IncRNA group = allLncRNA ::: data cols = Inpa (88 / 6668) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, full range 1000 -750 -500 250 factor(label) 0 0 40 30 20 10 0 0.5 1.0 1.5 0.0

rank

count