lncRNA group = IDRlessthan0\_1 ::: data cols = Inpa (16 / 231) = (func/total IncRNA) ::: biotype = processed\_transcript PageRank, full range 12.5 10.0 7.5 5.0 2.5 factor(label) 0.0 count 0 1 6 -4 2 0 0.5 0.0 1.0 1.5

rank