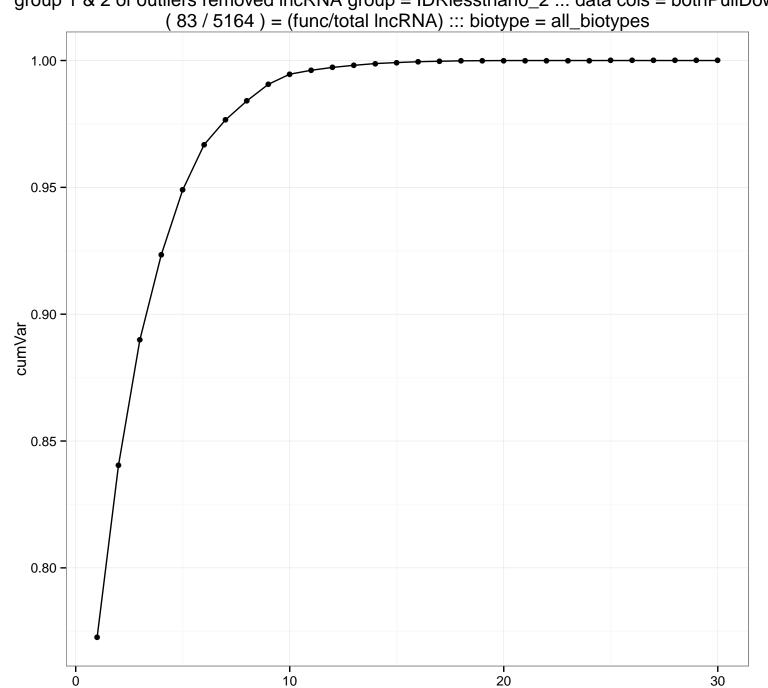variance of PCA components
group 1 & 2 of outliers removed lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDow
(83 / 5164) = (func/total lncRNA) ::: biotype = all_biotypes