IncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns (30 / 1701) = (func/total IncRNA) ::: biotype = antisense subset = 30 IncRNAs out of 1701 total IncRNAs

average features from Derrien 2012 datasheet LncRNA_tx_size phastcons_primate_intrphastcons_primate_transcript_score No Exons 6000 15 600 600 4000 10 400 400 2000 200 200 -5 0 false false false true true false true true phastcons primate promoterastcons mammal transcorpastcons mammal introductions mammal promoter score 800 800 600 600 600 600 withinSubset 400 false 400 400 true 200 -200 200 200 0 0 0 false true false true false true false true phastcons_vertebrate_trans@rijatstscns_vertebrate_iptrastcscns_vertebrate_promoter_score 800 750 600 600 400 500 400 200 250 200 0 0 false false true true false true value of facet label