lncRNA group = IDRlessthan0_01 ::: data cols = lpa (10/37) = (func/total lncRNA) ::: biotype = processed_transcript PageRank, range = (-0.1, 1.1)2.5 2.0 -1.5 factor(label) density 1.0 0.5 -0.0 -0.4 0.0 8.0

rank