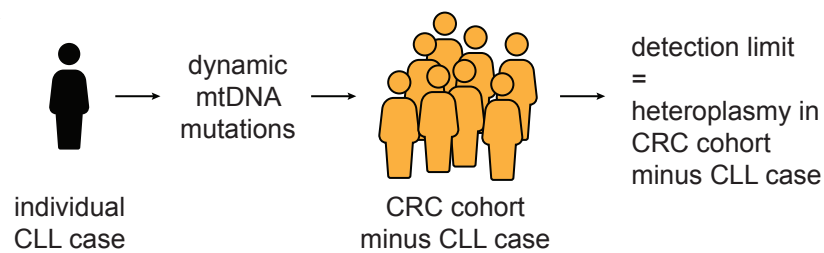
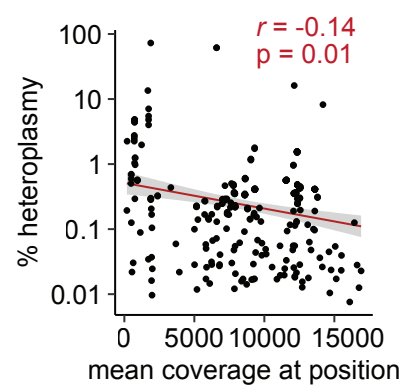
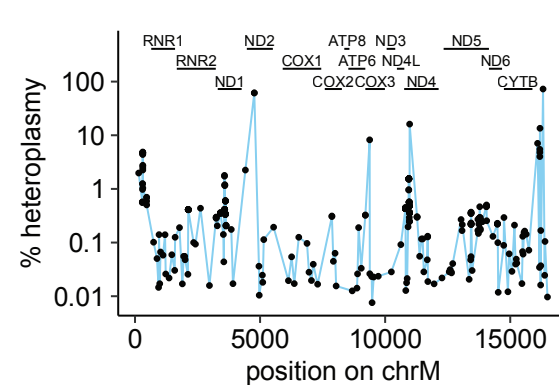
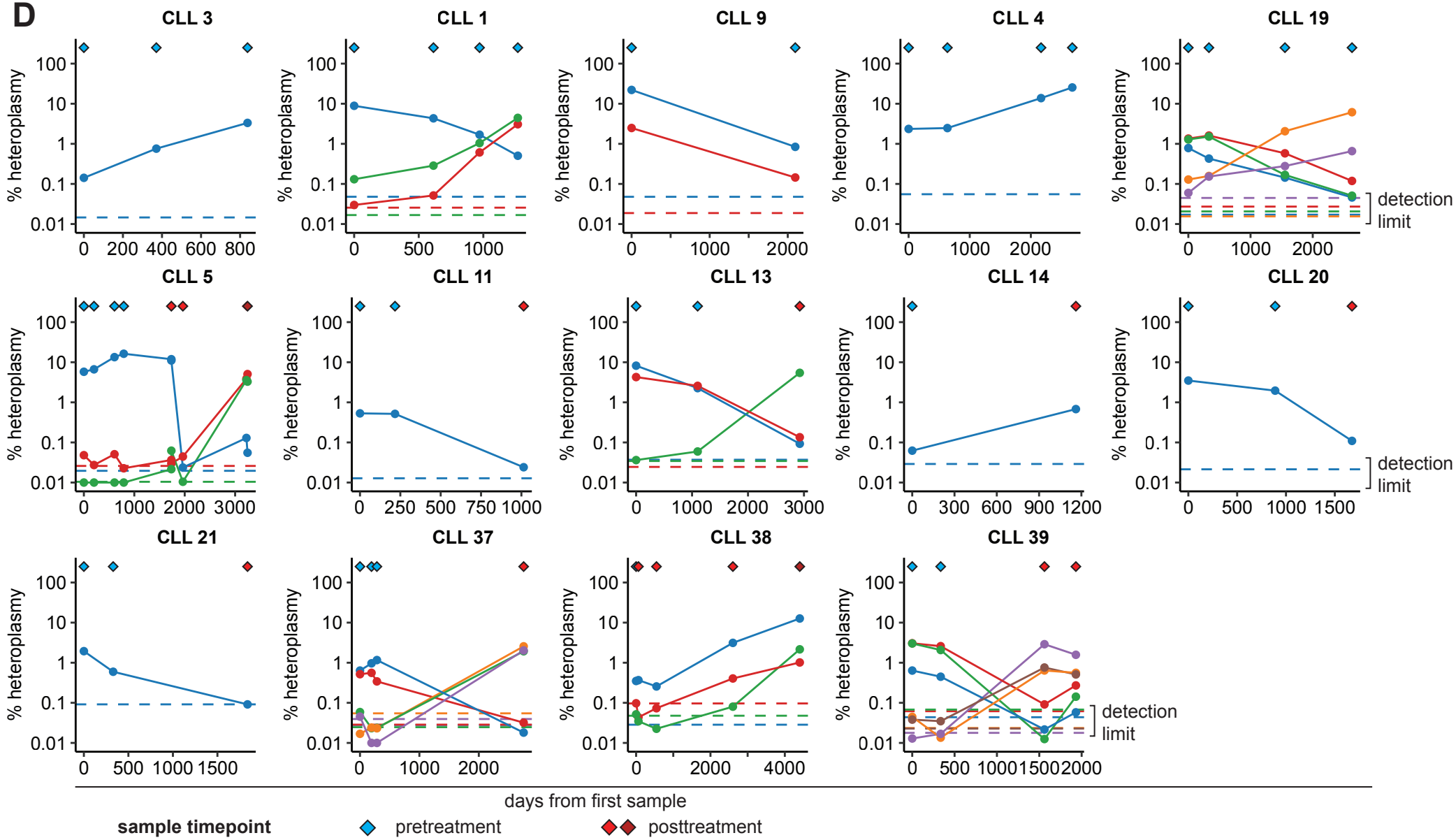


A**B****C****D**

Suppl. Fig. 6. Limit of detection of mitochondrial DNA mutations extracted from bulk RNA-seq data. A The limit of detection (LOD) for mtDNA mutations for each CLL case in bulk RNA-seq data were defined as the heteroplasmy for each mutation within the rest of the study cohort (i.e. background heteroplasmy), excluding samples from the respective CLL case. B LOD of mtDNA mutations is weakly correlated with the average sequencing coverage at that position on chrM. C LOD of mtDNA mutations is driven by their location on chrM. D Longitudinal tracking of mtDNA mutations alongside their respective LOD. Each line corresponds to one individual mtDNA mutation.