PCA of IncRNA expr :
no r1 group IncRNA group = IDRlessthan0_1 ::: data cols = bothPullDowns (16 / 232) = (func/total IncRNA) ::: biotype = processed_transcript 1.0 count 100 75 50 25 0.5 0.0 -0.5 0.0 1.0

log(Comp.2)

log(Comp.3)