IncRNA group = allLncRNA ::: data cols = lpa (87 / 5951) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, full range 0 1200 800 400 factor(label) 0 0 50 -40 -30 -20 10 0 1.5 0.0 1.0 0.5

rank

count