Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
17ave	4,195,238	1.94	1.95	979,580	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
17ave	64.4%	1.92	69.6%	108,616	2,368

- 17ave .. `/ngc/projects2/gm/data/archive/2022/variants/snv/17averfff-103913449613-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01755-220923_A01961_AH33KNDSX5-EXT_LAB`
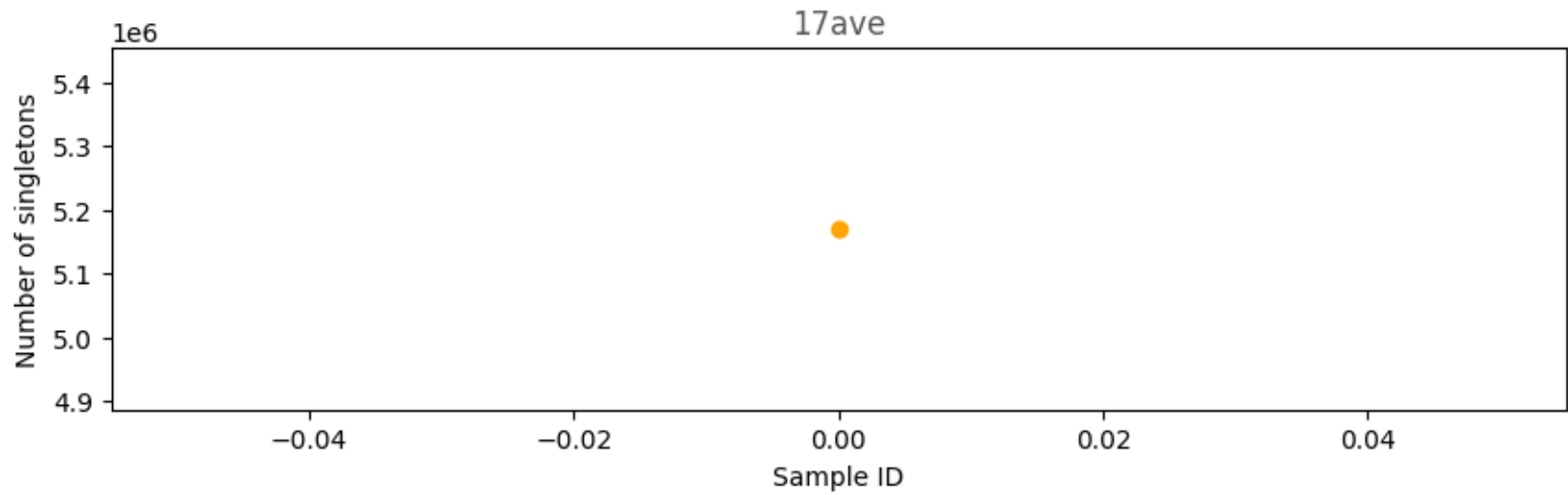 `KA_NGCWGS-NGCWGS05517_22RKG015807_snv_germline_raw.haplotype_caller.vcf.gz`
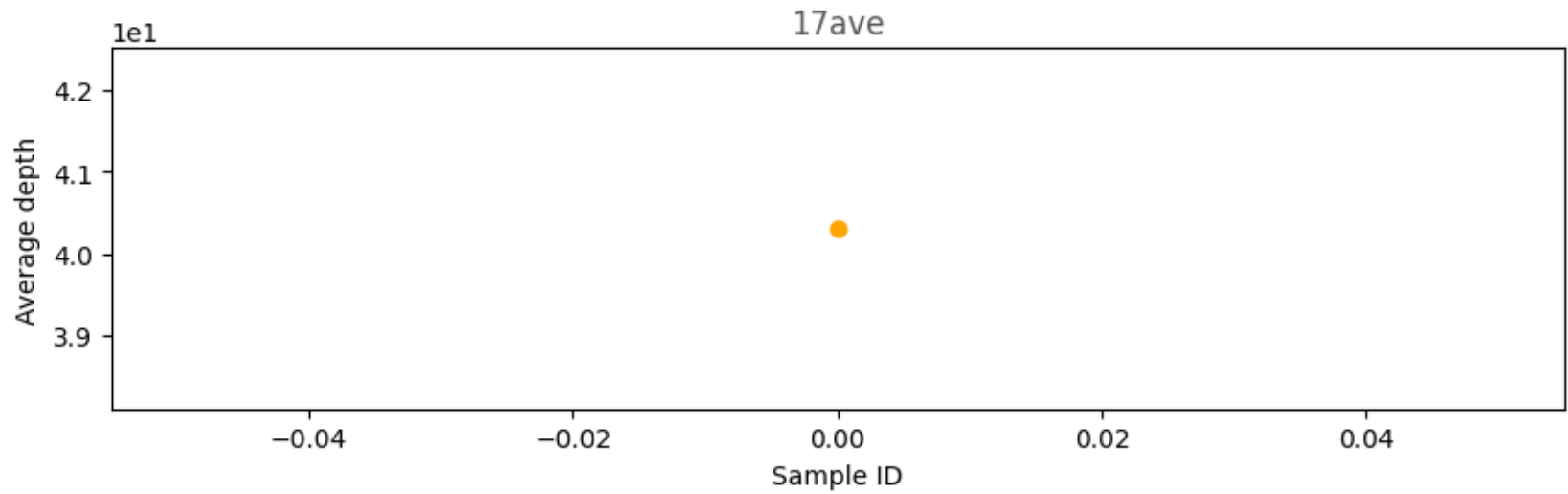
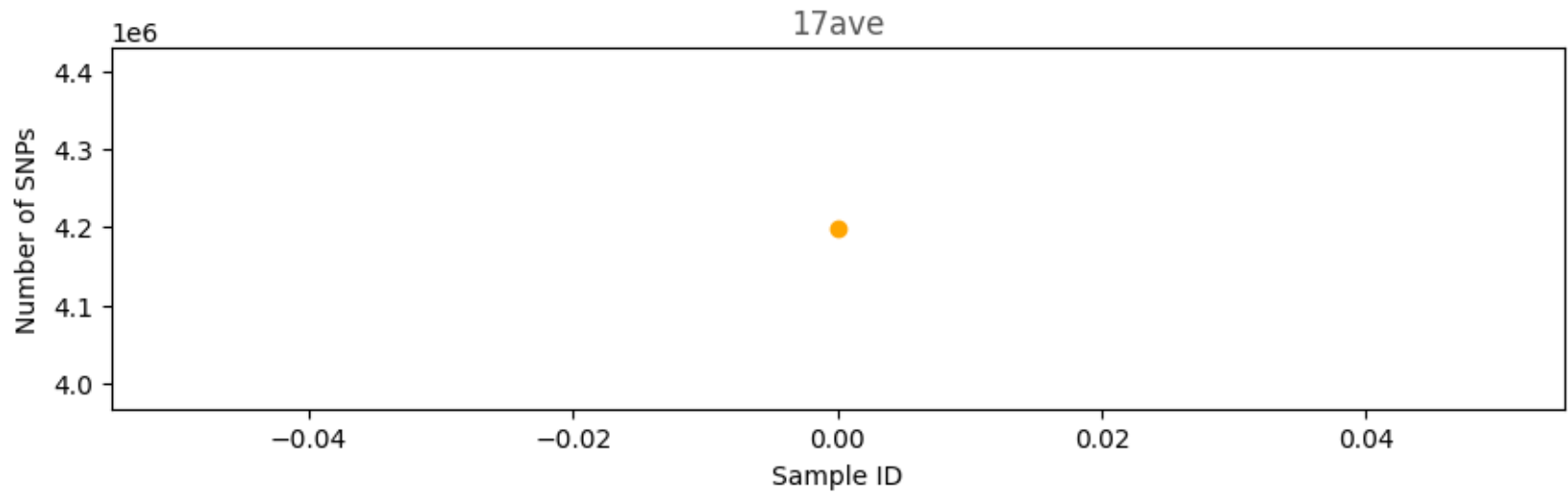# Ts/Tv by sample
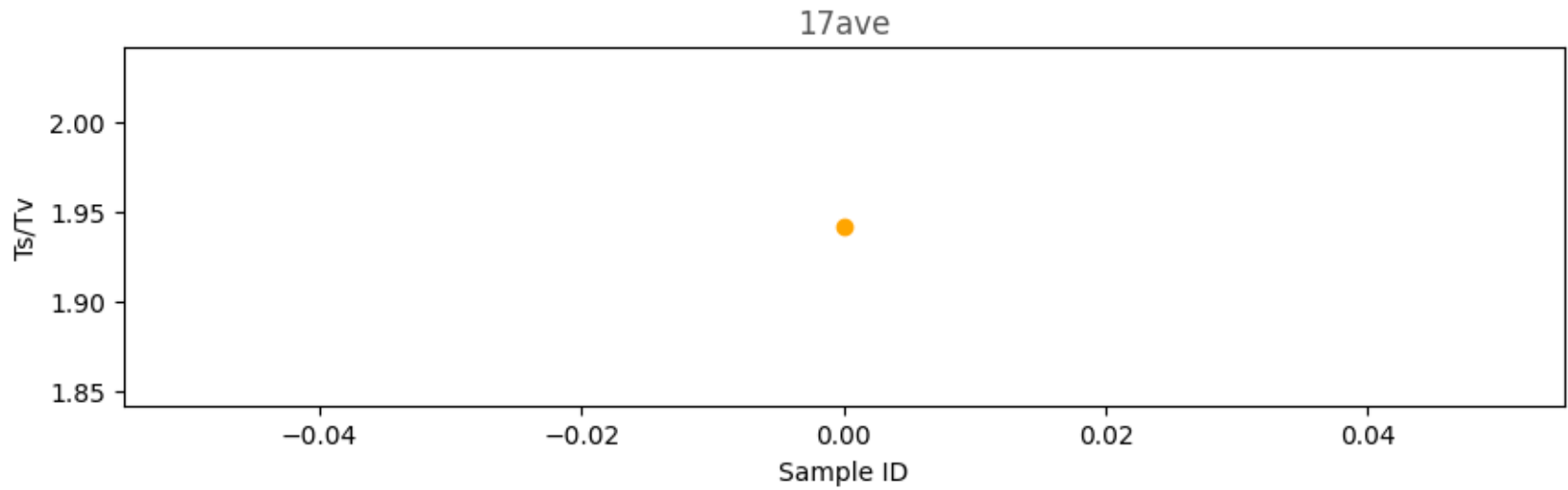
Hets vs non-ref Homs by sample
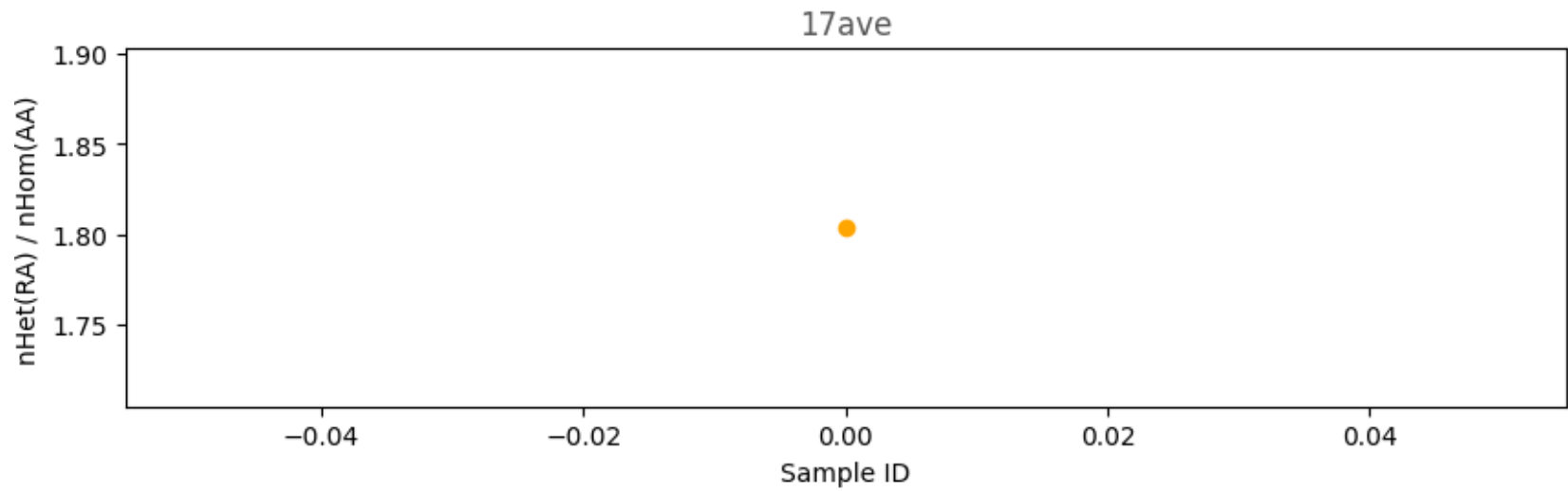
Singletons by sample (hets and homs)
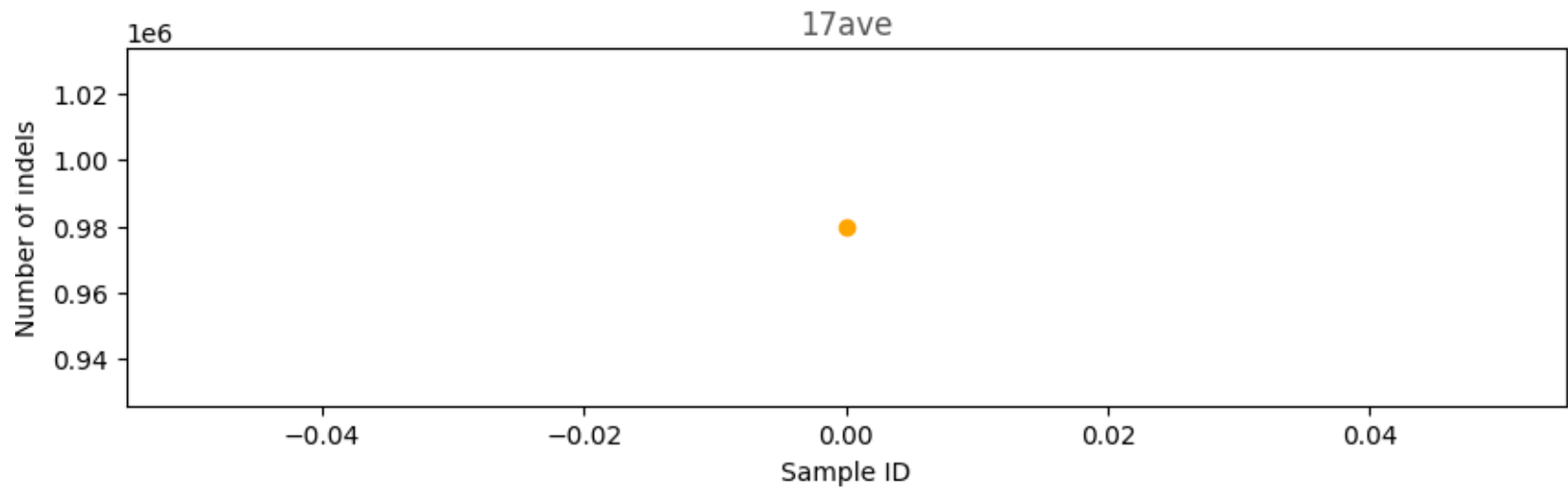
Average depth by sample
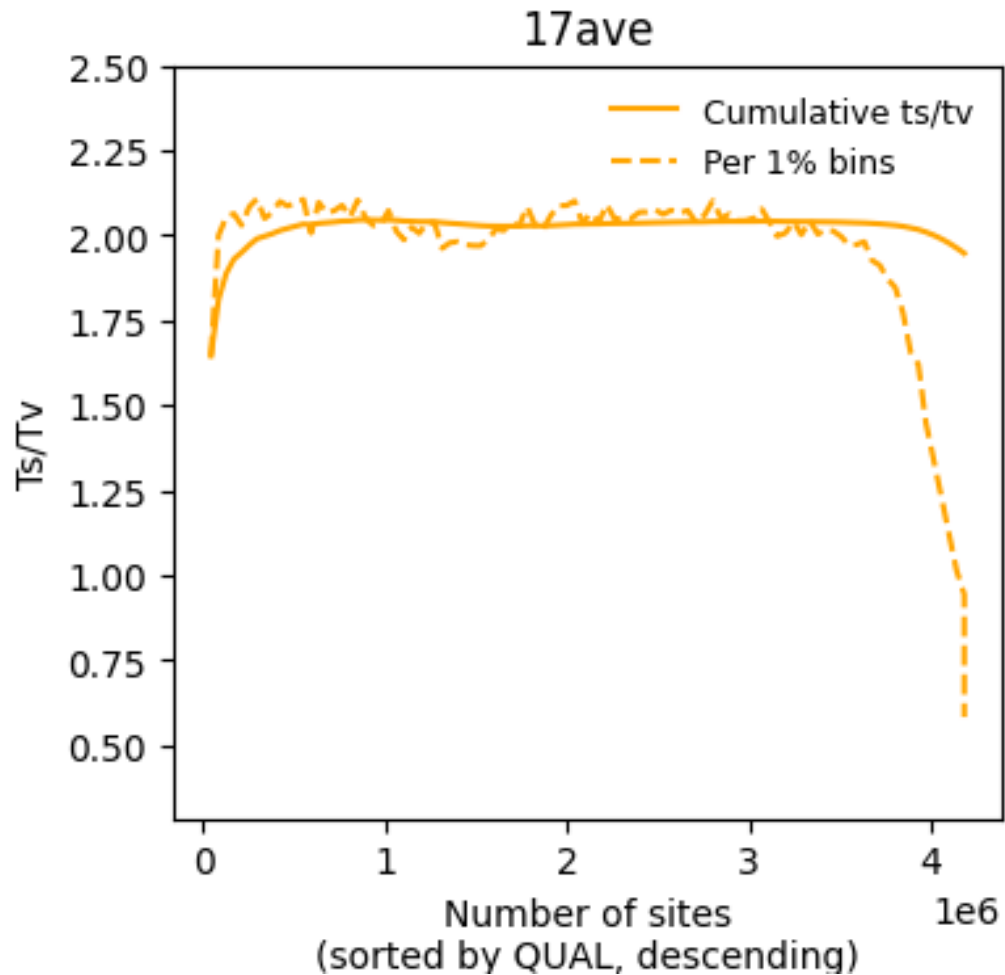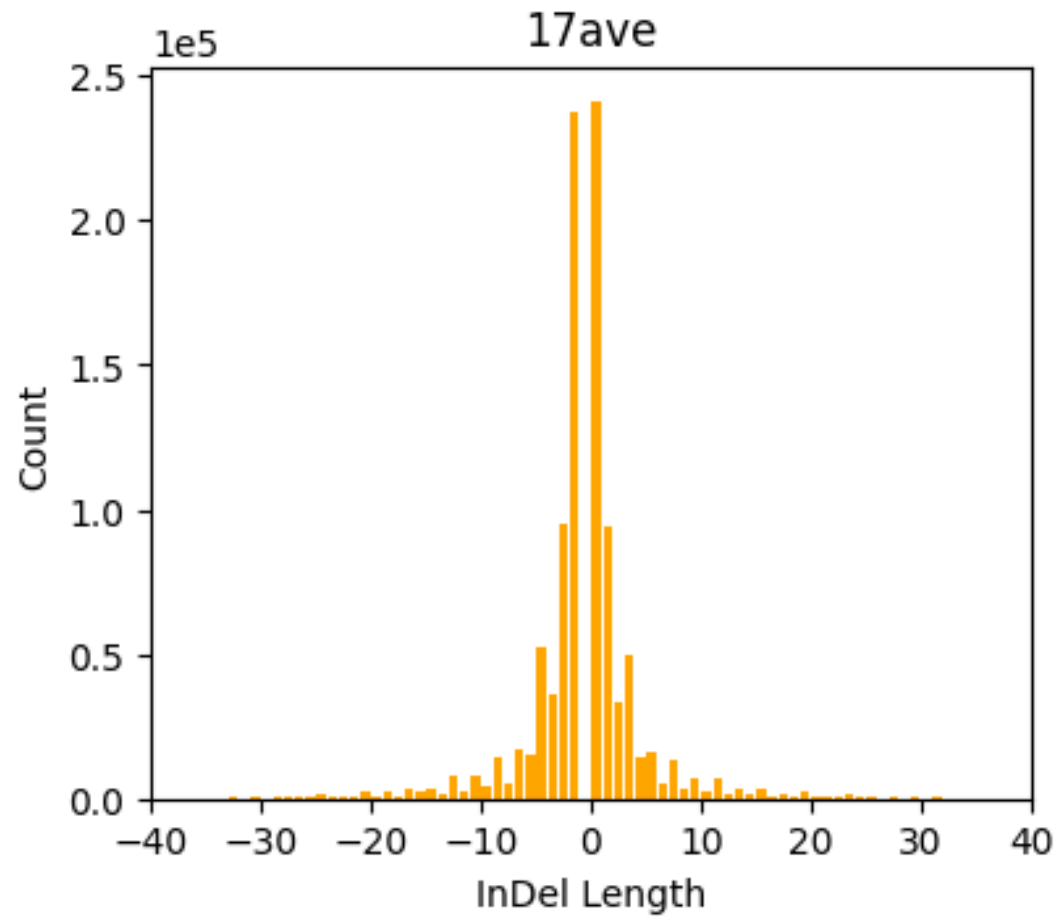
Number of SNPs by sample

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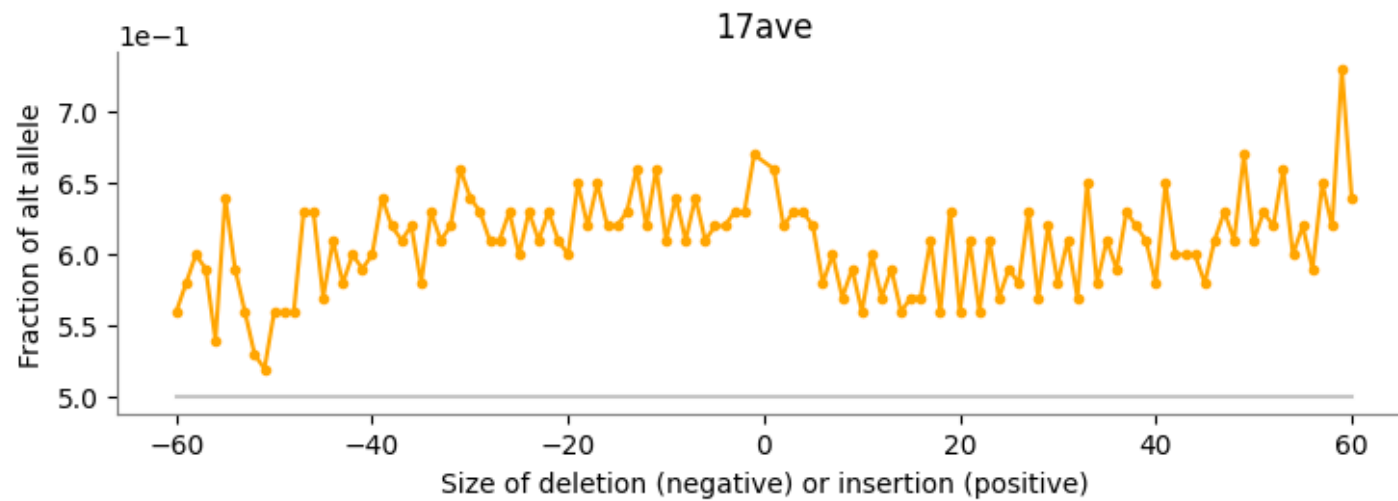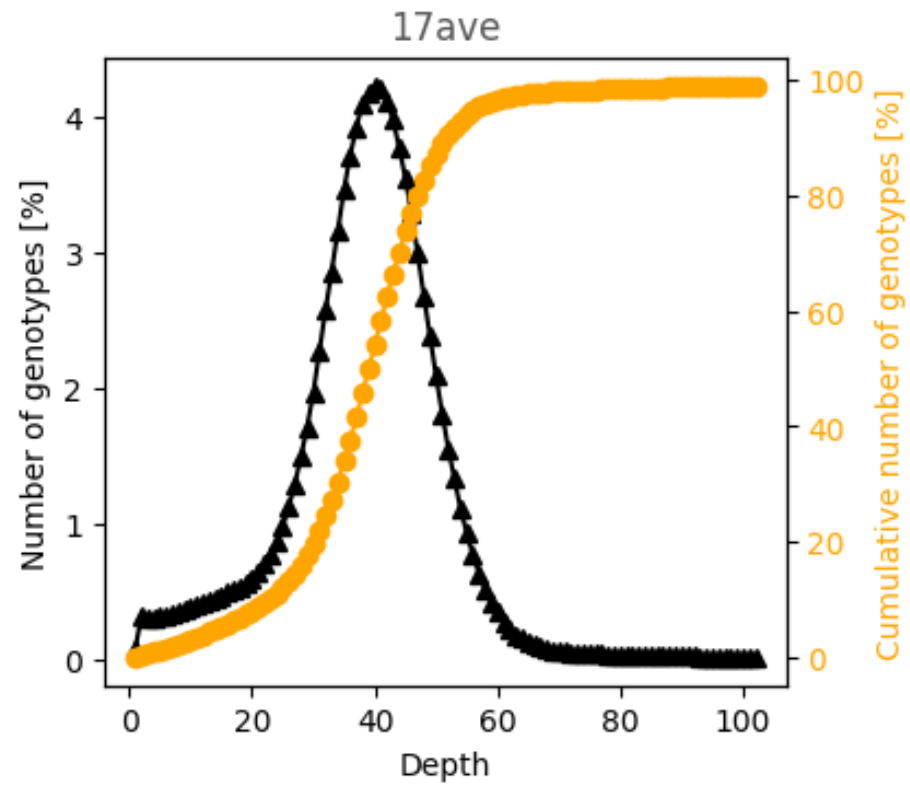
Ts/Tv stratified by QUAL
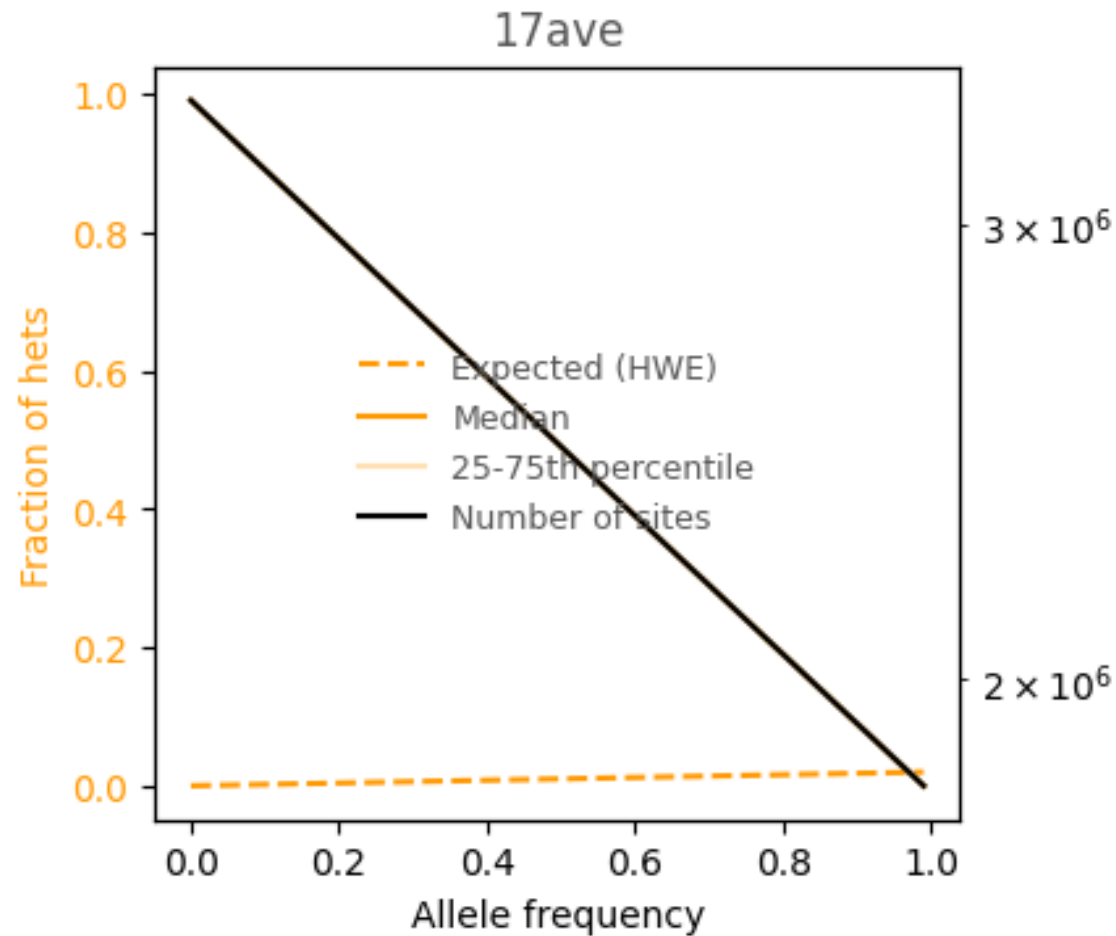
Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types