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