IncRNA group = allLncRNA ::: data cols = lpa (52 / 3884) = (func/total lncRNA) ::: biotype = remove_antisense PageRank, full range 0 750 · 500 250 factor(label) 0 count 0 1 30 20 10 -

1.0

rank

1.5

0

0.0

0.5