Comparison of peptide score distributions 2012_PGNEXUS_proteome_con.sp.canonical.isoforms_katana 2012_PGNEXUS_proteome_junc.exons_dragon 10000 -7500 -20000 5000 -10000 2500 -Frequency (absolute) Peptide Hit Type Swissprot hits 200 250 330 350 550 100 150 100 150 20 20 Exons from 3FT Junctions from 3FT 2020_phosphoproteome_OBseries_con_sp.hsa.canonical.isoforms 2020_phosphoproteome_OBseries_junc.exons Contaminant Reversed sequence 3e+05 1e+05 -2e+05 5e+04 · 1e+05

0e+00 **-**

Score_{MaxQuant} + 1

9

100

250 300 350 400 450 550

20

100

0e+00 -