PCA of IncRNA expression data all Inc RNAs IncRNA group = allLncRNA ::: data cols = Inpa ( 17 / 372 ) = (func/total IncRNA) ::: biotype = processed\_transcript 1.5 log(Third Component) withinSubset rest true 0.5 -0.0 -0.5 0.0 1.0 1.5 2.0 log(First Component)