Comparison of peptide score distributions 2012_PGNEXUS_proteome_con.sp.canonical.isoforms_katana 2012_PGNEXUS_proteome_junc.exons_dragon 7500 20000 5000 10000 2500 Frequency (absolute) Peptide Hit Type Swissprot hits 150 20 20 100 Exons from 3FT Junctions from 3FT 2020_phosphoproteome_OBseries_con_sp.hsa.canonical.isoforms 2020_phosphoproteome_OBseries_junc.exons Contaminant Reversed sequence 150000 3e+05 100000 2e+05 50000 1e+05 0e+00 ·

9

100

50

100

150

250 300 350 400 450 550 600

Score_{MaxQuant} + 1