IncRNA group = IDRIessthan0_2 ::: data cols = bothPullDowns (53 / 3463) = (func/total IncRNA) ::: biotype = remove_antisense subset = 53 IncRNAs out of 3463 total IncRNAs

average features from Derrien 2012 datasheet LncRNA_tx_sizephastcons_primate_introdnastcons_primate_transcript_score No Exons 600 600 10000 -20 -400 400 5000 10 -200 -200 0 0 false false false true false true true true phastcons primate promoterastcones mammal transplinations mammal inploastscons mammal promoter score 800 800 750 600 600 600 withinSubset value value 500 400 false 400 true 250 200 200 200 0 0 false true false true false true false true phastcons_vertebrate_transcriptastcons_vertebrate_iphastcons_vertebrate_promoter_score 800 800 -750 600 600 500 400 400 250 200 200 -0 0 false false false true true true

value of facet label