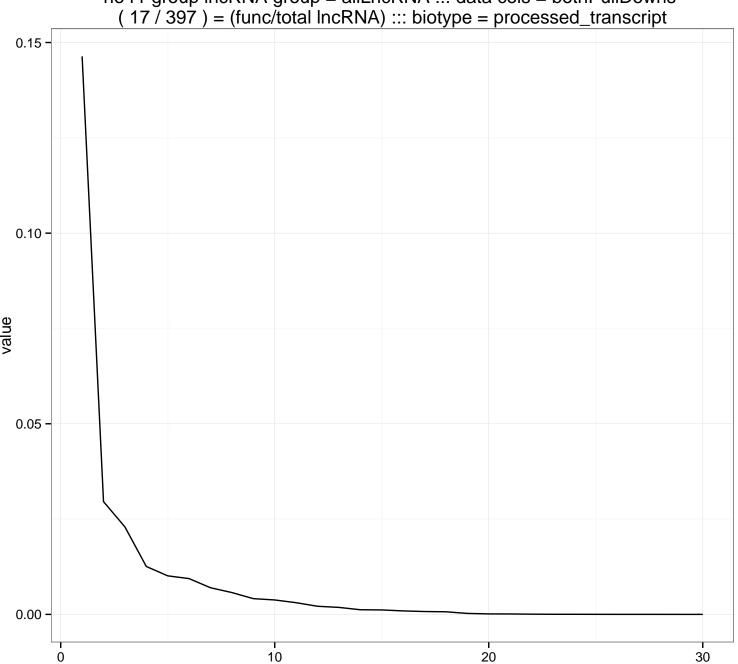
PCA IncRNA expressionmatrix

Component variation over all components
no r1 group lncRNA group = allLncRNA ::: data cols = bothPullDowns
(17 / 397) = (func/total lncRNA) ::: biotype = processed_transcript



component