

XXX\_YYY\_ZZZ

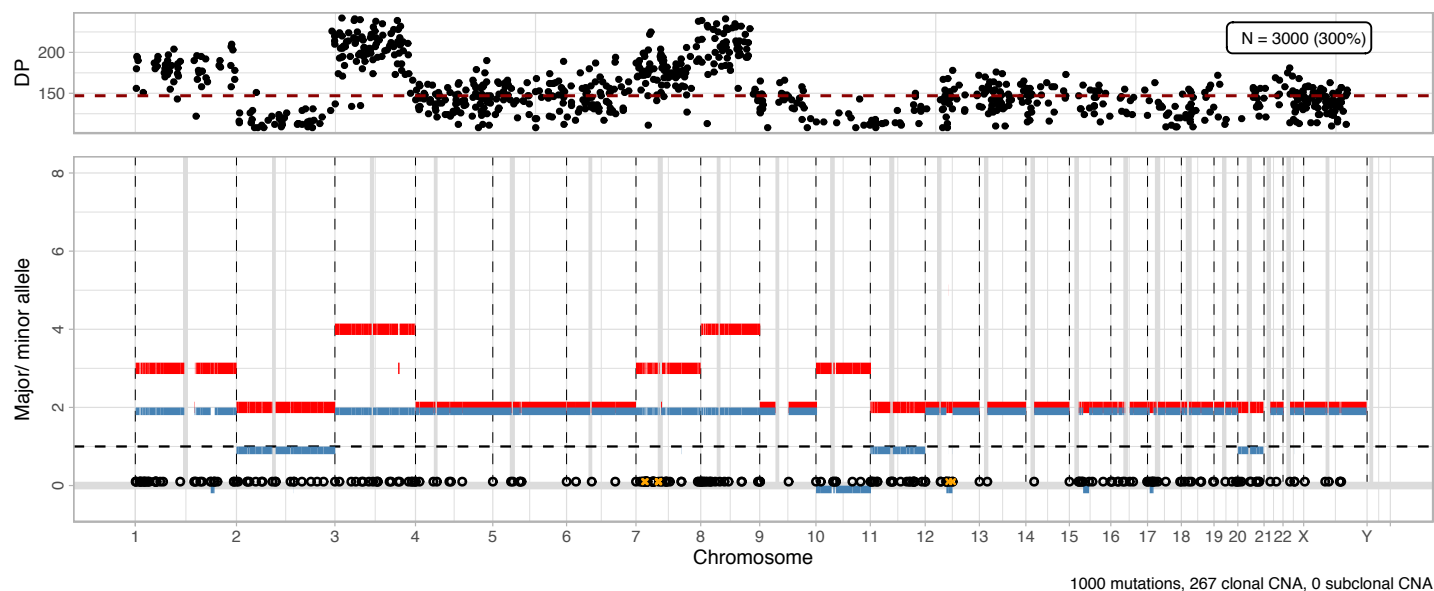
CNA segments (A), karyotype coverage (B) and summary QC (C).

Monoclonal architecture ~ QC PASS.

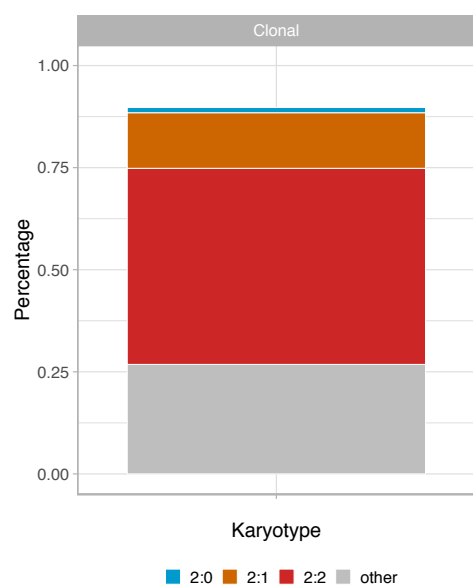
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CNA segments (A), karyotype coverage (B) and summary QC (C).

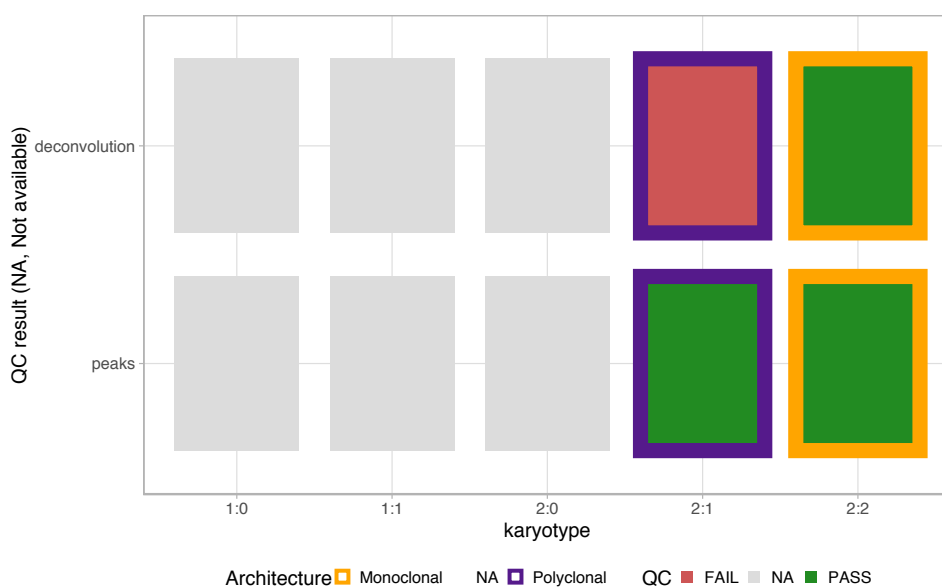
**A** Ploidy 4, most prevalent karyotype is 2:2 (Major:minor).



**B** Karyotypes coverage



**C** Summary QC results



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Subclonal deconvolution from raw VAF data, per karyotype.

## mobster and BMix fits per karyotype

