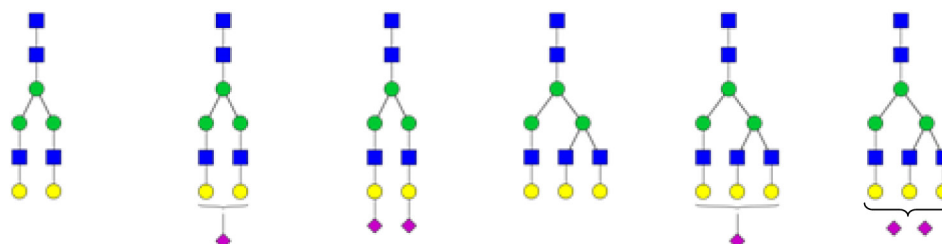
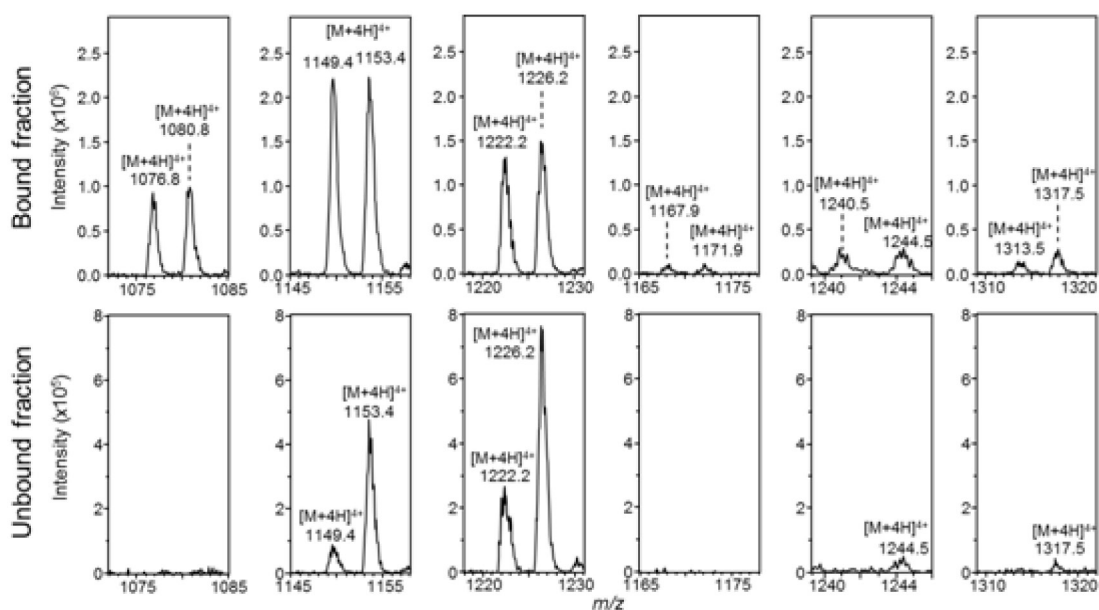


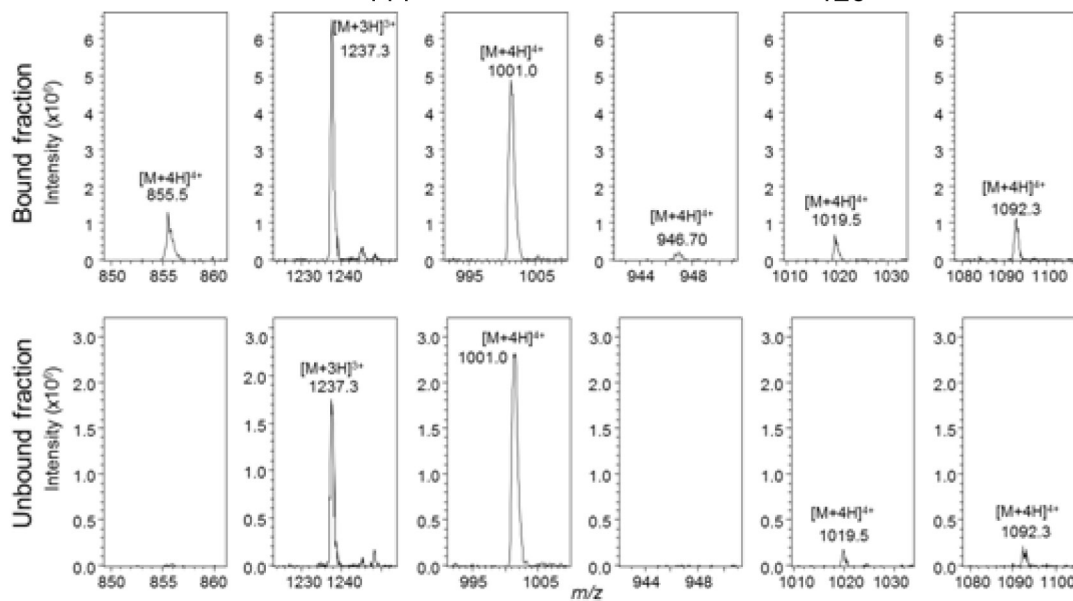
A



Site 1 M(ox)₅₄VSHH**N**LTTGATLINEQWLLTTAK₇₇



Site 4 V₁₁₁VLHP**N**YSQVDIGLIK₁₂₆



B

Site 2 and 3 $N_{78}LFLNHSENA_{90}TAK_{90}$

