PCA of IncRNA expression data all Inc RNAs IncRNA group = allLncRNA ::: data cols = Inpa (23 / 962) = (func/total IncRNA) ::: biotype = processed_transcript 2.0 log(Third Component) withinSubset rest true 0.5 -0.0 -2.0 0.0 0.5 1.0 1.5 log(First Component)