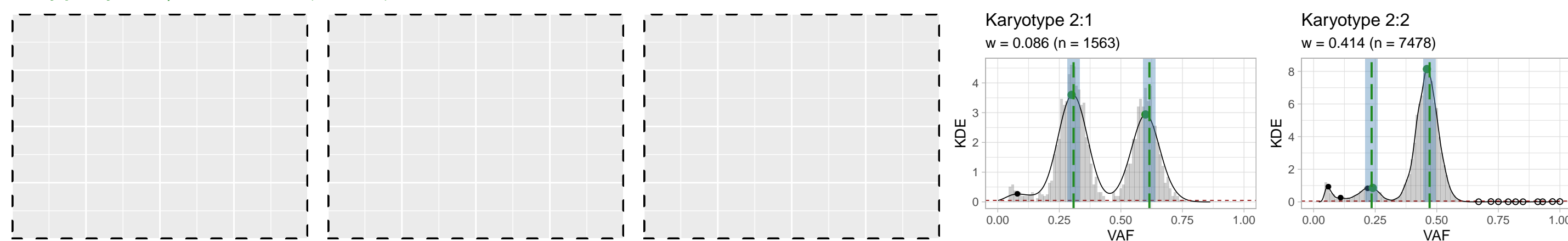


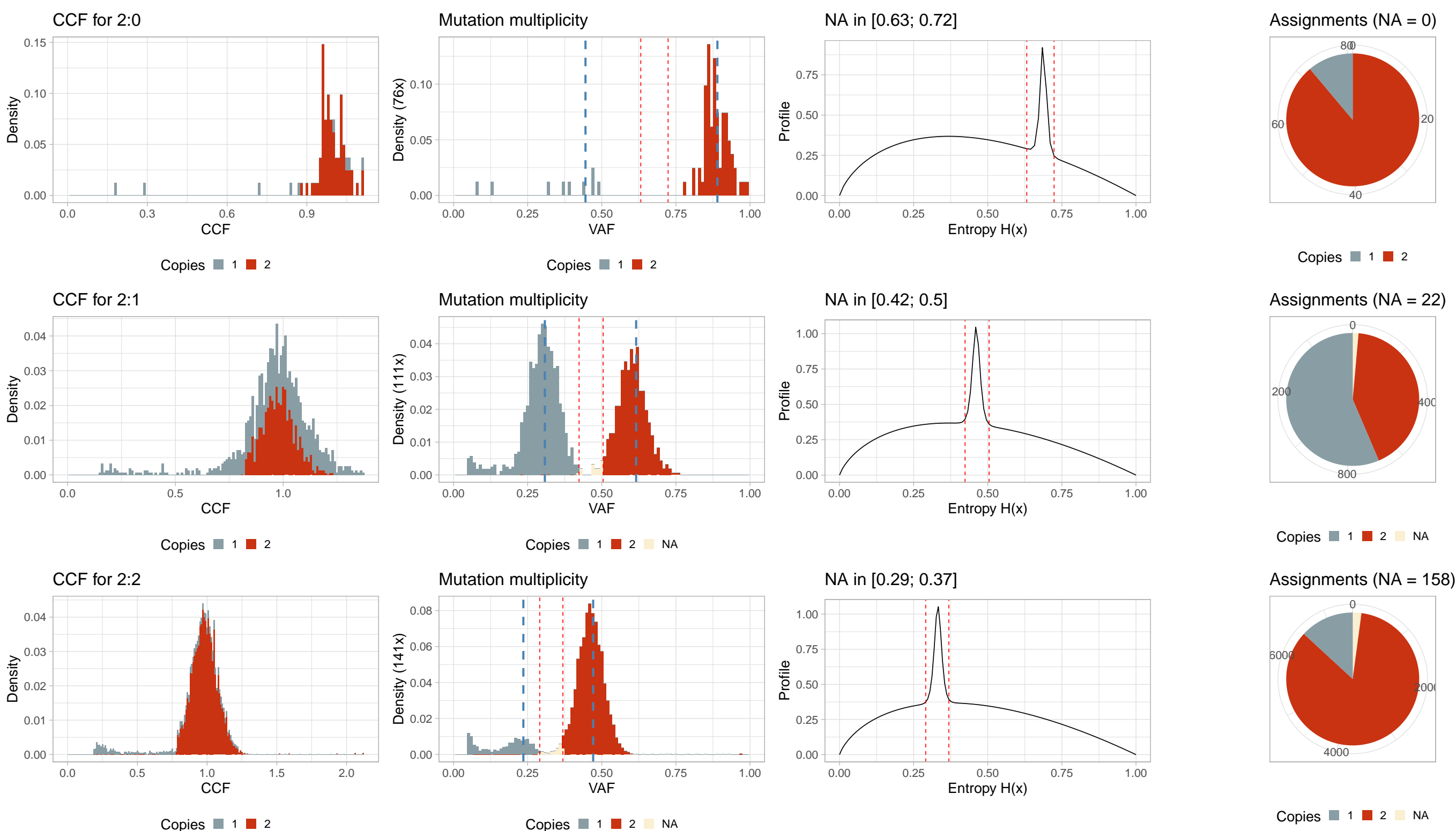
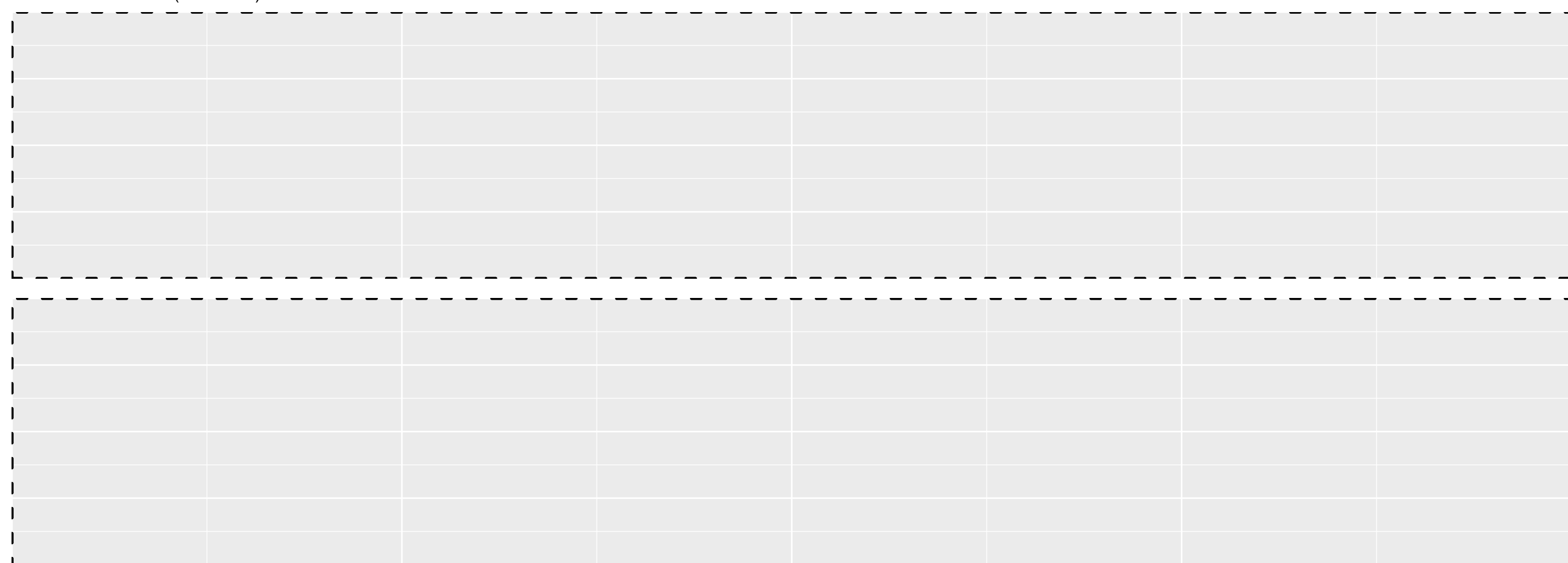
Input purity 0.89 ; CNAqc ploidy 4(2:2)

13141 mutations, 267 clonal CNA, 0 subclonal CNA

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

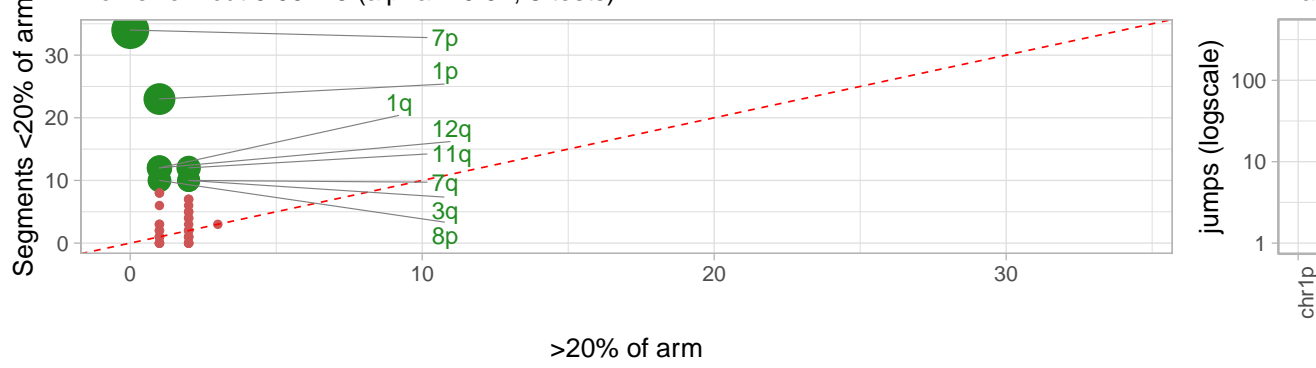


Cancer Cell Fractions (estimated



Segments overfragmentation in 8 arm

Bonferroni cut 0.00125 (alpha = 0.01, 8 tests)



Segments jumps

Maximum jump 414

