



Glycan Parameters	fucosylation / bisecting GlcNAc / sialylation / hybrid / high mannose / truncated	F	oF	B	S0	S>0	S1	S2	S3	S4	M	HB	trunc
		95.5	6.4	37.8	53.0	47.0	29.3	17.3	0.4	0.0	2.8	4.7	0.0
	galactosylation / terminal galactosylation	G0	G>0	G1	G2	G3	G4	tG0	tG>0	tG1	tG2	tG3	tG4
	GalNAc/terminal GalNAc	GalNAc0	GalNAc>0	GalNAc1	GalNAc2	GalNAc3	GalNAc4	tGalNAc0	tGalNAc>0	tGalNAc1	tGalNAc2	tGalNAc3	tGalNAc4
		96.8	3.2	3.2	0.0	0.0	0.0	97.2	2.8	2.8	0.0	0.0	0.0
	antennary / phosphorylation / sulfation	A1	A2	A3	A4	phos0	phos>0	phos1	phos2	sulf0	sulf>0	sulf1	sulf2
	NeuGc / alpha Gal (Galili epitope)	NeuGc0	NeuGc>0	NeuGc1	NeuGc2	NeuGc3	NeuGc4	Ga0	Ga>0	Ga1	Ga2	Ga3	Ga4
LacNAc repeats / acetylation / blood group antigens		100.0	0.0	0.0	0.0	0.0	0.0	100.0	0.0	0.0	0.0	0.0	0.0
		LacNAc0	LacNAc>0	LacNAc1	LacNAc2	LacNAc3	Le ^{A/X}	sLe ^{A/X}	Le ^{B/Y}	H-Type			% assigned glycans
		100.0	0.0	0.0	0.0	0.0	0.9	4.5	0.0	0.0			% unassigned glycans
		Ac0	Ac>0	Ac1	Ac2	Ac3							% impurities
Additional result information		100.0	0.0	0.0	0.0	0.0							total structure number
	total peak area	32568											109
	total compound number	109											
	total isomere number	28											
General Sample-information	einzelne Strukturen	38											
	Sample identification	21_0177	PM 1528										
	Additional sample information	Standard for RF											
	Organism / Cell line	H9D8											
Sample Preparation	Sample amount / concentration / volume	15µg											
	Sample buffer	PBS											
	Fluorescence Tag	RapiFluor™ (Waters Inc.)											
LC Settings	Sample preparation protocol	210528 CSH 987											
	LC equipment	ACQUITY UPLC i-Class Series (SM-FTN, BSM, CH, FLR)											
	LC Column	Acquity UPLC Glycan BEH Amide (PN: 186004742, Waters, #01453525816628)											
MS Settings	LC Eluents	100mM Ammoniumformiat-Puffer pH 4.5 / Acetonitril											
	LC method	glycans_rapifluor_classic_LC_01											
	MS equipment	Compact											
Quality Control Standard	MS method	glycans_rapifluor_classic_MS_02											
	External quality control standard												
	Glycan parameters	Std ID#	F	B	G1	G2	G3	G4	S1	S2	S3	S4	M
	Current standard	21_0177	95.5	37.8	34.3	45.6	3.0	0.0	29.3	17.3	0.4	0.0	2.8
		21_0153	96.0	35.1	33.7	45.4	3.3	0.0	28.8	17.3	0.4	0.0	2.8
	Previous standards	21_0134	95.4	37.3	34.3	45.4	3.9	0.0	28.7	18.3	0.6	0.0	2.4
		21_0111	96.7	37.2	33.6	47.2	3.2	0.0	28.6	18.7	0.6	0.0	2.8
		21_0102	95.2	35.3	33.6	44.0	3.9	0.0	30.2	15.3	0.6	0.0	3.3
		21_0058	95.2	36.9	34.4	46.3	3.1	0.0	28.6	19.0	0.6	0.0	2.8
		21_0040	94.9	35.0	33.8	45.2	3.9	0.0	30.4	15.7	0.7	0.0	3.5
MS Settings	Mean (previous stds)	95.6	36.1	33.9	45.6	3.6	0.0	29.2	17.4	0.6	0.0	0.0	2.9
	absolute Δ (current std - mean std)	0.1	1.7	0.4	0.1	0.6	0.0	0.0	0.0	0.1	0.0	0.0	0.1
	max absolute deviation of Δ (current std - mean std)	2.5	2.5	2.5	2.5	2.5	2.5	2.5	2.5	2.5	2.5	2.5	2.5
	compliance status (absolute deviation)	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓	✓



Certificate of Analysis