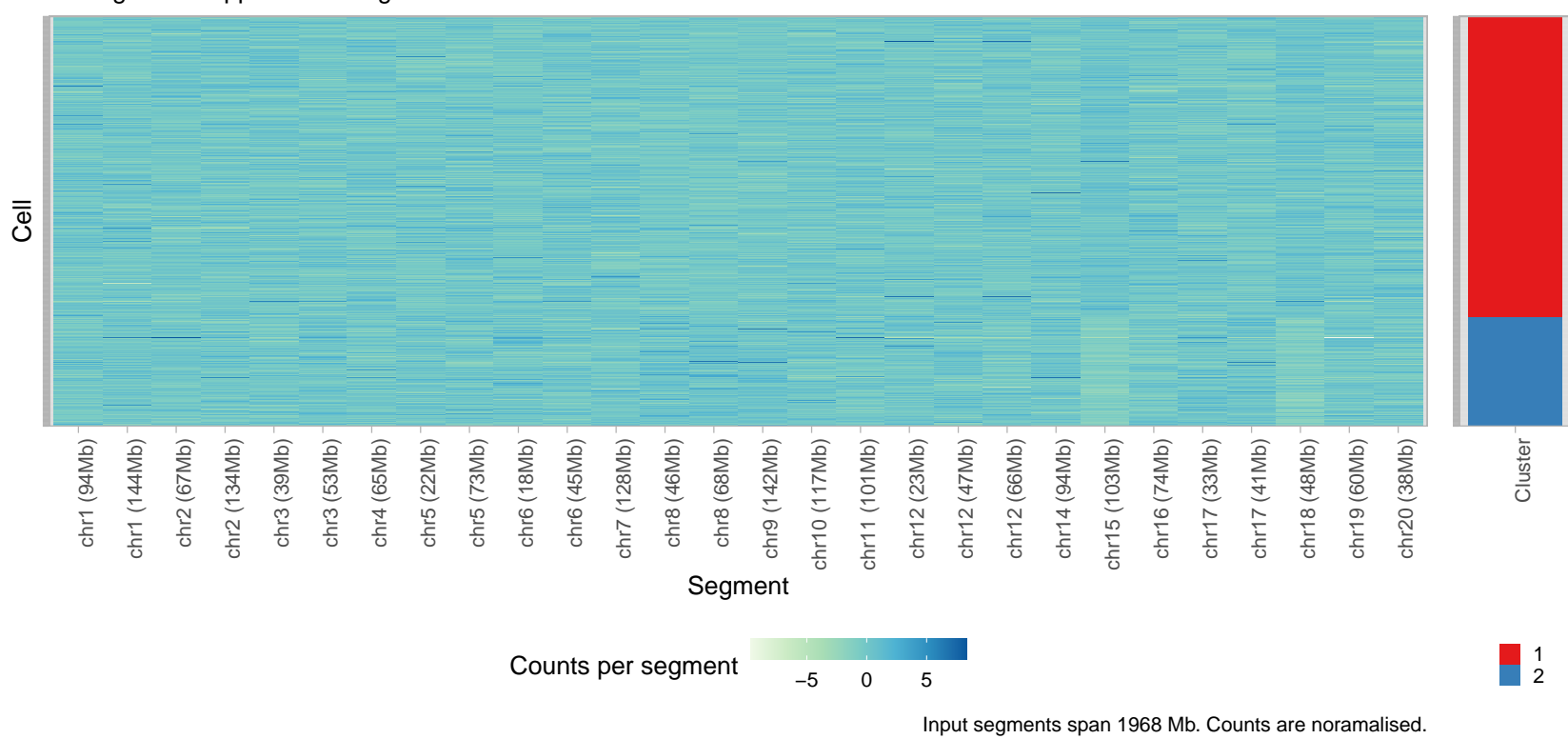
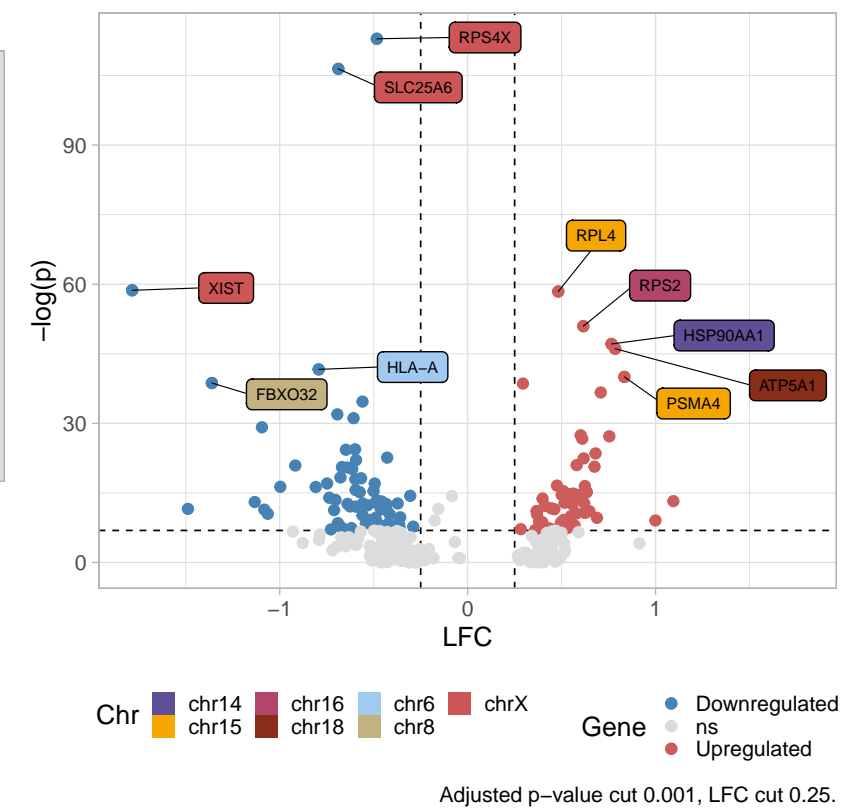


**B** Raw CONGAS dataset (k =2)  
2153 genes mapped to 28 segments.



**C** DE (1 vs 2 with DESeq2)  
Cutoffs: p-value  $p < 0.001$  LFC  $< 0.25$ .



**D** Counts per chromosome

