lncRNA group = IDRlessthan0_1 ::: data cols = lpa (69 / 2581) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, full range 0 100 50 factor(label) 0 0 1 20 15 10 5 0 -

1.0

rank

1.5

0.5

0.0

count