IncRNA group = allLncRNA ::: data cols = Inpa (52 / 4449) = (func/total lncRNA) ::: biotype = remove\_antisense PageRank, full range 0 600 -400 -200 factor(label) 0 count 0 1 30 20 10 -0 0.0 1.0 1.5 0.5

rank