lncRNA group = IDRlessthan0_1 ::: data cols = Inpa (23/935) = (func/total IncRNA) ::: biotype = lincRNAPageRank, range = (-0.1, 1.1)factor(label) 0.0 0.4 8.0 rank

3 ·

2 ·

1 ·

0

density