IncRNA group = allLncRNA ::: data cols = Inpa (86 / 5711) = (func/total lncRNA) ::: biotype = all\_biotypes PageRank, full range 0 800 600 400 200 factor(label) 0 0 1 40 30 20 10 0 1.5

1.0

rank

count

0.0

0.5