S7 Fig. Hexosylation of the peptide ADLESLISSKEK of MYPU_3200. Orbitrap MS1 showing the doubly and triply charged ions. The 81.0268 shift for $z = 2$ between the non-glycosylated and hexose peptides equates to a mass shift of 162.0536 Da with a mass accuracy of 0.0008 Da. The 54.0170 shift for $z = 3$ between non-glycosylated and hexose forms equates to a mass shift of 162.0510 Da with a mass accuracy of 0.0018 Da. The theoretical and experimental calculated values for $m/z$ are given in bold. The images presented were obtained from an LC peak of MS scans and are expanded to show the charge states of each form.