K] ⁺	892.27					194.5 ± 0.4				
Lacto-N-fucopentaose III	853.77	[M + Rb] ⁺	938.22					195.3 ± 0.5				
Lacto-N-fucopentaose III	853.77	[M + Cs] ⁺	986.21					195.7 ± 0.3				
Lacto-N-fucopentaose V	853.77	[M + Be] ²⁺	431.16					197.3 ± 0.3				
Lacto-N-fucopentaose V	853.77	[M + Mg] ²⁺	438.65					197.4 ± 0.5				
Lacto-N-fucopentaose V	853.77	[M + Ca] ²⁺	446.63					196.7 ± 0.2				
Lacto-N-fucopentaose V	853.77	[M + Sr] ²⁺	470.61					197.8 ± 1.1				
Lacto-N-fucopentaose V	853.77	[M + Ba] ²⁺	495.61					197.6 ± 0.9				
Lacto-N-fucopentaose V	853.77	[M + H] ⁺	854.31	195								
Lacto-N-fucopentaose V	853.77	[M + Li] ⁺	860.32				198.0 ± 0.3					
Lacto-N-fucopentaose V	853.77	[M + Na] ⁺	876.30	201.8 ± 0.7	194		198.4 ± 0.3					

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Fenn11	^{DT} CCS _{He} Williams10	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang15	^{DT} CCS _{N2} May14	TIMS ^{DT} CCS _{N2} Pu16 §	^{TW} CCS _{N2} Hofmann17	^{TW} CCS _{N2} Struwe16
Lacto-N-fucopentaose V	853.77	[M + K] ⁺	892.27			199.4 ± 0.3						
Lacto-N-fucopentaose V	853.77	[M + Rb] ⁺	938.22			199.9 ± 0.4						
Lacto-N-fucopentaose V	853.77	[M + Cs] ⁺	986.21			201.7 ± 1.0						
Lacto-N-difucohexaose I	999.91	[M + Be] ²⁺	504.19					226.5 ± 0.5				
Lacto-N-difucohexaose I	999.91	[M + Mg] ²⁺	511.67					226.0 ± 0.5				
Lacto-N-difucohexaose I	999.91	[M + Ca] ²⁺	519.66					22.1 ± 0.6				
Lacto-N-difucohexaose I	999.91	[M + Sr] ²⁺	543.63					223.1 ± 0.5				
Lacto-N-difucohexaose I	999.91	[M + Ba] ²⁺	568.63					220.2 ± 1.1				
Lacto-N-difucohexaose I	999.91	[M + Li] ⁺	1006.38			221.5 ± 0.8			301.4			
Lacto-N-difucohexaose I	999.91	[M + K] ⁺	1038.33			223.7 ± 0.6			303.5			
Lacto-N-difucohexaose I	999.91	[M + Rb] ⁺	1084.28			223.7 ± 0.7			303.2			
Lacto-N-difucohexaose I	999.91	[M + Cs] ⁺	1132.27			223.7 ± 0.6			301.2			
Lacto-N-difucohexaose I (peak 1)	999.91	[M + Na] ⁺	1022.35	225.6 ± 1.1	217	223.5 ± 0.7			290.6			
Lacto-N-difucohexaose I (peak 2)	999.91	[M + Na] ⁺	1022.35						304.2			
Lacto-N-difucohexaose II	999.91	[M + Be] ²⁺	504.19					226.5 ± 0.4				
Lacto-N-difucohexaose II	999.91	[M + Mg] ²⁺	511.67					227.2 ± 0.8				
Lacto-N-difucohexaose II	999.91	[M + Ca] ²⁺	519.66					225.5 ± 0.5				
Lacto-N-difucohexaose II	999.91	[M + Sr] ²⁺	543.63					225.7 ± 1.4				
Lacto-N-difucohexaose II	999.91	[M + Ba] ²⁺	568.63					225.5 ± 0.9				
Lacto-N-difucohexaose II	999.91	[M + Li] ⁺	1006.38			217.7 ± 0.3	220.6±0.4					
Lacto-N-difucohexaose II	999.91	[M + K] ⁺	1038.33			219.9 ± 0.3	222.9±0.4		305.8			
Lacto-N-difucohexaose II	999.91	[M + Rb] ⁺	1084.28			219.8 ± 0.2	222.8±0.4					
Lacto-N-difucohexaose II	999.91	[M + Cs] ⁺	1132.27			220.4 ± 0.2	223.5±0.3					
Lacto-N-difucohexaose II (peak 1)	999.91	[M + Na] ⁺	1022.35	220.6 ± 1.0	214	218.7 ± 0.3	221.6±0.5		291.2			
Lacto-N-difucohexaose II (peak 2)	999.91	[M + Na] ⁺	1022.35						306.3			
Lacto-N-hexaose (LNH)	1072.96	[M - H] ⁻	1071.37							255.2 ± 1.6		
Lacto-N-hexaose (LNH)	1072.96	[M + H] ⁺	1073.39							225.5 ± 0.4		
Lacto-N-hexaose (LNH)	1072.96	[M + Na] ⁺	1095.37							232.6 ± 0.2		
Lacto-N-hexaose (LNH)	1072.96	[M + Cl] ⁻	1107.35							244.5 ± 2.4		
Lacto-N-neohexaose (LNnH)	1072.96	[M - H] ⁻	1071.37							227.6 ± 1.6		

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Fenn11	^{DT} CCS _{He} Williams10	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang15	^{DT} CCS _{N2} May14	TIMS ^{DT} CCS _{N2} Pu16 §	^{TW} CCS _{N2} Hofmann17	^{TW} CCS _{N2} Struwe16
Lacto-N-neohexaose (LNnH)	1072.96	[M + H] ⁺	1073.39									228.6 ± 1.0
Lacto-N-neohexaose (LNnH)	1072.96	[M + Na] ⁺	1095.37									224.9 ± 0.2
Lacto-N-neohexaose (LNnH)	1072.96	[M + Cl] ⁻	1107.35									244.3 ± 1.2

‡ Values before and after the vertical bar | are derived from a calibration with carbohydrates and peptides, respectively.

§ Values separated by the vertical bar | are values from the same reference but different instruments.

Oligosaccharides

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Gaye15	^{DT} CCS _{He} Lee97	^{DT} CCS _{He} Fenn11	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Huang15	^{TW} CCS _{He} Yang16	^{DT} CCS _{N2 200°C} Clowers05	^{DT} CCS _{N2} May14	^{TW} CCS _{N2} Paglia14
Sucrose	342.3	[M + Li] ⁺	349.13	104.5				107.1 ± 0.3					
Sucrose	342.3	[M + Na] ⁺	365.11					108.9 ± 0.3				170	
Sucrose	342.3	[M + K] ⁺	381.08					110.3 ± 0.3					
Sucrose	342.3	[M + Rb] ⁺	427.03					111.3 ± 0.2					
Sucrose	342.3	[M + Cs] ⁺	475.02					112.9 ± 0.3					
Trehalose	342.30	[M - H] ⁻	341.11									166	
Trehalose	342.3	[M + Li] ⁺	349.13	105.1				107.9 ± 0.3					
Trehalose	342.3	[M + Na] ⁺	365.11					110.6 ± 0.4				173	
Trehalose	342.3	[M + K] ⁺	381.08					114.0 ± 0.4					
Trehalose	342.3	[M + Rb] ⁺	427.03					115.0 ± 0.4					
Trehalose	342.3	[M + Cs] ⁺	475.02					116.5 ± 0.4					
Cellobiitol	344.31	[M + Na] ⁺	367.12							146.09			
Melibiitol	344.31	[M + Na] ⁺	367.12							146.52			
N-acetyl-lactoseamine	383.35	[M + Na] ⁺	406.13			129.2 ± 2.1							
Cellobiose (PHN label)	432.43	[M + Na] ⁺	455.16						139.4 ± 0.3				
Gentiobiose (PHN label)	432.43	[M + Na] ⁺	455.16						134.3 ± 0.2				
Isomaltose (PHN label)	432.43	[M + Na] ⁺	455.16						134.7 ± 0.3				
Kojibiose (PHN label)	432.43	[M + Na] ⁺	455.16						134.0 ± 0.2				
Lactose (PHN label)	432.43	[M + Na] ⁺	455.16						137.3 ± 0.3				
Laminaribiose (PHN label)	432.43	[M + Na] ⁺	455.16						128.6 ± 0.5				
Maltose (PHN label)	432.43	[M + Na] ⁺	455.16						138.1 ± 0.2				
Nigerose (PHN label)	432.43	[M + Na] ⁺	455.16						126.5 ± 0.6				
Sophorose (PHN label)	432.43	[M + Na] ⁺	455.16						133.4 ± 0.7				
Isomaltotriose	504.44	[M + Mg] ²⁺	264.08					138.3 ± 0.3					
Isomaltotriose	504.44	[M + Ca] ²⁺	272.07					136.0 ± 1.0					
Isomaltotriose	504.44	[M + Sr] ²⁺	296.04						142.2 ± 0.3				
Isomaltotriose	504.44	[M + Ba] ²⁺	321.04						145.5 ± 0.4				
Isomaltotriose	504.44	[M + Li] ⁺	511.19					140.8 ± 0.1					
Isomaltotriose	504.44	[M + Na] ⁺	527.16					142.3 ± 0.1		184.14			
Isomaltotriose	504.44	[M + K] ⁺	543.13					143.4 ± 0.1					
Isomaltotriose	504.44	[M + Rb] ⁺	589.08					143.4 ± 0.1					

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Gaye15	^{DT} CCS _{He} Lee97	^{DT} CCS _{He} Fenn11	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Huang15	^{TW} CCS _{He} Yang16	^{DT} CCS _{N2 200°C} Clowers05	^{DT} CCS _{N2} May14	^{TW} CCS _{N2} Paglia14
Isomaltotriose	504.44	[M + Cs] ⁺	637.07					144.2 ± 0.1					
Maltotriose	504.44	[M + Mg] ²⁺	264.08						136.5 ± 0.7				
Maltotriose	504.44	[M + Ca] ²⁺	272.07						136.0 ± 0.6				
Maltotriose	504.44	[M + Sr] ²⁺	296.04						139.4 ± 0.7				
Maltotriose	504.44	[M + Ba] ²⁺	321.04						142.8 ± 1.1				
Maltotriose	504.44	[M + H] ⁺	505.18									216.3	
Maltotriose	504.44	[M + Li] ⁺	511.19	138				138.6 ± 0.2					
Maltotriose	504.44	[M + Na] ⁺	527.16				140.6 ± 0.9 145.3 ± 1.5 ‡	142.9 ± 0.2				212.8	
Maltotriose	504.44	[M + K] ⁺	543.13					144.4 ± 0.2				216.8	
Maltotriose	504.44	[M + Rb] ⁺	589.08					147.0 ± 0.3					
Maltotriose	504.44	[M + Cs] ⁺	637.07					150.8 ± 0.2					
Melezitose	504.44	[M + Mg] ²⁺	264.08						137.9 ± 0.5				
Melezitose	504.44	[M + Ca] ²⁺	272.07						137.7 ± 0.6				
Melezitose	504.44	[M + Sr] ²⁺	296.04						142.6 ± 0.4				
Melezitose	504.44	[M + Ba] ²⁺	321.04						146.5 ± 0.3				
Melezitose	504.44	[M + H] ⁺	505.18								202.6		
Melezitose	504.44	[M + Li] ⁺	511.19	132.6				134.1 ± 0.2				202.9	
Melezitose	504.44	[M + Na] ⁺	527.16					133.5 ± 0.2		173.57	202.7		
Melezitose	504.44	[M + K] ⁺	543.13					134.2 ± 0.2			221.9		
Melezitose	504.44	[M + Cs] ⁺	637.07					136.9 ± 0.2			205.8		
Melezitose (peak 1)	504.44	[M + Rb] ⁺	294.54					134.7 ± 0.3			204.1		
Melezitose (peak2)	504.44	[M + Rb] ⁺	294.54								219.2		
Raffinose	504.44	[M + Mg] ²⁺	264.08						136.2 ± 0.3				
Raffinose	504.44	[M + Ca] ²⁺	272.07						133.6 ± 0.4				
Raffinose	504.44	[M + Sr] ²⁺	296.04						138.4 ± 0.6				
Raffinose	504.44	[M + Ba] ²⁺	321.04						141.3 ± 1.1				
Raffinose	504.44	[M - H] ⁻	503.16								201		
Raffinose	504.44	[M + Li] ⁺	511.19	137.4				137.3 ± 0.4					
Raffinose	504.44	[M + Na] ⁺	527.16					138.8 ± 0.4		180.91	210.7	210	
Raffinose	504.44	[M + K] ⁺	543.13					140.0 ± 0.4			212.7		

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Gaye15	^{DT} CCS _{He} Lee97	^{DT} CCS _{He} Fenn11	^{TW} CCS _{He} Gelb14 ‡	^{TW} CCS _{He} Huang13	^{TW} CCS _{He} Huang15	^{TW} CCS _{He} Yang16	^{DT} CCS _{N2 200°C} Clowers05	^{DT} CCS _{N2} May14	^{TW} CCS _{N2} Paglia14
Raffinose	504.44	[M + Cs] ⁺	637.07					141.8 ± 0.4					
Raffinose (peak1)	504.44	[M + Rb] ⁺	589.08					140.5 ± 0.4*			218.7		
Raffinose (peak2)	504.44	[M + Rb] ⁺	589.08								229.7		
Cellotetraose	666.58	[M + Na] ⁺	689.21		160								
Isomaltotetraose	666.58	[M + Na] ⁺	689.21		163								
Maltotetraose	666.58	[M + H] ⁺	665.21								238.3		
Maltotetraose	666.58	[M + Na] ⁺	689.21		159						235.3		
Cellobiose (PMP label)	672.69	[M + H] ⁺	673.27					177.1 ± 0.2					
Gentibiose (PMP label)	672.69	[M + H] ⁺	673.27					176.2 ± 0.3					
Isomaltose (PMP label)	672.69	[M + H] ⁺	673.27					174.7 ± 0.2					
Kojibiose (PMP label)	672.69	[M + H] ⁺	673.27					180.0 ± 0.5					
Lactose (PMP label)	672.69	[M + H] ⁺	673.27					177.9 ± 0.6					
Laminaribiose (PMP label)	672.69	[M + H] ⁺	673.27					184.4 ± 0.3					
Maltose (PMP label)	672.69	[M + H] ⁺	673.27					182.5 ± 0.3					
Nigerose (PMP label)	672.69	[M + H] ⁺	673.27					180.1 ± 0.4					
Sophorose (PMP label)	672.69	[M + H] ⁺	673.27					179.2 ± 0.2					
Maltohexaose	990.86	[M + Na] ⁺	1013.32		206						286.4		
Maltohexaose	990.86	[M + K] ⁺	1029.29								293.3		
Maltoheptaose	1153.00	[M + H] ⁺	1153.40								303.3		
Maltoheptaose	1153.00	[M + Na] ⁺	1175.37								303.1		
Maltoheptaose	1153.00	[M + K] ⁺	1191.34								303.4		

‡ Values before and after the vertical bar | are derived from a calibration with carbohydrates and peptides, respectively.

*Peak 2 is not reported here.

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{N2} Clowers05	^{TW} CCS _{N2} Hinneburg16	^{TW} CCS _{N2} Guttman16
α -D-GalNAc-(1-3)-D-GalNAc-ol	426.42	[M + Na] ⁺	449.17	160.98		
α -D-GalNAc-(1-6)-D-GalNAc-ol	426.42	[M + Na] ⁺	449.17	164.28		
β -D-GlcNAc-(1-3)-D-GalNAc-ol	426.42	[M + Na] ⁺	449.17	163.49		
β -D-GlcNAc-(1-4)-D-GlcNAc-ol	426.42	[M + Na] ⁺	449.17	165.44		
β -D-GlcNAc-(1-6)-D-GalNAc-ol	426.42	[M + Na] ⁺	449.17	169.70		
α NeuAc-2,3- β -Gal-1,4-GlcNAc	fragment	[M - H ₂ O + H] ⁺	657.23		246	247.2 ± 0.6
α NeuAc-2,6- β -Gal-1,4-GlcNAc	fragment	[M - H ₂ O + H] ⁺	657.23		236	236.9 ± 0.4
β Gal-1,3-(α NeuAc-2,6-)GlcNAc	fragment	[M - H ₂ O + H] ⁺	657.23			231 ± 0.6

Cyclodextrin

molecule	MW in Da	type	<i>m/z</i>	^{DT} CCS _{He} Lee97	^{DT} CCS _{He} Fenn11	^{DT} CCS _{N2} May14
Alpha-Cyclodextrin	972.84	[M + H] ⁺	973.32			285.2
Alpha-Cyclodextrin	972.84	[M + K] ⁺	1011.28			287.7
Alpha-Cyclodextrin	972.84	[M + Na] ⁺	995.31	204	200.7 ± 0.5	285.5
Beta-Cyclodextrin	1134.98	[M + K] ⁺	1173.33			320.3
Beta-Cyclodextrin	1134.98	[M + Na] ⁺	1157.36		231.4 ± 0.6	319.7
Beta-Cyclodextrin (peak 1)	1134.98	[M + H] ⁺	1135.38			301.3
Beta-Cyclodextrin (peak 2)	1134.98	[M + H] ⁺	1135.38			319.6
Gamma-Cyclodextrin	1297.12	[M + Cs] ⁺	1429.33			338.2
Gamma-Cyclodextrin	1297.12	[M + H] ⁺	1297.43			322.6
Gamma-Cyclodextrin	1297.12	[M + K] ⁺	1335.39			324.8
Gamma-Cyclodextrin	1297.12	[M + Li] ⁺	1303.44			317.7
Gamma-Cyclodextrin	1297.12	[M + Na] ⁺	1319.41			322.1
Gamma-Cyclodextrin	1297.12	[M + Rb] ⁺	1381.33			327.3