## Comparison of peptide score distributions 2012\_PGNEXUS\_proteome\_con.sp.canonical.isoforms\_katana 2012\_PGNEXUS\_proteome\_junc.exons\_katana 2012\_PGNEXUS\_proteome\_sp.canonical.isoforms\_junc.exons\_katana 25000 20000 900 20000 15000 600 10000 10000 300 5000 Frequency (absolute) Peptide Hit Type Swissprot hits 051 2 200 050 250 000 000 000 000 000 000 20 100 100 20 Exons from 3FT Junctions from 3FT 2020\_phosphoproteome\_OBseries\_con\_sp.hsa.canonical.isoforms 2020\_phosphoproteome\_OBseries\_junc.exons Contaminant Reversed sequence 15000 3e+05 10000 2e+05 5000 1e+05 0e+00 -150 200 250 300 450 9 100 $Score_{MaxQuant} + 1$