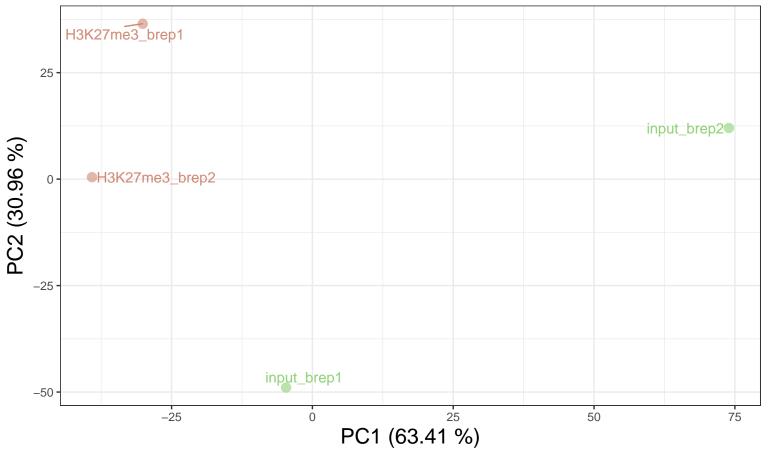
## PCA based on counts.all.genome.bw10000.tab

Unit variance scaling per sample



Condition - ChIP - input