IncRNA group = IDRlessthan0\_01 ::: data cols = lpa ( 11 / 109 ) = (func/total IncRNA) ::: biotype = lincRNA logRank, range = median +/- 2 ds factor(label) 0 1 1.00 -0.75 -

0.6

logRank

0.8

0.4

6

4

2 ·

0

0.50

0.25

0.00

0.2

count