lncRNA group = IDRlessthan0\_1 ::: data cols = lpa ( 42 / 1356 ) = (func/total lncRNA) ::: biotype = lincProcTrans PageRank, range = median +/- 2 ds 0 60 -40 -20 factor(label) 0 0 15 -10 -5 -0 -0.5 1.0 0.0 1.5

rank

count