Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78lye	4,165,112	1.94	1.95	965,199	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78lye	64.3%	1.92	69.1%	103,330	2,283

- 78lye .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78lyeresf-103801978847-Normal_B`
    `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211110_A01411_AHT7VJDSX2-RHGM_LABKA_WGSA`
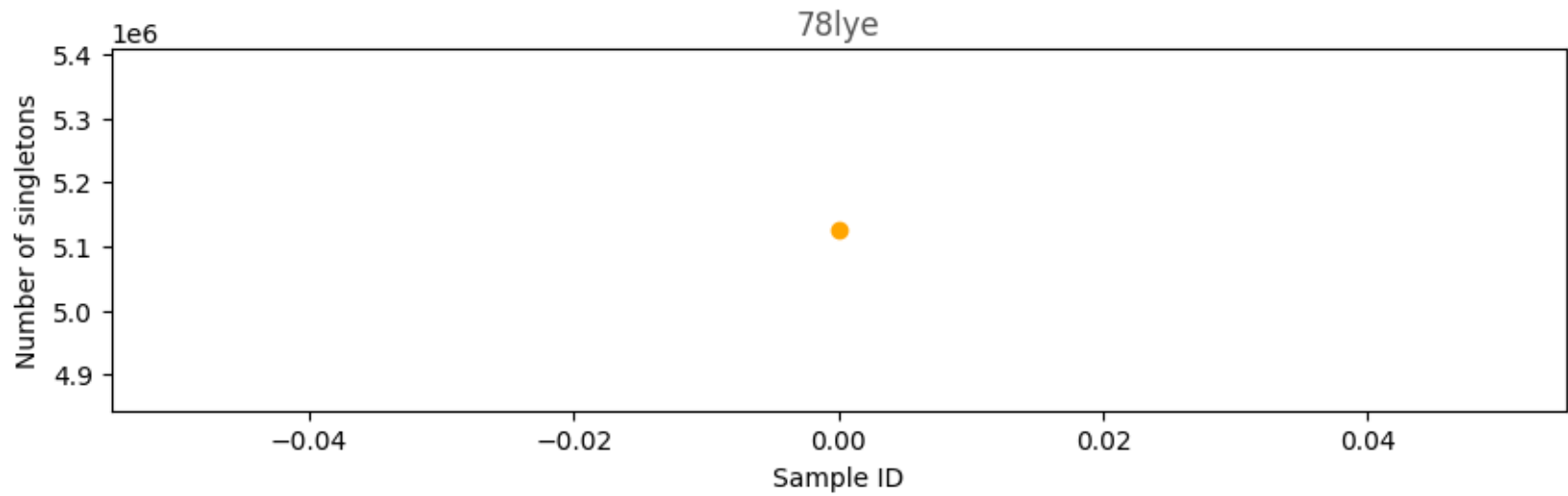   `KUT-WGSAKUT03300_21RKG027263x01_snv_germline_raw.haplotype_caller.vcf.gz`
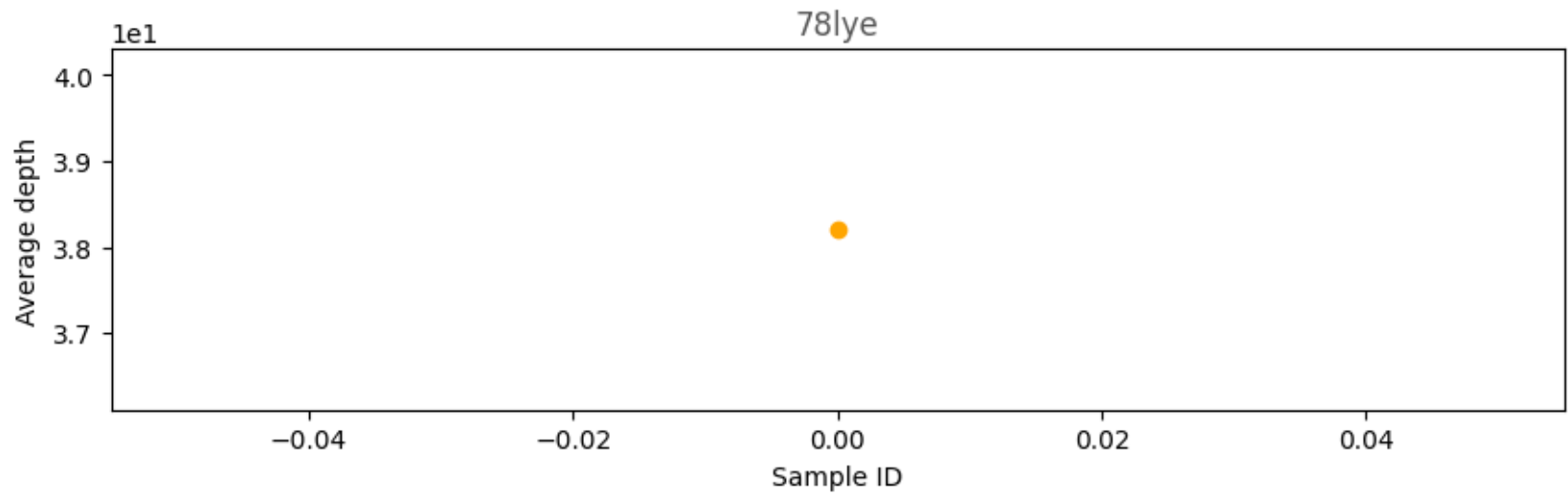
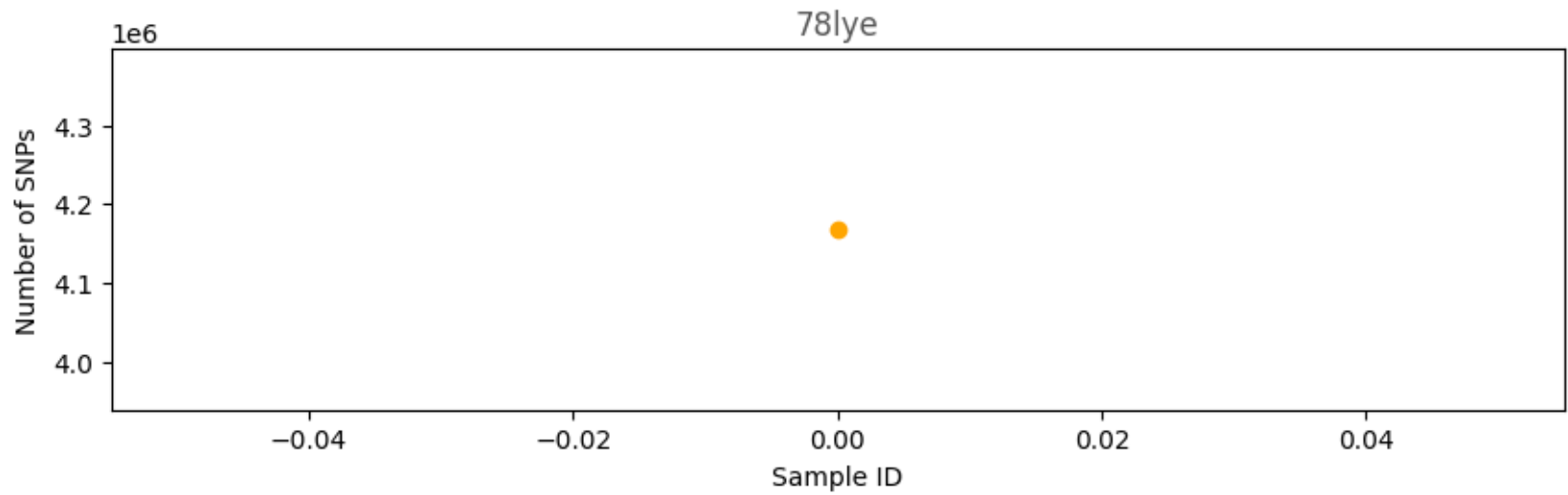# Ts/Tv by sample
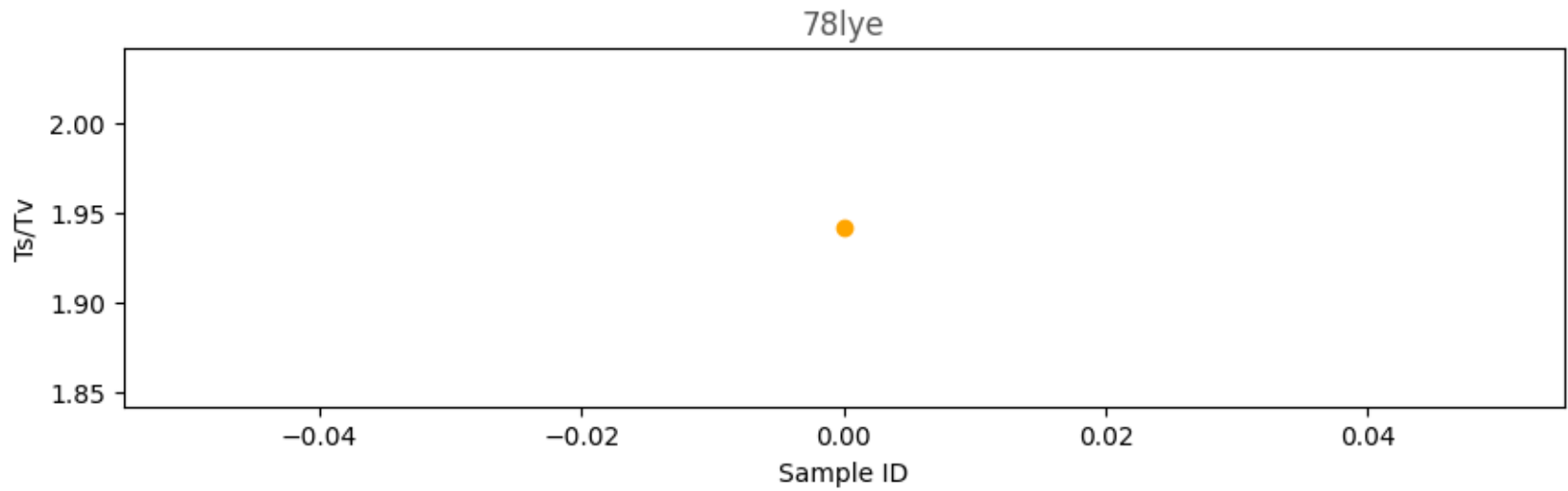
Hets vs non-ref Homs by sample
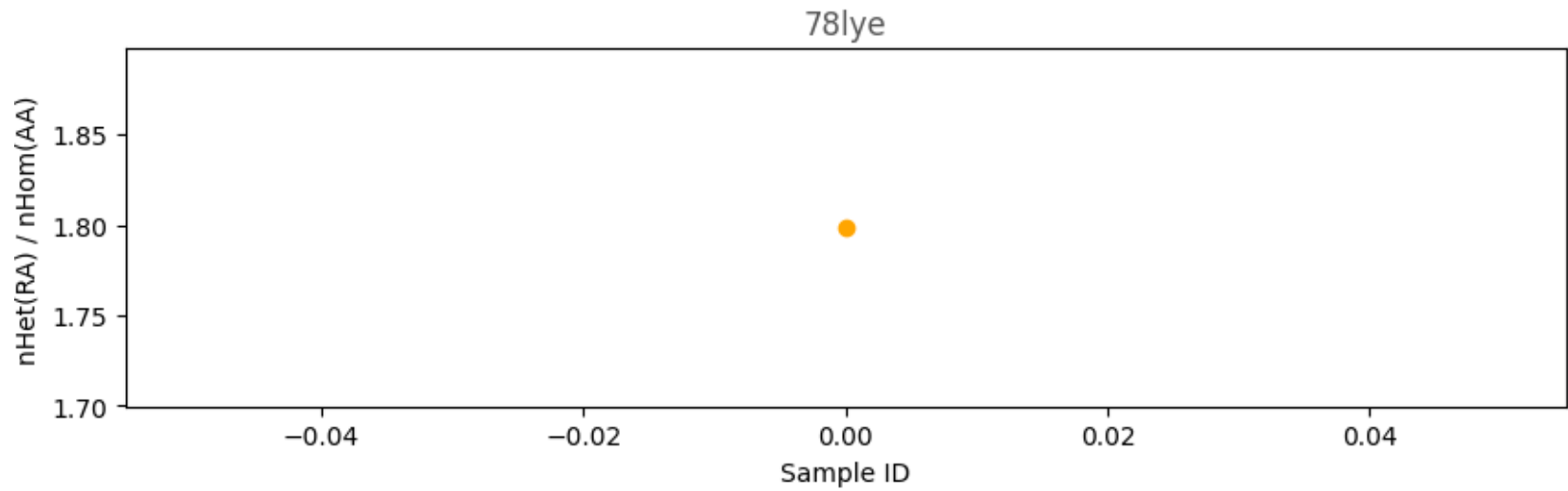
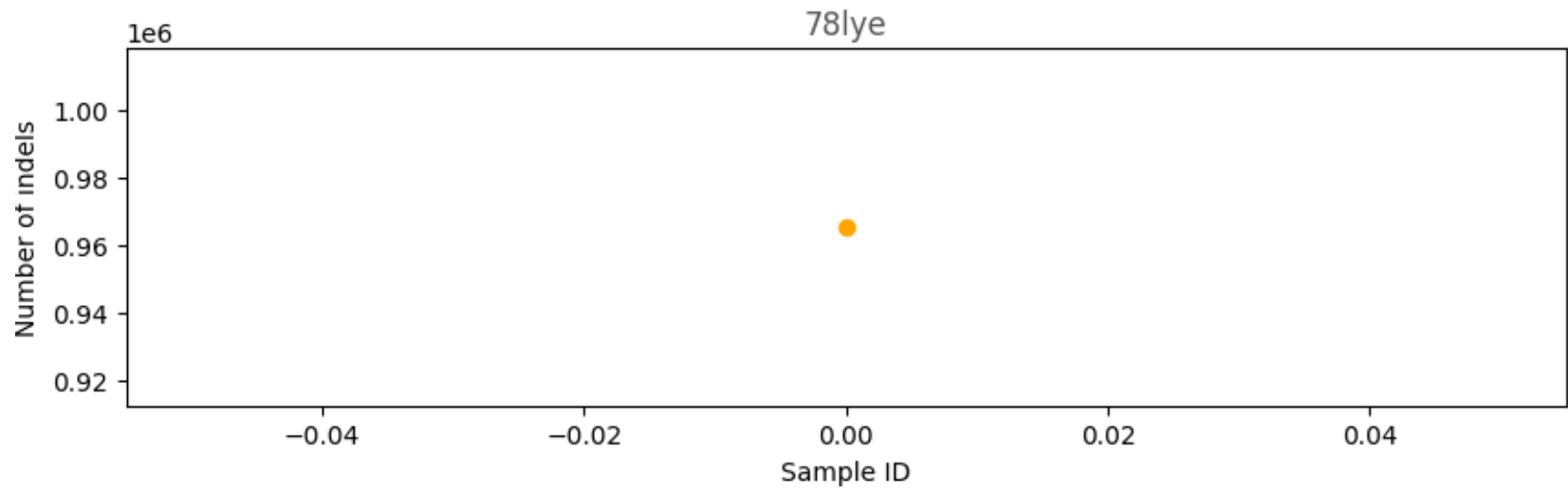# Singletons by sample (hets and homs)
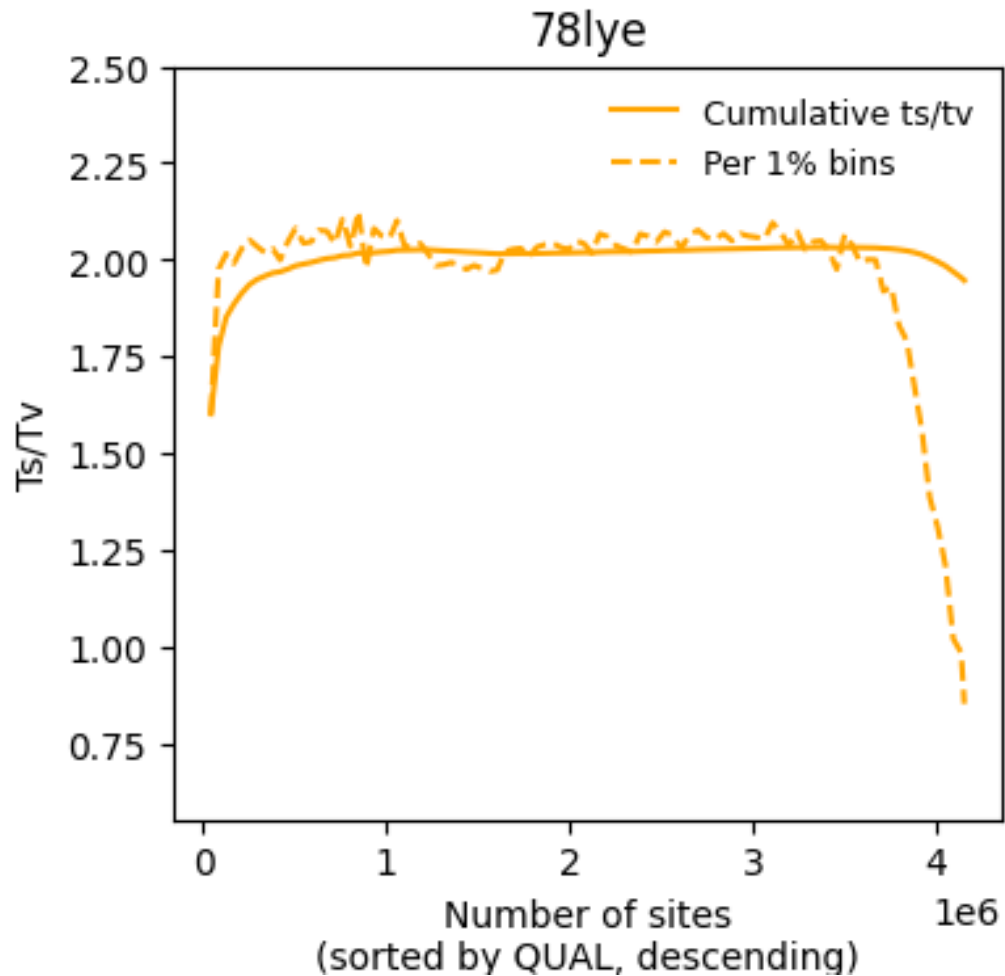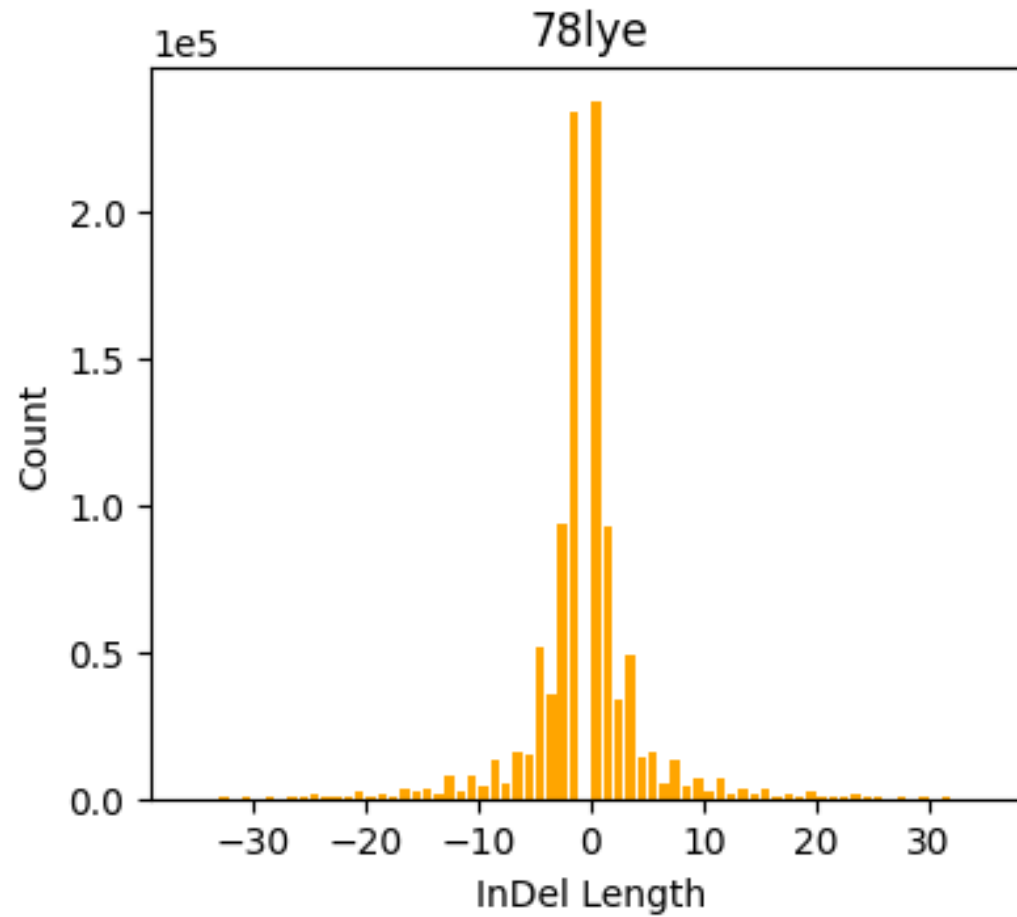
Average depth by sample
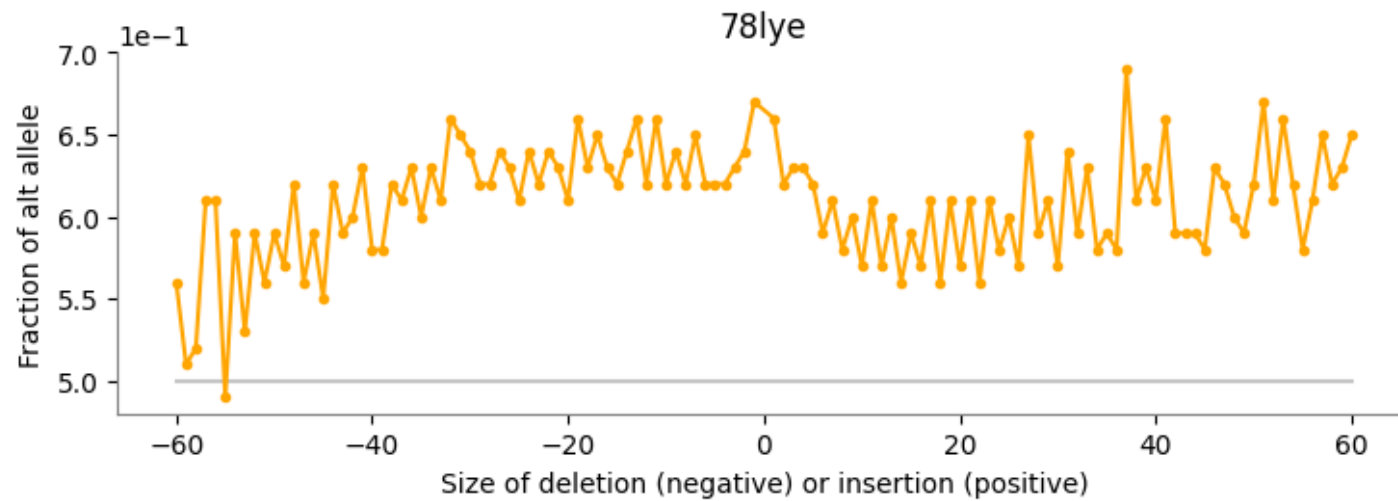
Number of SNPs by sample

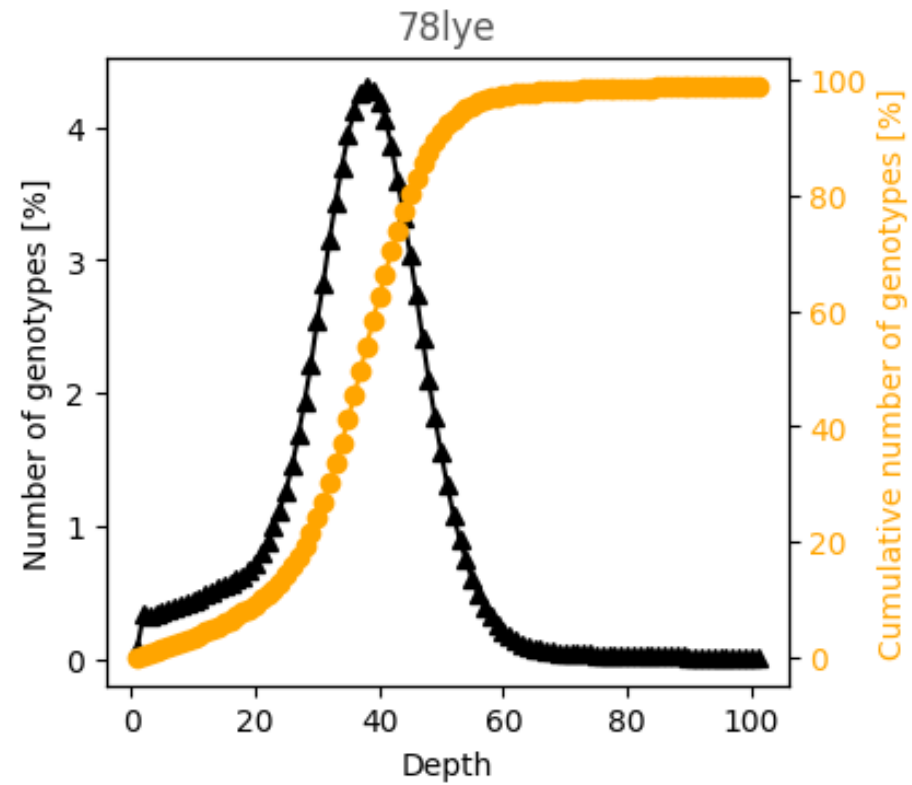# Number of indels by sample
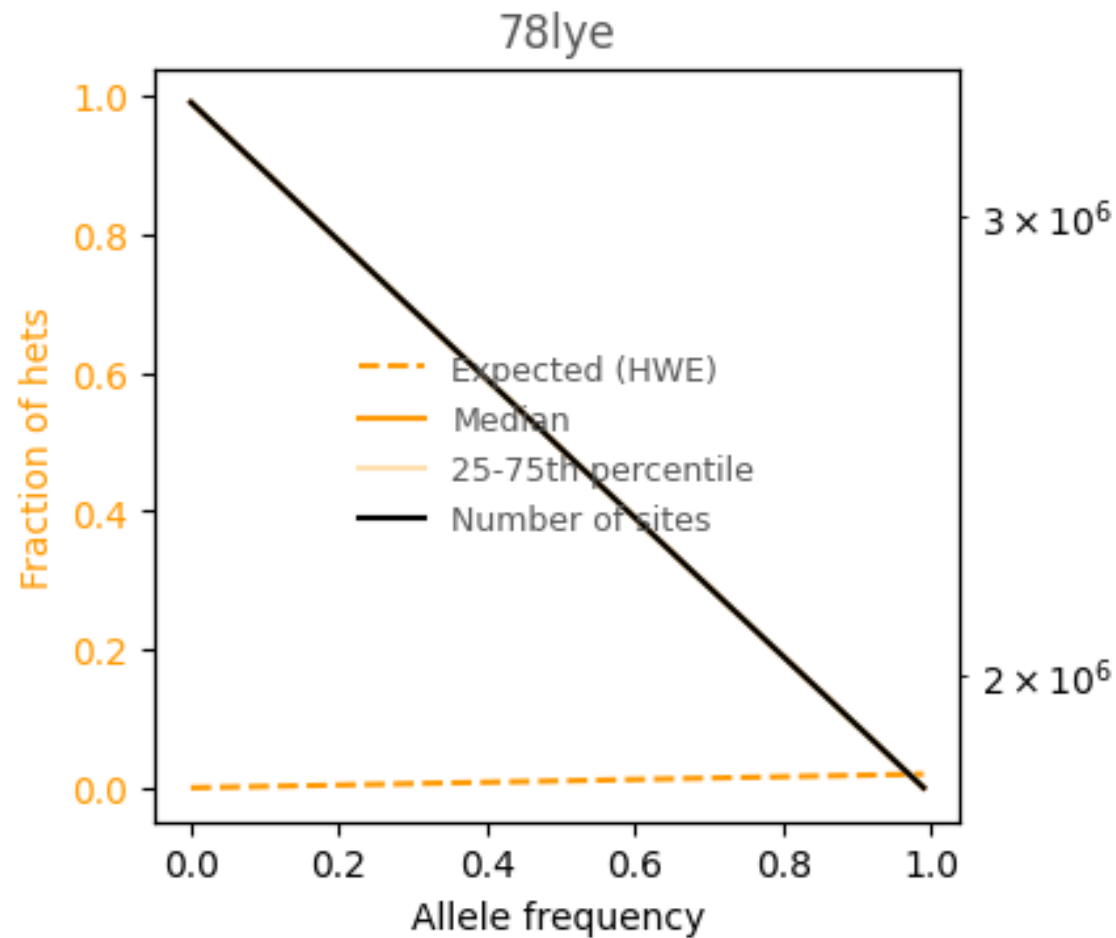
Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

78lye

Substitution types