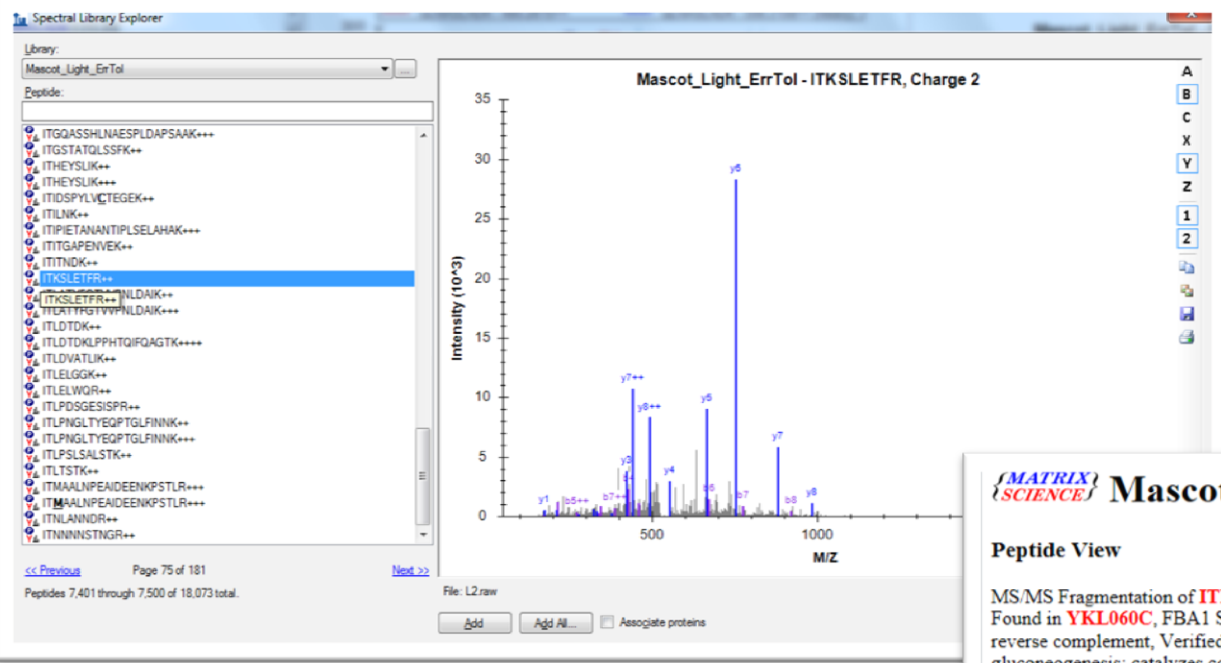




Mascot Search Results

Peptide View

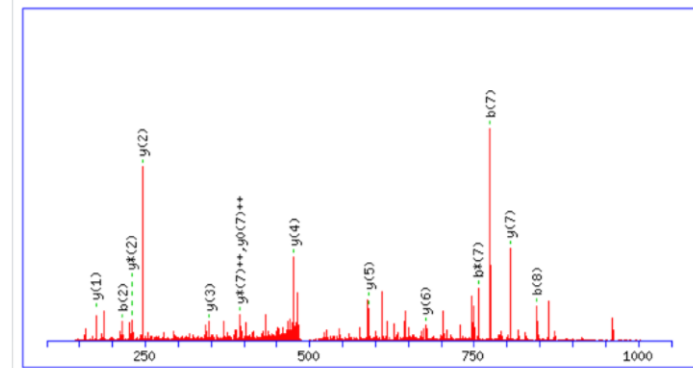
MS/MS Fragmentation of **ITKSLETFR**
 Found in **YKL060C**, FBA1 SGDID:S000001543, Chr XI from 327487-326408, Genome Release 64-1-1, reverse complement, Verified ORF, "Fructose 1,6-bisphosphate aldolase, required for glycolysis and gluconeogenesis; catalyzes conversion of fructose 1,6 biphosphate to glycera

Match to Query 8784: 1017.581152 from(509.797852,2+) intensity(204502.0000)
 Title: L2.12425.2 File:"L2.raw", NativeID:"controllerType=0 controllerNumber=1 scan=12425"
 Data file L:\Salvo_25\UnbiasedPTM_newhopes\2_LongColumn\2_DB_search\2_MGF\L2.mgf

Click mouse within plot area to zoom in by factor of two about that point

Or, Plot from 100 to 1050 Da Full range

Label all possible matches Label matches used for scoring



Monoisotopic mass of neutral peptide Mr(calc): 1017.5819
 Fixed modifications: Carbamidomethyl (C) (apply to specified residues or termini only)
 Variable modifications:
 F8 : Phe->Ala (F)
 Ions Score: 33
 Matches : 14/86 fragment ions using 36 most intense peaks (help)