Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
22tys	4,079,790	1.94	1.94	939,430	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
22tys	63.3%	1.91	67.4%	93,100	1,950

- 22tys .. `/ngc/projects2/gm/data/archive/2022/variants/snv/22tysbyff-103888872213-Normal_B`
`lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220706_A01176_BHMWWFDSX3-RHGM_LABKA_WGSA`
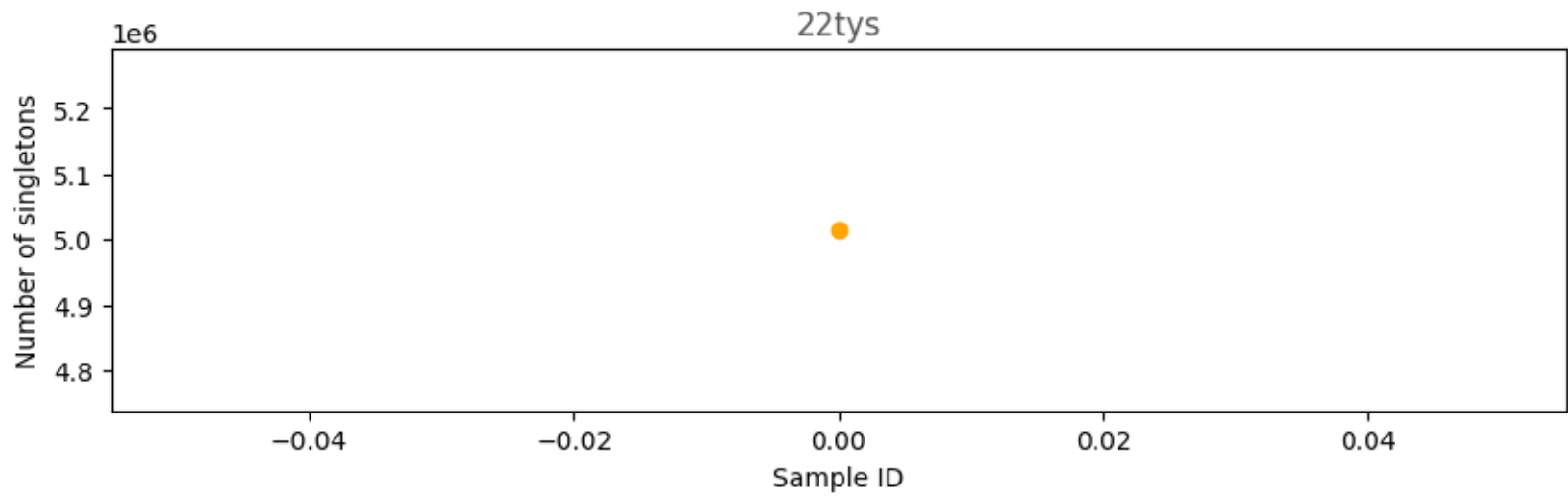`KUT-WGSAKUT04593_22RKG016361x01_snv_germline_raw.haplotype_caller.vcf.gz`
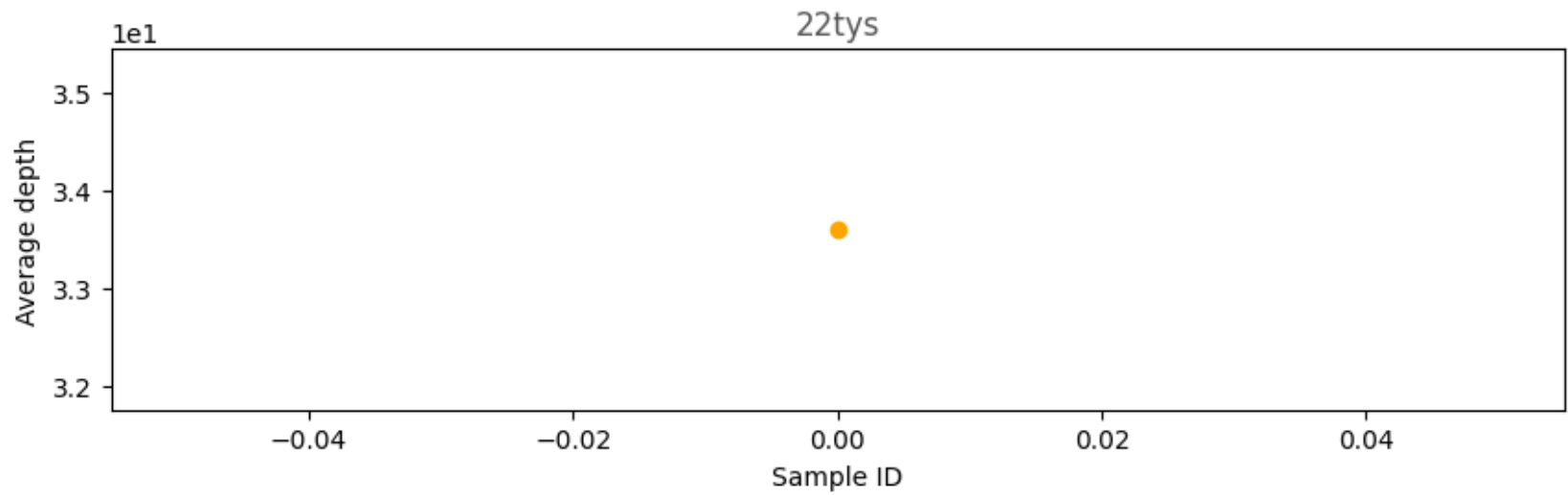
Ts/Tv by sample
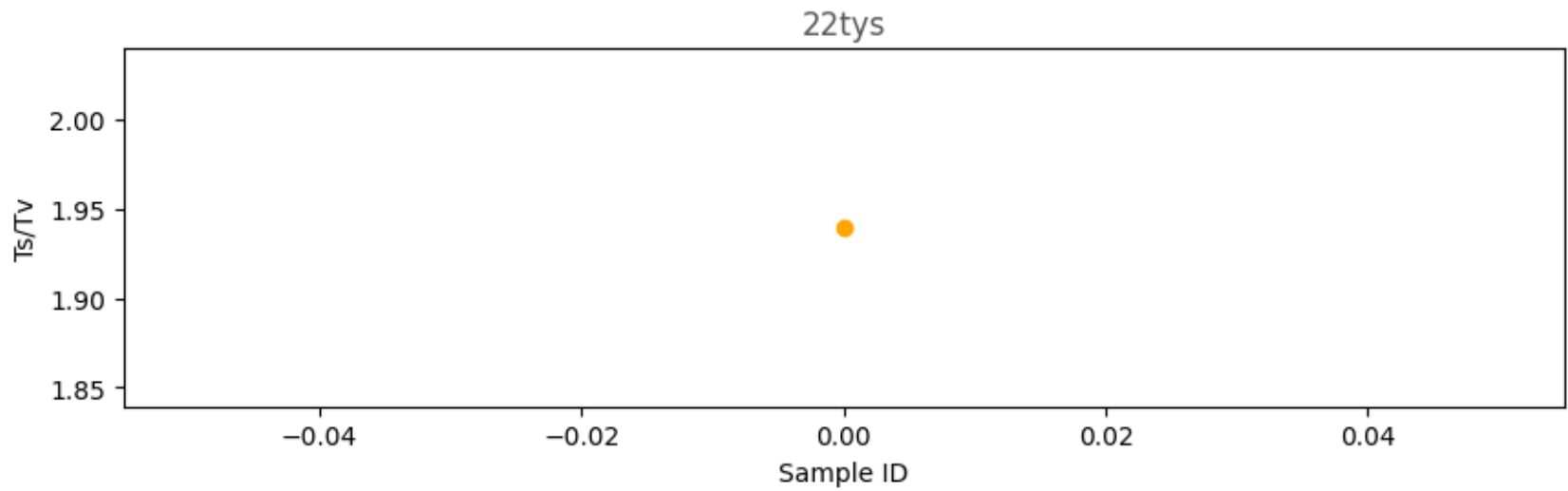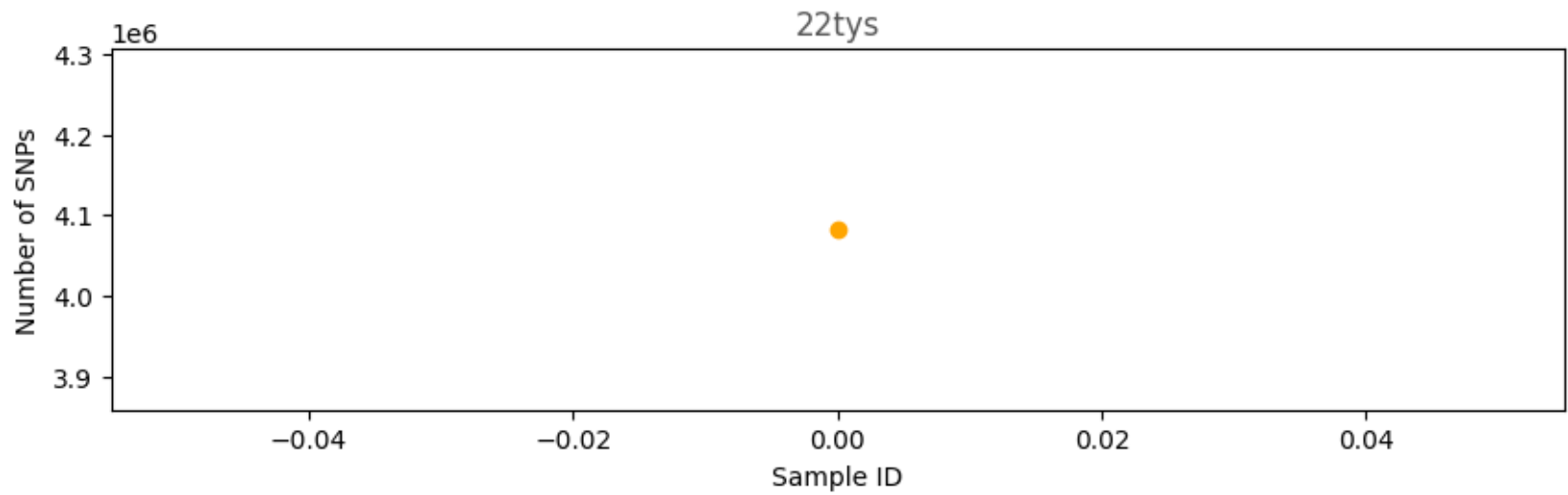
Hets vs non-ref Homs by sample
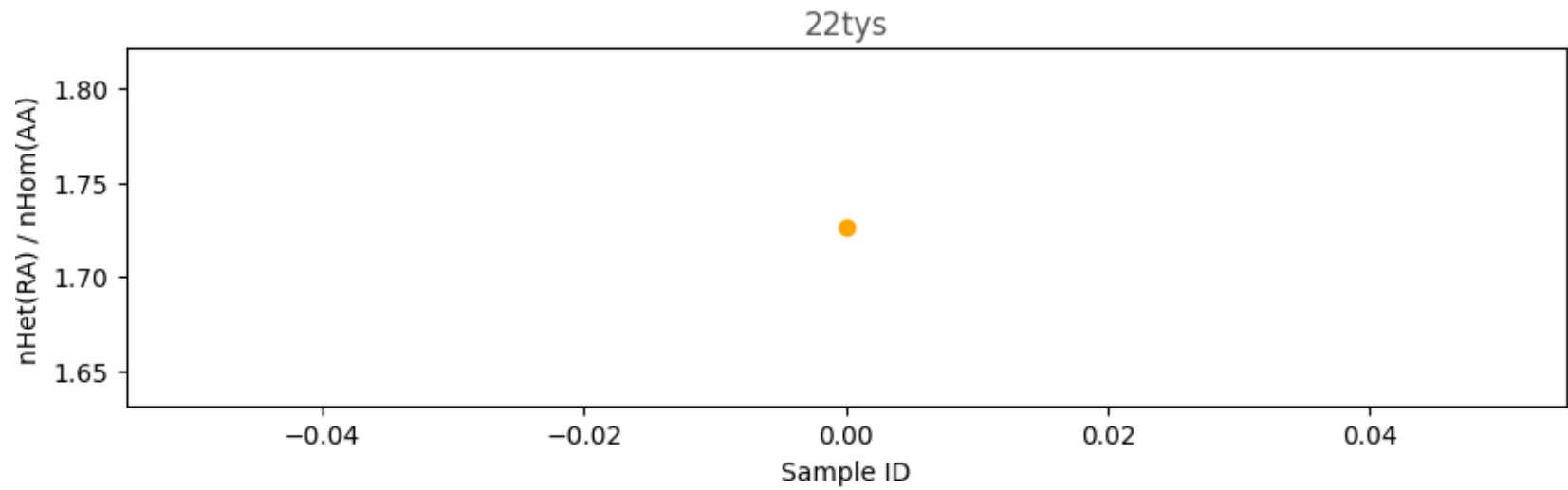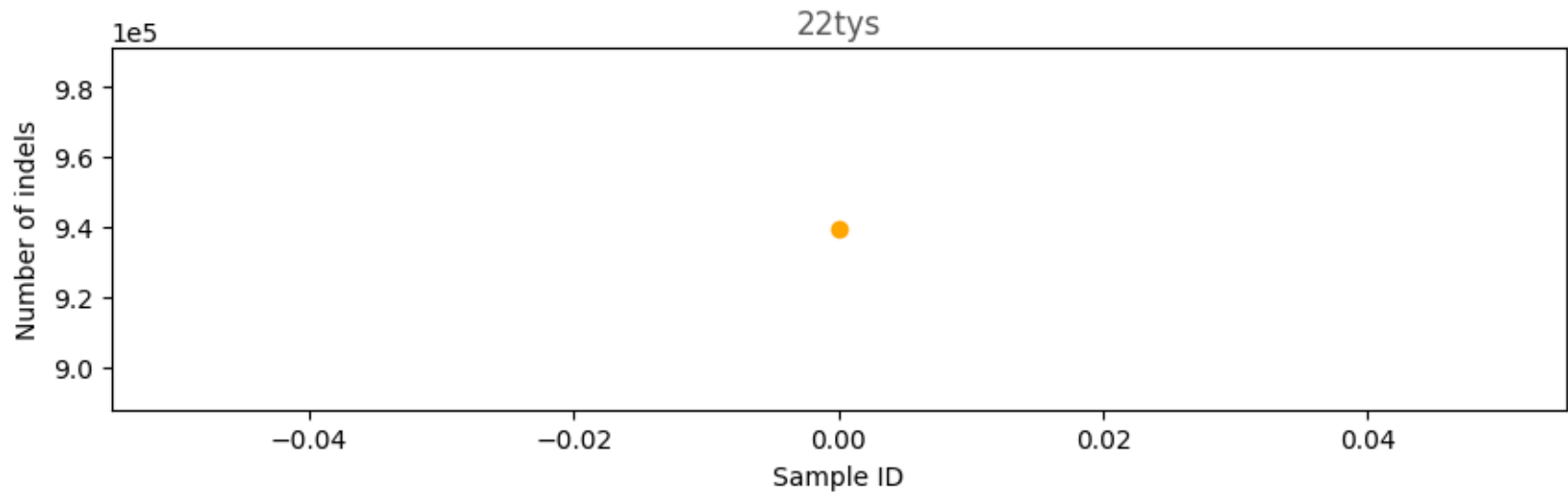
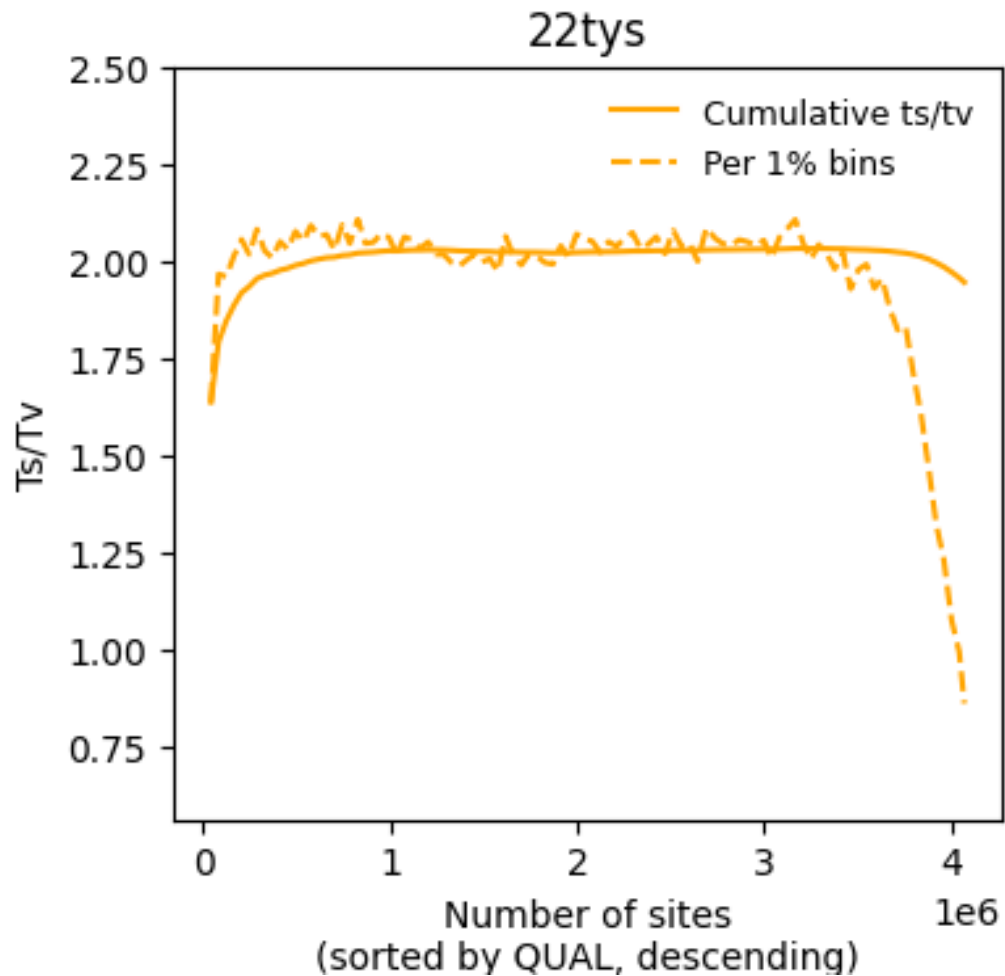# Singletons by sample (hets and homs)
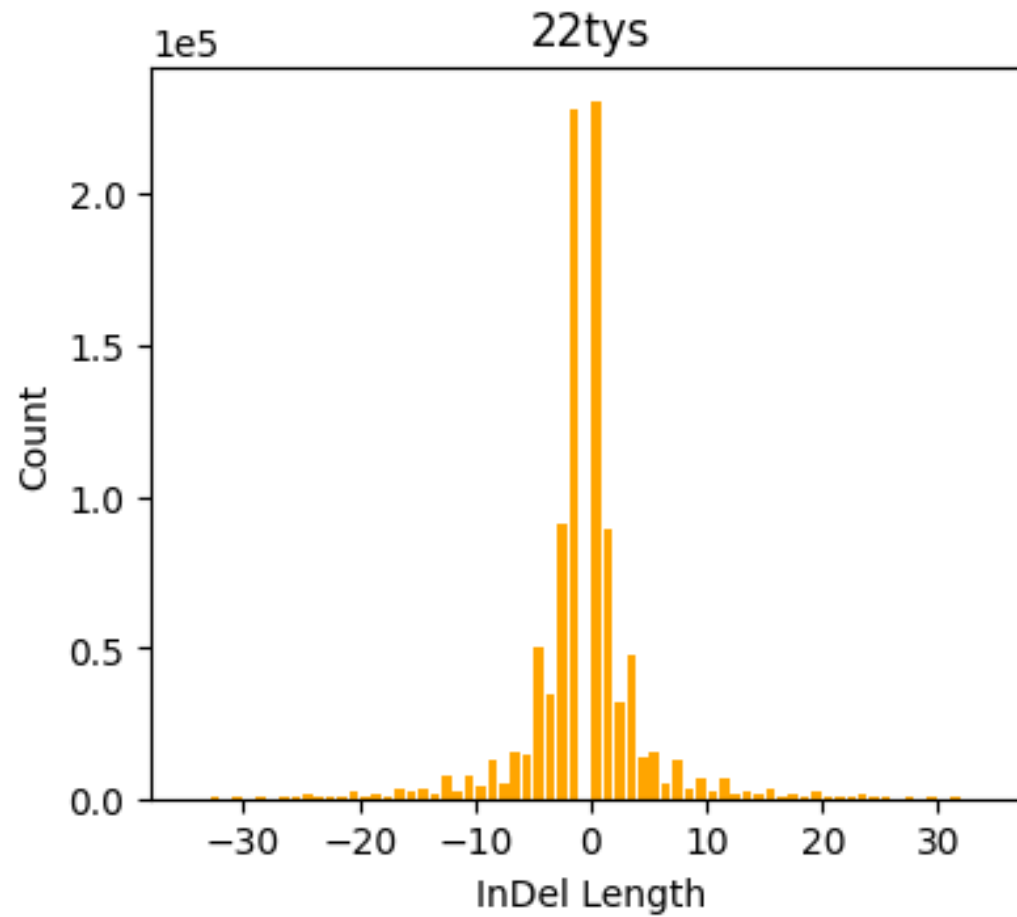
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

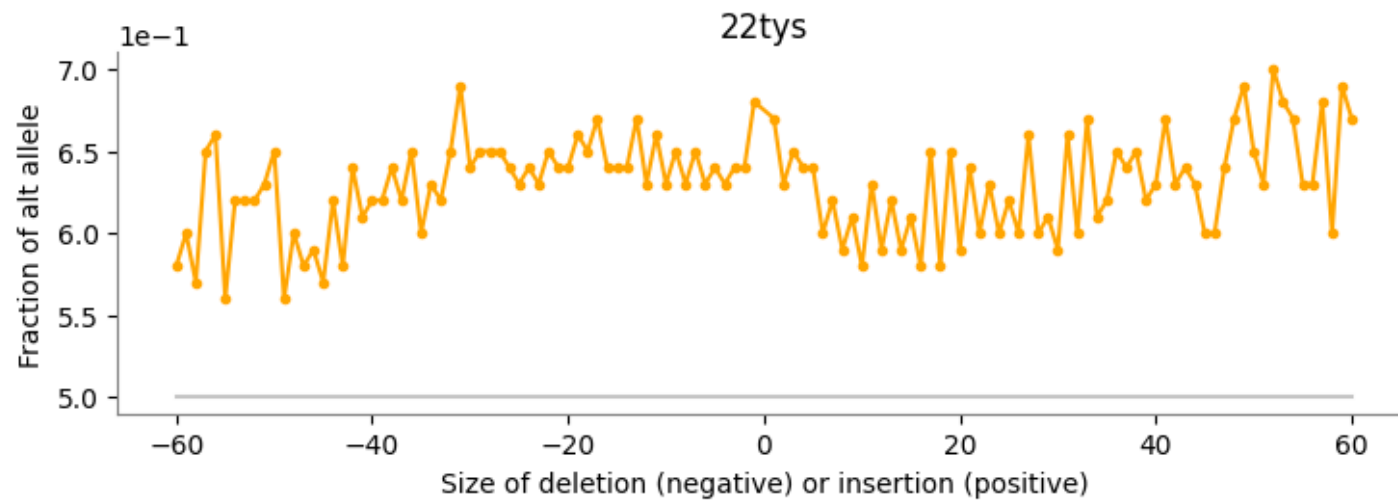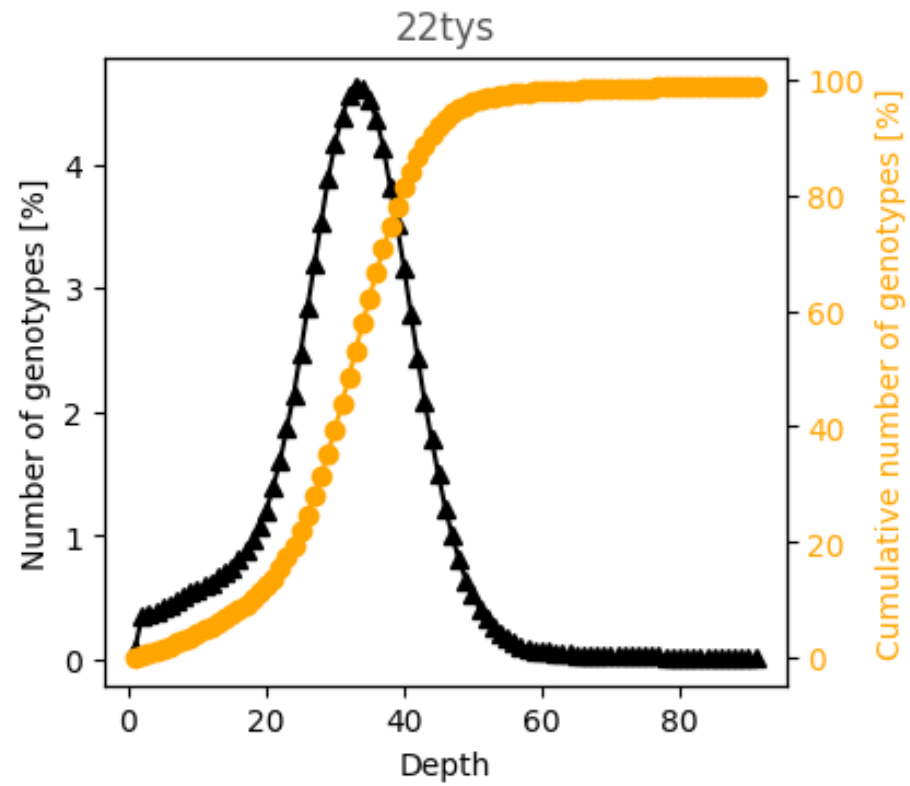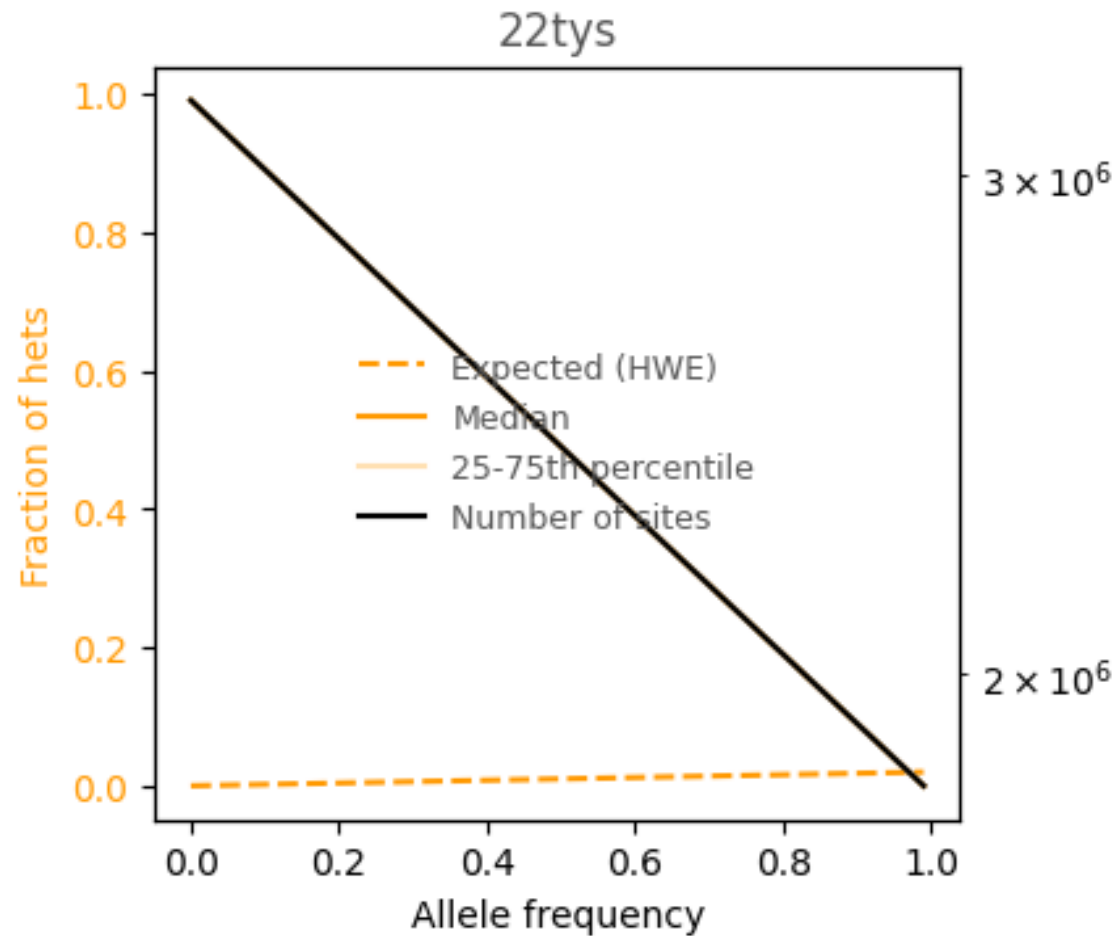# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types