Figure S5. Upper spectra illustrate the precursor mass scan at 28.6-28.9 min of the sample from non-stimulated Jurkat cells found in the IMAC un-bound fractions. For illustration purposes, only the m/z 400-750 region is shown. Blue inset highlights the m/z 414.35 ion. MS2 scan of this ion identifies it as the 161-170 unphosphorylated RNSVRLVIRK peptide. Black lines indicate peptide cleavage.