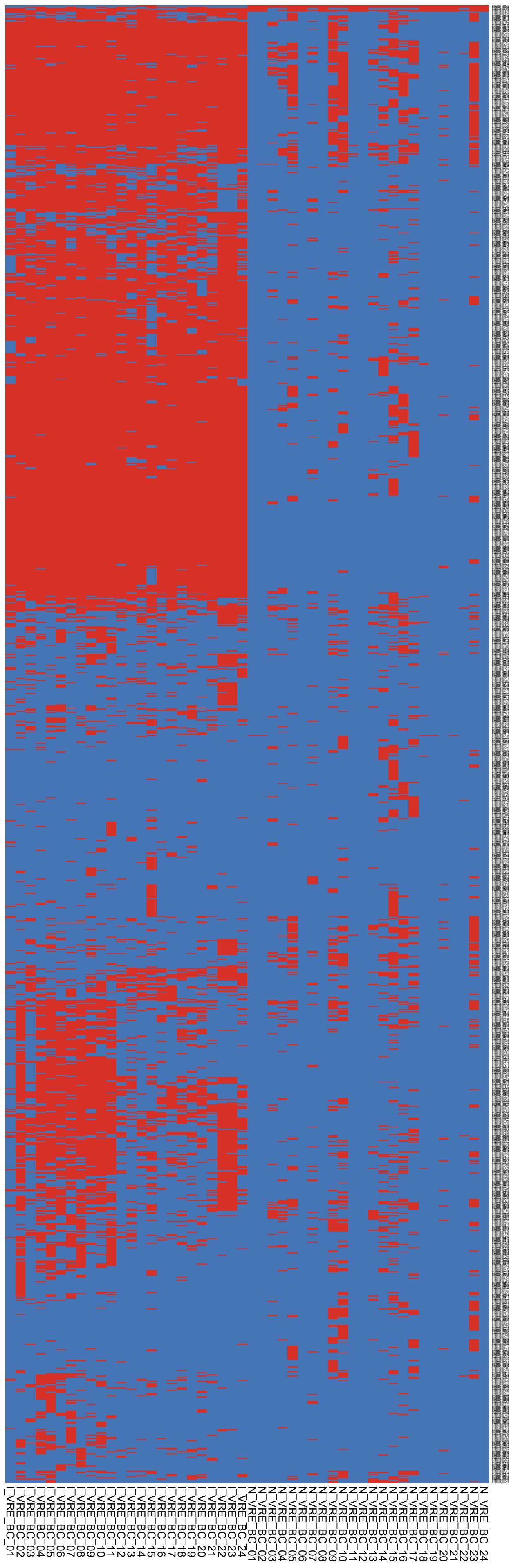




Genes with coverage that differs $\pm 25\%$ from the mean (VRE, Hybrid)



Supplementary Figure 6: Heatmaps of eye-catching and potentially due to different filters excluded genes in the hybrid validation experiment 3 part 1 on VRE data. Shown are genes with more than 50% of heterozygous variant-calls, genes with below 55 mapping quality and genes with a coverage that differs more than 25% from the mean.