lncRNA group = IDRlessthan0_1 ::: data cols = lpa (69 / 2848) = (func/total lncRNA) ::: biotype = all_biotypes PageRank, full range 0 150 100 50 factor(label) 0 0 1 20 -15 -10 5 0 -0.5 1.5 0.0 1.0

rank

count