IncRNA group = IDRlessthan0_1 ::: data cols = Ipa
(42 / 1356) = (func/total IncRNA) ::: biotype = lincProcTrans PageRank, full range 0 80 60 -40 20 factor(label) 0 0 15 -10 -5 -

1.0

rank

1.5

count

0 -

0.0

0.5