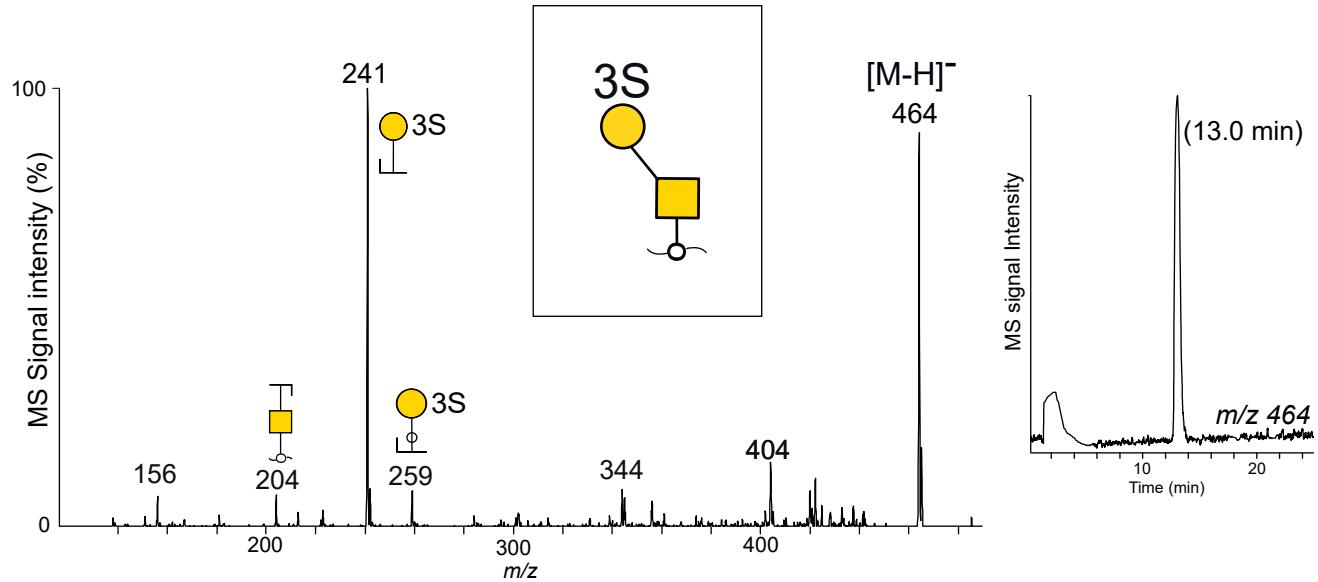
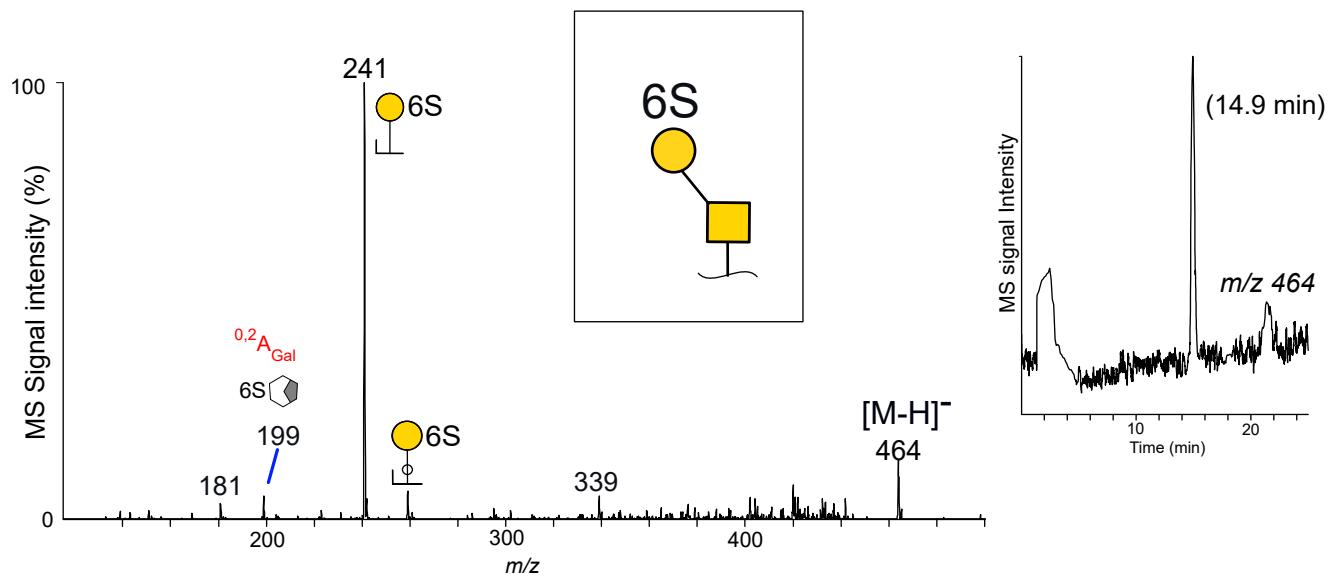


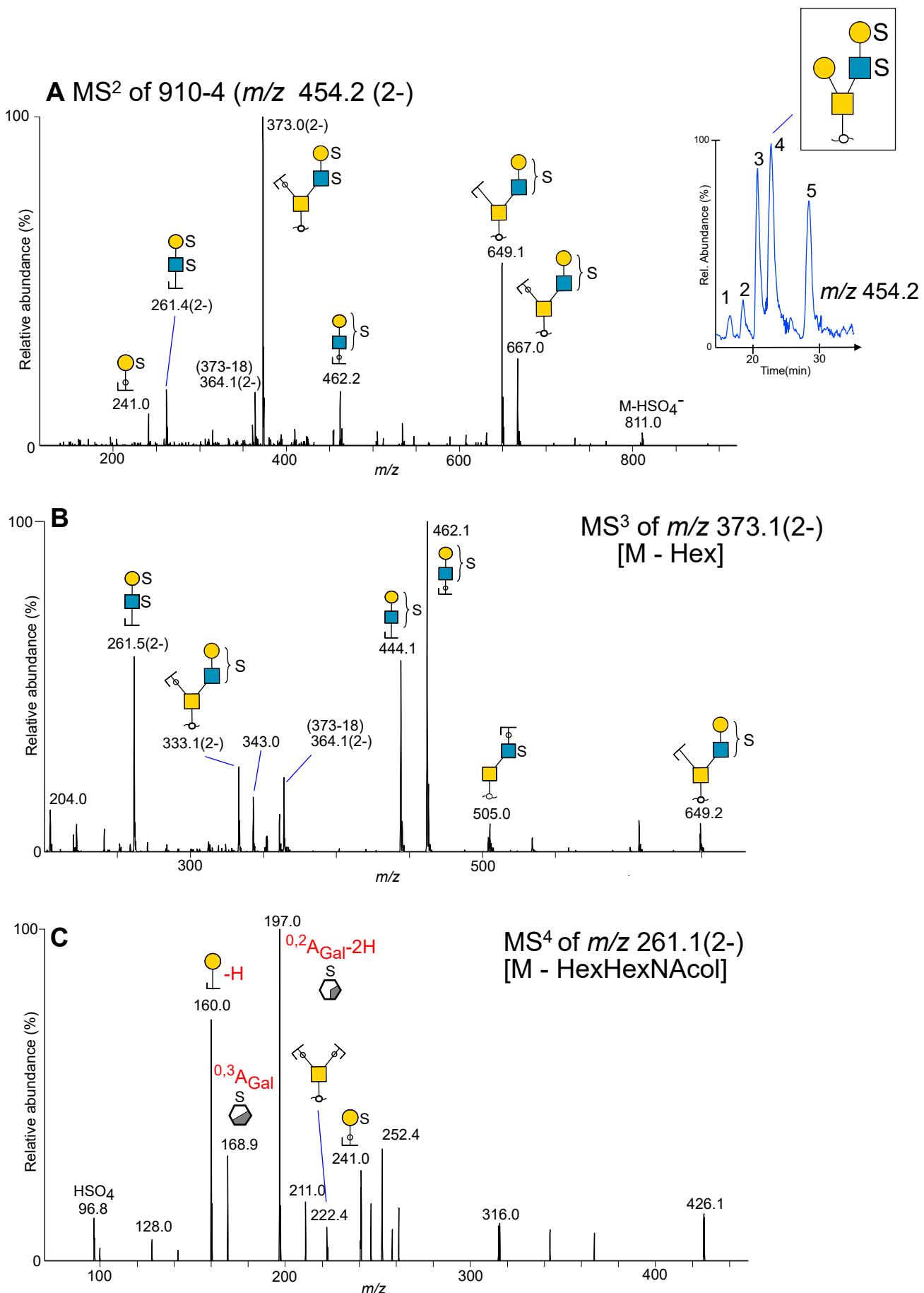
A Gal3ST4



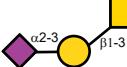
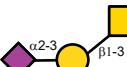
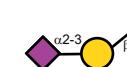
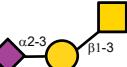
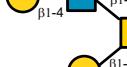
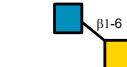
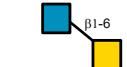
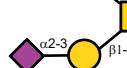
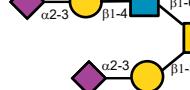
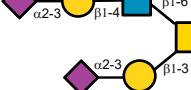
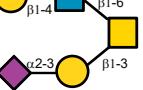
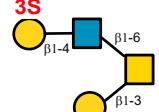
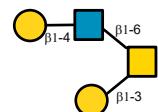
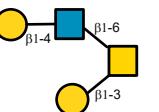
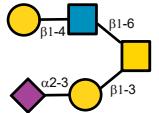
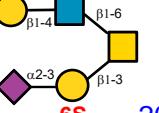
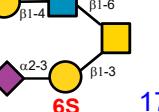
B CHST1



Supplemental Figure 1 LC- MS² spectra of sulfated core 1 O-glycans collected after fragmentation of the precursor ion at m/z 464 ([M-H]⁻) from recombinantly produced PSGL-1/mlgG2b protein purified from CHO cells. PSGL-1 was co-transfected with Gal3ST4 (A) or CHST1 (B). Included in each figure is EIC of m/z 464 from LC-MS run. For key of symbols, see Figure 1.



Supplemental Figure 2 Fragmentation of the disulfated isomer 910-4. Glycan sequence was interpreted from MS2, MS3 and MS4 experiments. (A) MS² spectrum of parent detected as $[M-2H]^{2-}$ ions at m/z 454.2²⁻ (inserted EIC). Fragment ions are annotated with proposed compositions. For some fragment ions, more than one composition may be possible. (B) MS³ spectrum of the m/z 373.0²⁻ from the MS² experiment shown in (A). (C) MS⁴ spectra of the ions at m/z 261.1²⁻ from the MS³ experiment shown in (B). For key of symbols, see Figure 1.

m/z [M-xH] ^{x-}	PSGL-1/mlgG2b + GCNT1 ¹	+ GAL3ST2	+ GAL3ST4	+ CHST1
675	 2%	 8%	 11%	 10%
749	 1%		 6%	 6%
878	 2%	 6%	 8%	 5%
1040	 39%		 40%	 24%
1331	 54%		 30%	 16%
829		 6%	 5%	 3%
1120		 79%		 20%
599				 17%

Supplemental Figure 3 Table of glycans detected after transfection of CHO-cells with PSGL-1/mlgG2b and GCNT1 constructs together with constructs for sulfotransferases. Included is also the relative abundance as measured by their ion intensity of parent ions from the LC-MS runs. For key of symbols, see Figure 1.