PCA of IncRNA expression data all Inc RNAs IncRNA group = allLncRNA ::: data cols = Inpa (17 / 372) = (func/total IncRNA) ::: biotype = processed_transcript 2 · withinSubset rest true 0 · 0.0 0.5 1.0 1.5 2.0 log(First Component)

log(Second Component)