Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
91bes	4,104,160	1.93	1.93	954,524	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
91bes	63.0%	1.90	67.9%	101,516	2,133

- 91bes .. `/ngc/projects2/gm/data/archive/2022/variants/snv/91bessatf-103892326325-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220720_A01176_AHMY7JDSX3-RHGM_LABKA_WGS-`
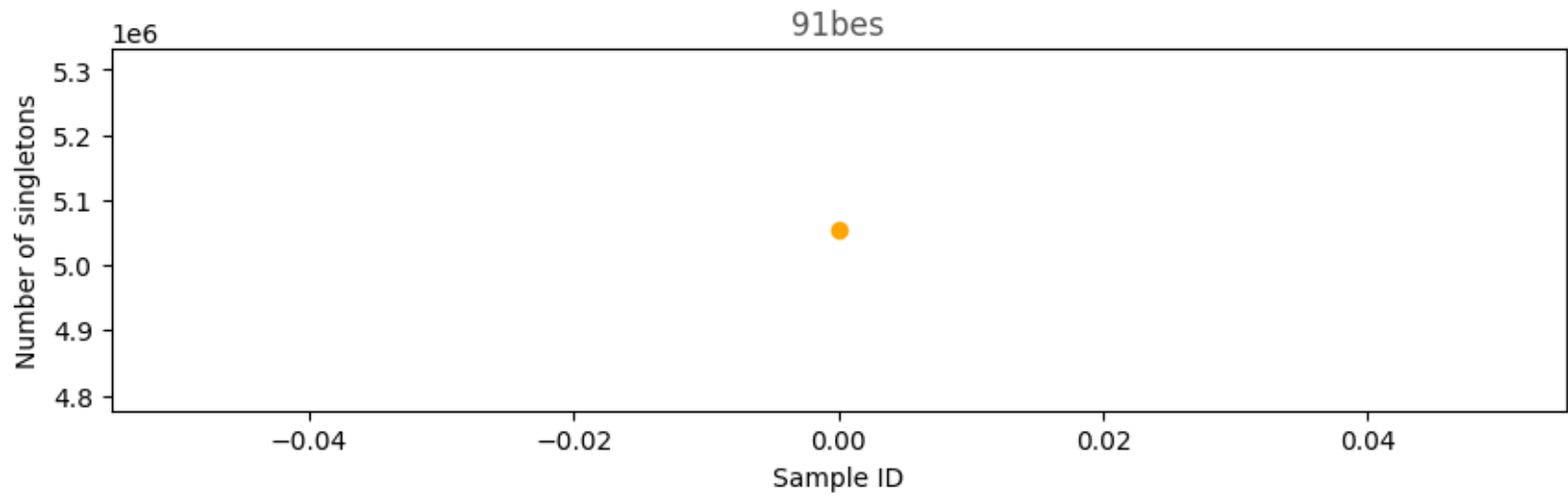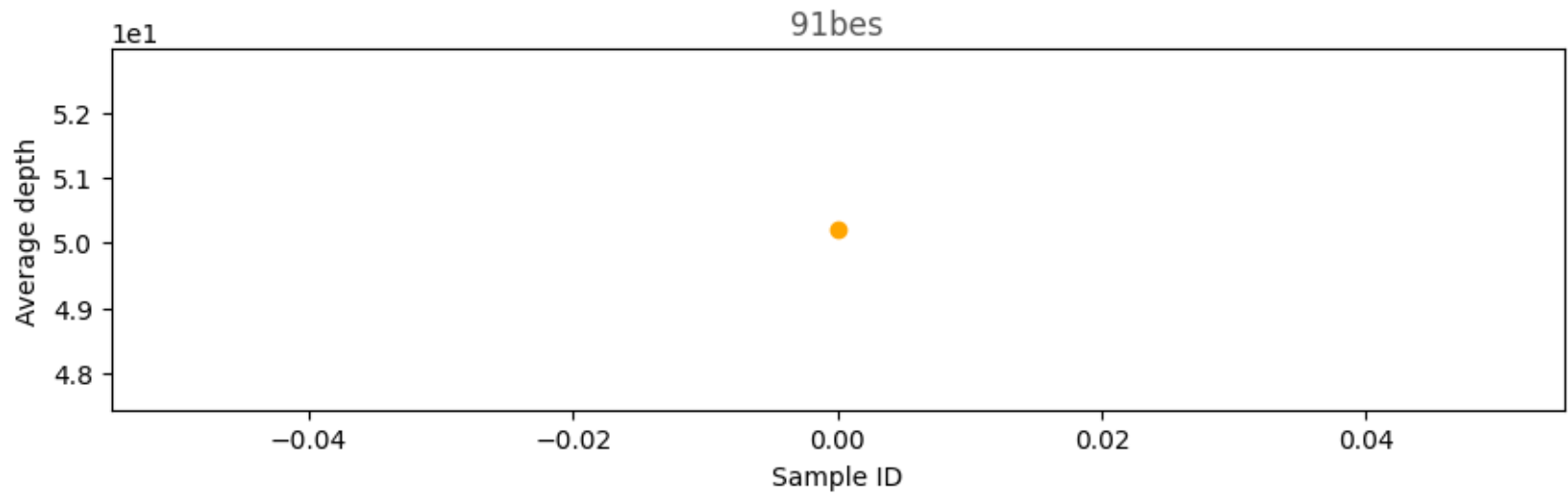 `WGS04683_snv_germline_raw.haplotype_caller.vcf.gz`
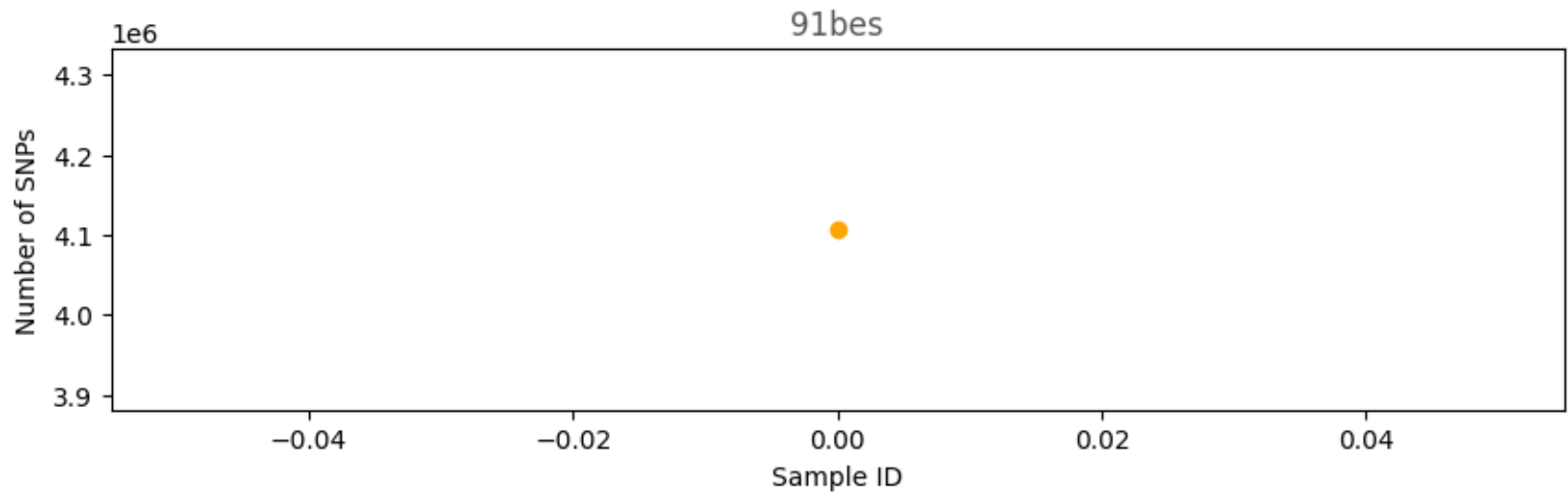
Ts/Tv by sample
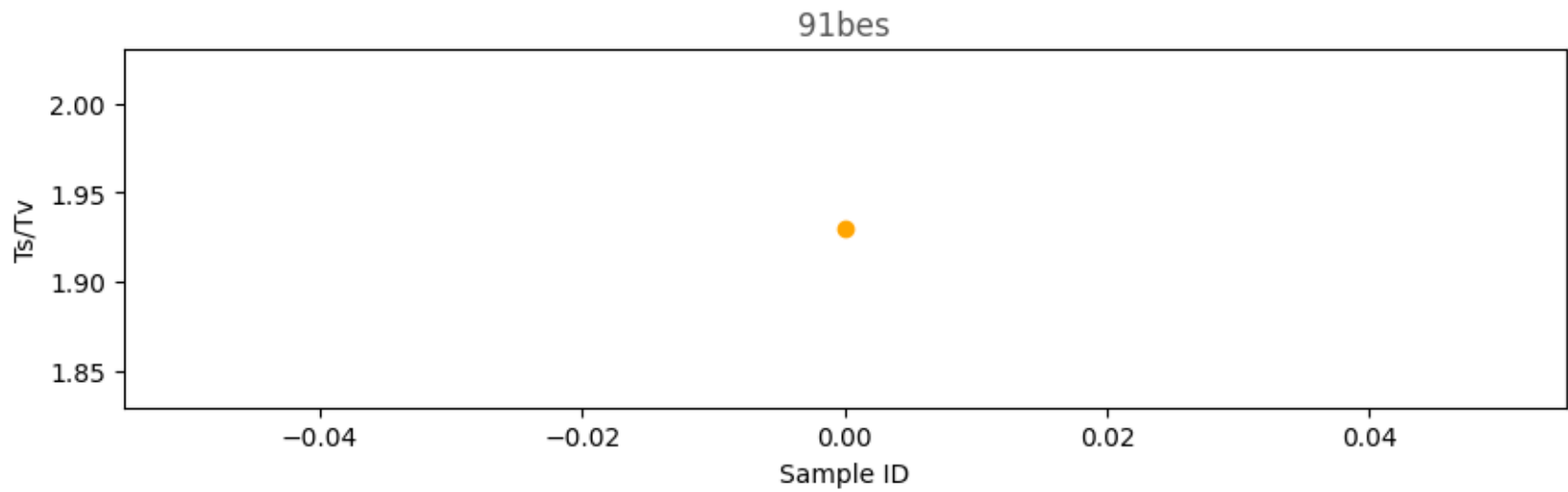
Hets vs non-ref Homs by sample

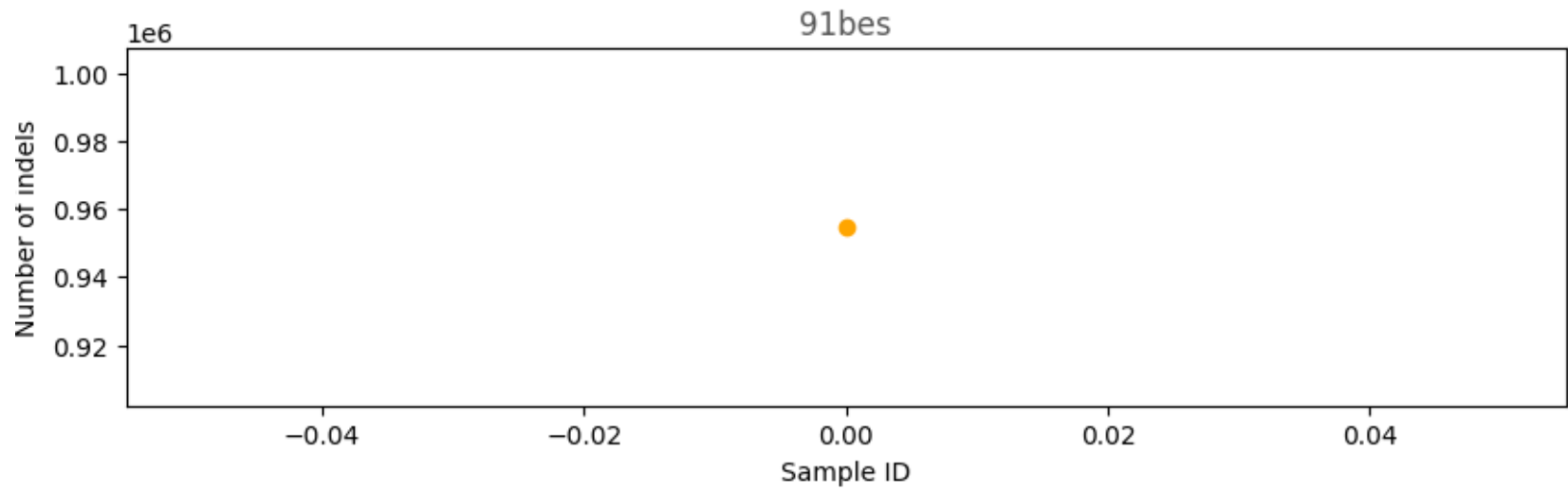91bes
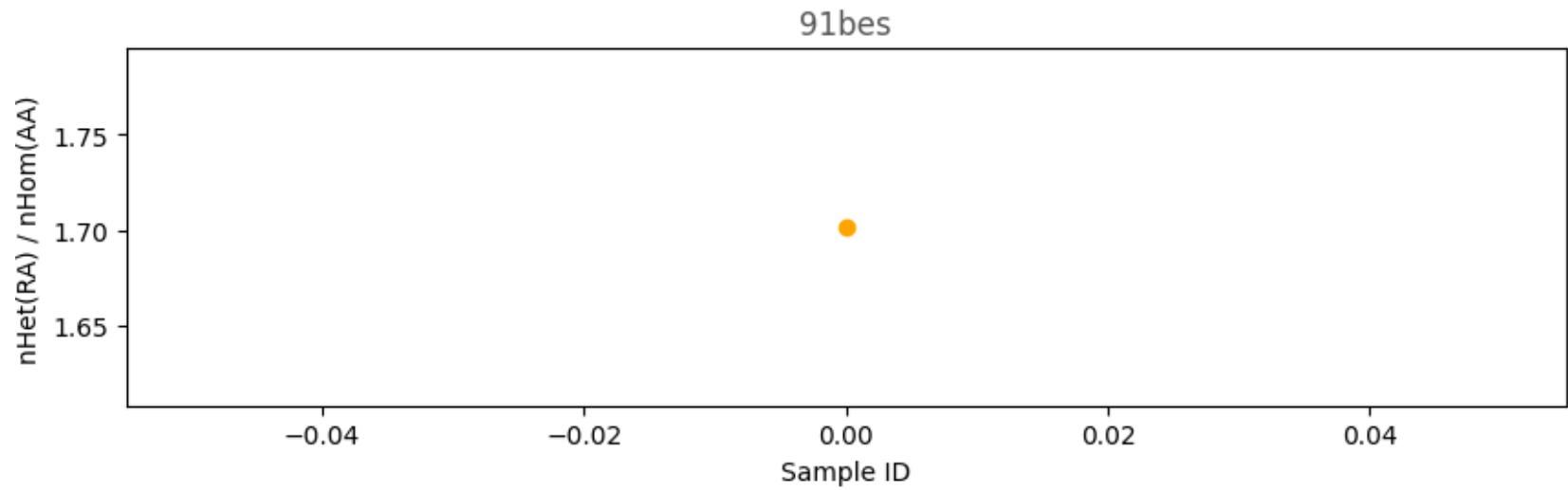
Singletons by sample (hets and homs)
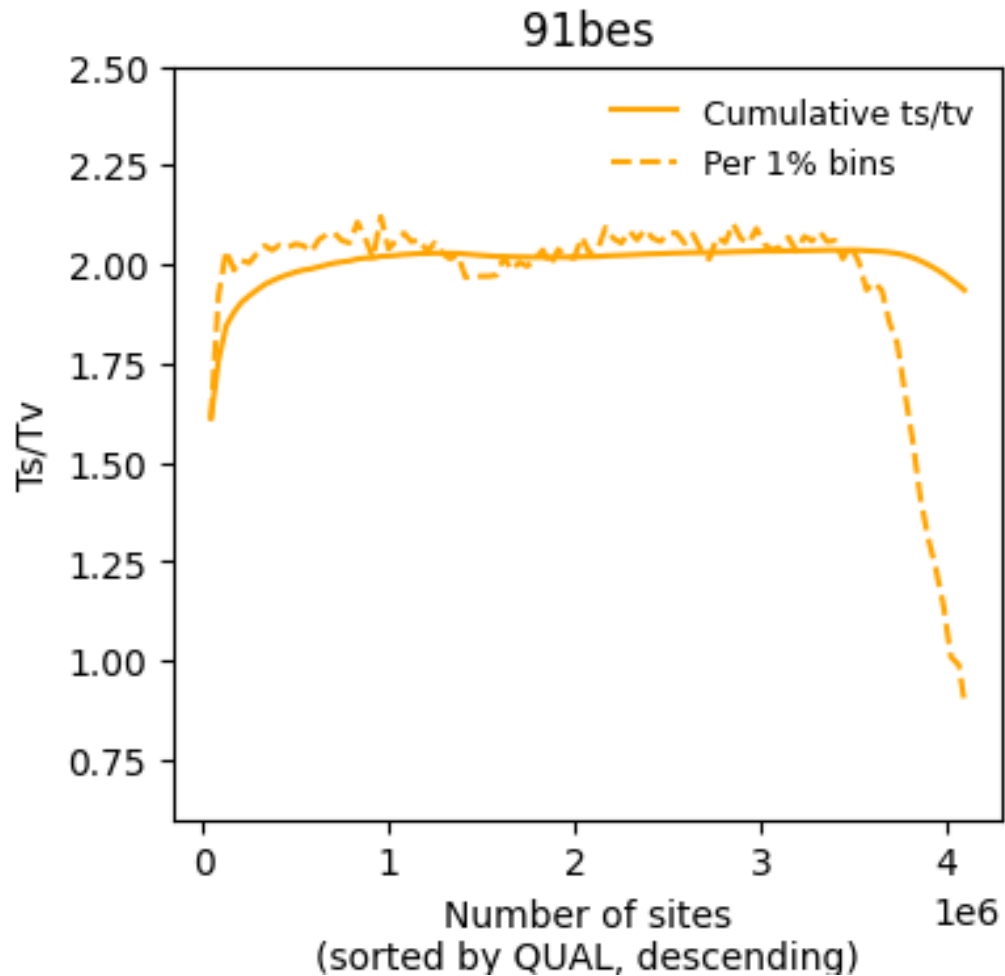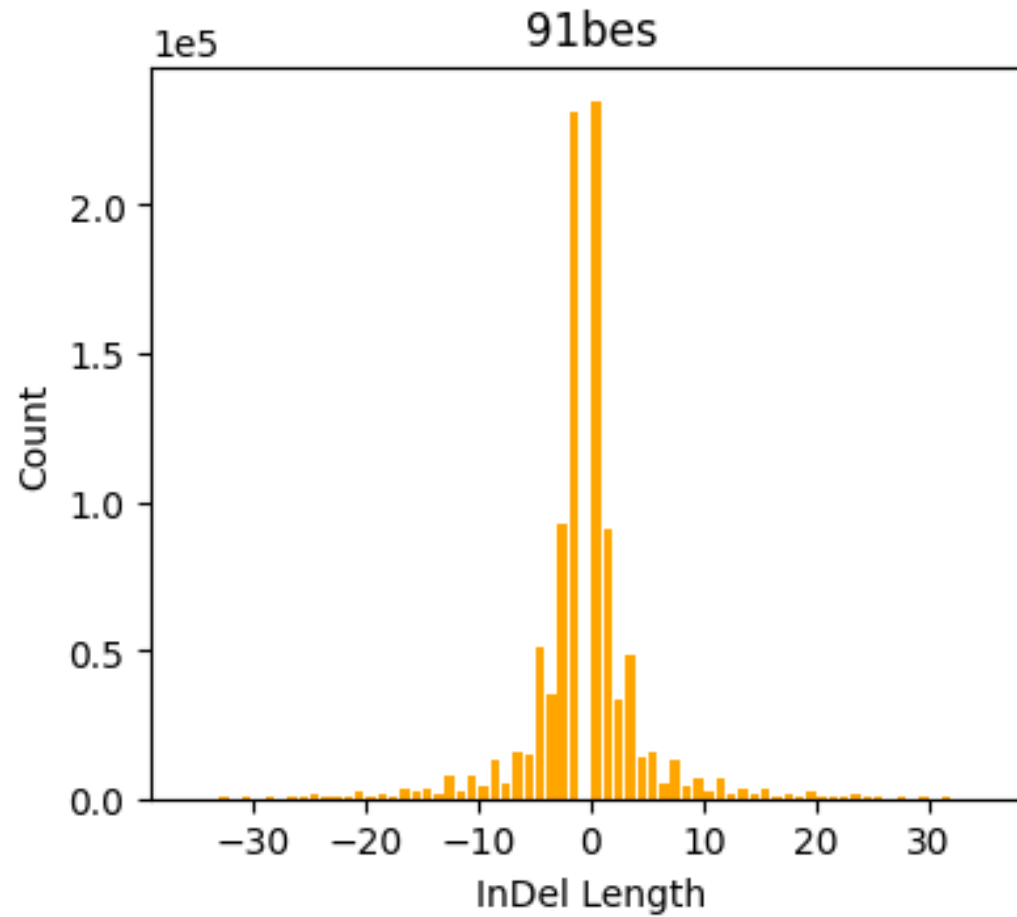
Average depth by sample
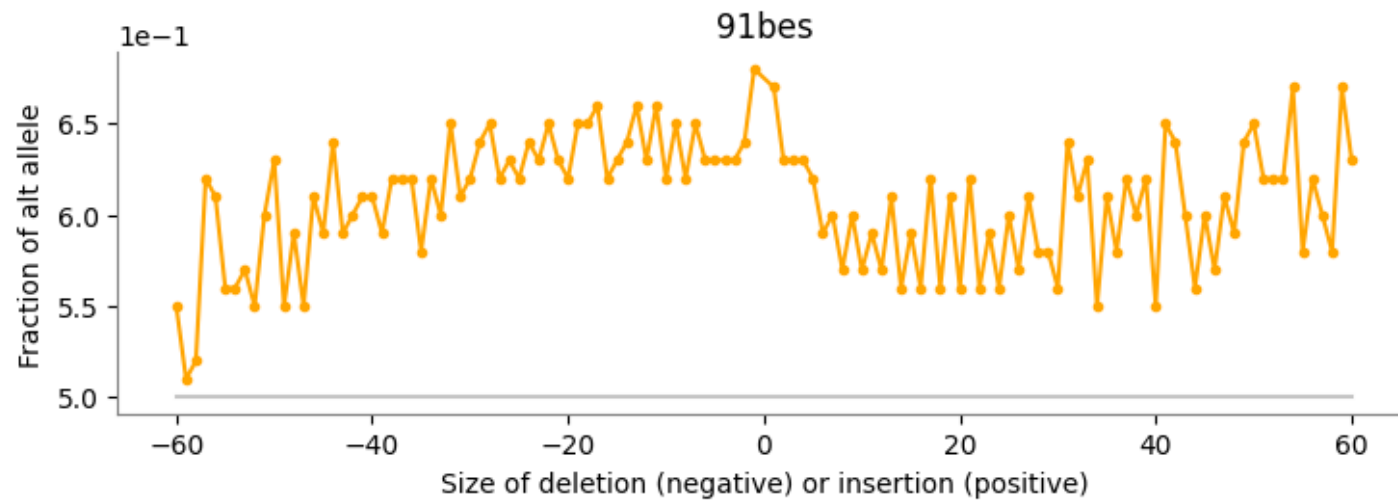
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

91bes

Indel distribution

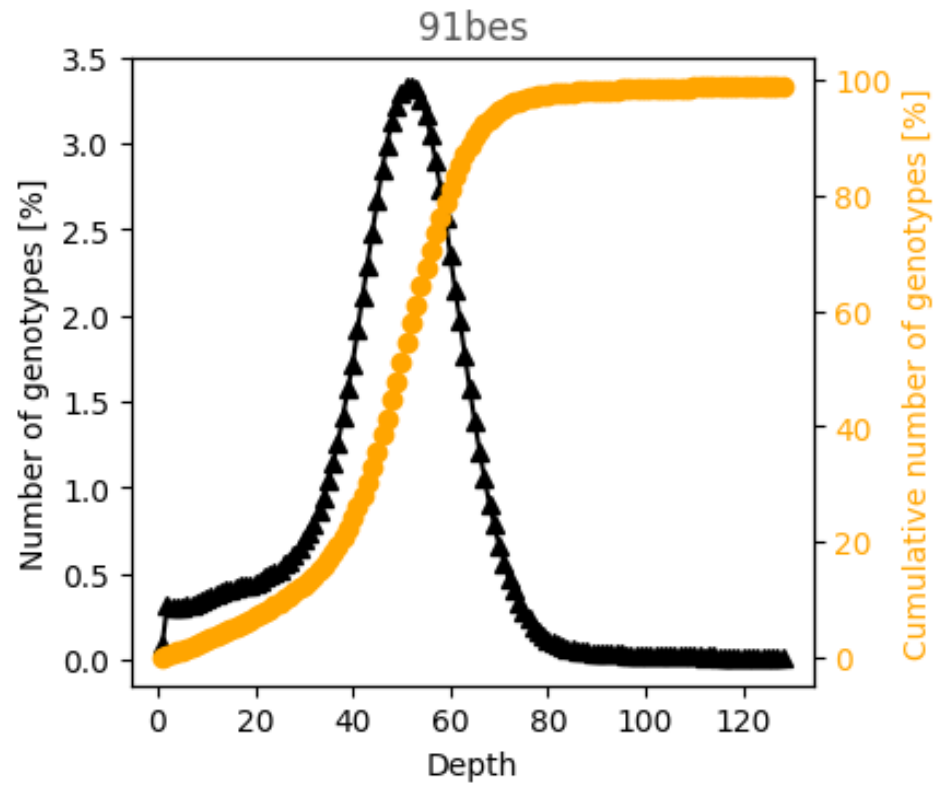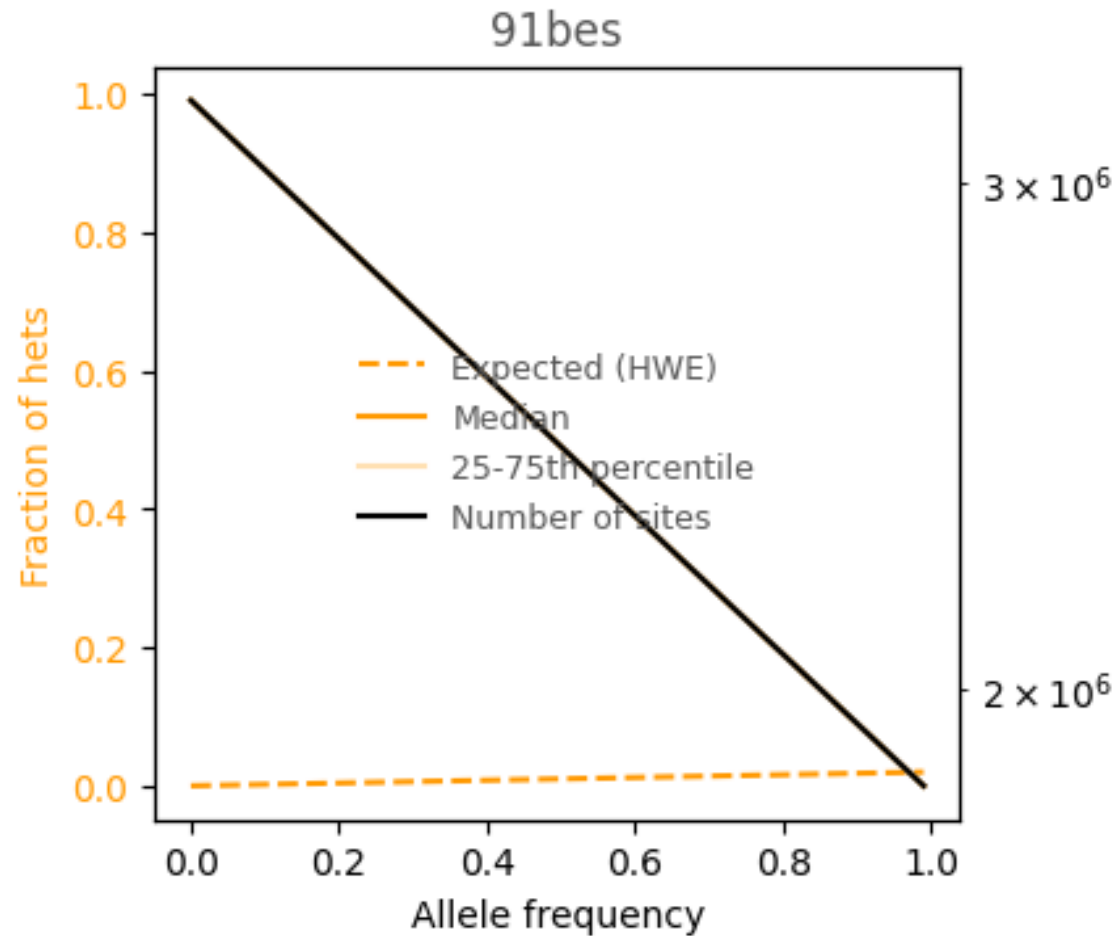# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types