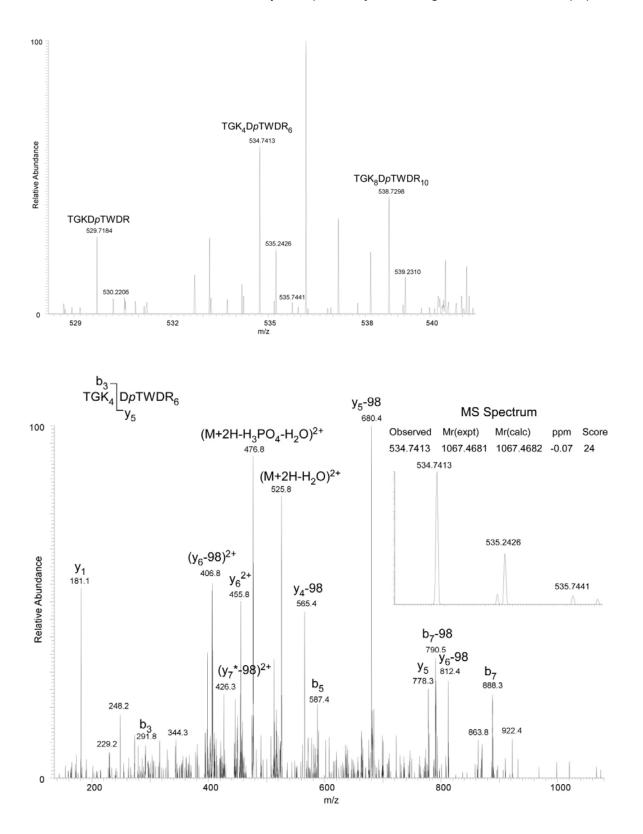
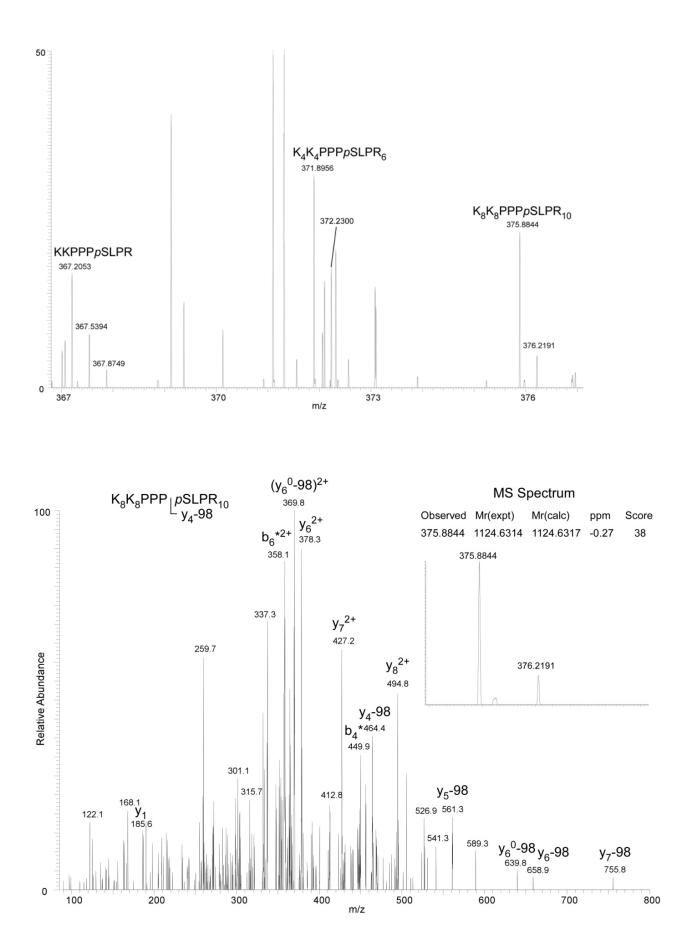
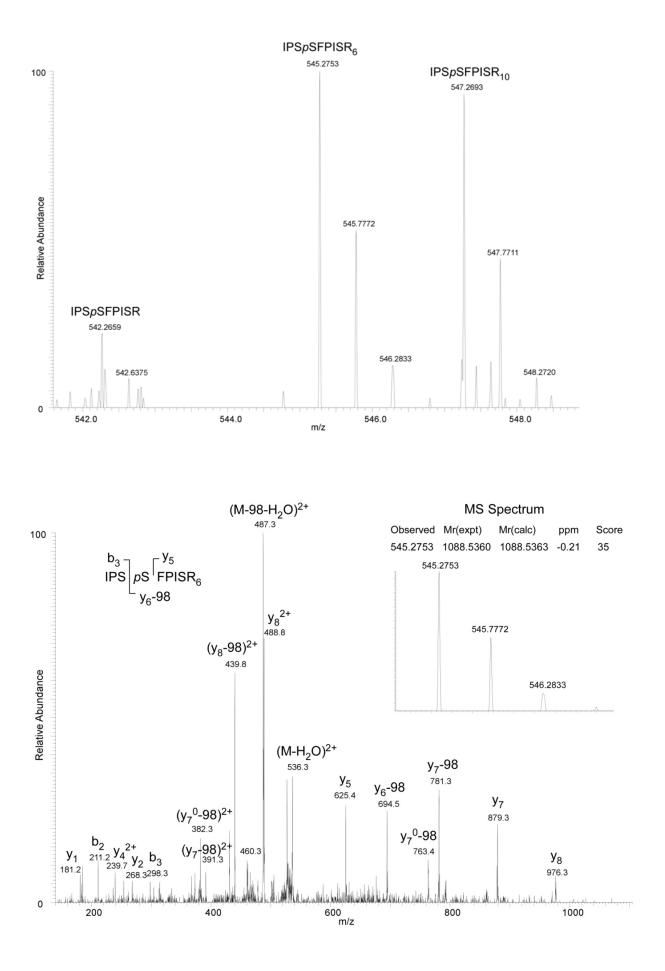
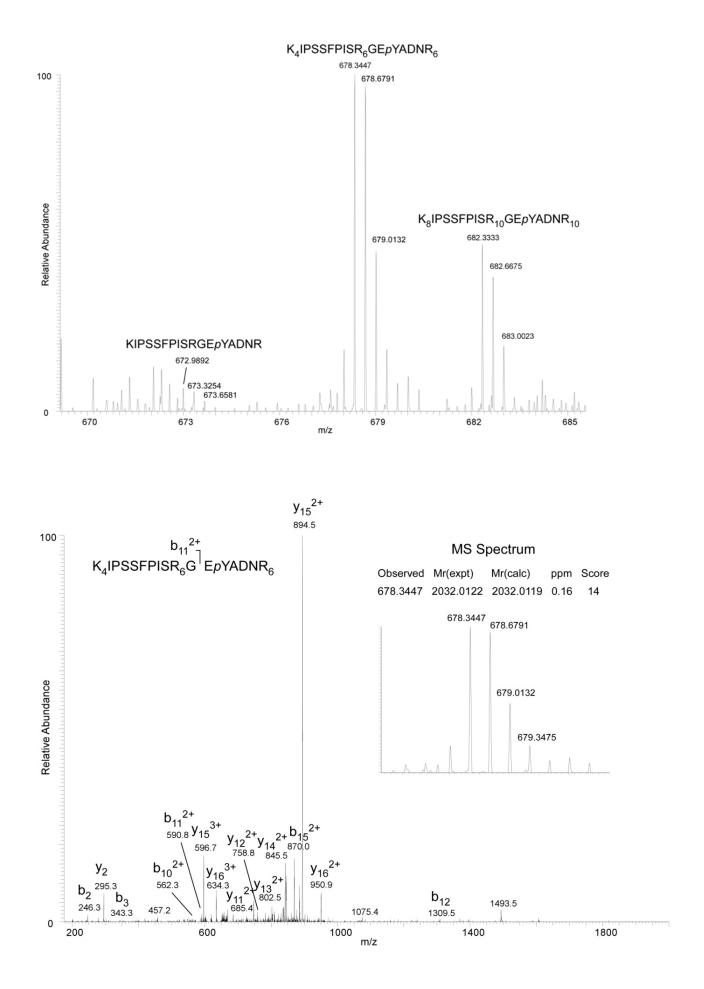
Supplementary Data 2

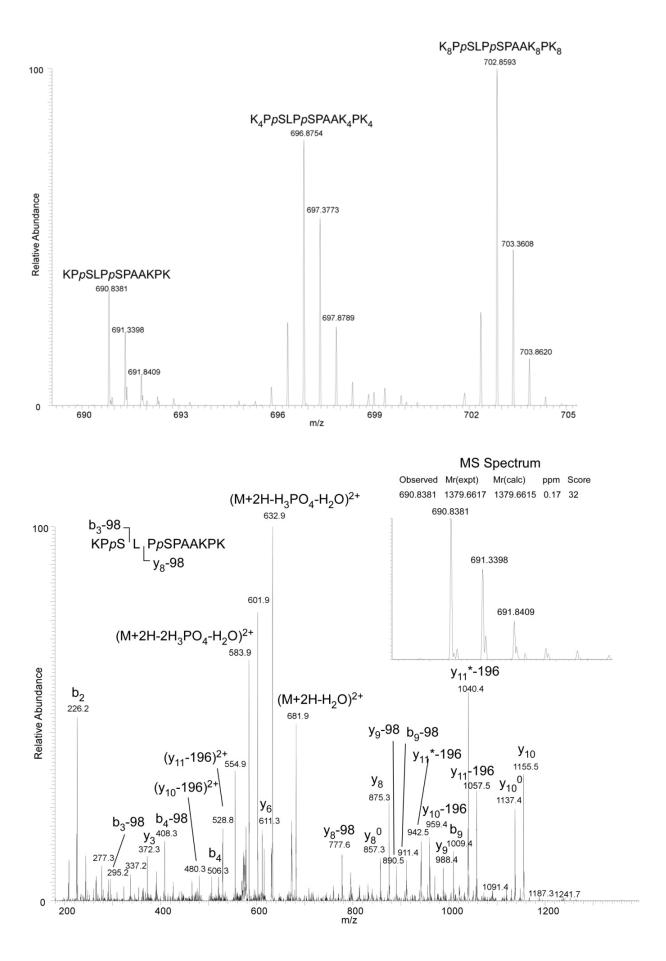
Annotated ESI-MSMS spectra of phosphorylated and quantified peptides derived from SLP-65 after digestion with trypsin and enrichment. The annotation clearly shows the phosphorylation site. Unambigiously identified phosphorylation sites are in capitals (pS/pT/pY). Sites in peptides that could not be unambiguously identified are in small letters (s/t/y). Spectra were recorded either on a Q-ToF MS (Waters) and/or an Orbitrap MS (ThermoFisherScientific). The precursor in MS showed a mass accuracy of \leq 15 ppm or \leq 2 ppm, respectively. Y-type and b-typ ions are listed within the spectra and the given sequence. Subscribed numbers indicate the number of heavy isotopes in Lys- and Arg-residues within the peptide.

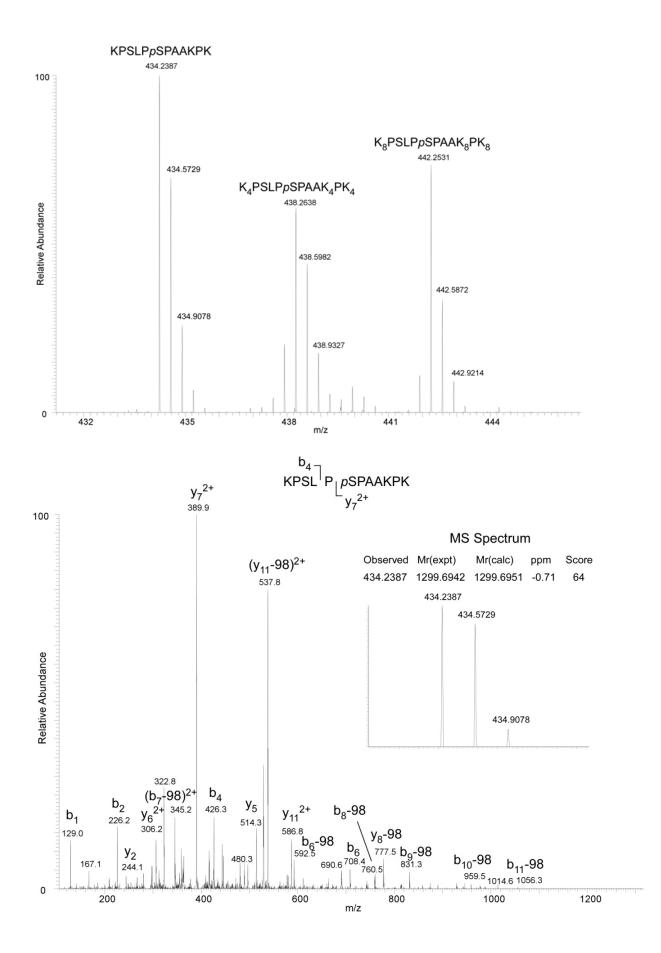




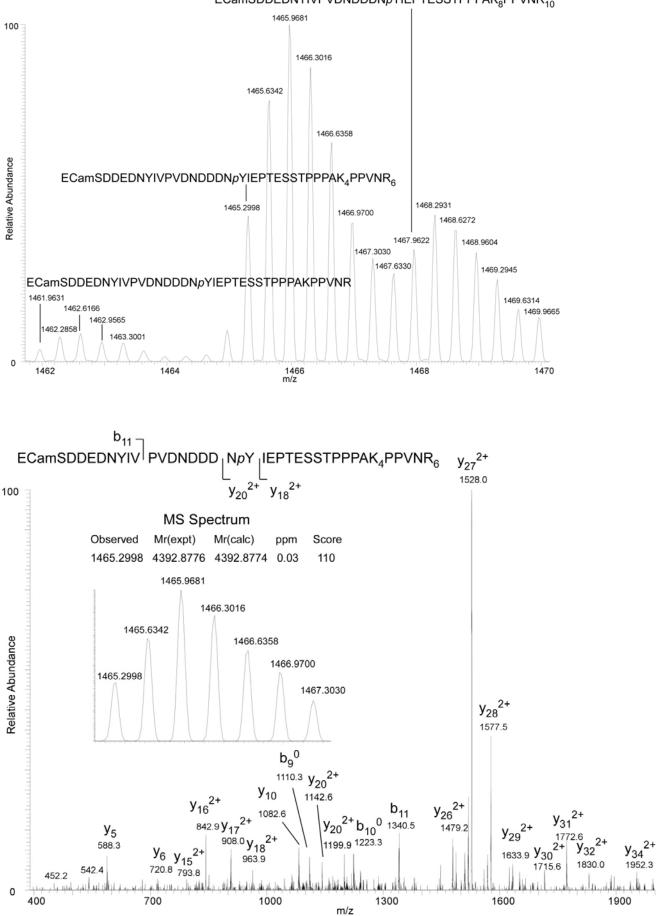


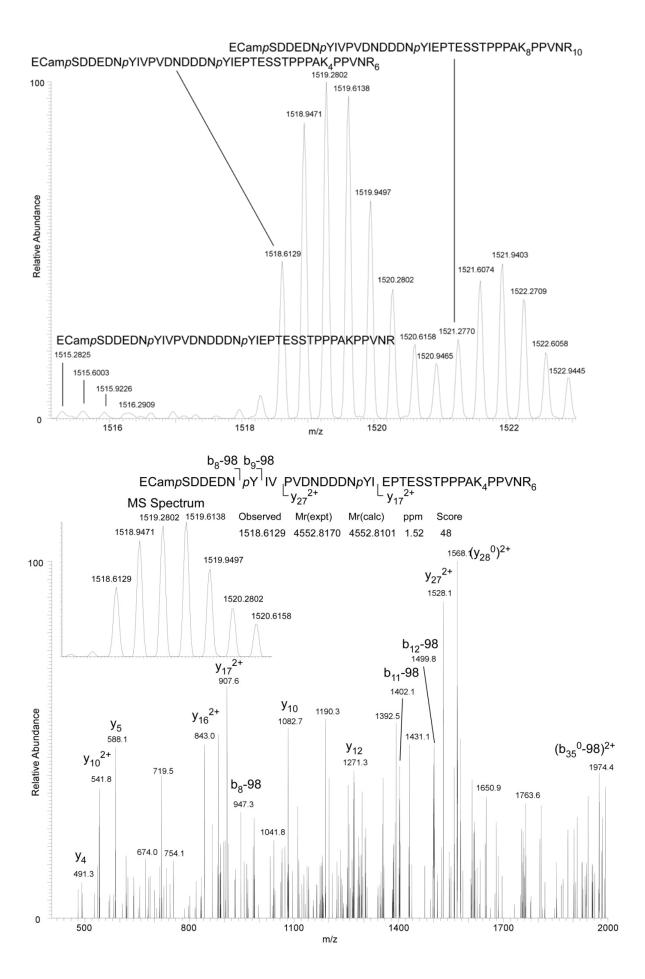


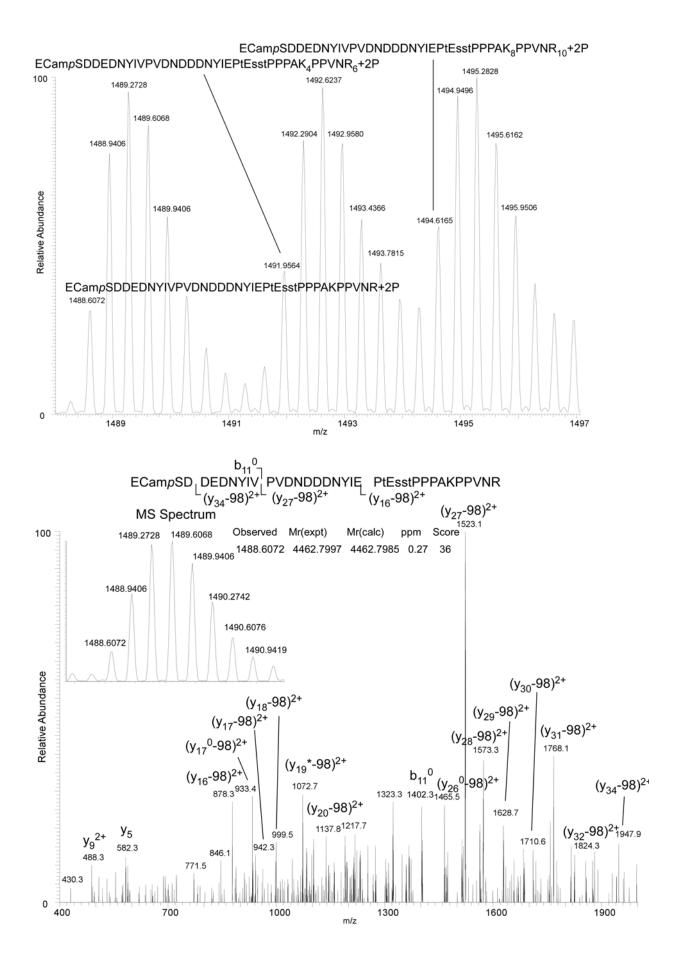


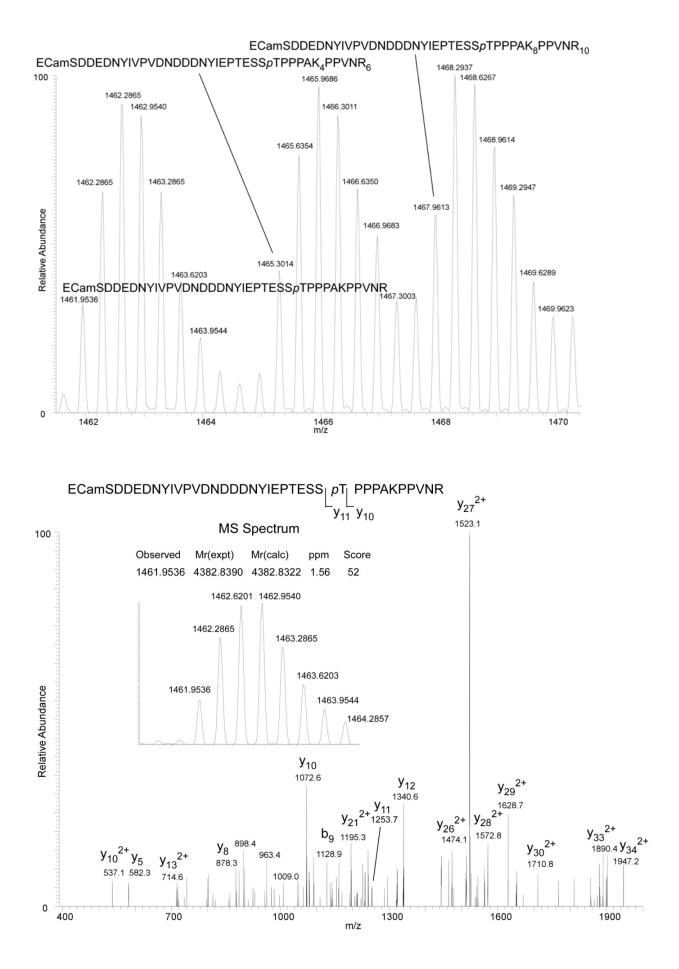


ECamSDDEDNYIVPVDNDDDNpYIEPTESSTPPPAK8PPVNR10

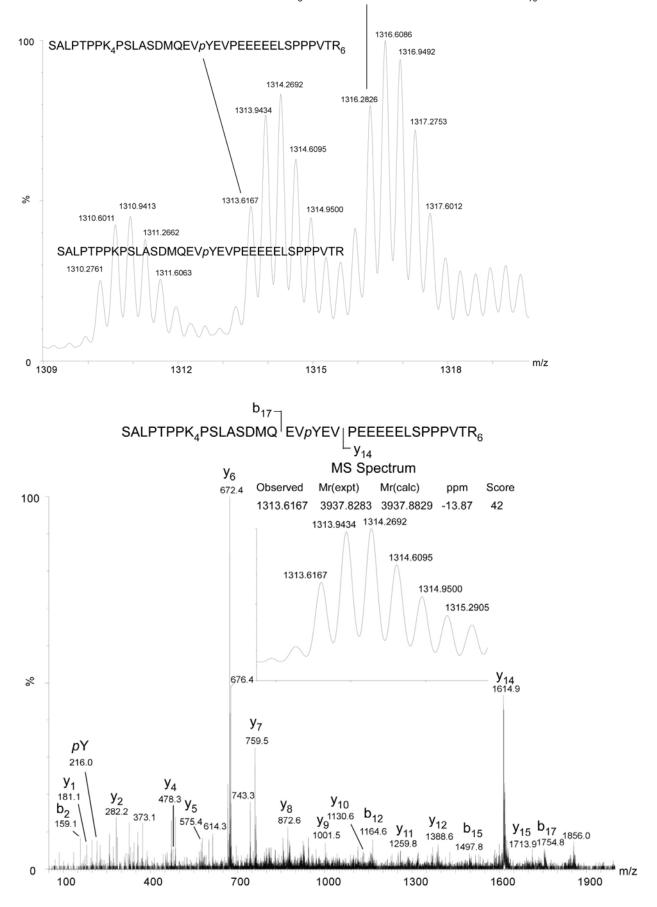


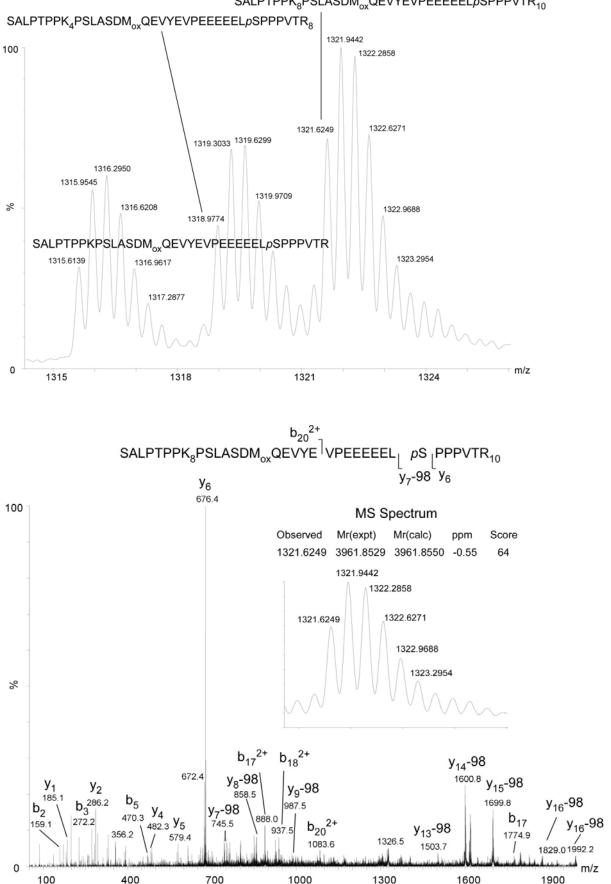






SALPTPPK₈PSLASDMQEVpYEVPEEEEELSPPPVTR₁₀





SALPTPPK₈PSLASDM_{ox}QEVYEVPEEEEELpSPPPVTR₁₀

SALPpTPPKPSLASDM_{ox}QEVYEVPEEEEELpSPPPVTR

