IncRNA group = allLncRNA ::: data cols = Inpa (36 / 2293) = (func/total IncRNA) ::: biotype = antisense logRank, range = median +/- 2 ds 2.5 2.0 -1.5 factor(label) 1.0 0.5 -0.0 -0.00 0.25 0.50 0.75

logRank

density