IncRNA group = allLncRNA ::: data cols = lpa (50 / 3553) = (func/total lncRNA) ::: biotype = remove_antisense PageRank, full range 0 800 -600 -400 200 factor(label) 0 count 0 1 30 20 10 -0 0.0 0.5 1.5 1.0

rank