lncRNA group = IDRlessthan0_01 ::: data cols = lpa (11/109) = (func/total IncRNA) ::: biotype = lincRNAPageRank, range = (-0.1, 1.1)factor(label) 0.0 0.4 8.0

rank

2.0

1.5 -

1.0

0.5 -

0.0

density