PCA of IncRNA expression data all Inc RNAs IncRNA group = allLncRNA ::: data cols = Inpa (23 / 962) = (func/total IncRNA) ::: biotype = processed\_transcript 130600 2.5 2.0 log(Second Component) withinSubset rest true 175061 0.5 226950 0.0 2.0 0.0 0.5 1.0 1.5 log(First Component)