Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
77twe	4,063,273	1.93	1.94	944,714	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
77twe	62.1%	1.90	66.8%	97,635	2,093

- 77twe .. `/ngc/projects2/gm/data/archive/2022/variants/snv/77twelltm-110305470116-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02377-221116_A01176_BHCYHYDSX5-EXT_LAB`
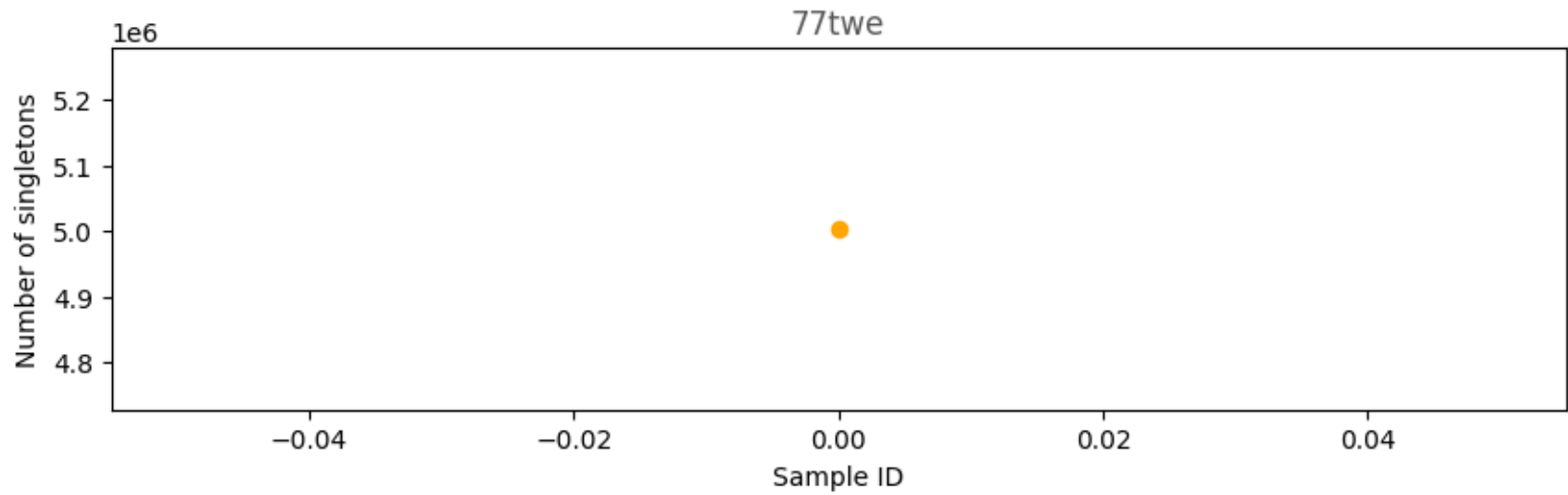 `KA_NGCWGS-NGCWGS06277_snv_germline_raw.haplotype_caller.vcf.gz`
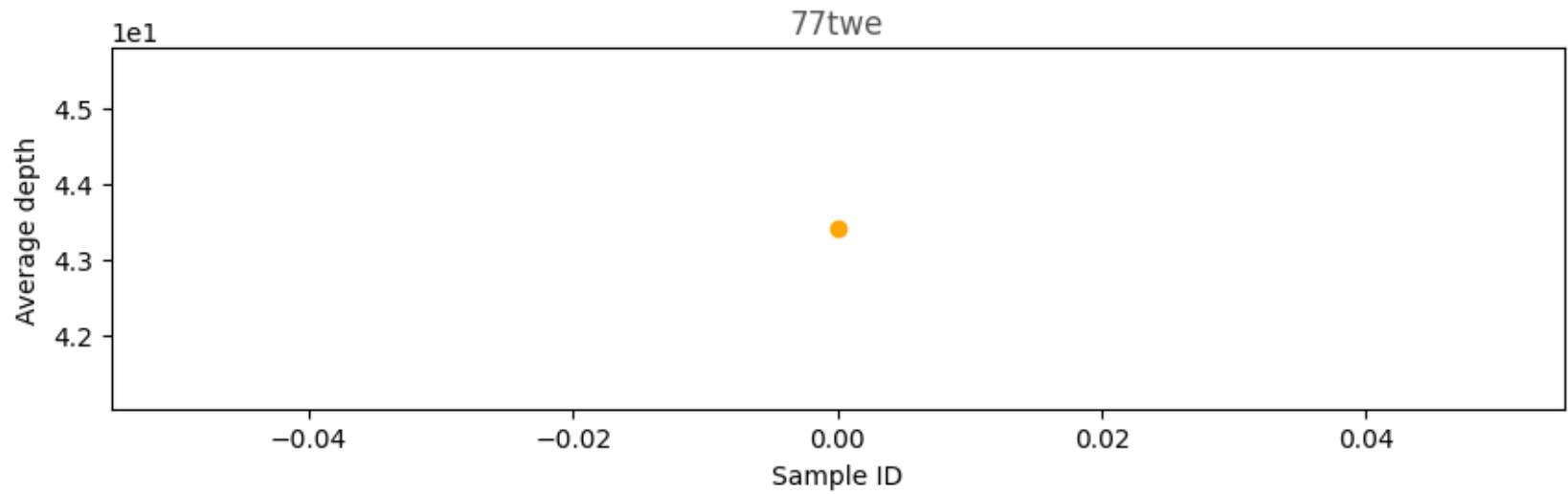
Ts/Tv by sample
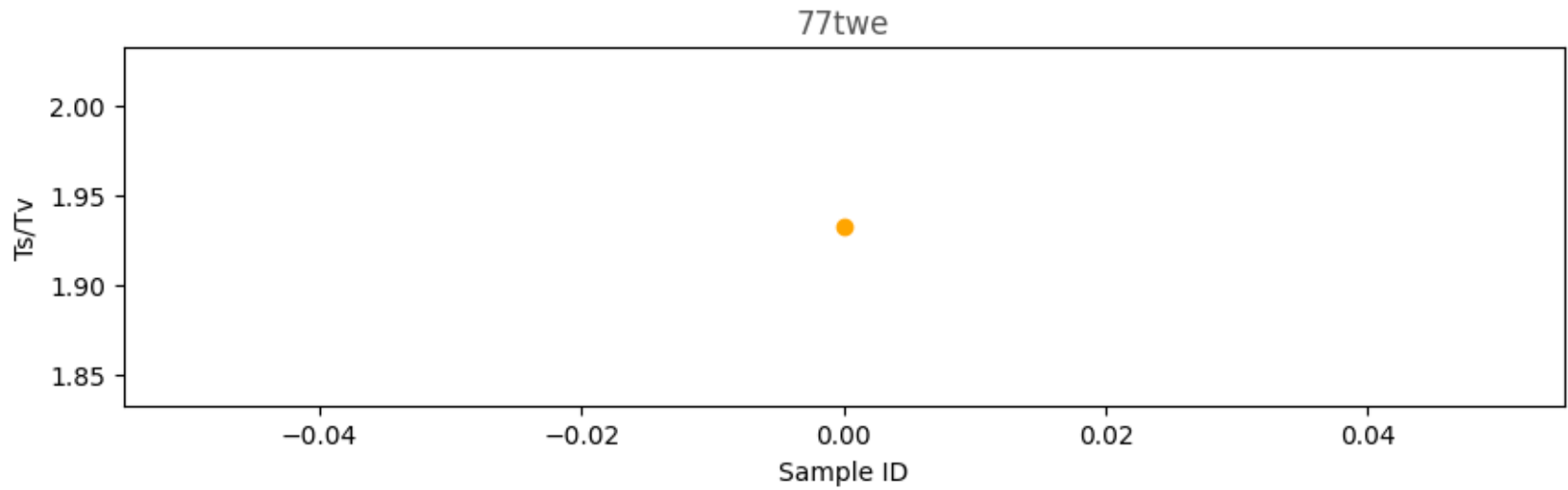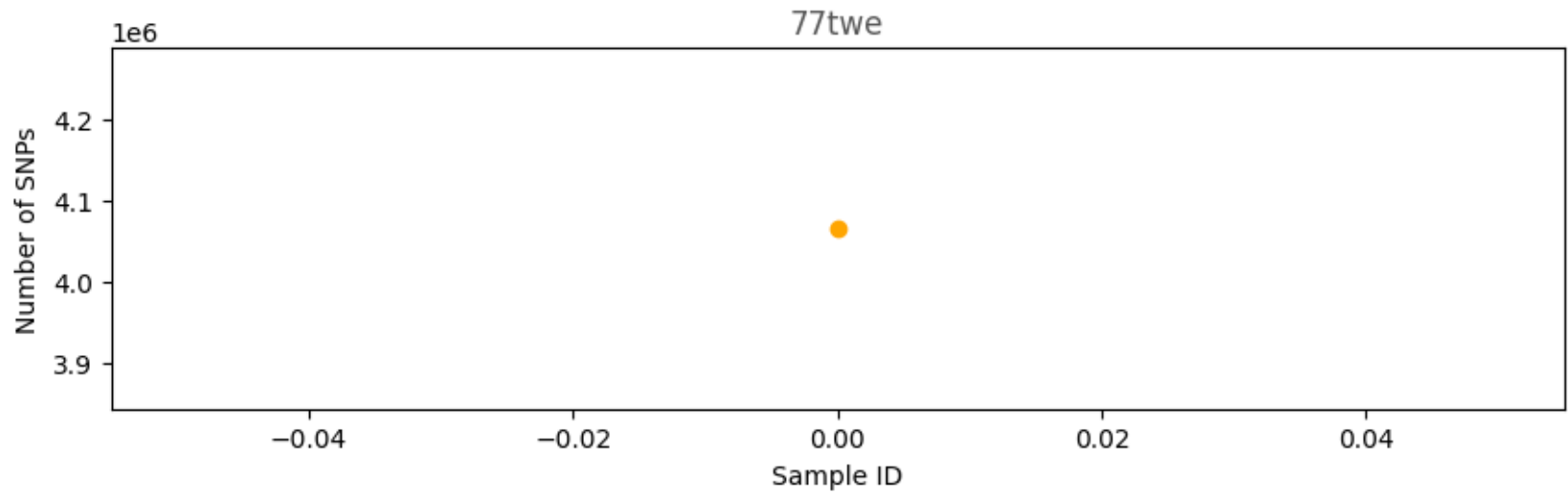
Hets vs non-ref Homs by sample
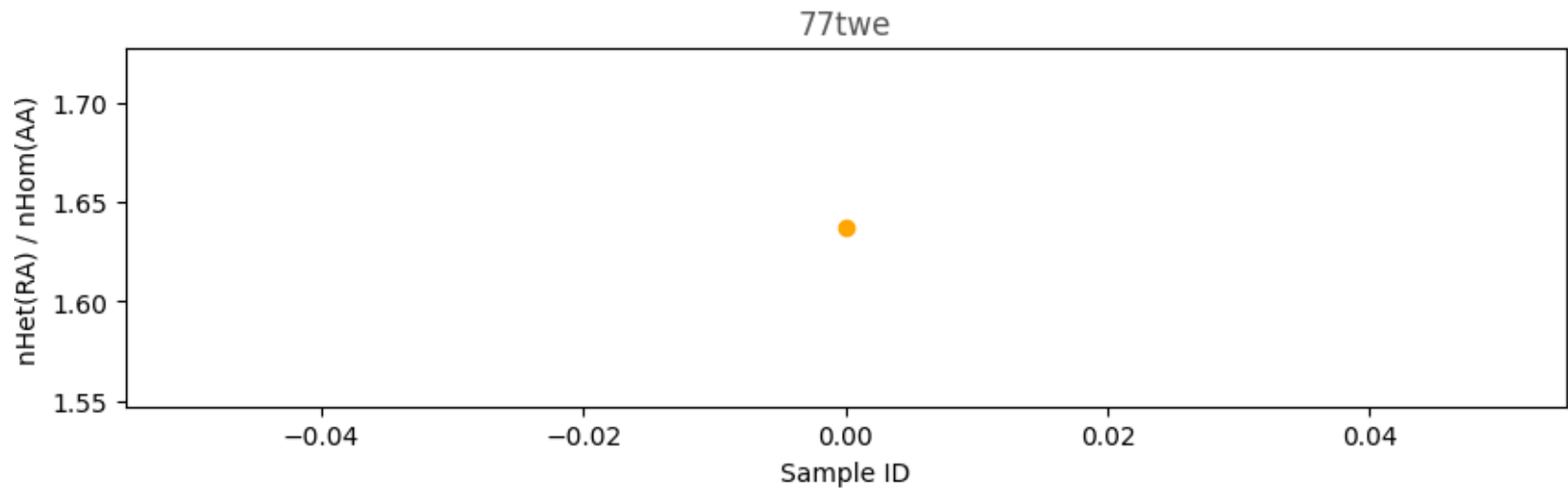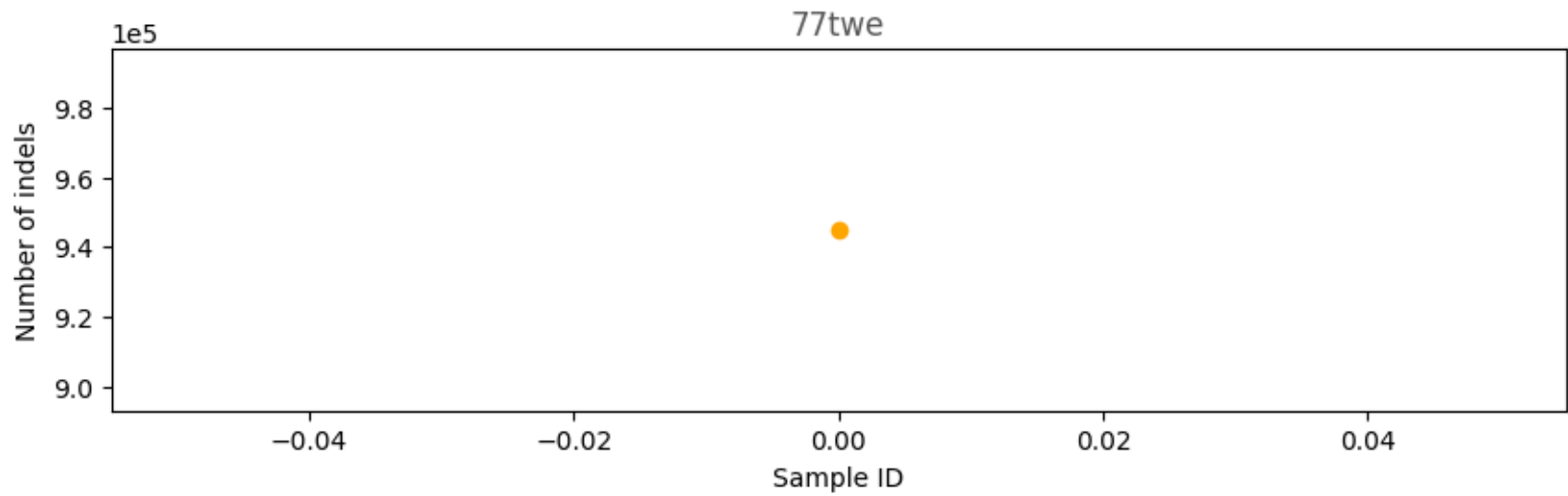
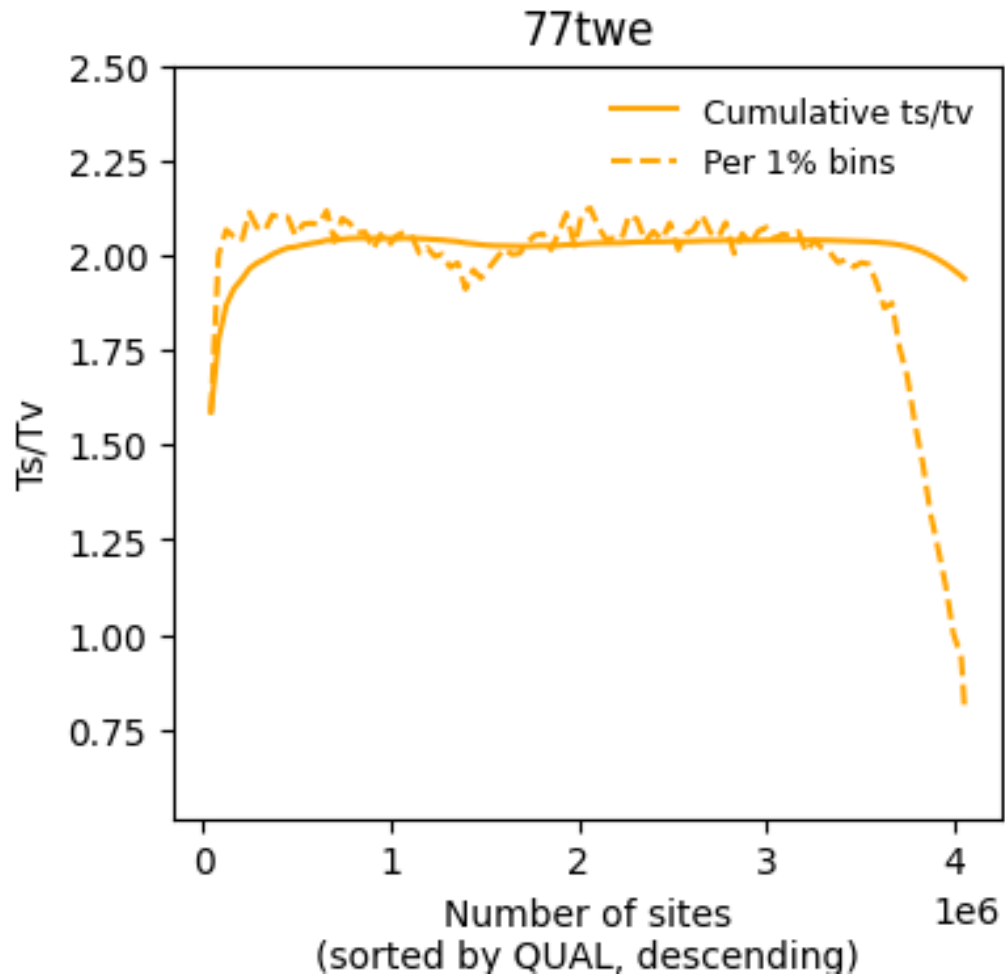# Singletons by sample (hets and homs)

Average depth by sample
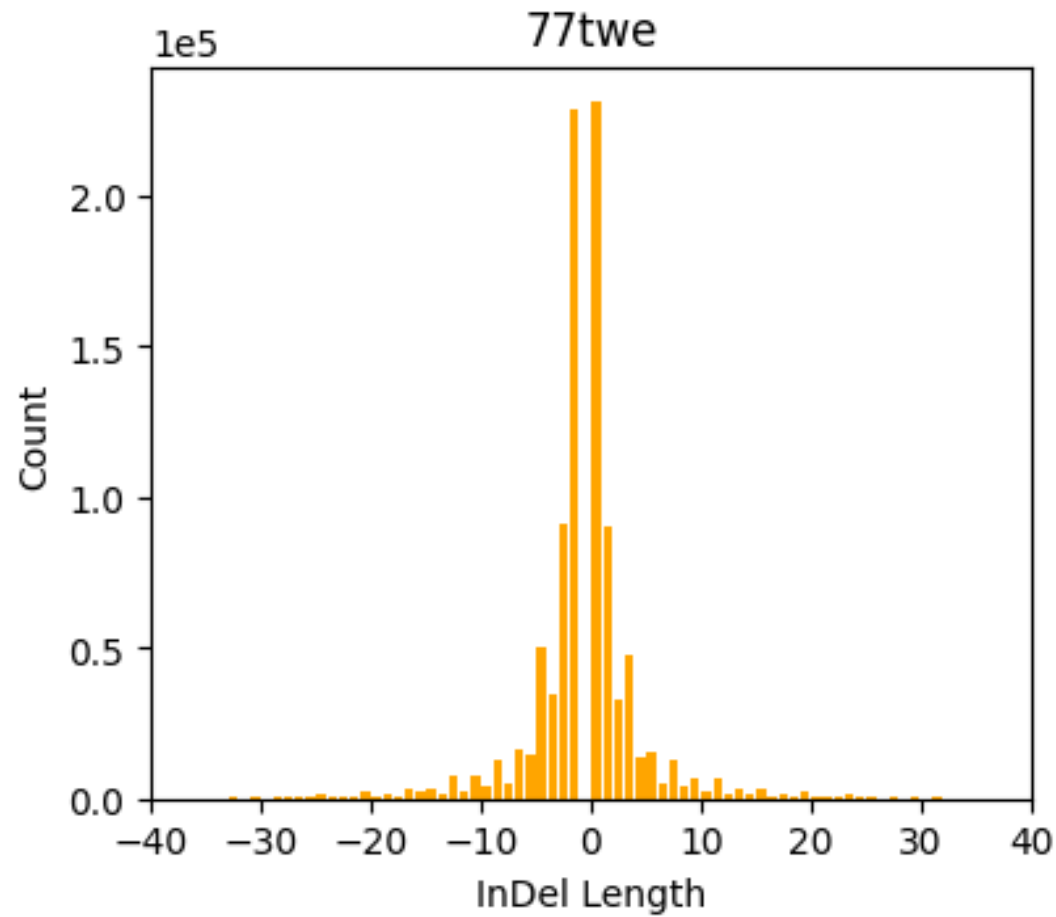
Number of SNPs by sample

Number of indels by sample
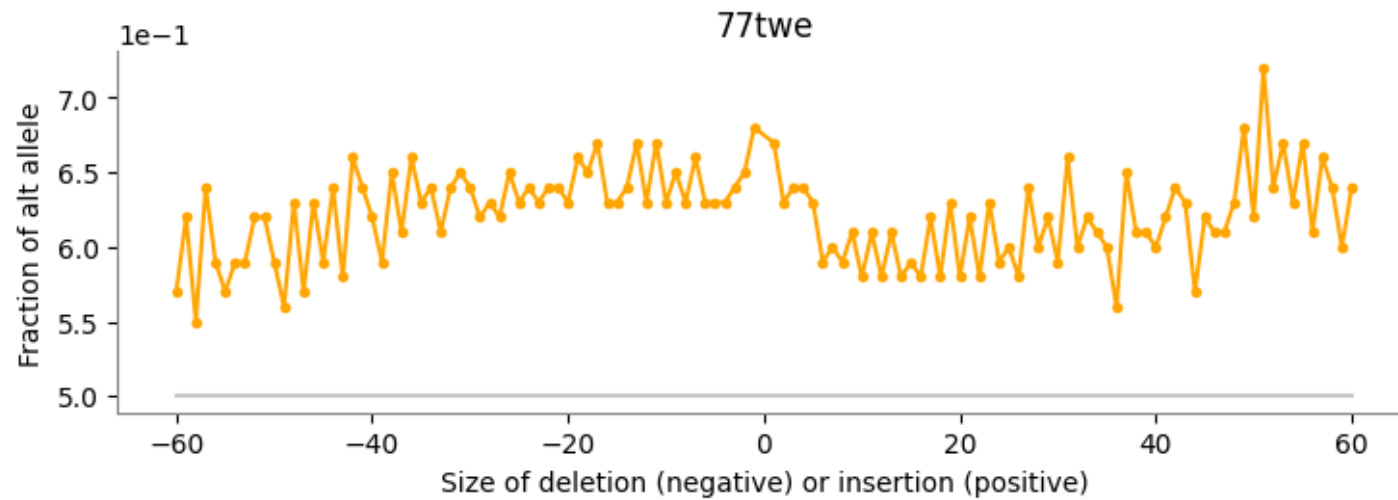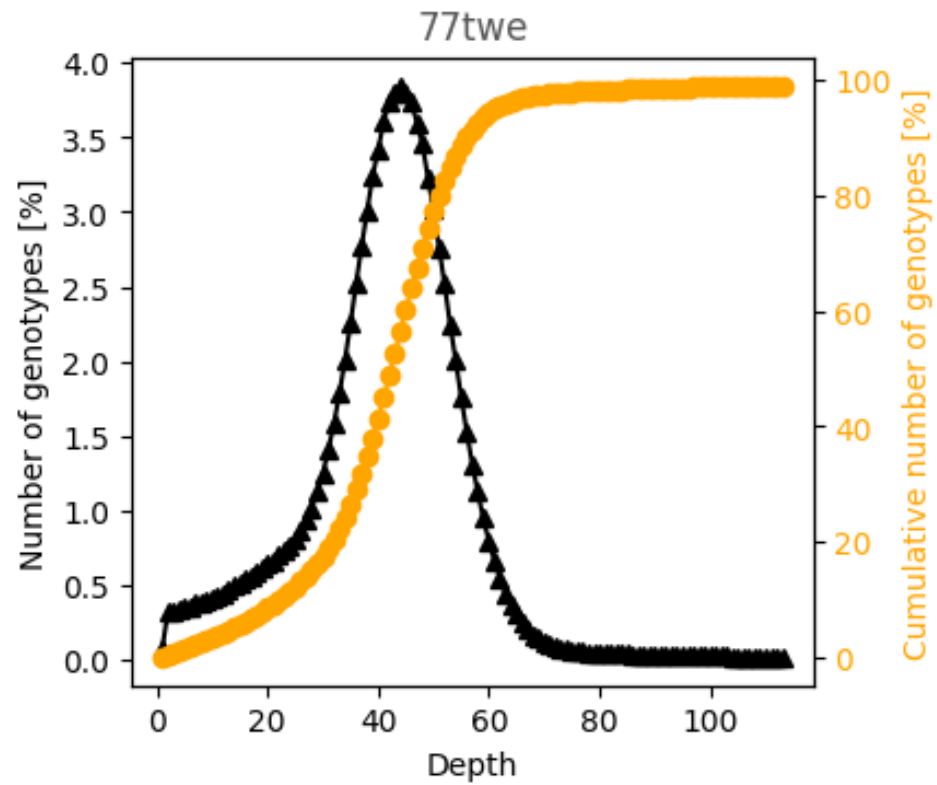
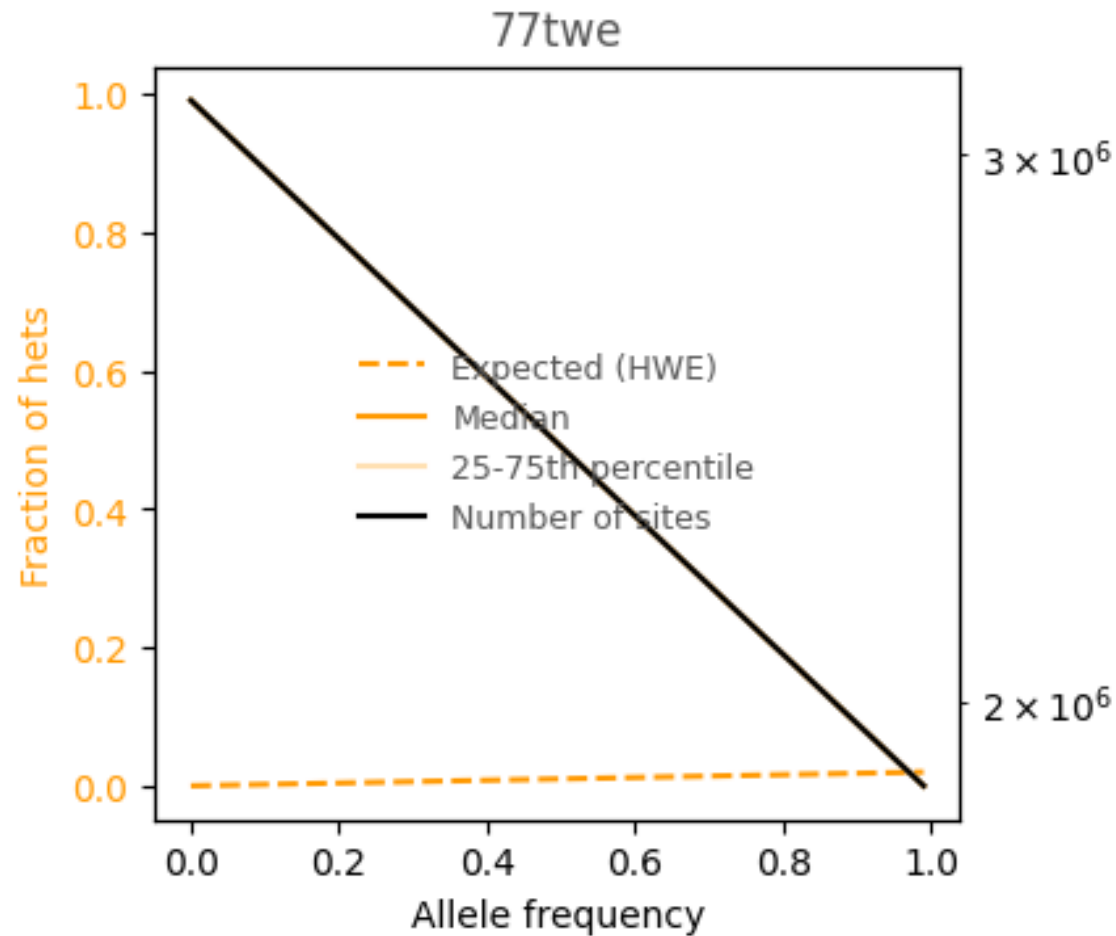# Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types