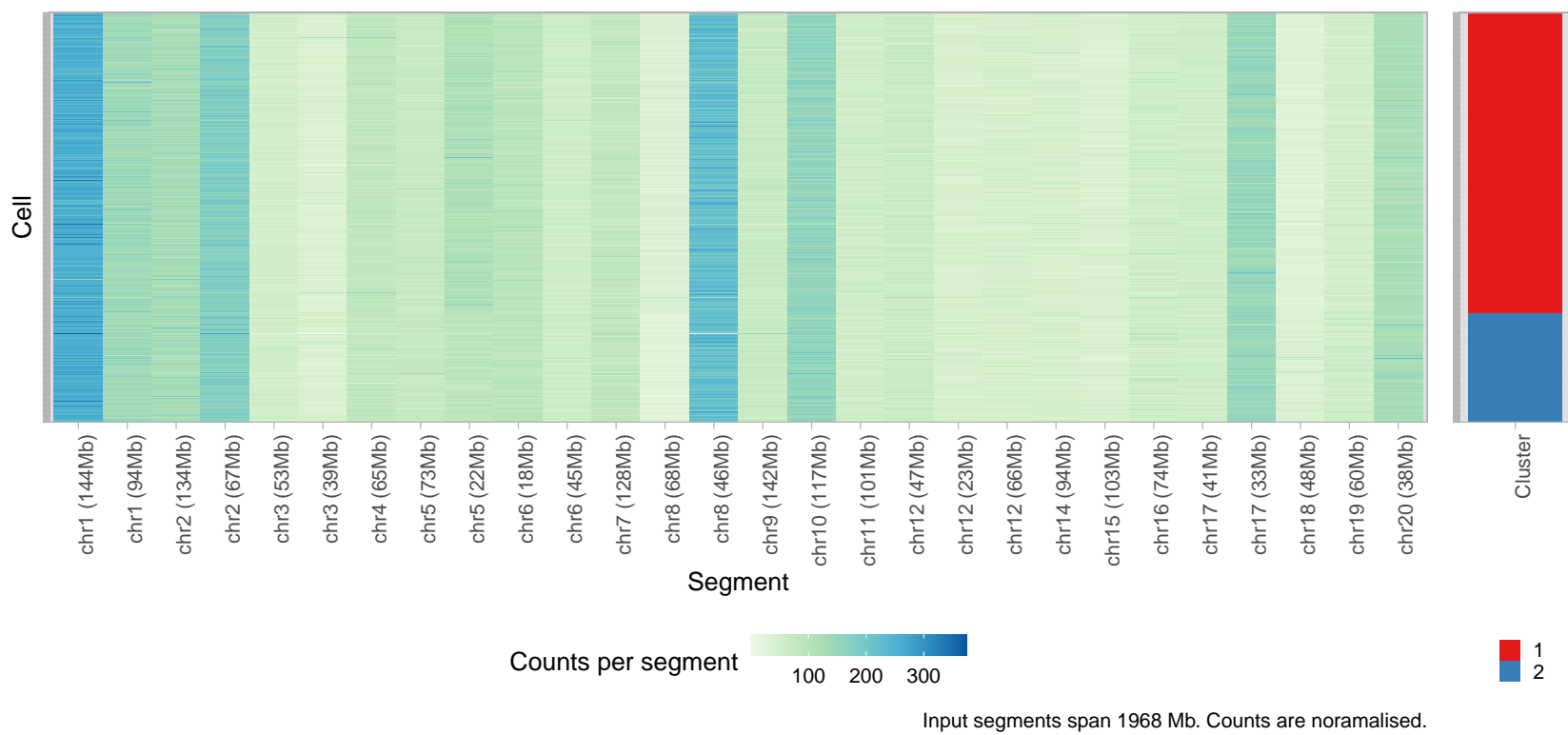
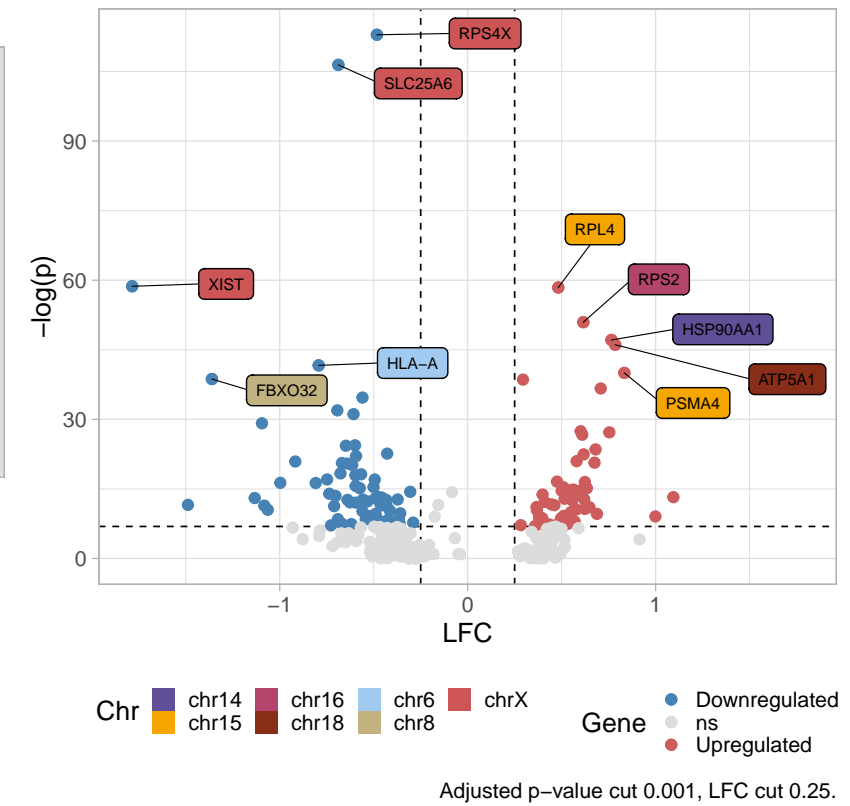


B Raw CONGAS dataset (k =2)
2153 genes mapped to 28 segments.



C DE (1 vs 2 with DESeq2)



D Counts per chromosome

