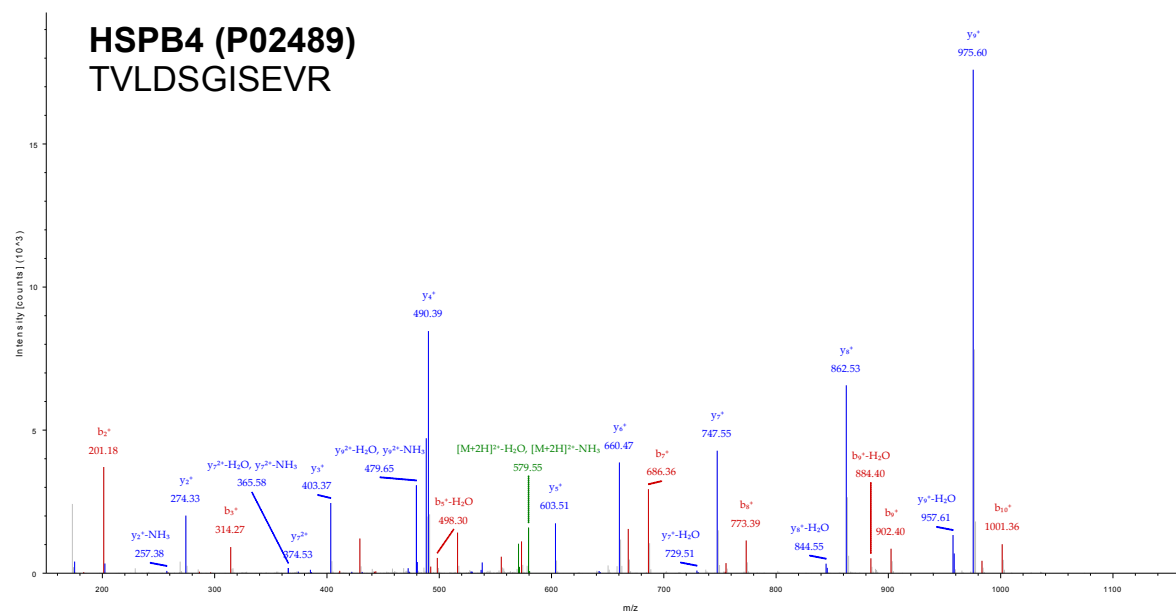
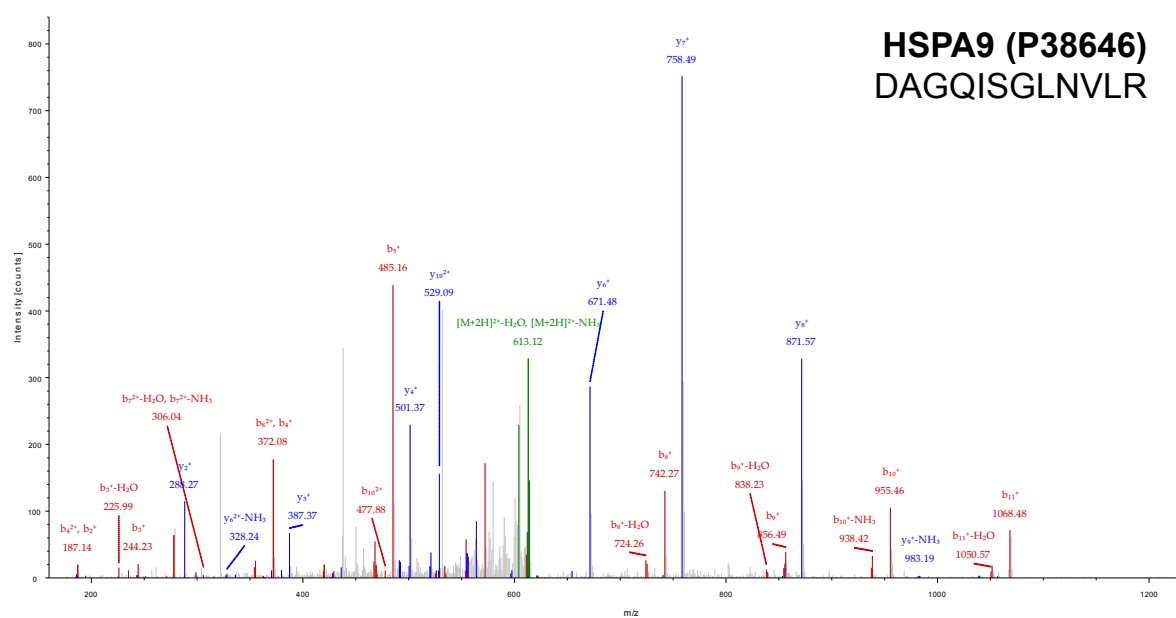
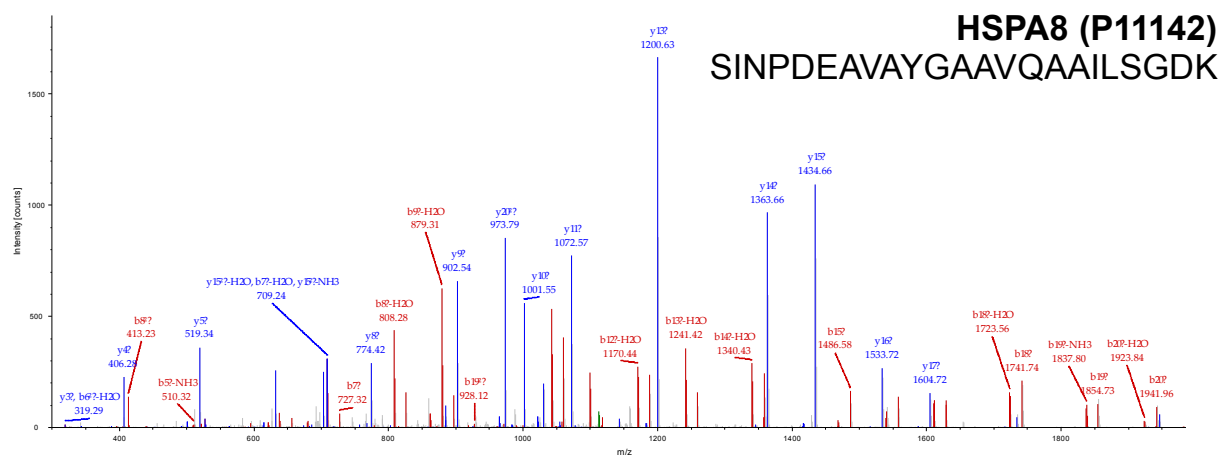
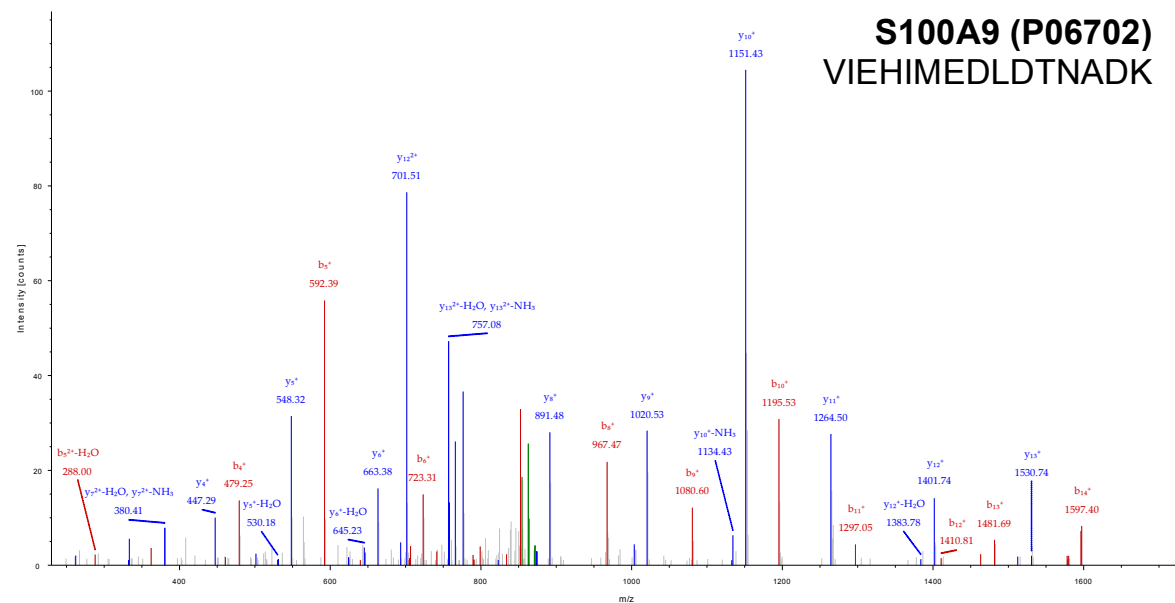
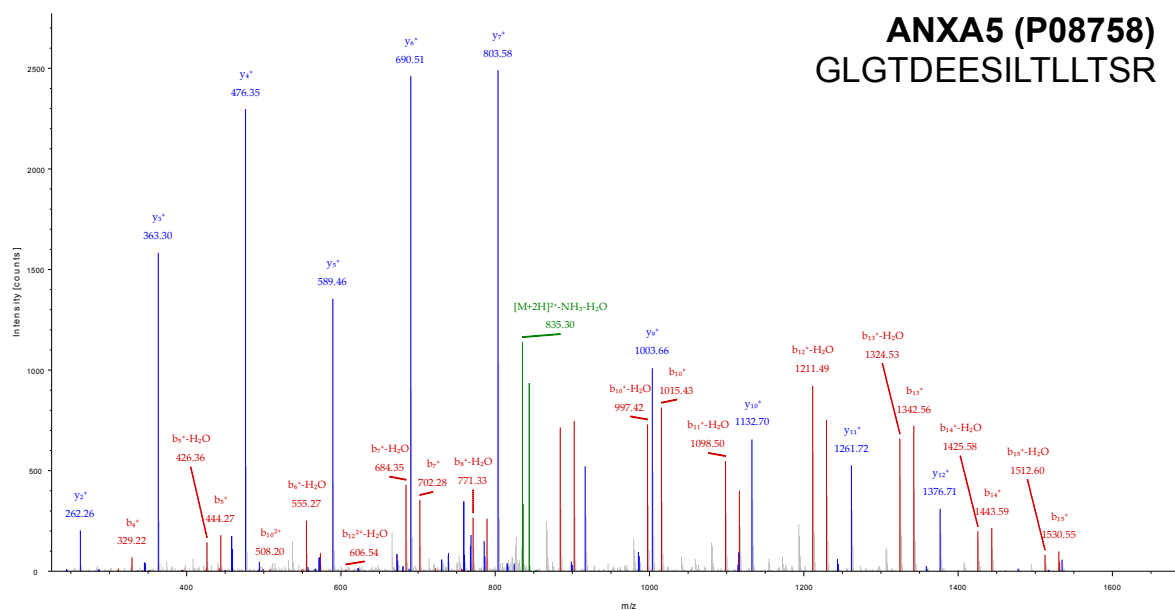






S3 Fig. Representative MS/MS spectra for the peptides matched to proteins identified in this study (see Table 1). The spectra were produced via collision-induced dissociation (CID) of the corresponding mass-selected precursor ions in the ion trap mass spectrometer. The MS/MS spectra contain sequence-determining product ions of the b- (shown in red) and y-series (shown in blue).