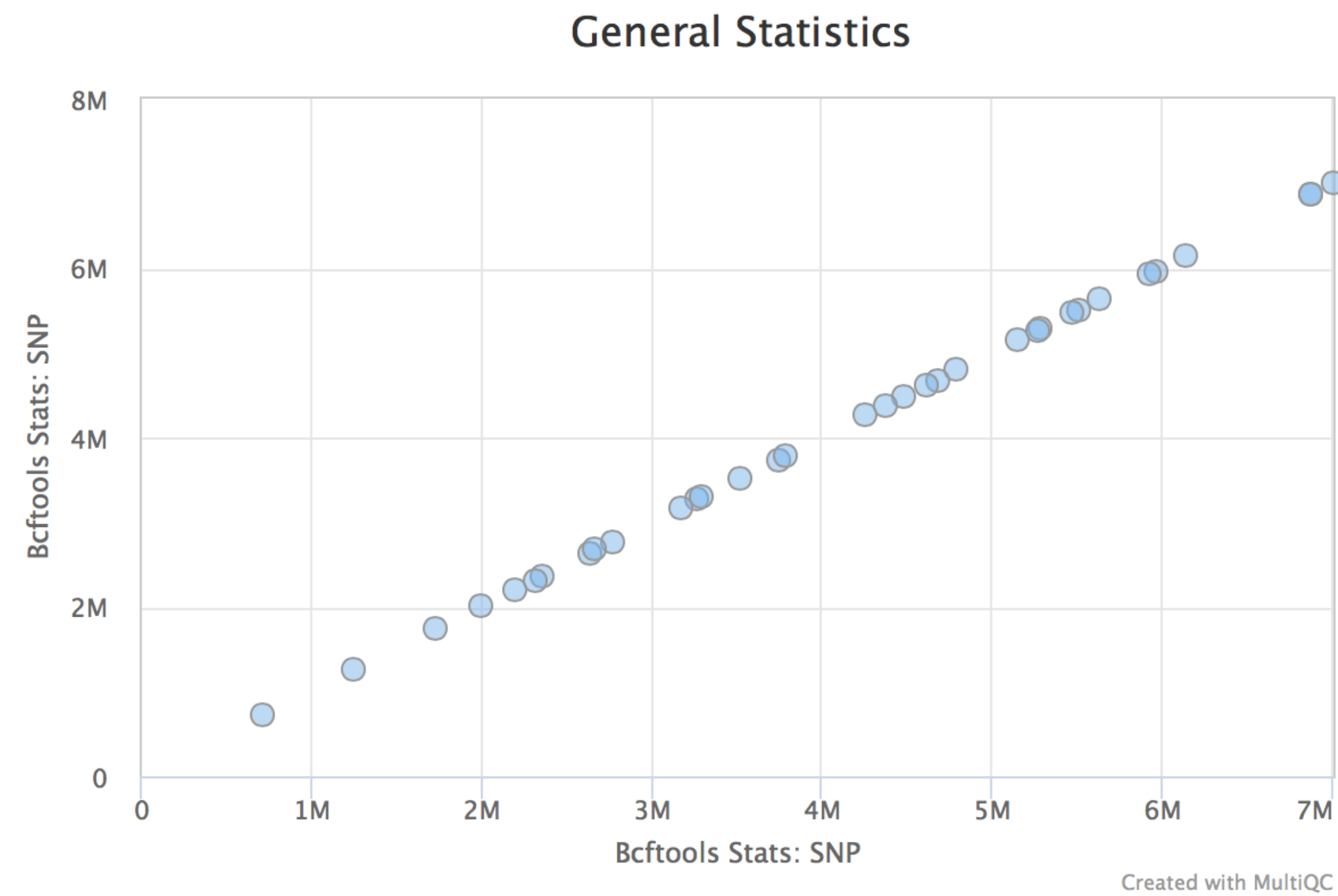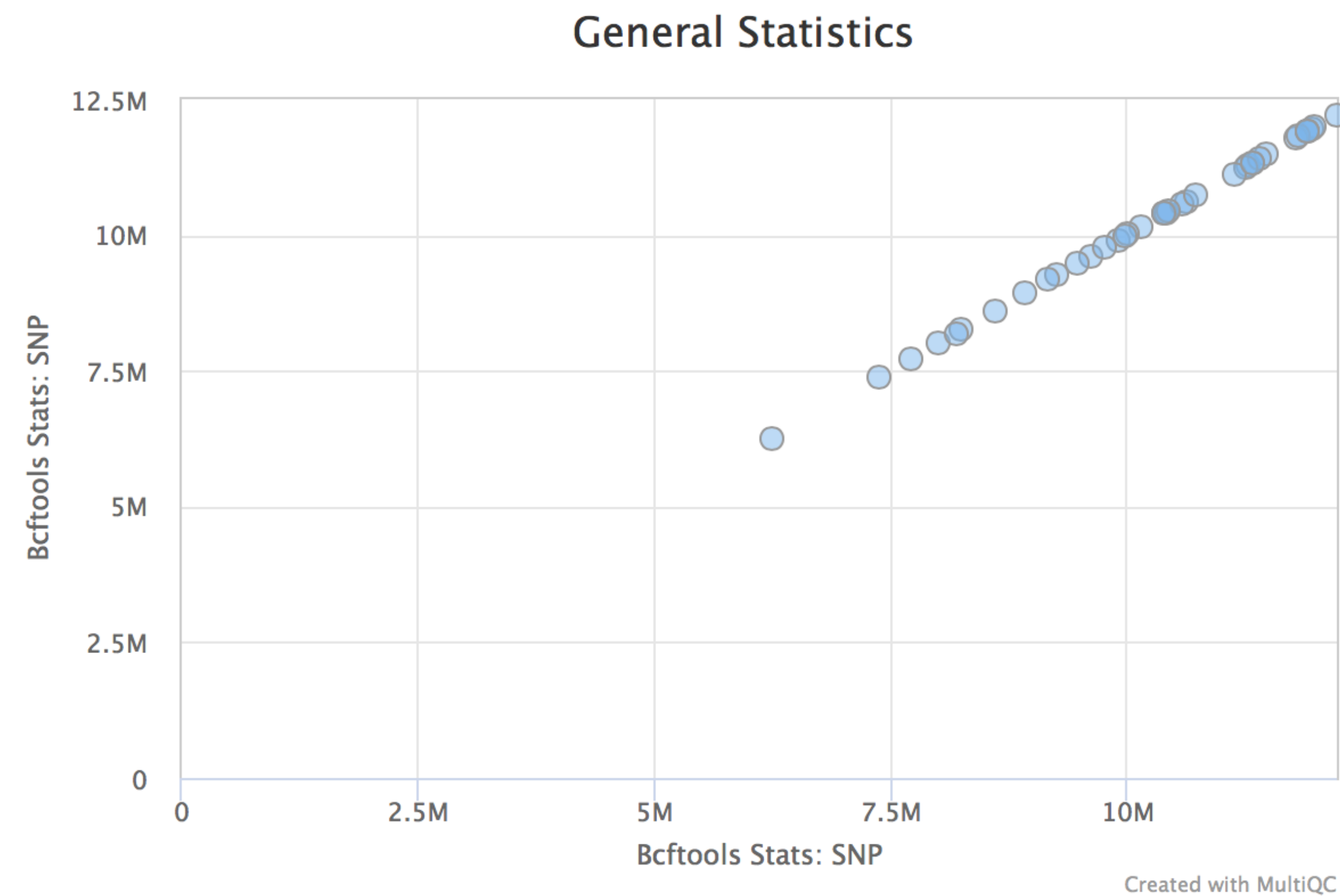# Summary of SNP calling with freebayes
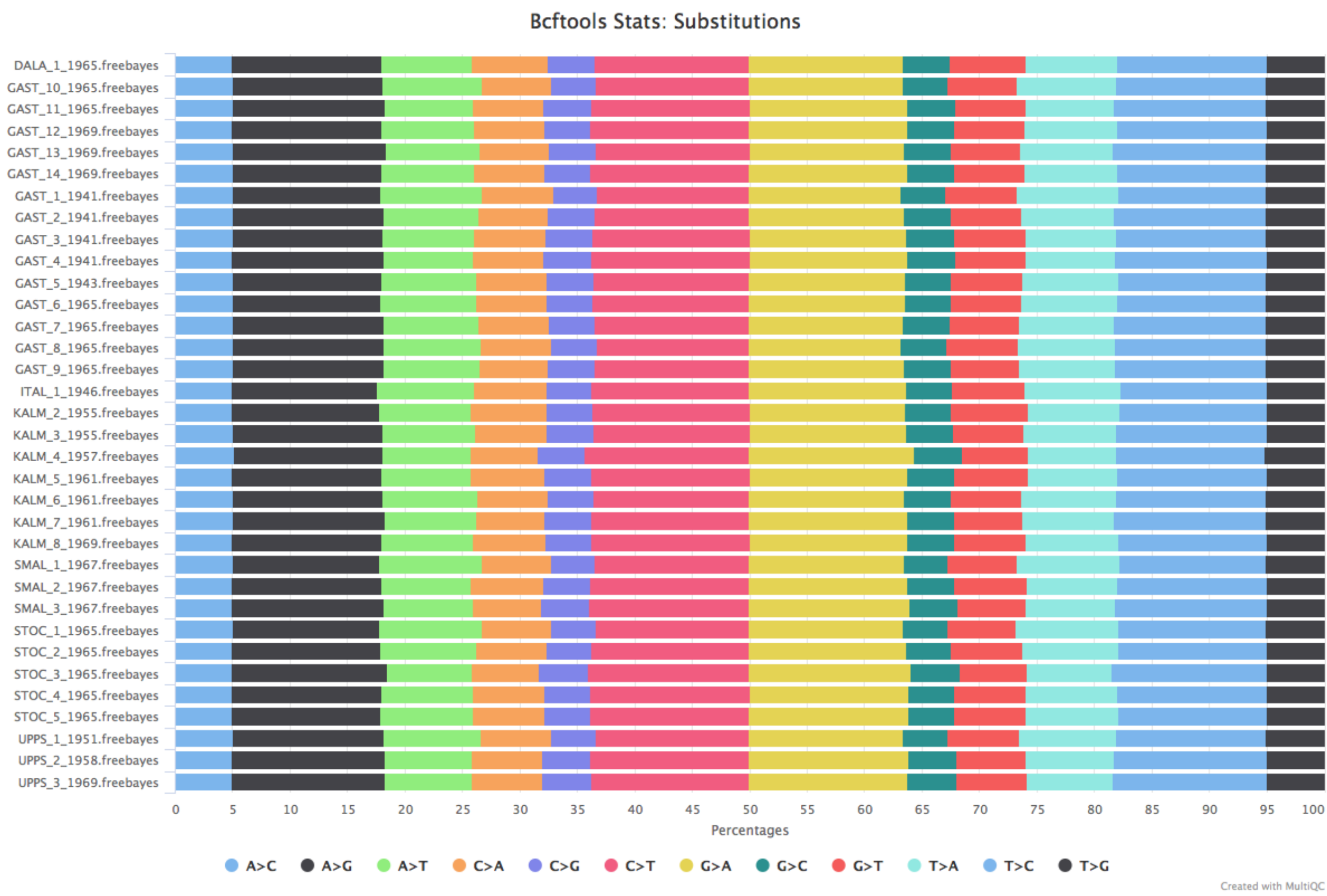

"Historical" samples
input GC > 50%
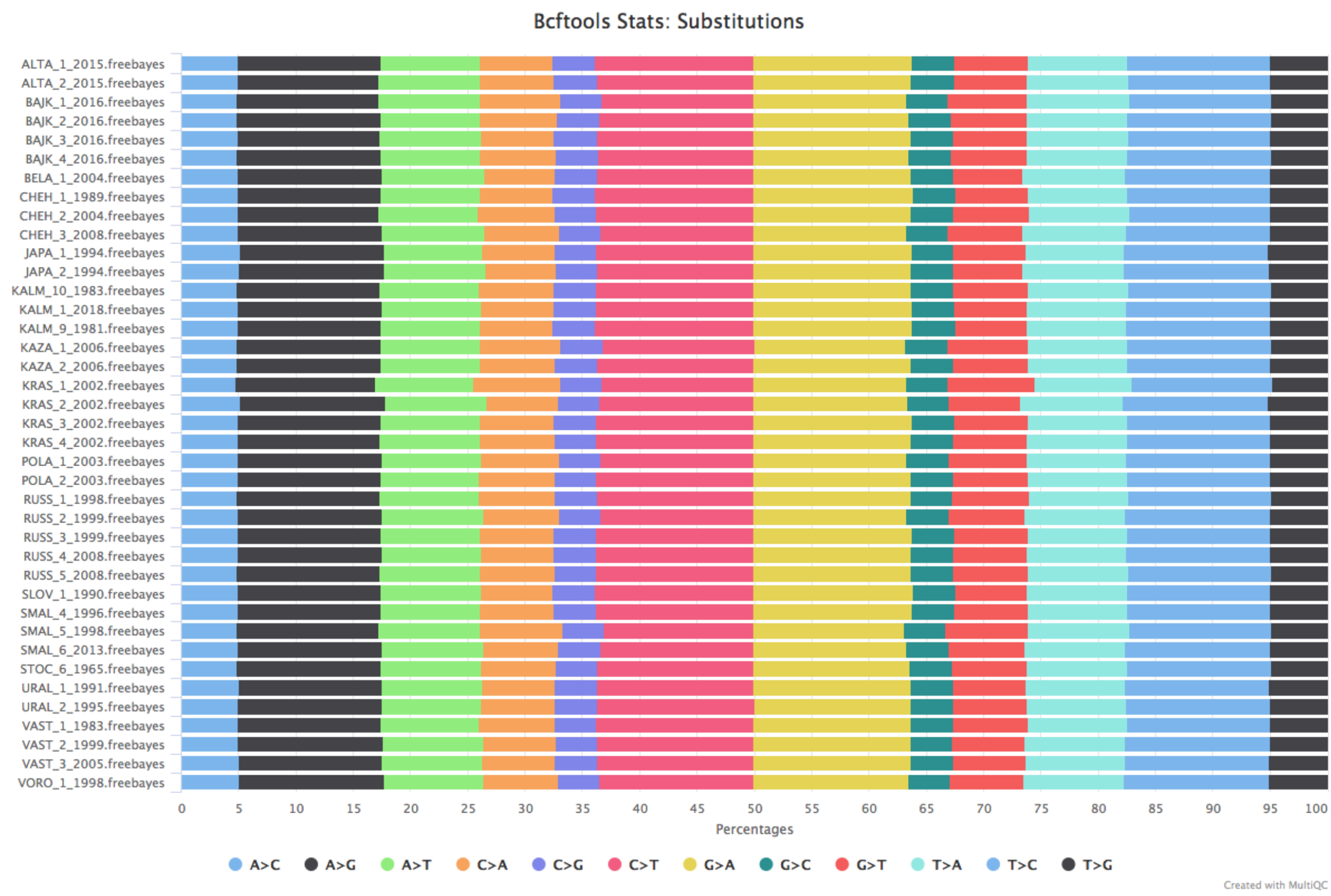
"Contemporary" samples
input GC > 50%

Summary of SNP calling with freebayes
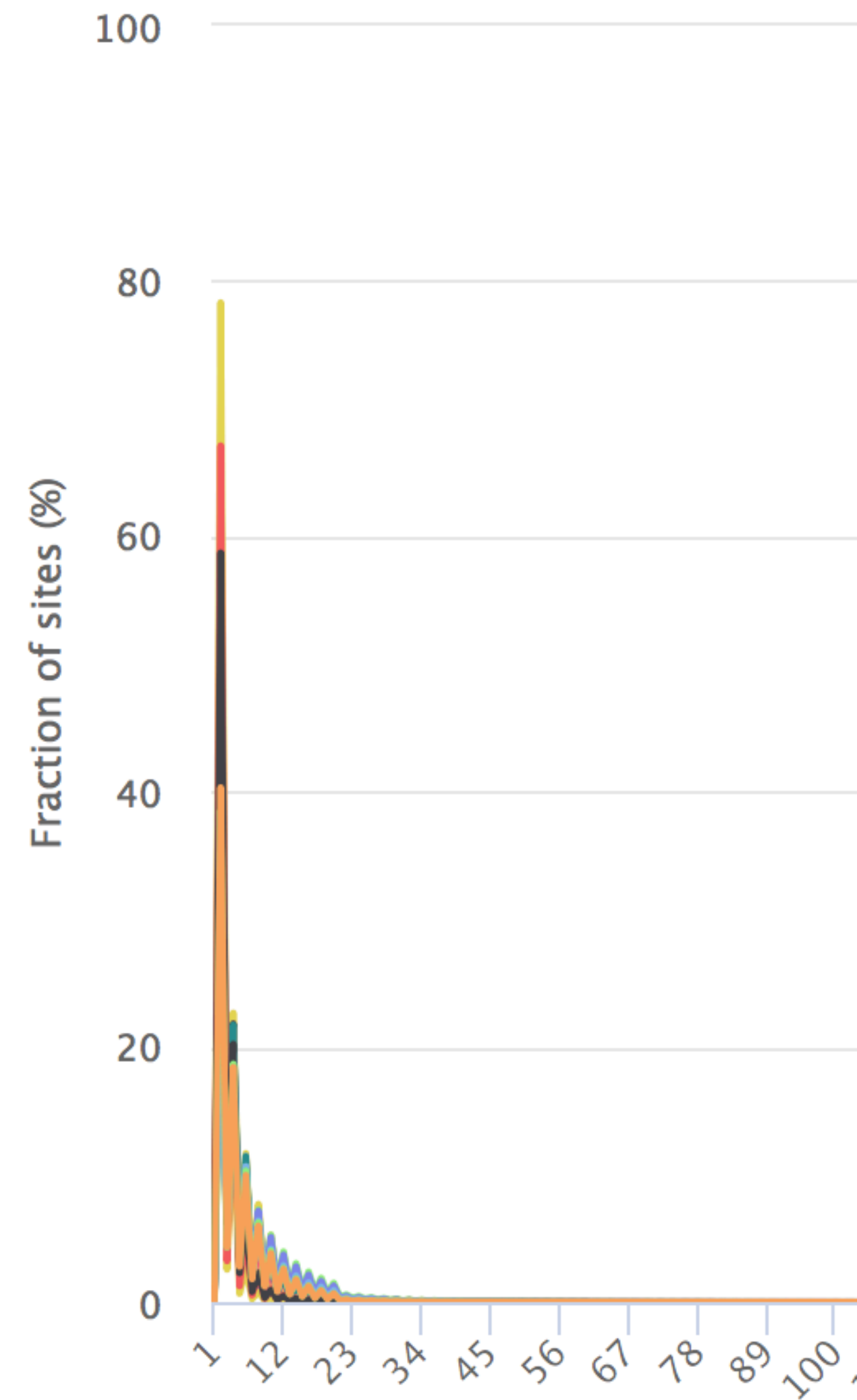
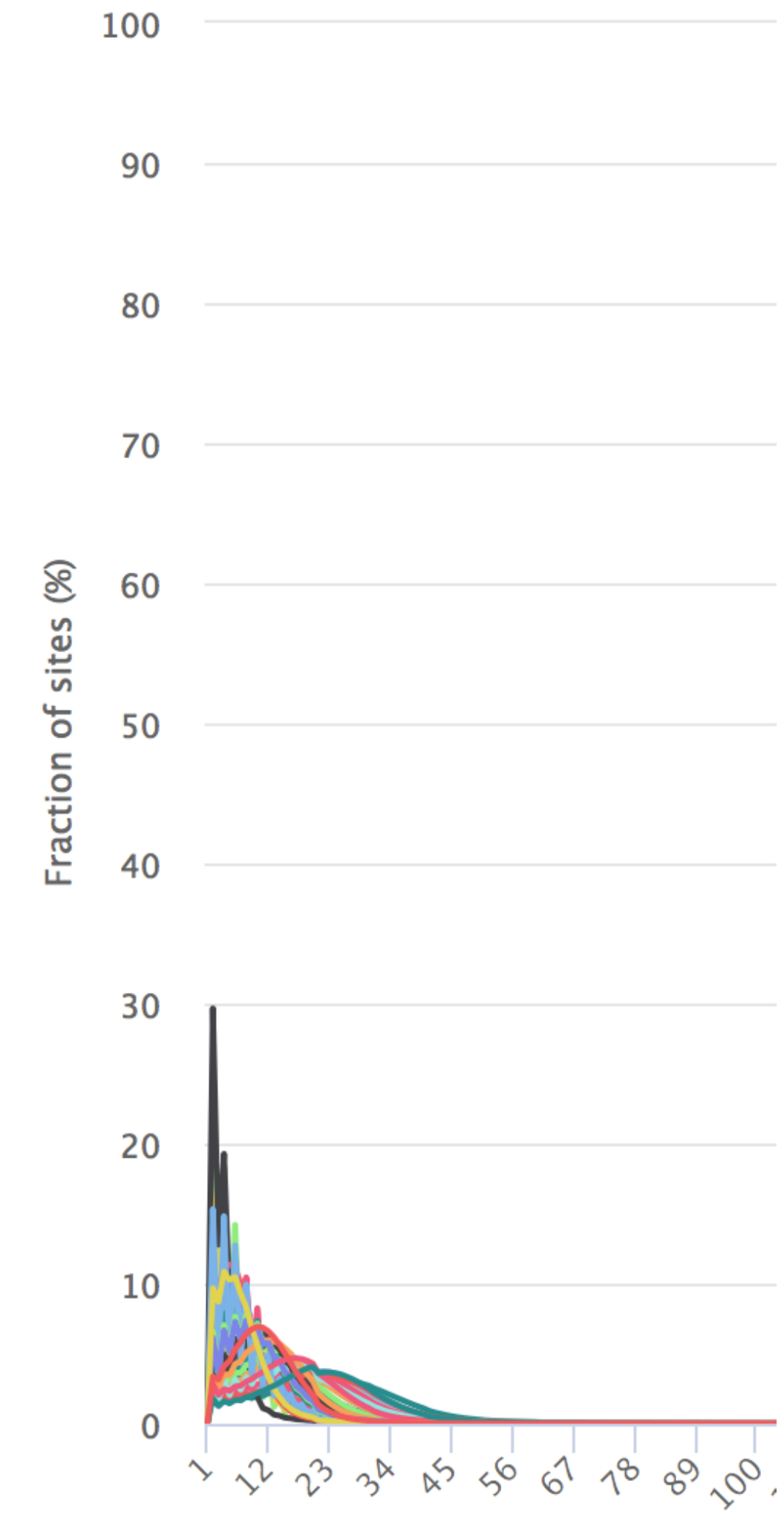## Type of substitutions

"Historical" samples
input GC > 50%

"Contemporary" samples
input GC > 50%

"Historical" samples
input GC > 50%

"Contemporary" samples
input GC > 50%

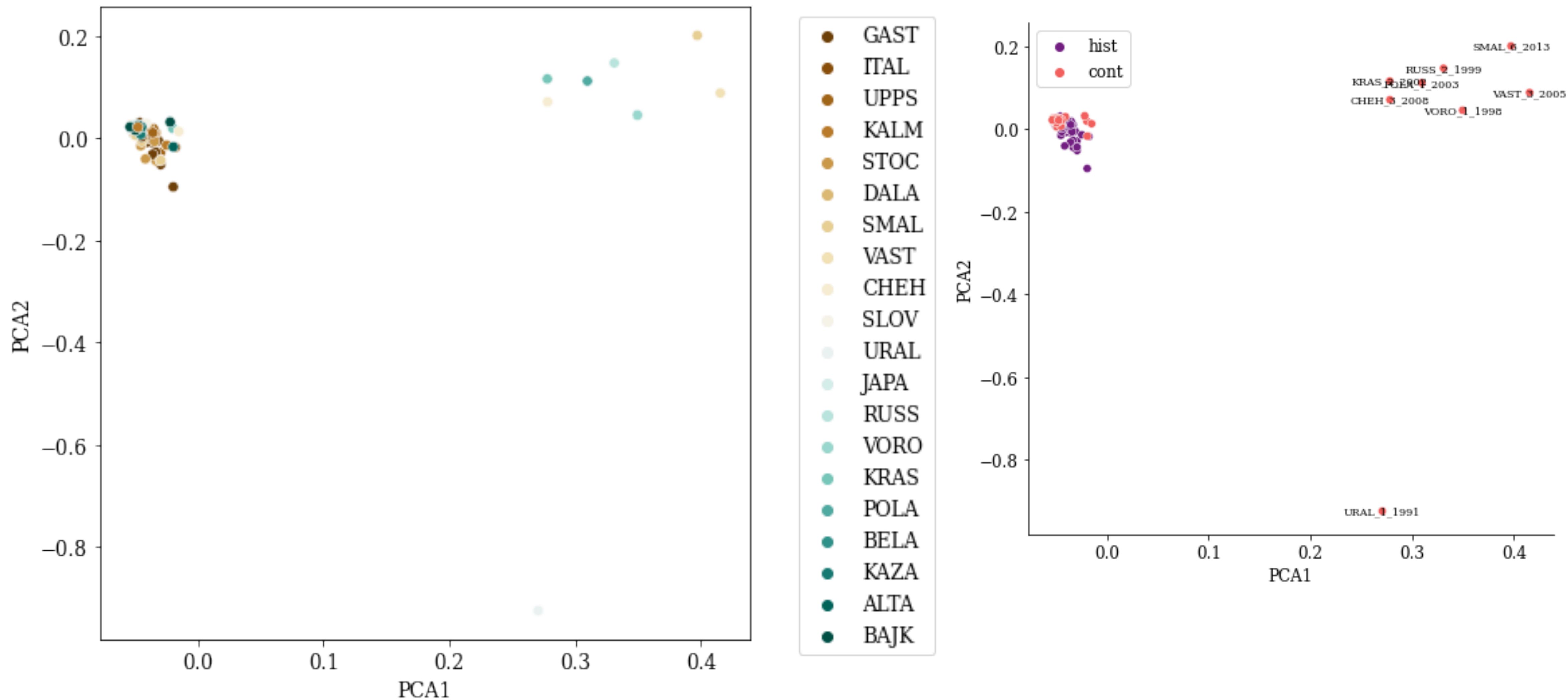Fraction of sites (%)

Variant depth

Fraction of sites (%)

Variant depth

Summary of hard filtering

number of samples:	73
number of records:	2203
number of no-ALTs:	0
number of SNPs:	2203
number of MNPs:	0
number of indels:	0
number of others:	0
number of multiallelic sites:	334

--remove-indels
--max-missing-count 10
--minQ 30
--min-meanDP 10
--max-meanDP 40

All data plotted

Outlier group removed