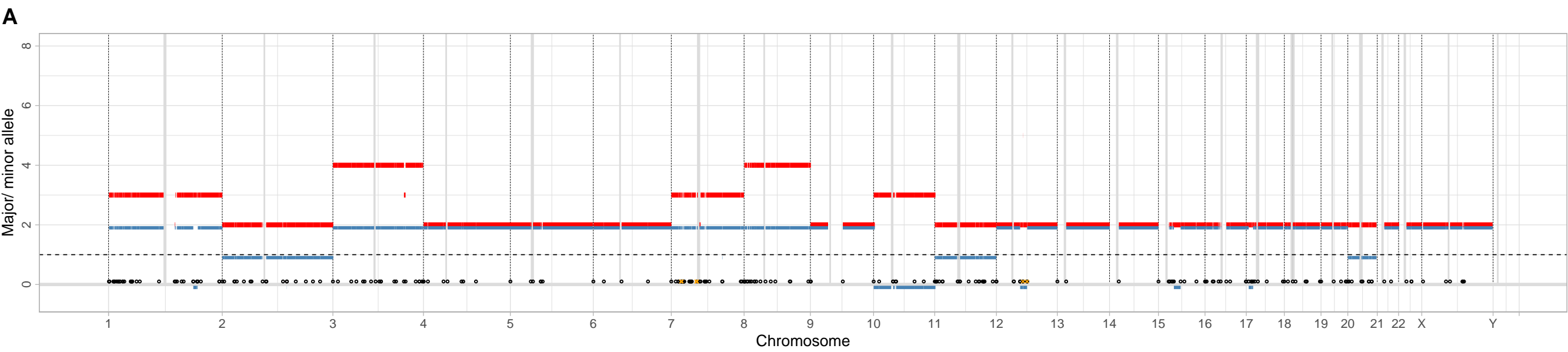


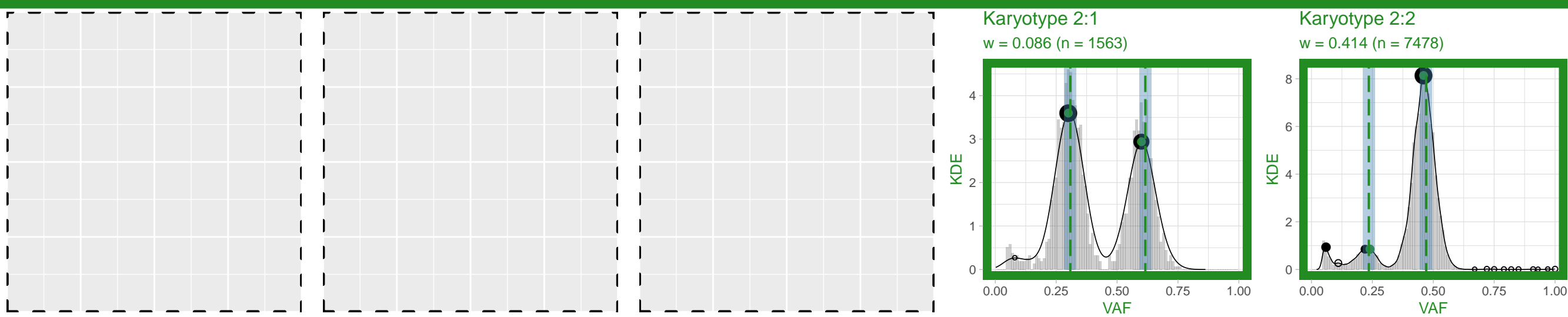
evoverse CNA/ mutation report : My Sample

Input purity 0.89 ; CNAqc ploidy 4 (2:2 karyotype)

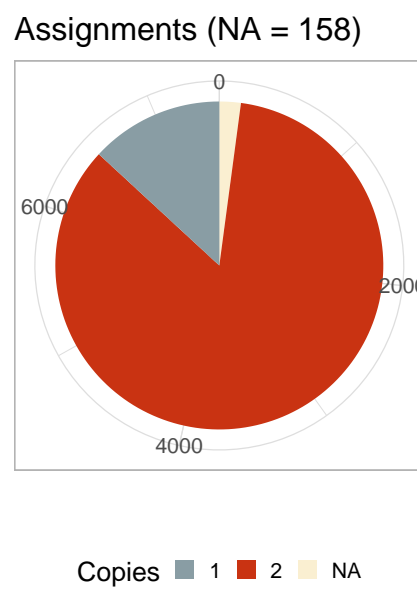
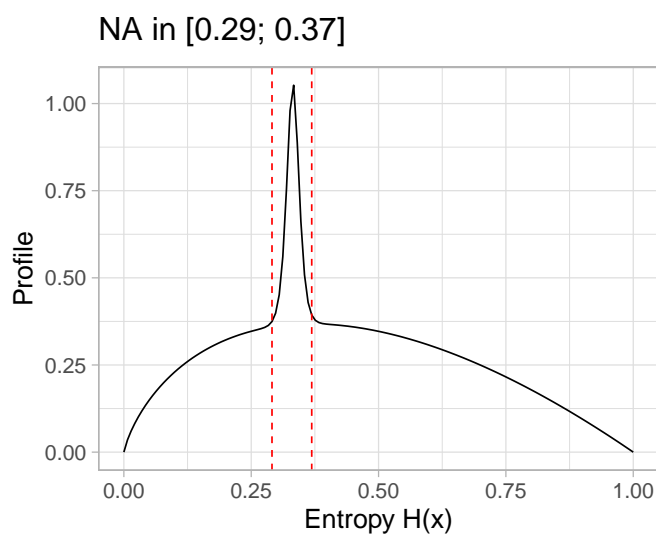
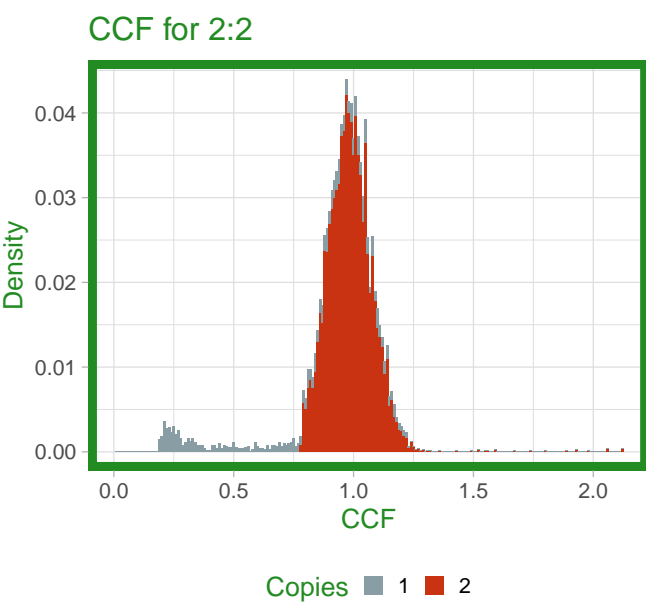
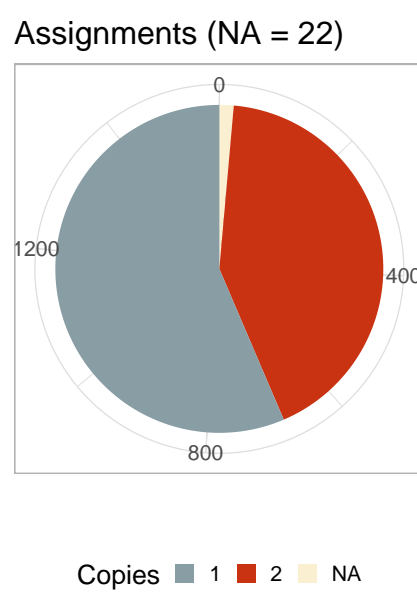
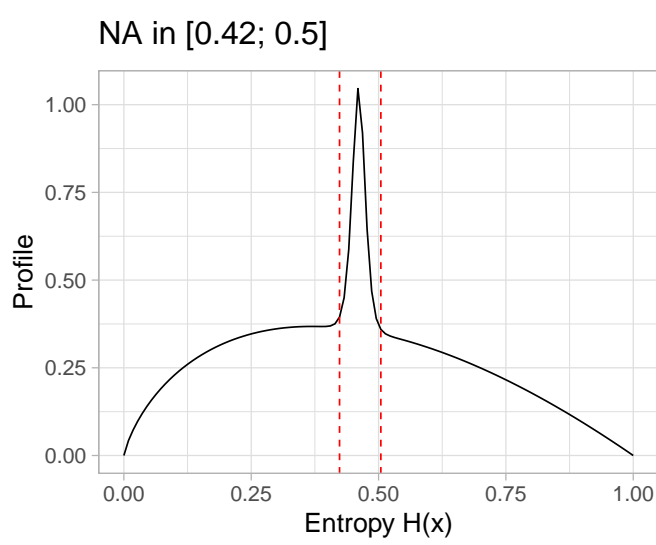
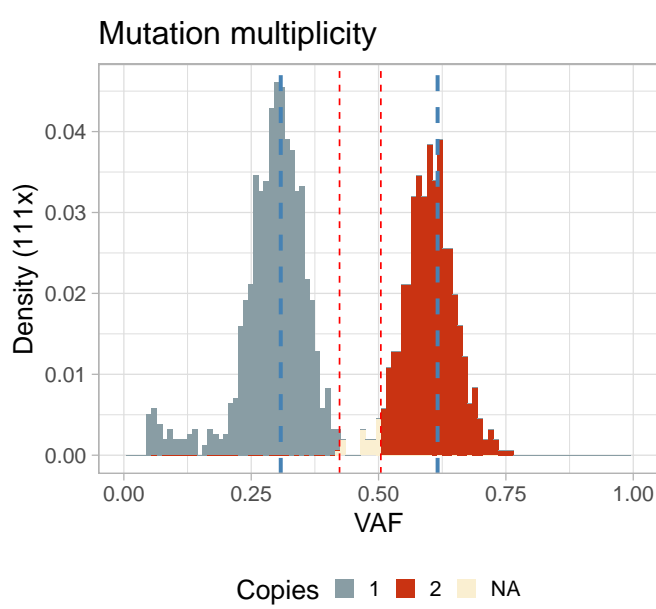
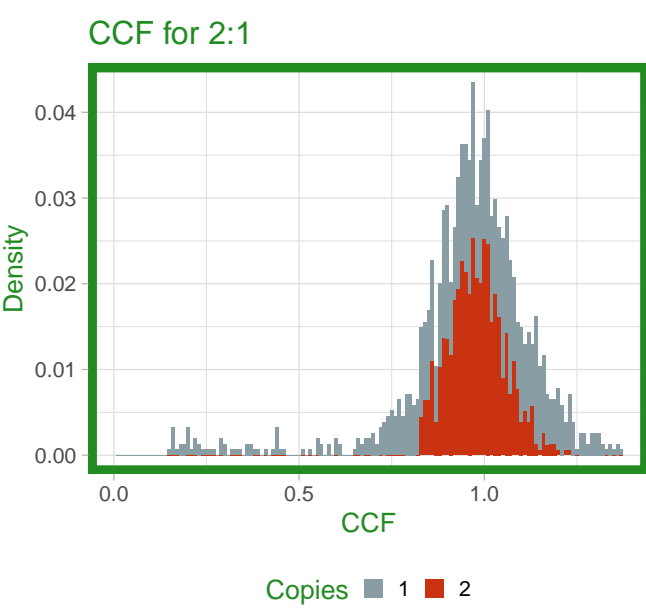
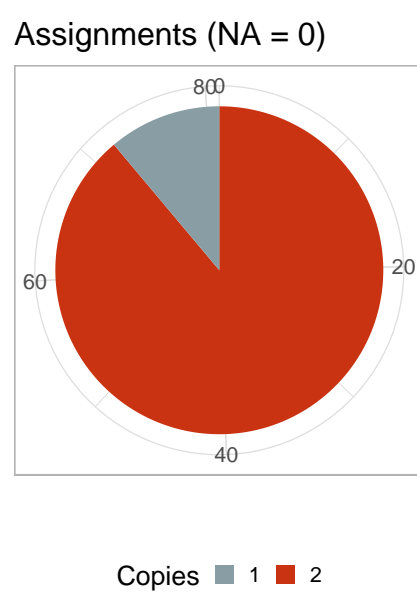
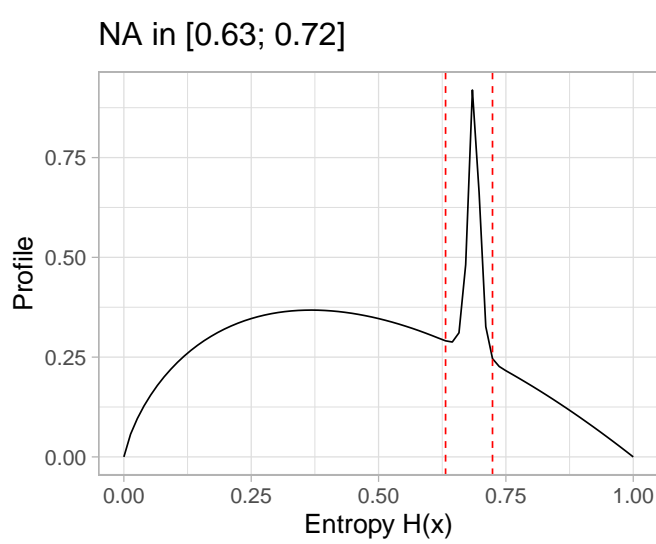
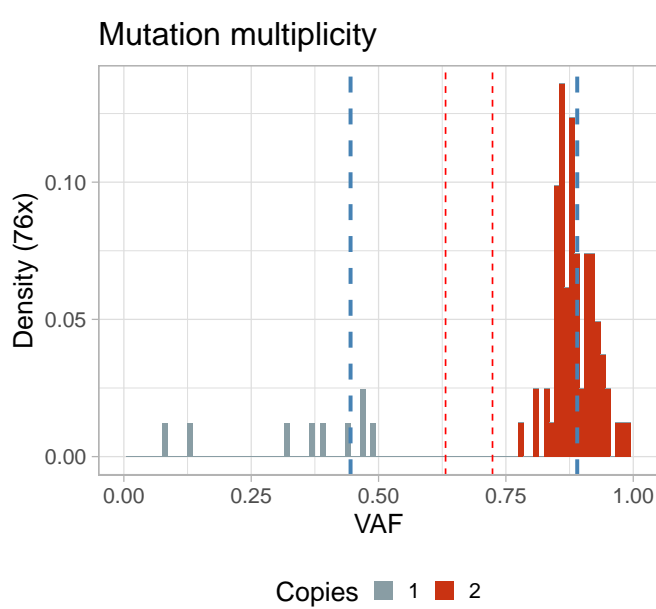
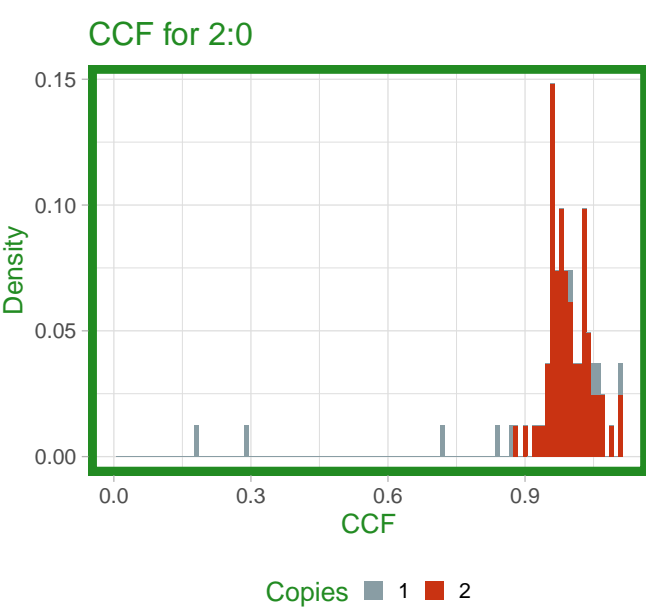
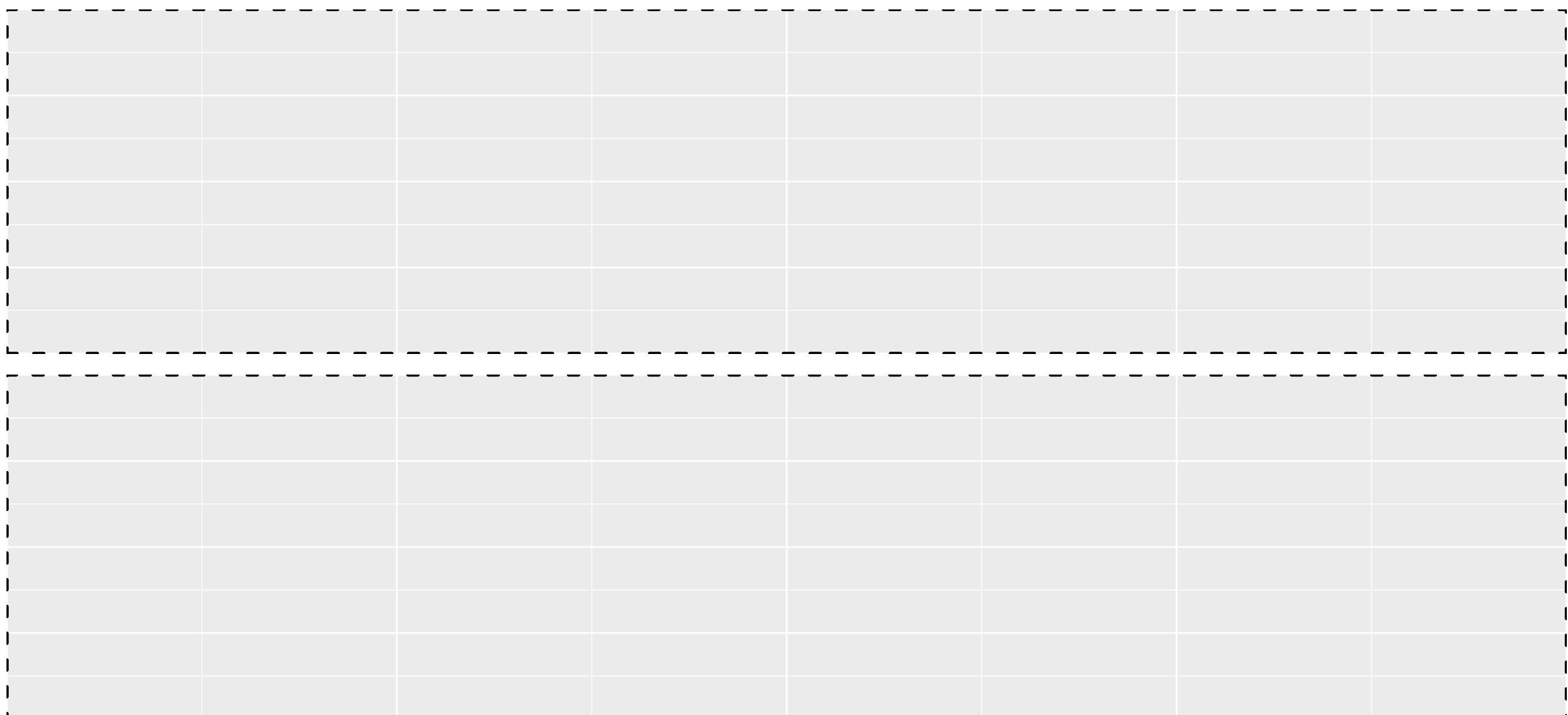


13141 mutations, 267 clonal CNA, 0 subclonal CNA

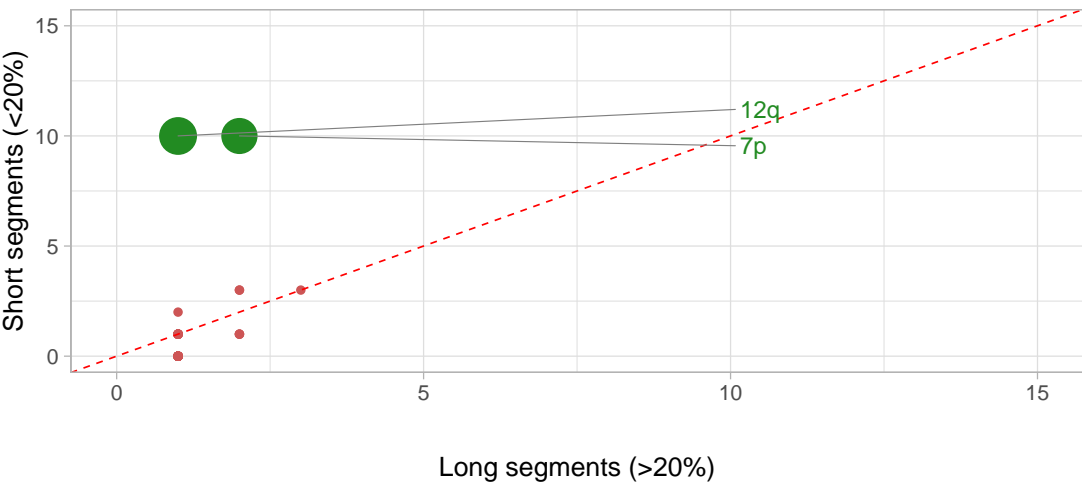
B
Purity/ploidy QC : $\rho = 0.004689562$ ($\varepsilon = 0.025$)



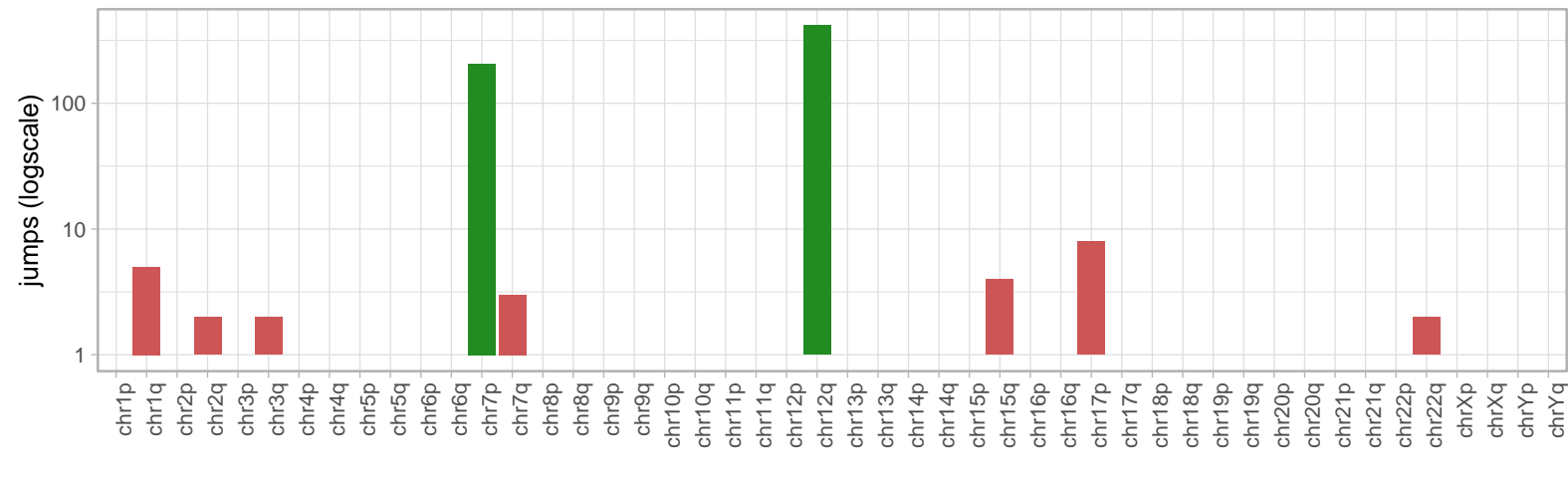
C
Cancer Cell Fractions (estimated)



D
2 overfragmented arms (Bonferroni alpha = 0.005)



Segments jumps



Fragmented ● FALSE ● TRUE

$-\log(p)$ • 0 ● 5 ● 10