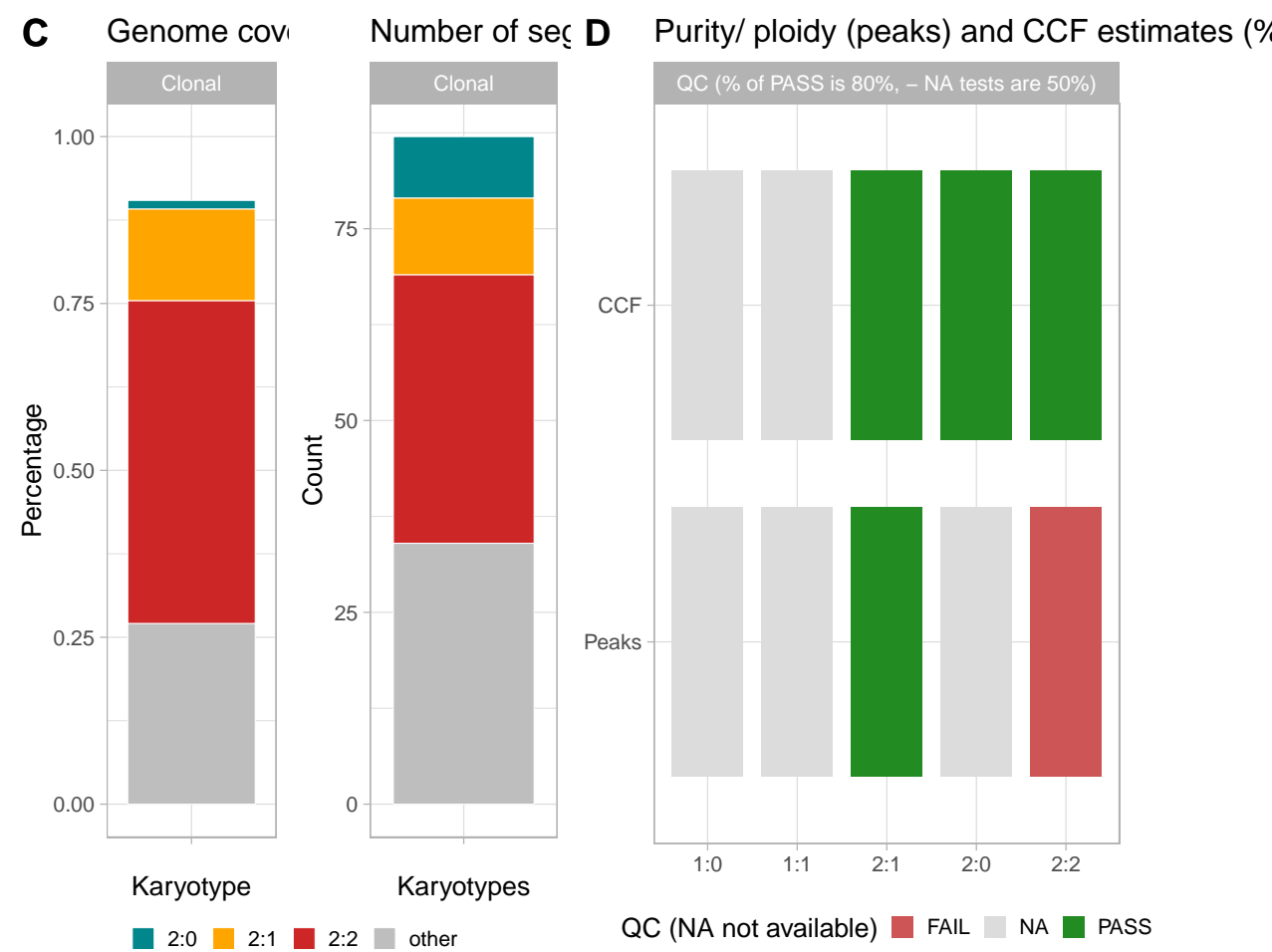
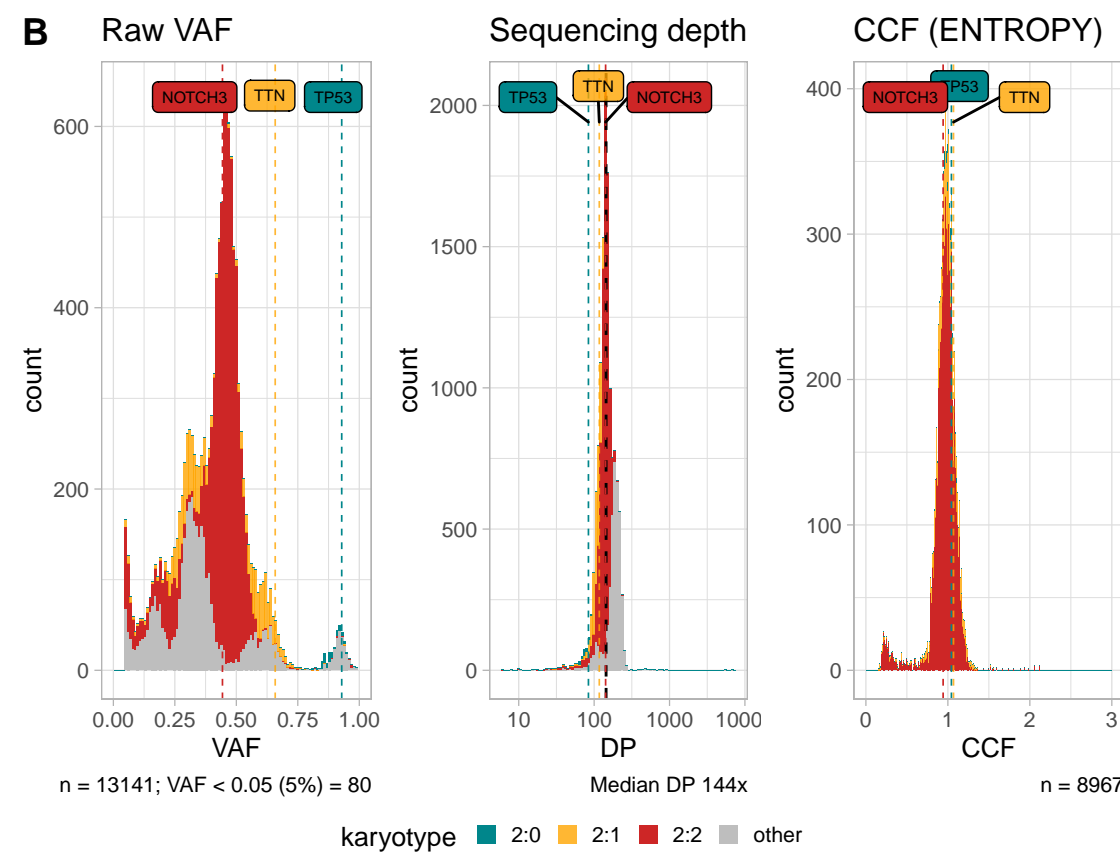
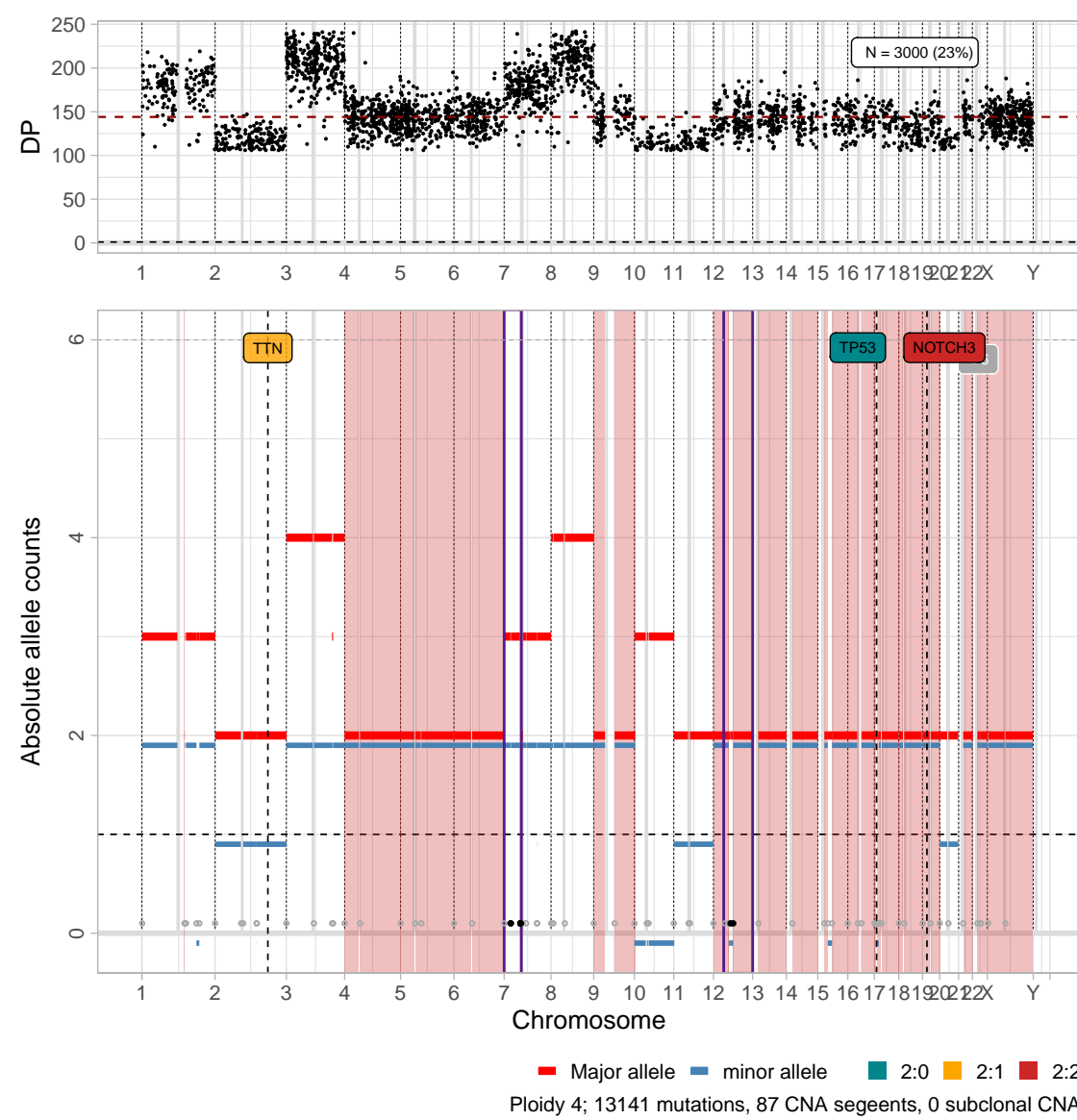
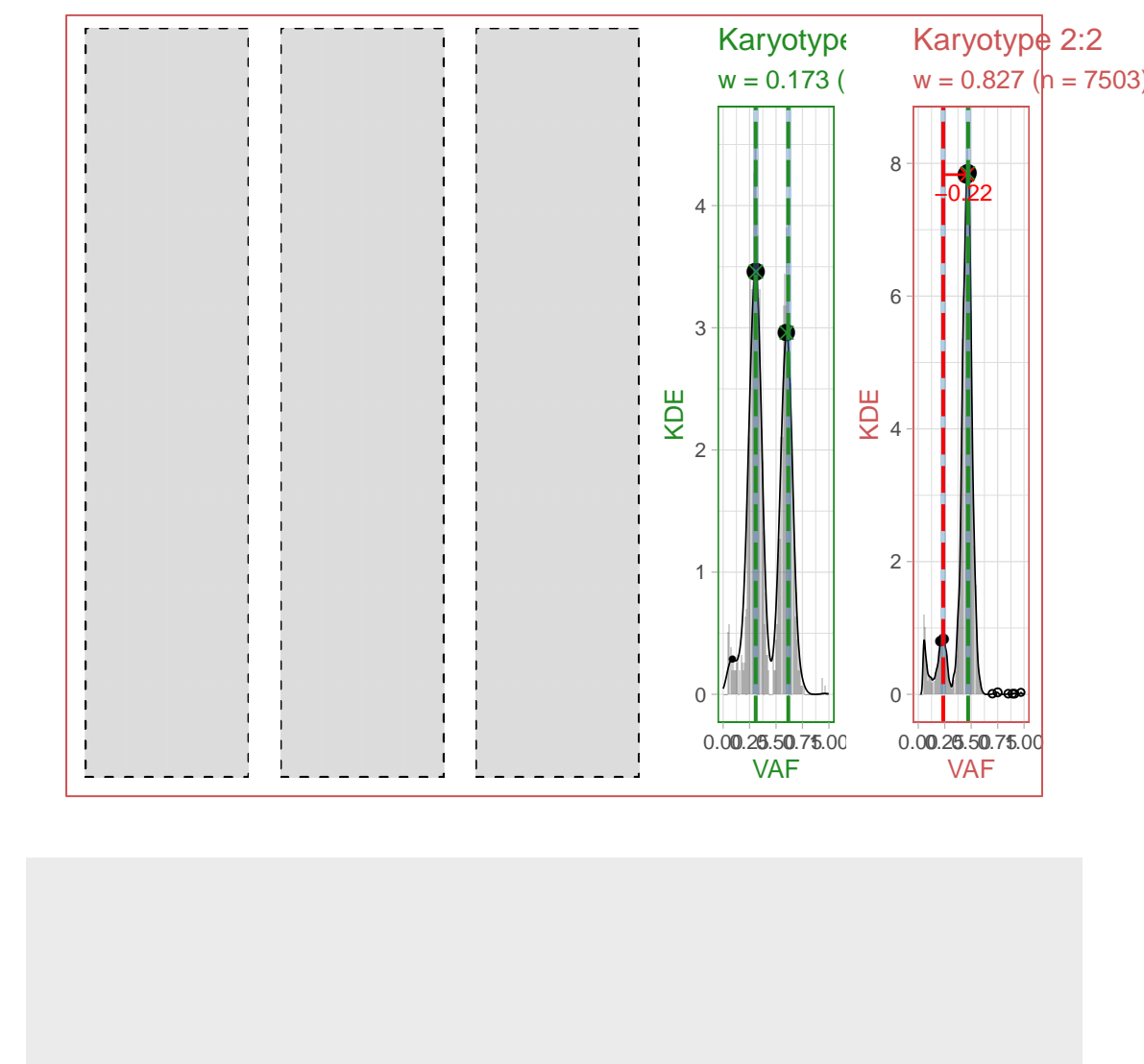
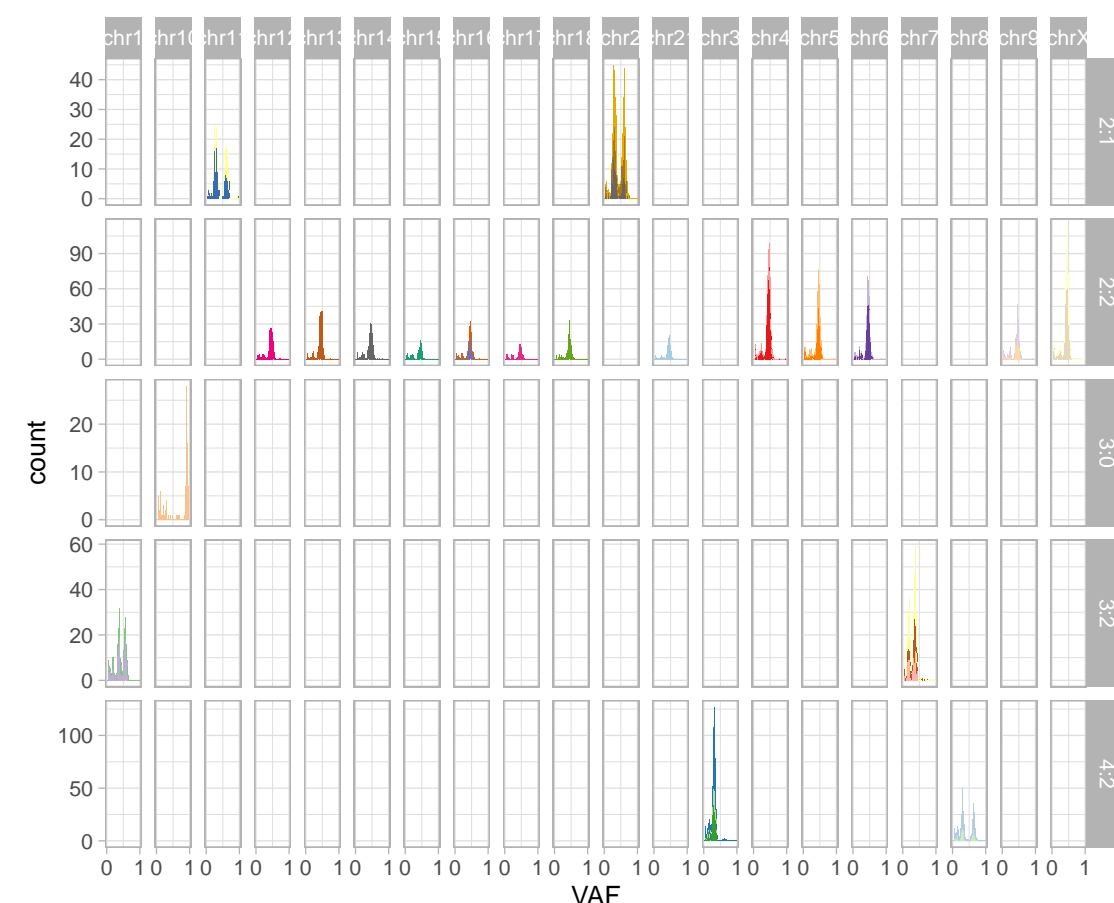
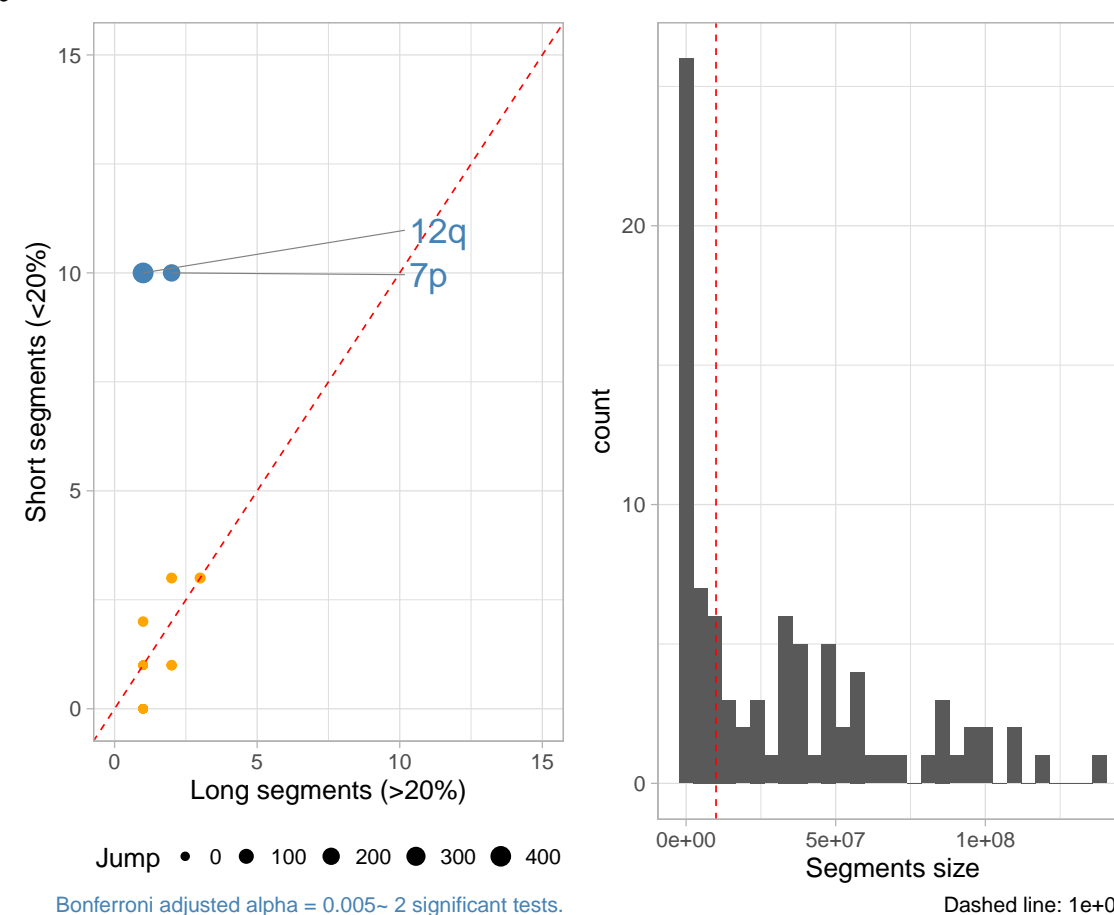


**A** Ploidy 4, most prevalent karyotype is 2:2 (Major:minor).**A** Purity and ploidy QC via peak detection (purity p = 89%)**B** Segments with >131.41 mutations (1% of total) and >2129860 bases (10% of media**C** Arm-level segments fragmentation

## QC PASS (per karyotype) if we can assign &gt;90% of mutation burden

