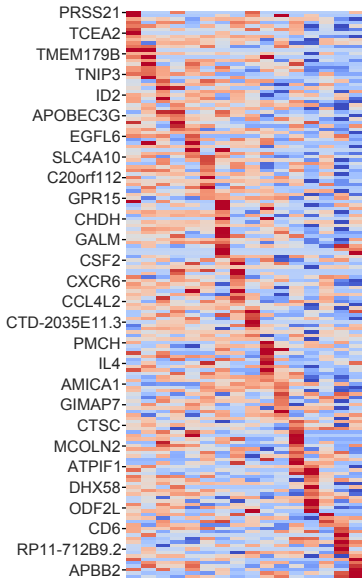
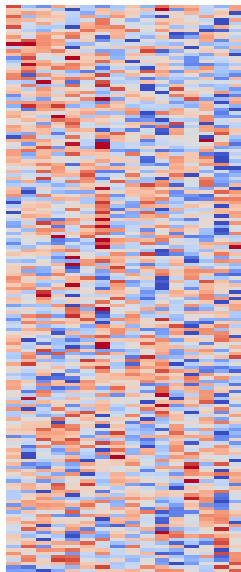


165 clonal ( $p < 0.01$ ) genes.



16 clones

Peaks near each genes  
height aggregated by gene.



16 clones