lncRNA group = IDRlessthan0_1 ::: data cols = lpa (69 / 2581) = (func/total lncRNA) ::: biotype = all_biotypes logRank, range = median +/- 2 ds 1.5 -1.0 factor(label) 0.5 0.0 -0.2 0.4 0.6 8.0 1.0

logRank

density