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 0.5 0.0 1.0 2.0 2.5 log(Second Component)