lncRNA group = allLncRNA ::: data cols = bothPullDowns (92 / 7622) = (func/total IncRNA) ::: biotype = all_biotypes PageRank, range = (-0.1, 1.1)10.0 -7.5 factor(label) 5.0 2.5 0.0 0.0 0.4 8.0

rank

density