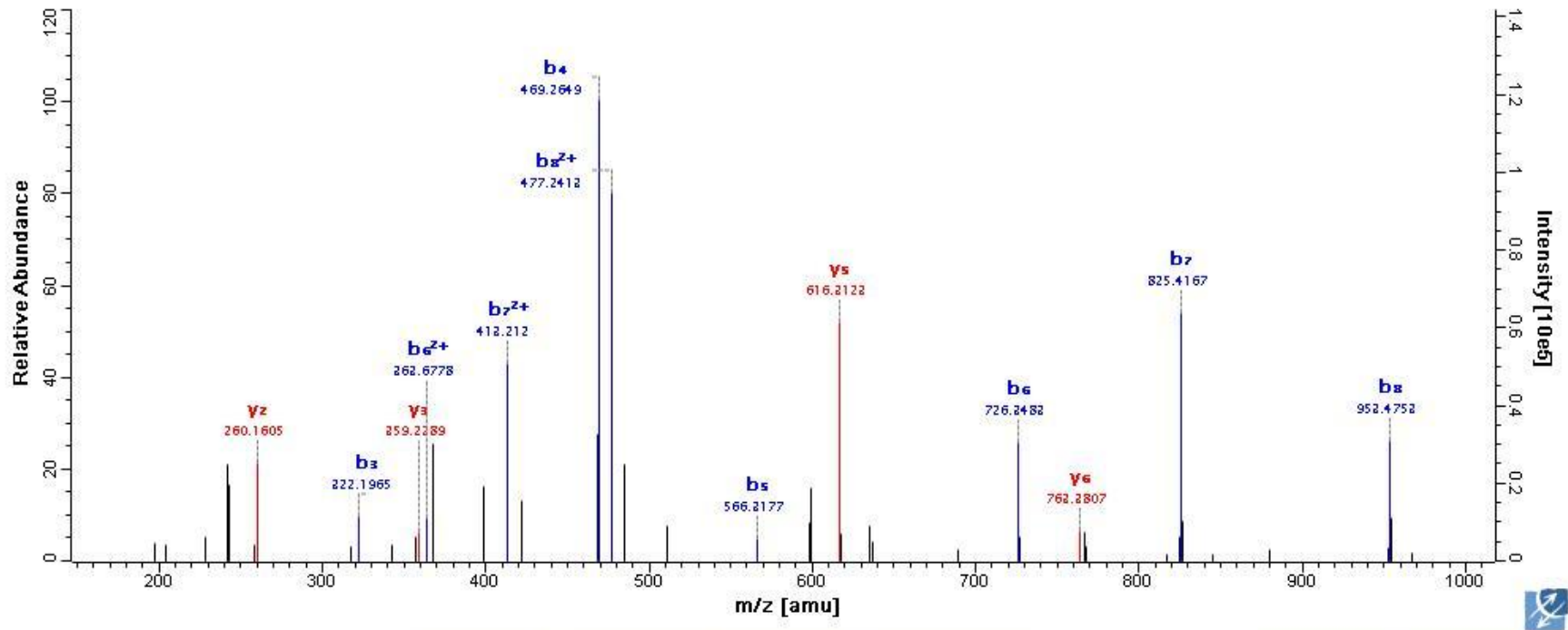
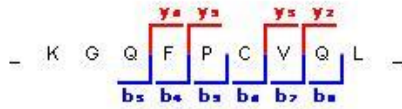


Source: 20121207_CO_0375ViVi_Exp1_Mix2_R01R04R05_GeLC_24
 Scannumber: 19010
 Protein: PB2delta
 Peptide Score: 100.15
 Method: ITMS; CID; 3



S2 Fig. MS/MS spectrum of the tryptic C-terminal unique peptide KGQFPCVQL of the H5N1 KAN-1 virus protein PB2_Δ. 100 μg of protein mixtures from KAN-1-infected A549 cells were separated on a one-dimensional gel followed by in-gel digest by trypsin. LC-MS/MS analyses were performed on an EasyLC nano-HPLC (Proxeon Biosystems) coupled to an LTQ Orbitrap Elite mass spectrometer (Thermo Scientific). The MS data were processed using default parameters of the MaxQuant software (v1.2.2.9).