Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78ada	4,058,917	1.92	1.93	950,237	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78ada	59.3%	1.88	64.3%	95,163	2,164

- 78ada .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78adavinf-103907650175-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01270-220907_A01176_BHYN5VDSX3-EXT_LAB`
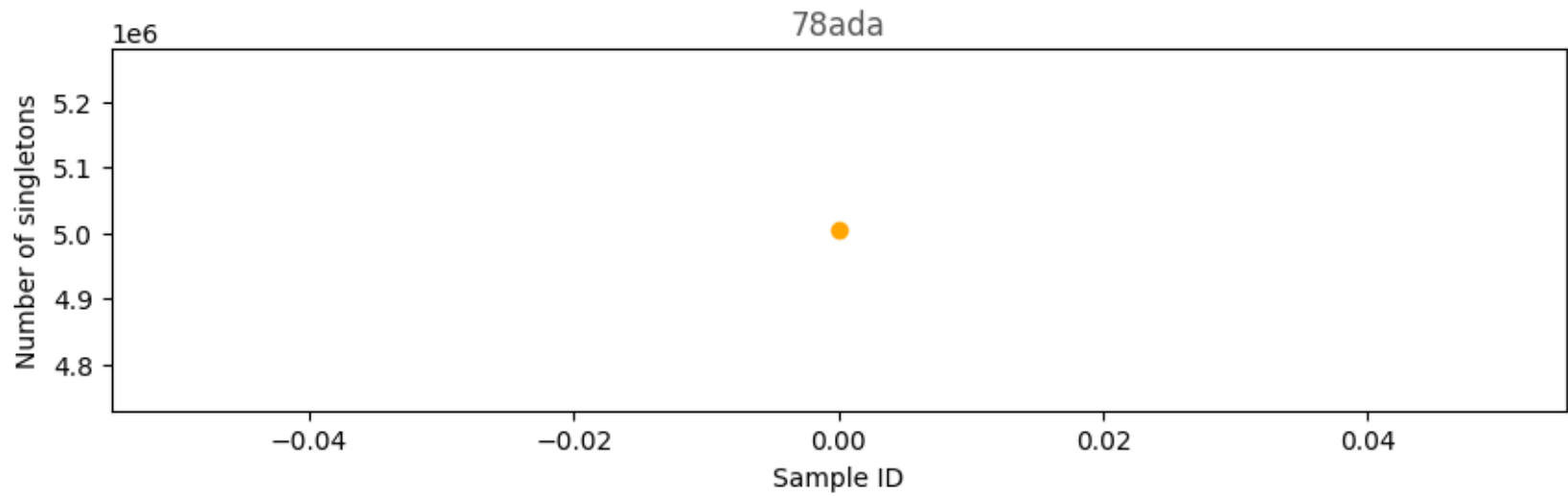 `KA_NGCWGS-NGCWGS05043_22RKG011401x01_snv_germline_raw.haplotype_caller.vcf.gz`
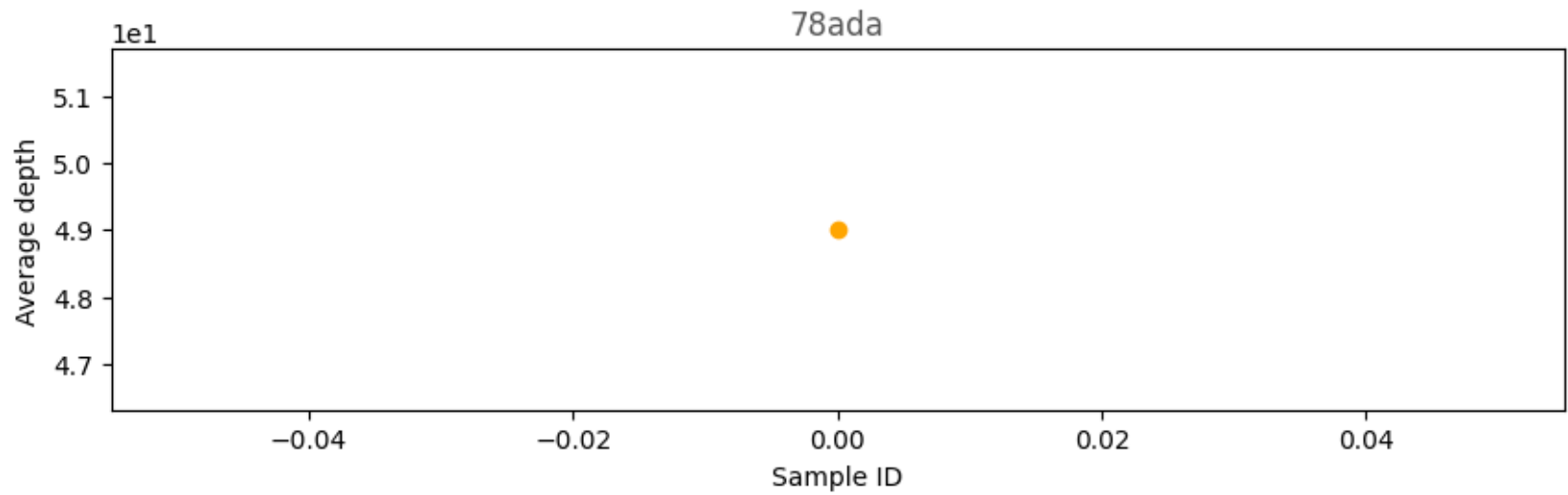
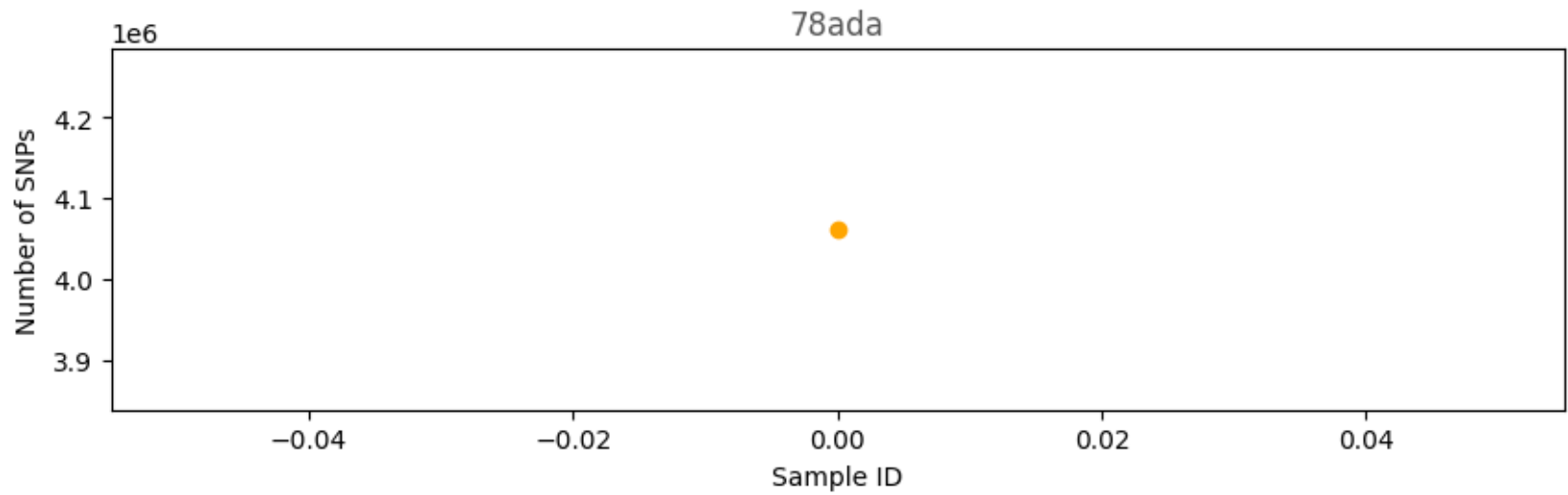# Ts/Tv by sample
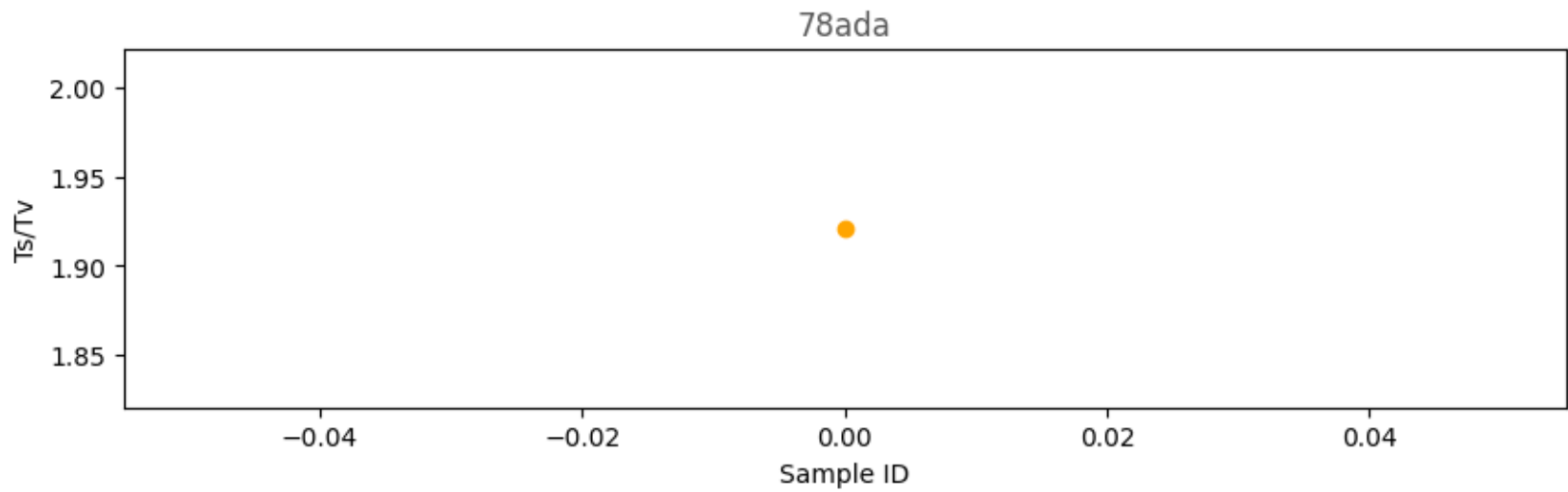
Hets vs non-ref Homs by sample
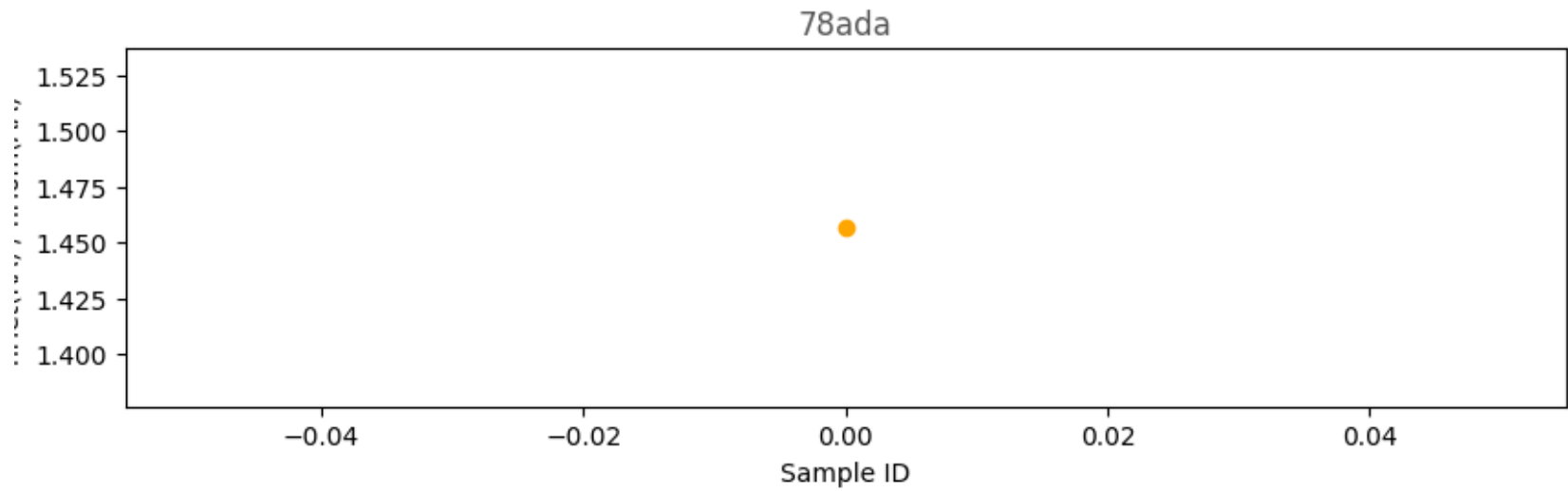
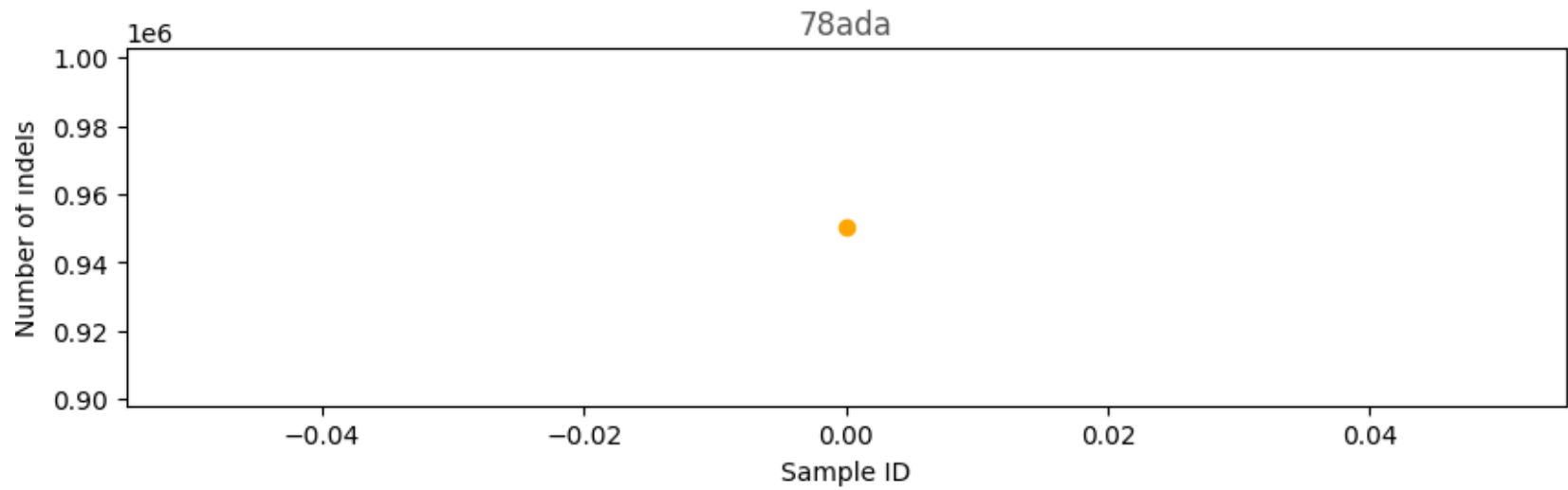# Singletons by sample (hets and homs)
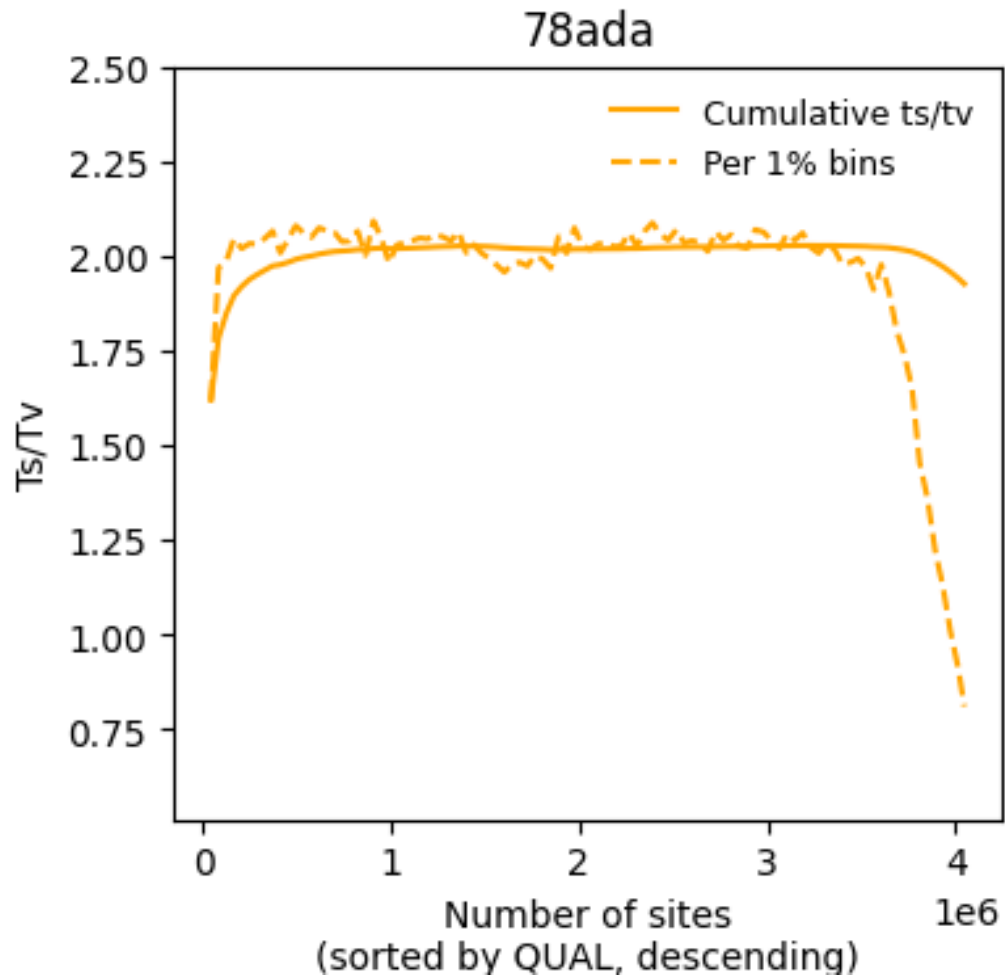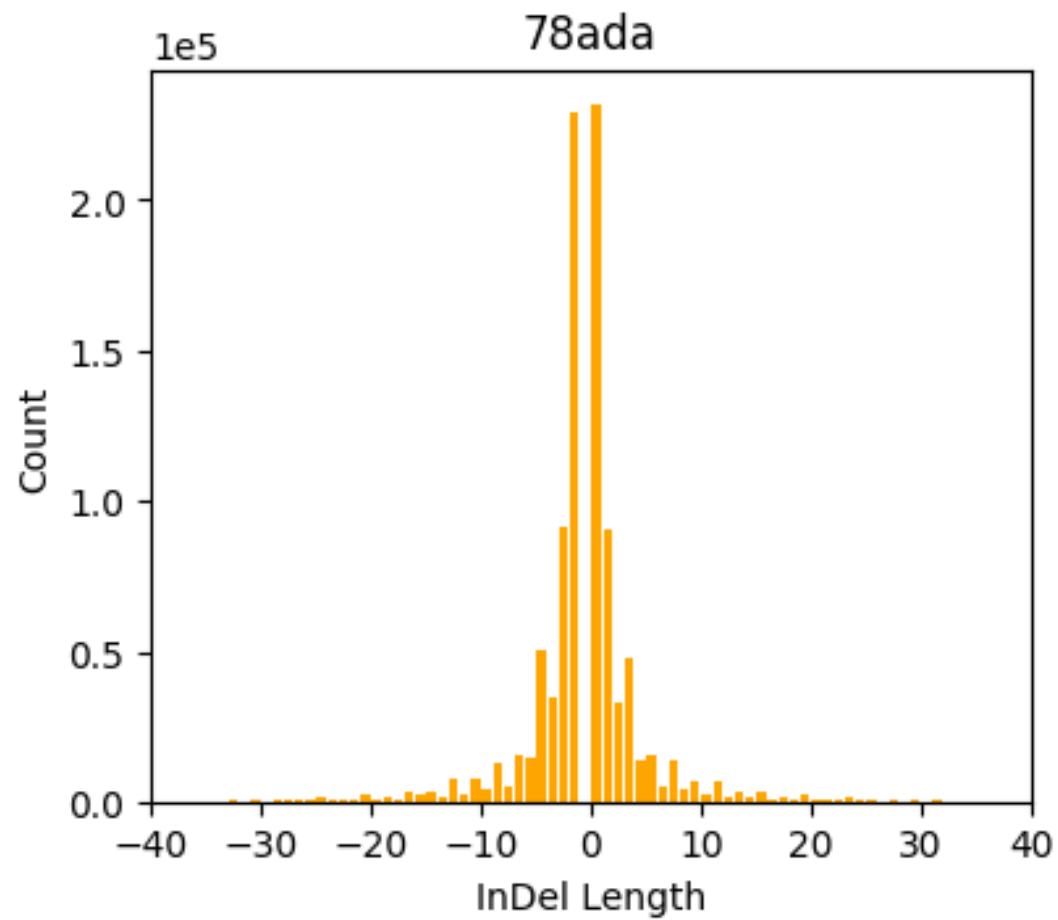
Average depth by sample
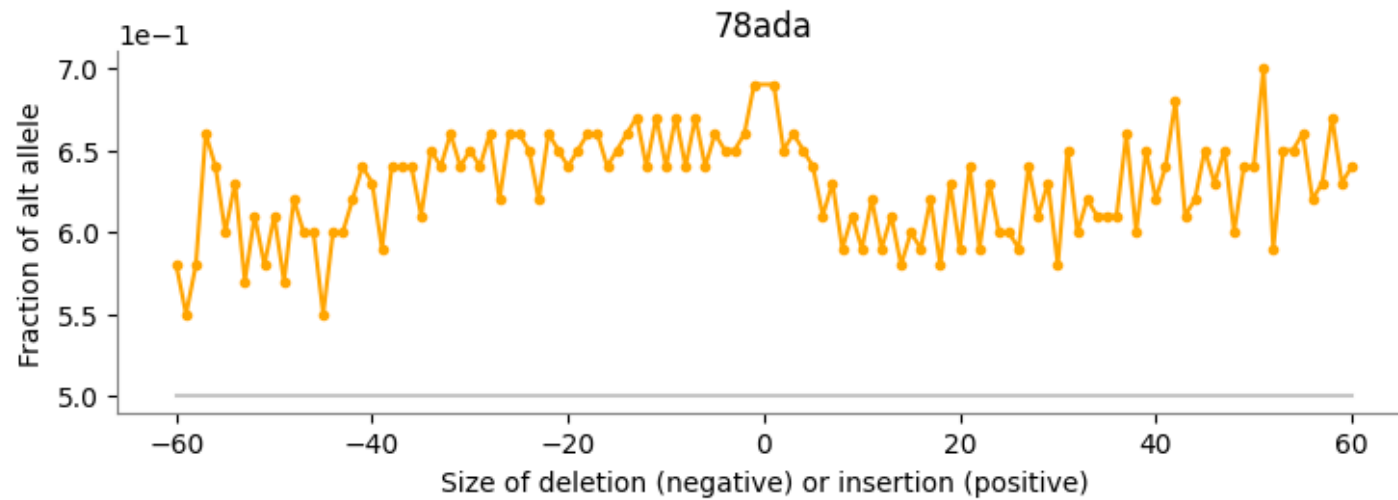
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

Indel distribution

78ada

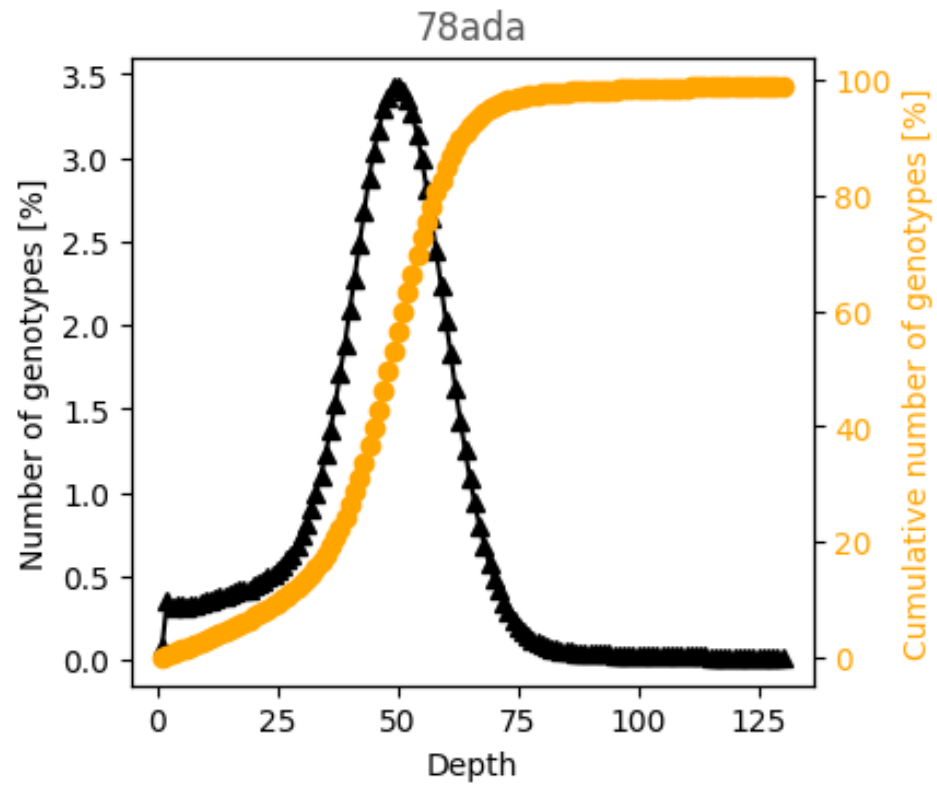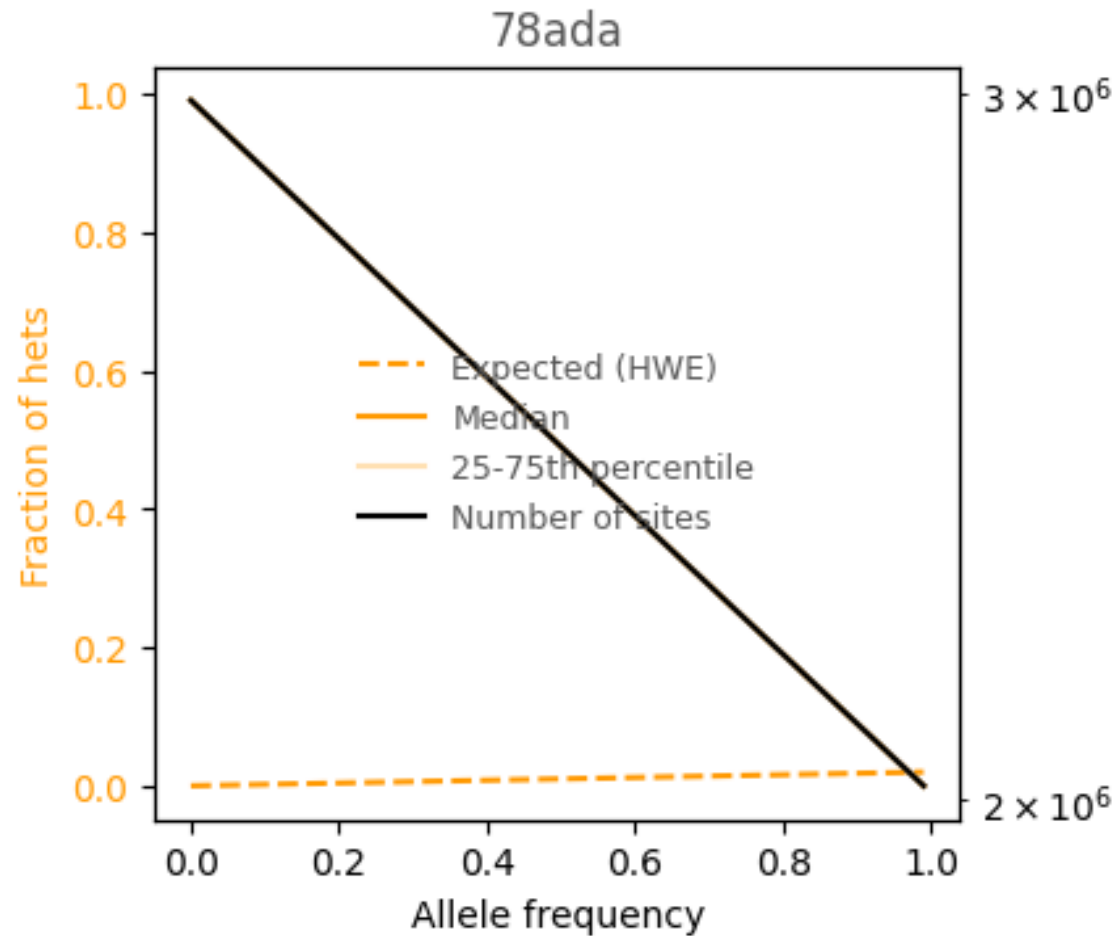# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types