Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78mar	4,173,072	1.93	1.93	976,365	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78mar	64.0%	1.90	69.7%	108,252	2,163

- 78mar .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78marsowf-103807551489-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00435-211206_A01466_AH2CT3DSX3-EXT_LAB`
  `KA_NGCWGS-NGCWGS03457_21RKG024433x01_snv_germline_raw.haplotype_caller.vcf.gz`
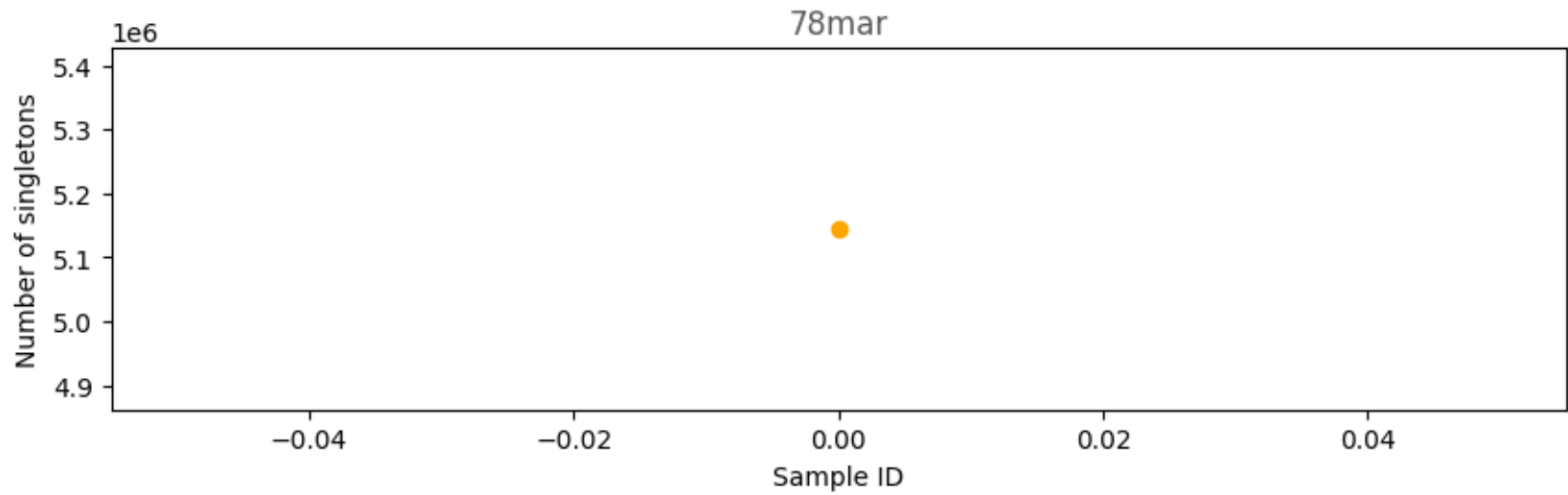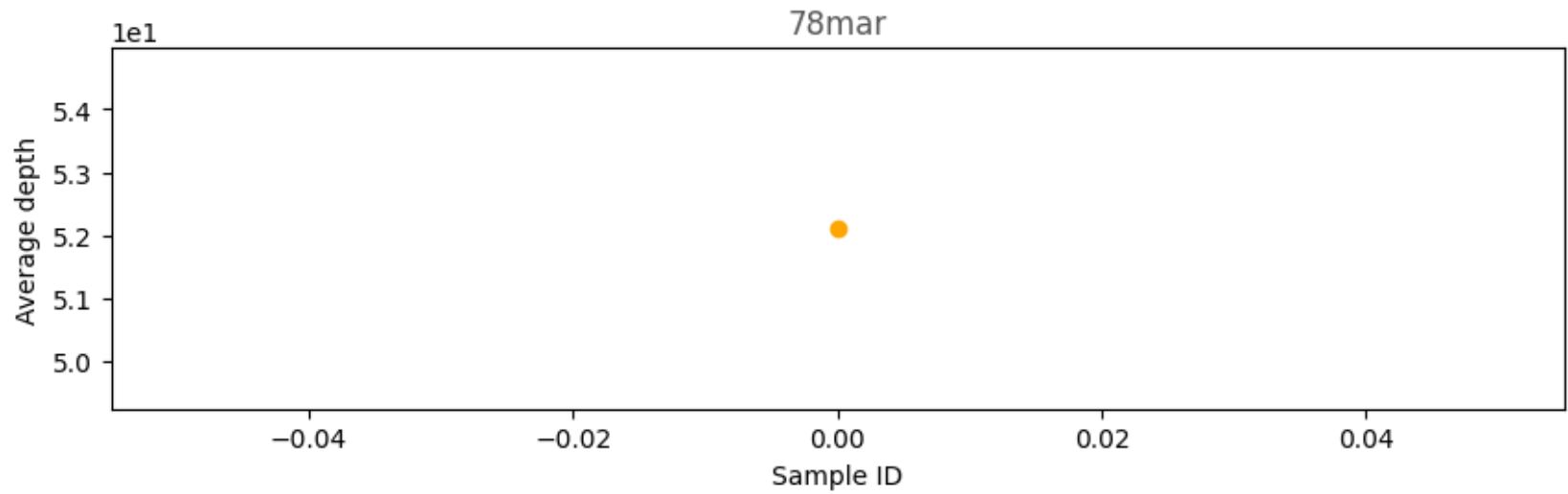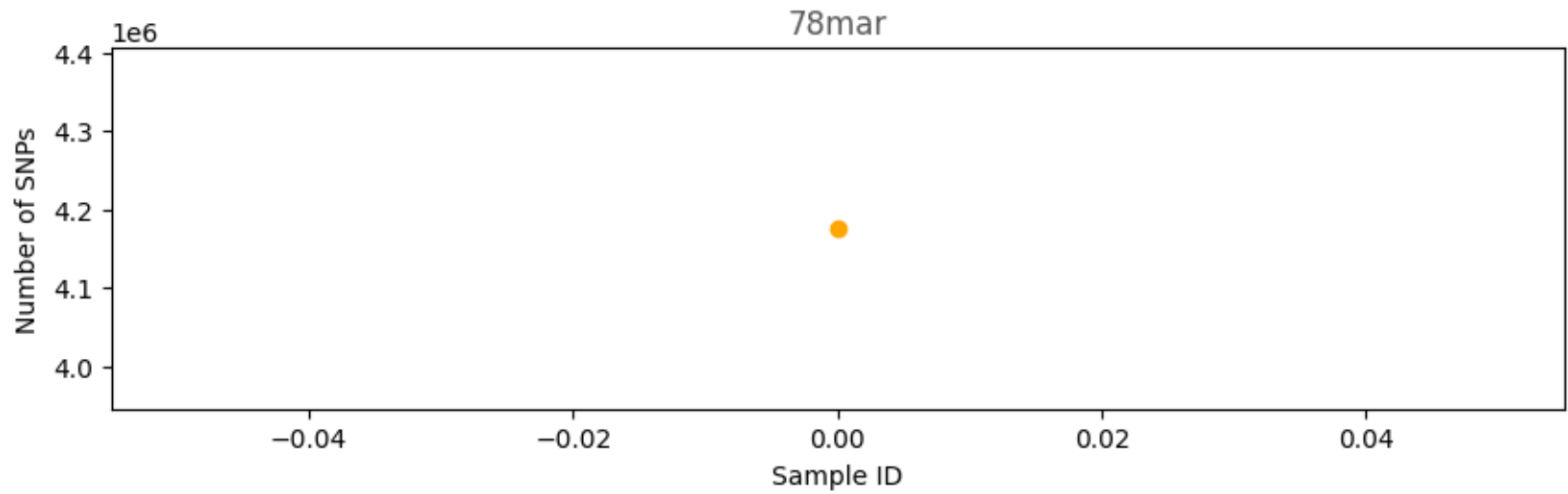
Ts/Tv by sample
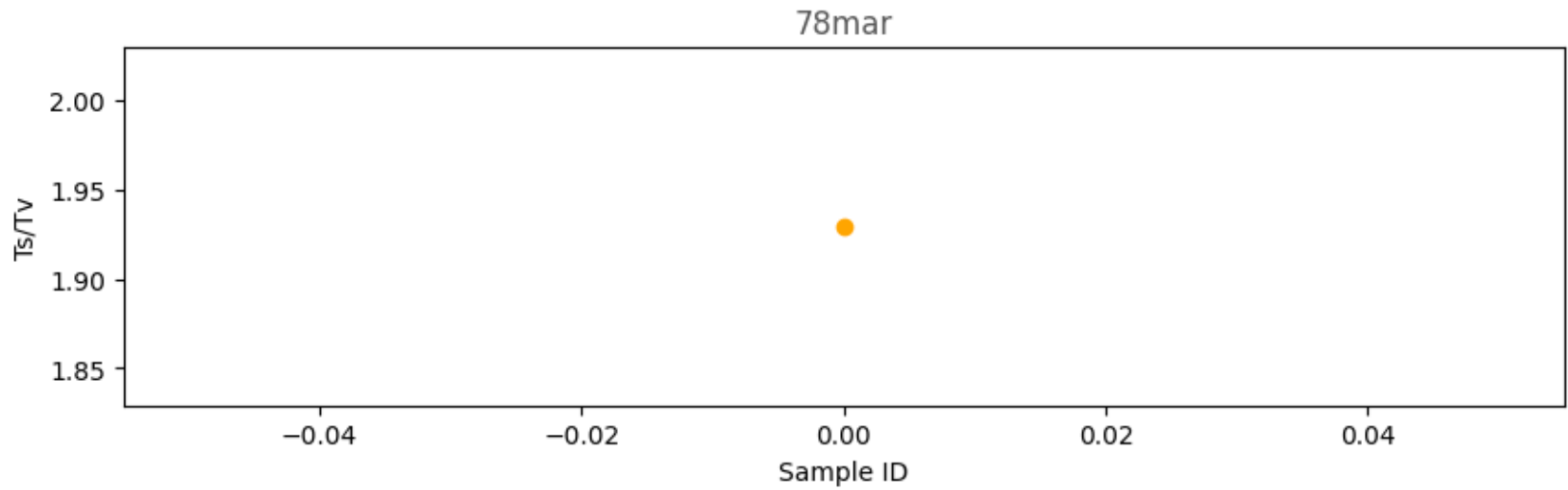
Hets vs non-ref Homs by sample
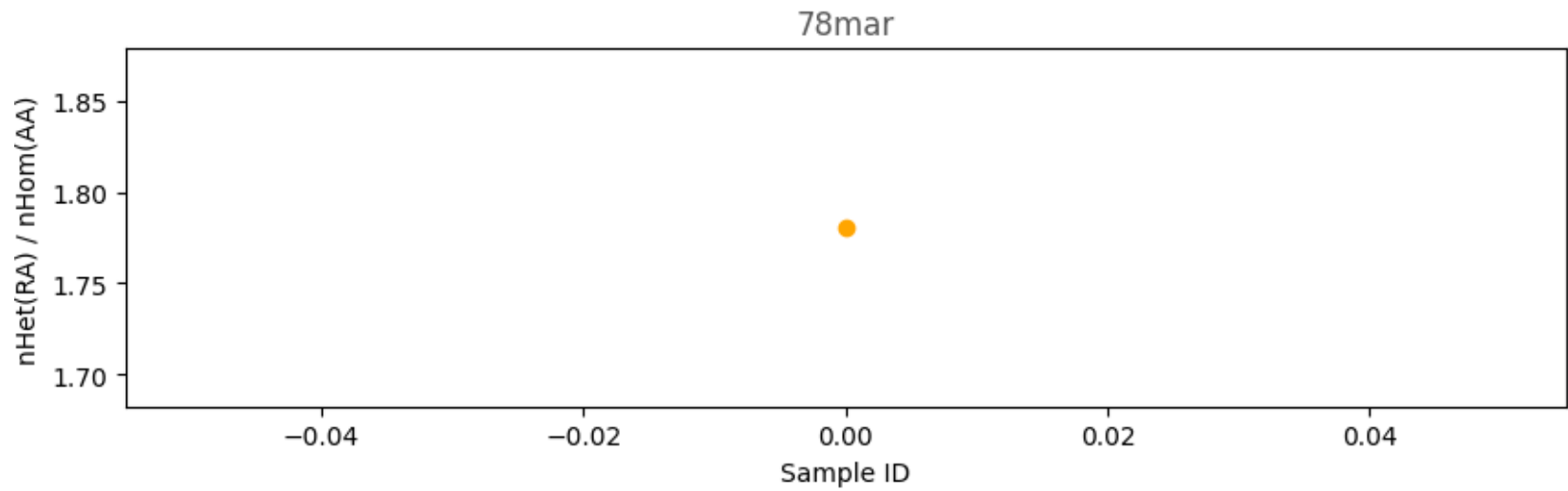
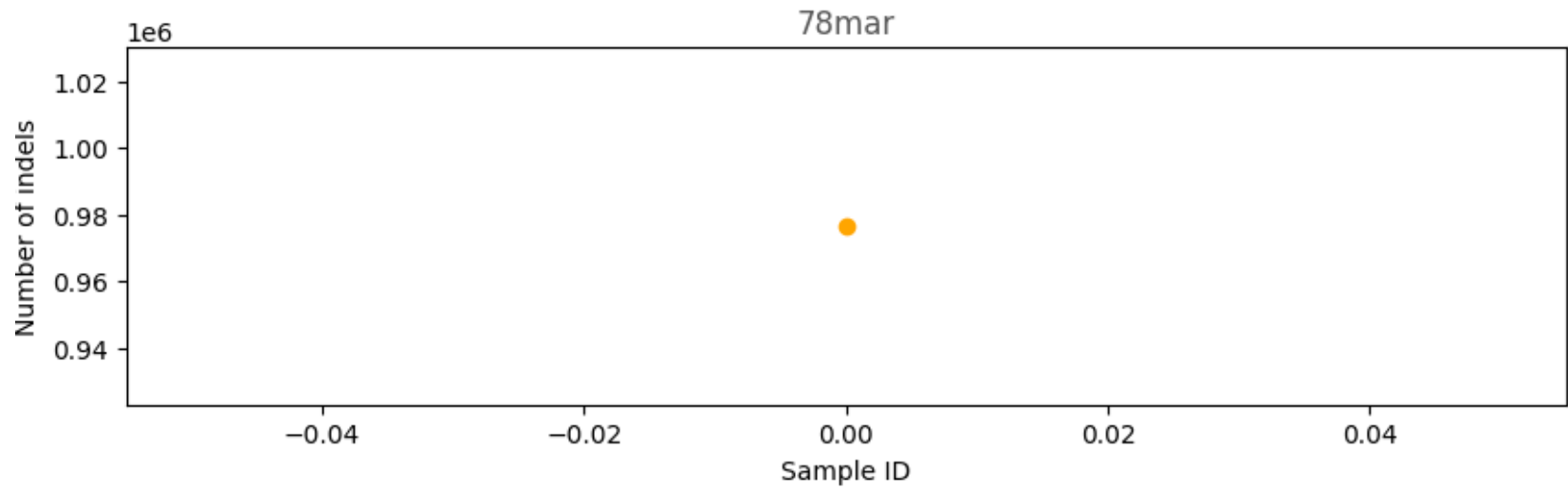# Singletons by sample (hets and homs)

Average depth by sample
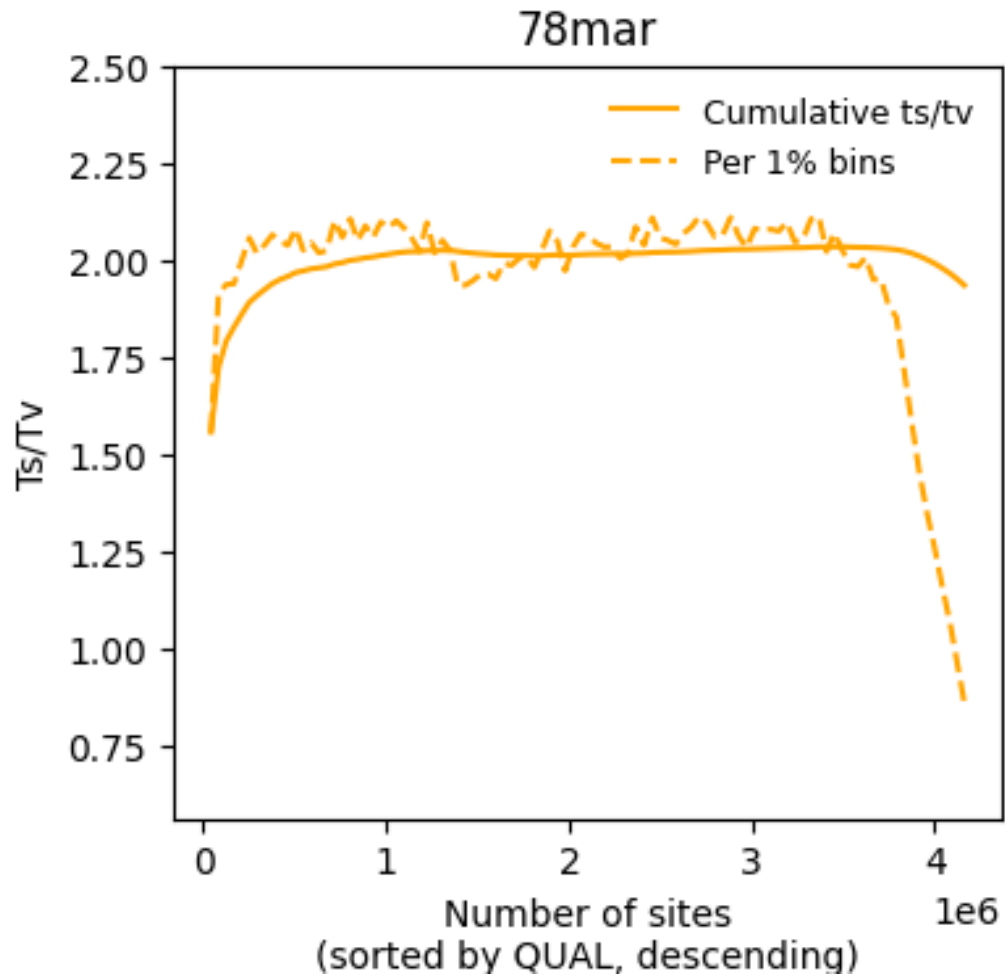
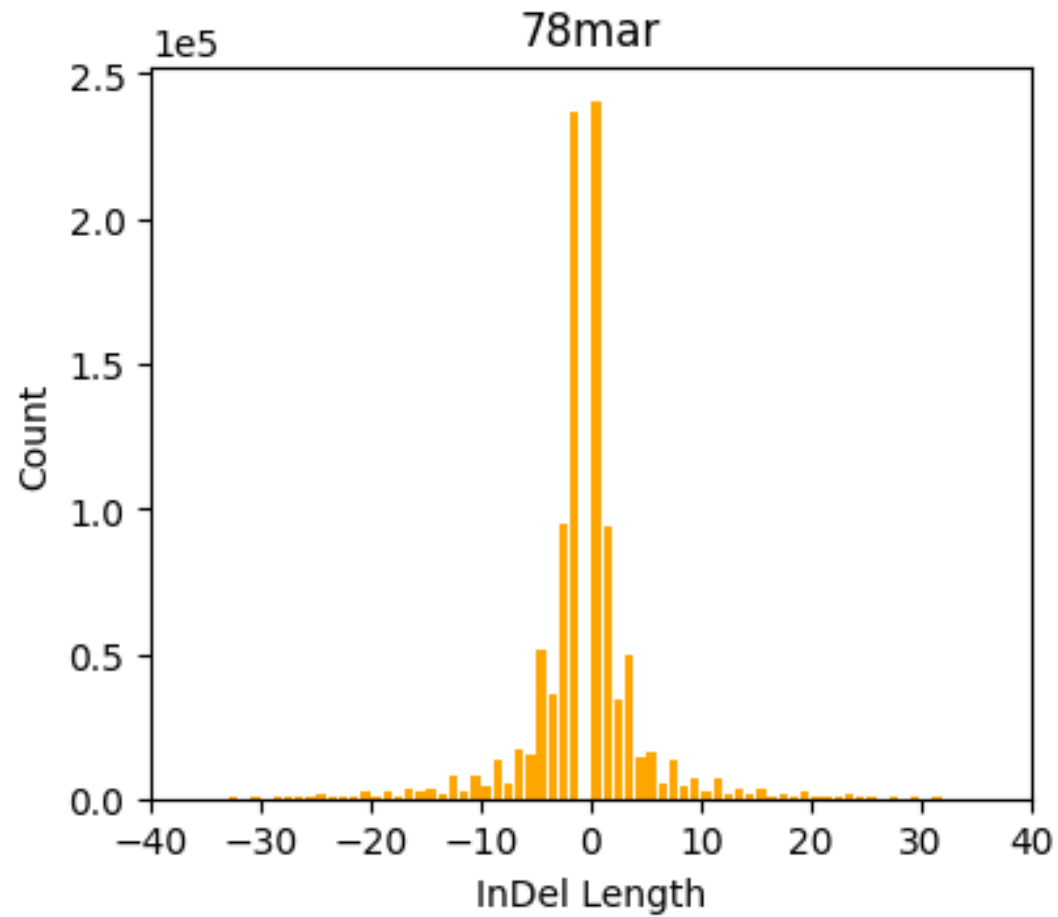# Number of SNPs by sample

Number of indels by sample
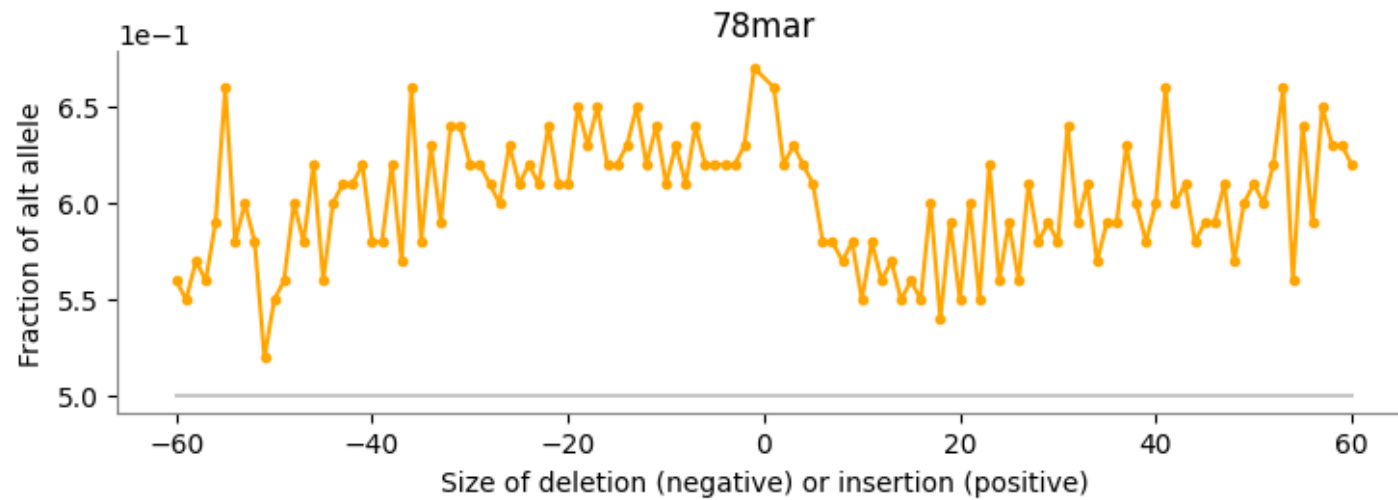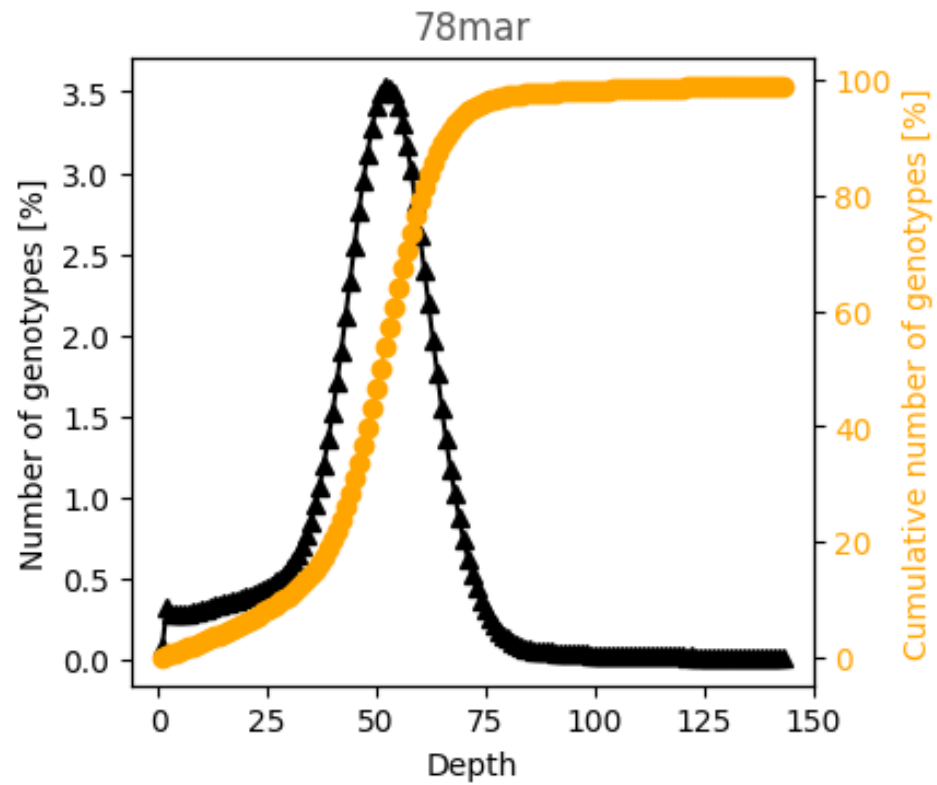
Ts/Tv stratified by QUAL
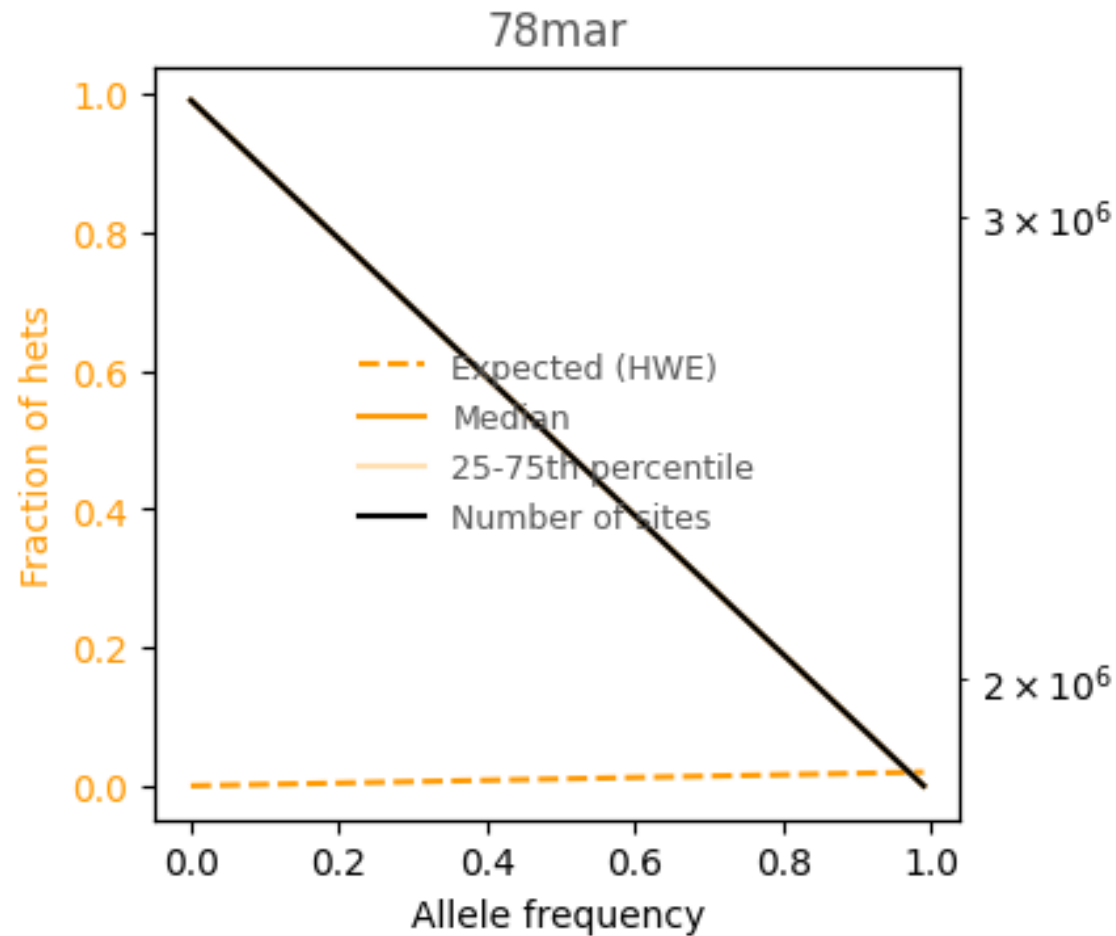
Indel distribution

78mar

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types