

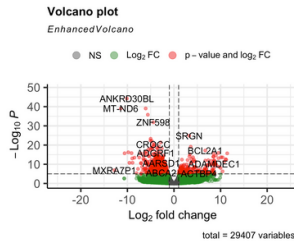
Sorteren en
indexeren BAM
bestanden



Count matrix
opstellen per gen



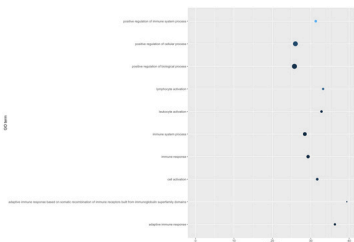
Bekijken welke genen
veranderd zijn



Volcano plot maken



GO-analyse



KEGG pathway (hsa04660)

