IncRNA group = allLncRNA ::: data cols = lpa (91 / 7106) = (func/total lncRNA) ::: biotype = all\_biotypes PageRank, full range 1500 1000 -500 factor(label) 0 0 50 -40 30 20 10 0 0.5 1.5 1.0 0.0 rank

count