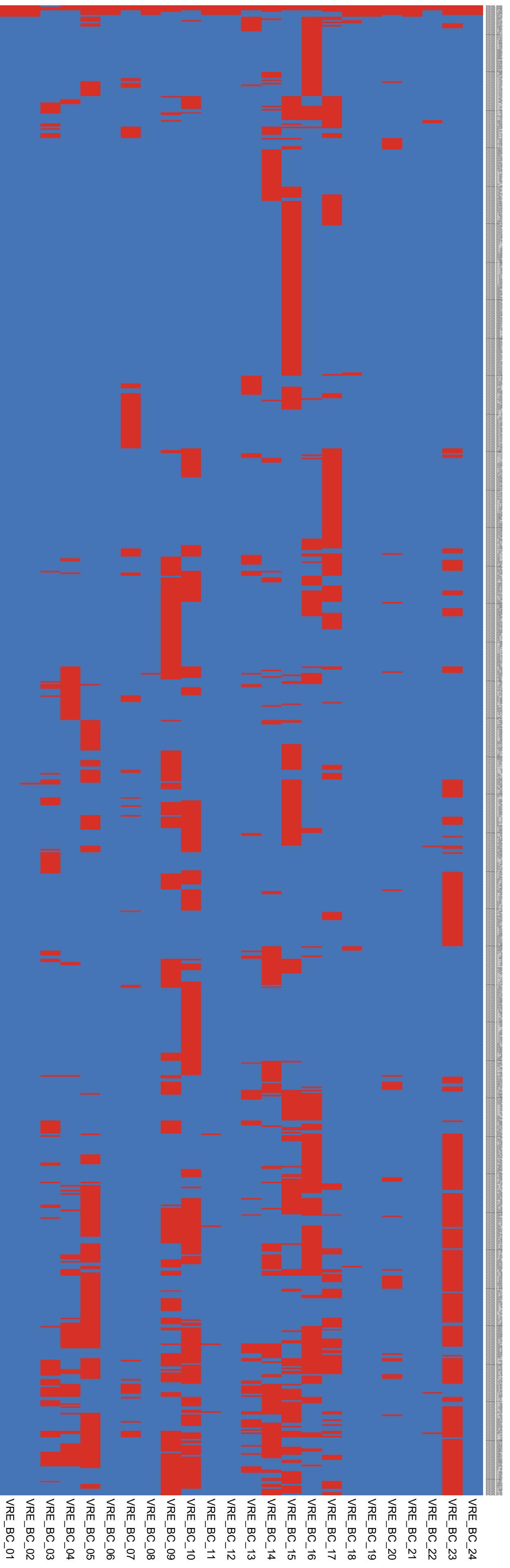


1
0.8
0.6
0.4
0.2
0
Genes with coverage that differs $\pm 25\%$ from the mean (VRE, Nanopore-only)



Supplementary Figure 4: Heatmaps of eye-catching and potentially due to different filters excluded genes in the Nanopore-only validation experiment 2 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.