lncRNA group = IDRlessthan0_2 ::: data cols = lpa (30/2025) = (func/total IncRNA) ::: biotype = lincRNAPageRank, range = (-0.1, 1.1)factor(label) 0.0 0.4 8.0

rank

5

4

3 -

2 ·

1 ·

0

density