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 6 · 5.0 8 2 4 2.5 7 -2 1 0.0 6 0 -2.55 0 true false true false true false false true phastcons_primate_promotestcoms_mammal_transcriptiastcons_mammal_introbastcons_mammal_promoter_score 5.0 log(value) 0.0 5.0 5.0 -5.0 withinSubset 2.5 false 2.5 2.5 true 0.0 0.0 0.0 --2.5 false true false true false true false true phastcons_vertebrate_transbristcons_vertebrate_inprbastscors_vertebrate_promoter_score 5.0 5.0 5.0 2.5 2.5 2.5 -0.0 0.0 0.0 false true false true false true

value of facet label