Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
62twh	4,115,092	1.92	1.93	955,122	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
62twh	63.4%	1.89	68.4%	101,810	2,108

- 62twh .. `/ngc/projects2/gm/data/archive/2022/variants/snv/62twhecef-103883427688-Normal_B`
`lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00951-220629_A01176_AHMWKCDSX3-EXT_LAB`
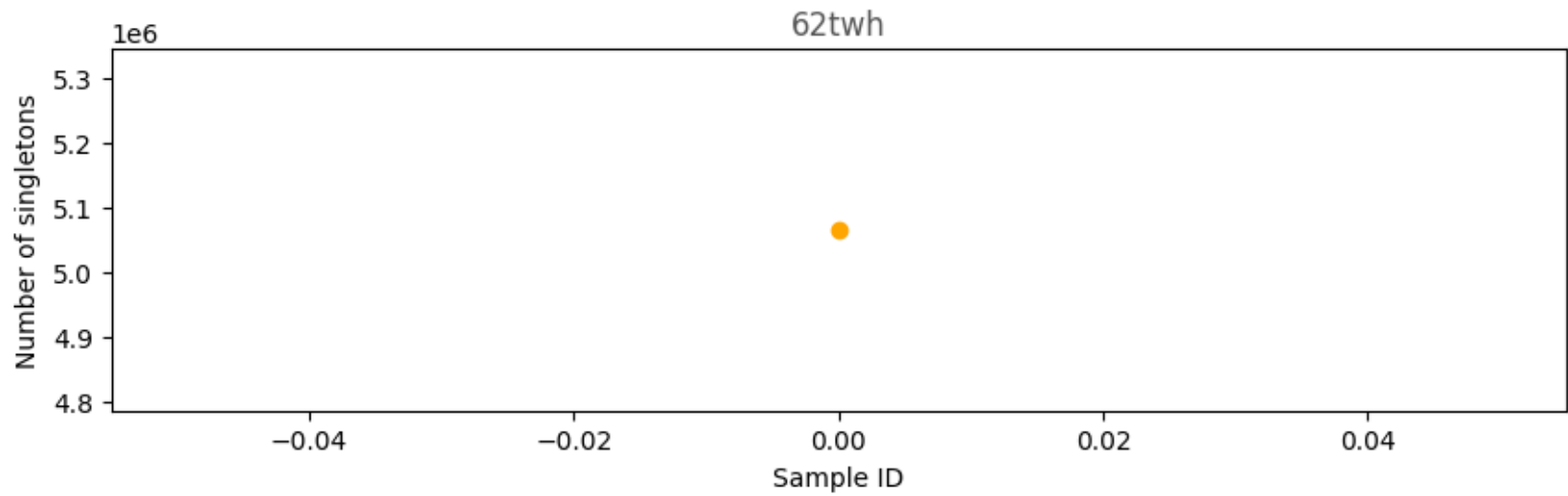`KA_NGCWGS-NGCWGS04550_snv_germline_raw.haplotype_caller.vcf.gz`
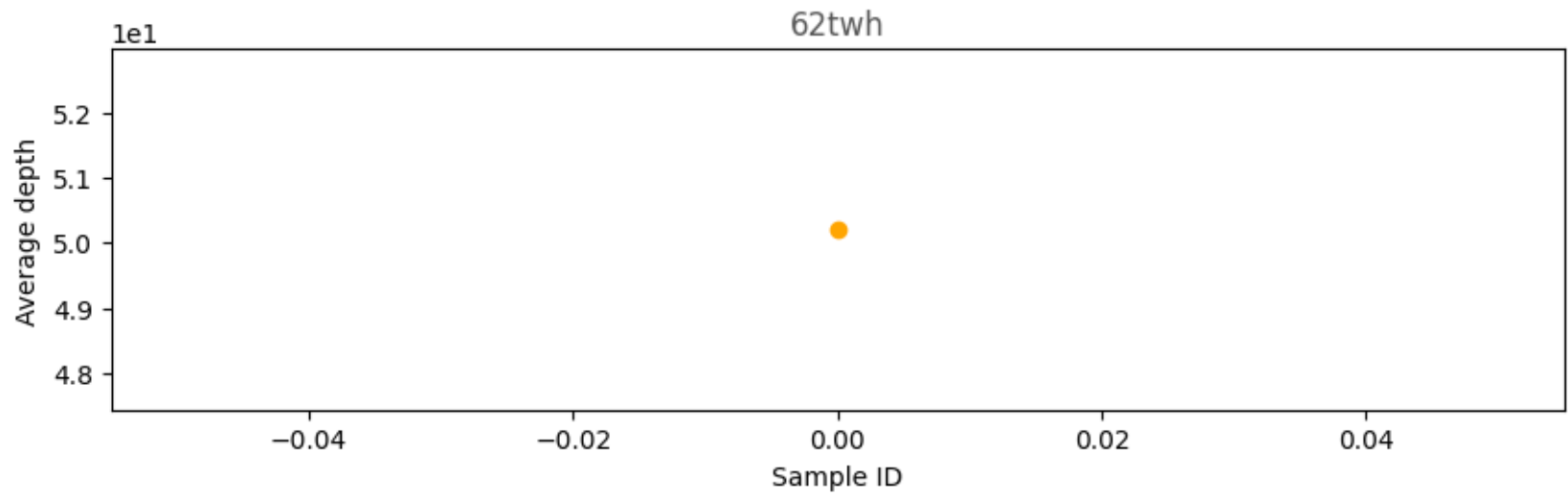
Ts/Tv by sample
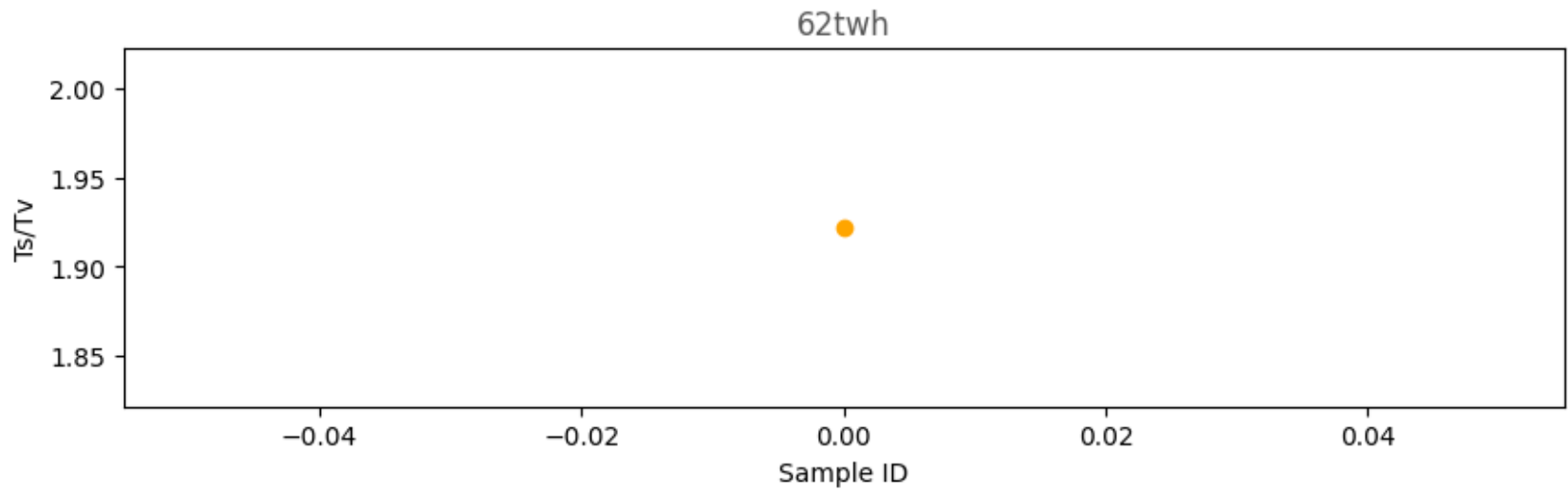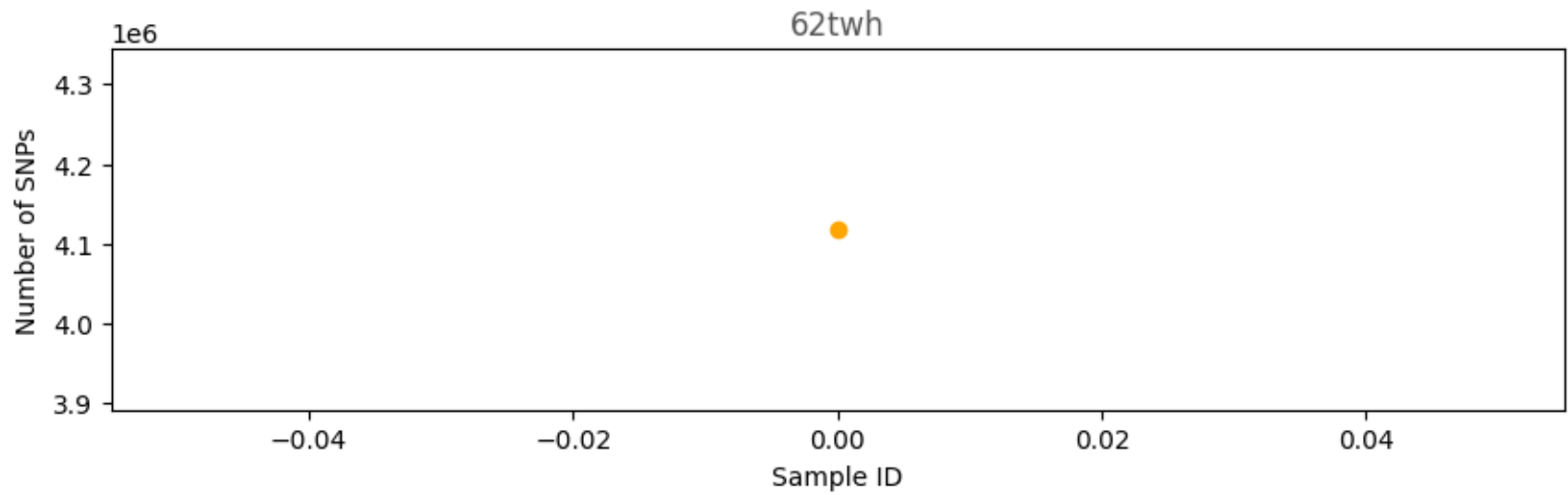
Hets vs non-ref Homs by sample
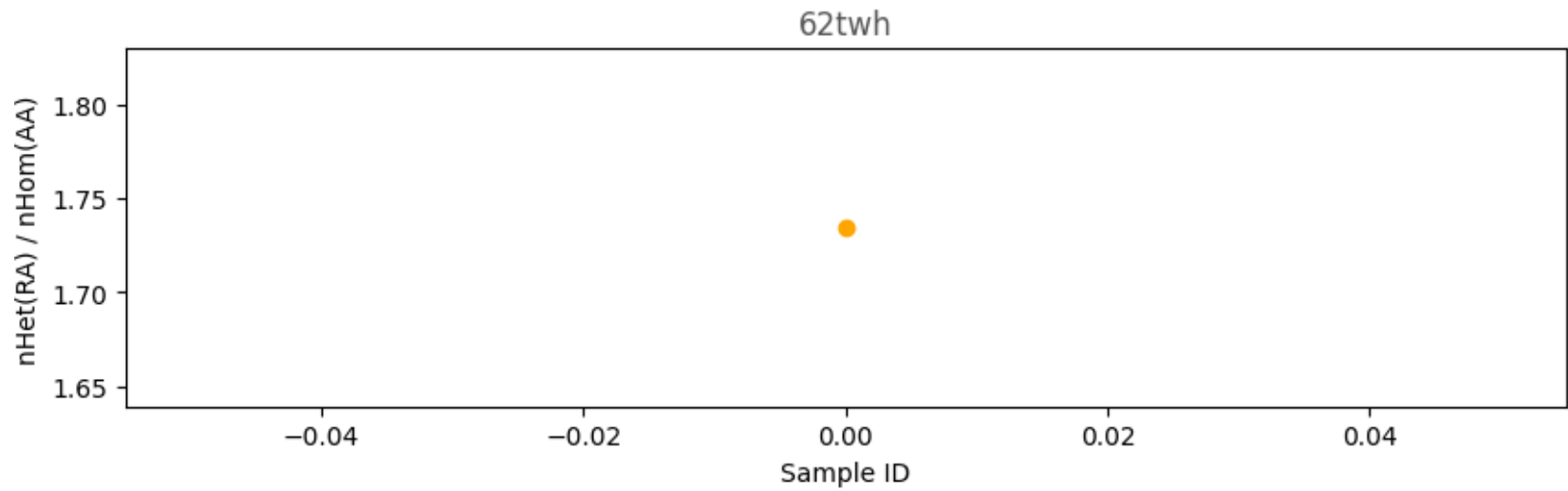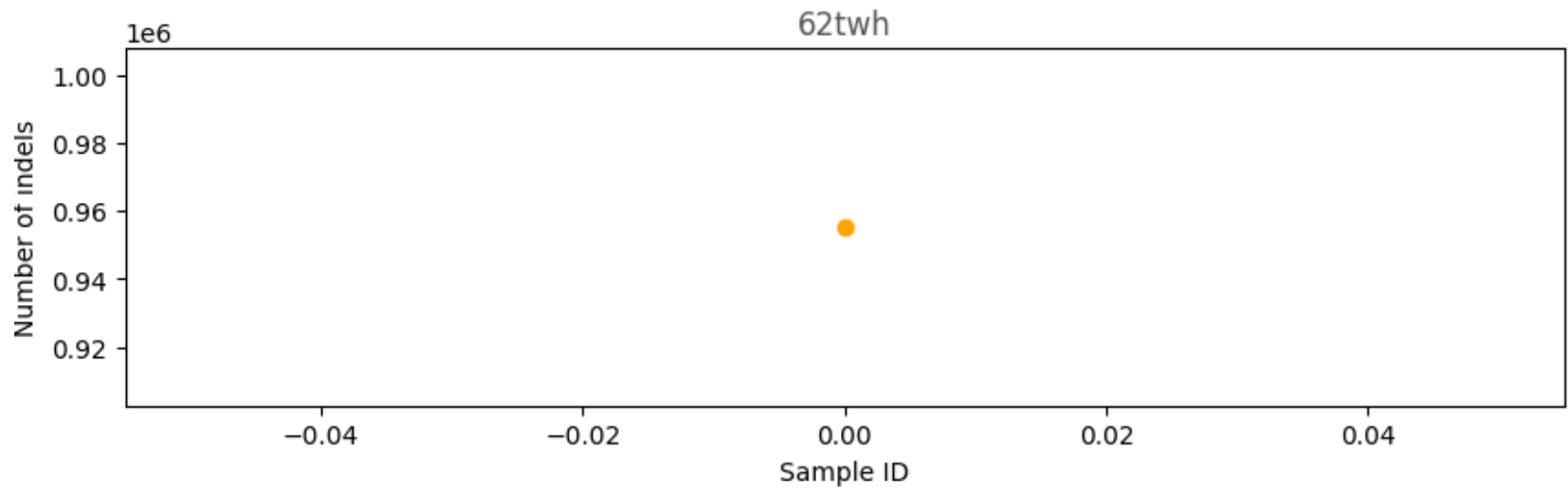
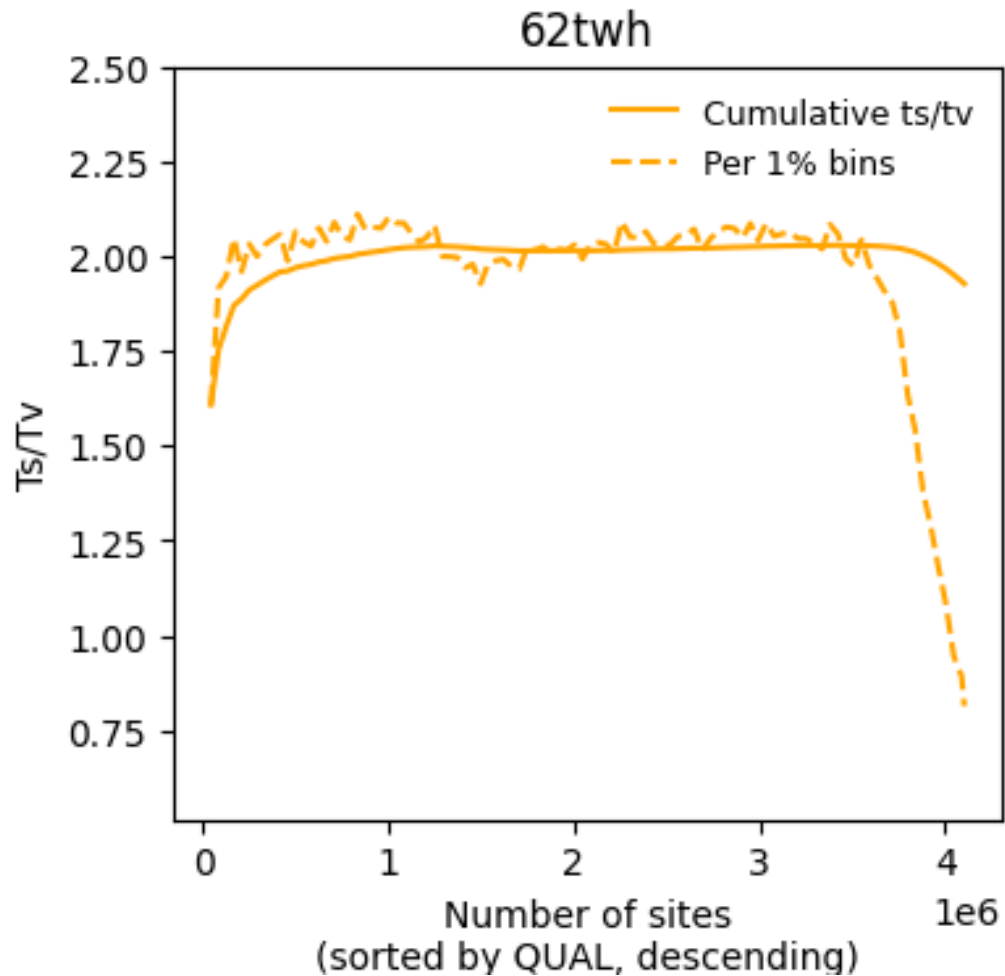# Singletons by sample (hets and homs)

Average depth by sample
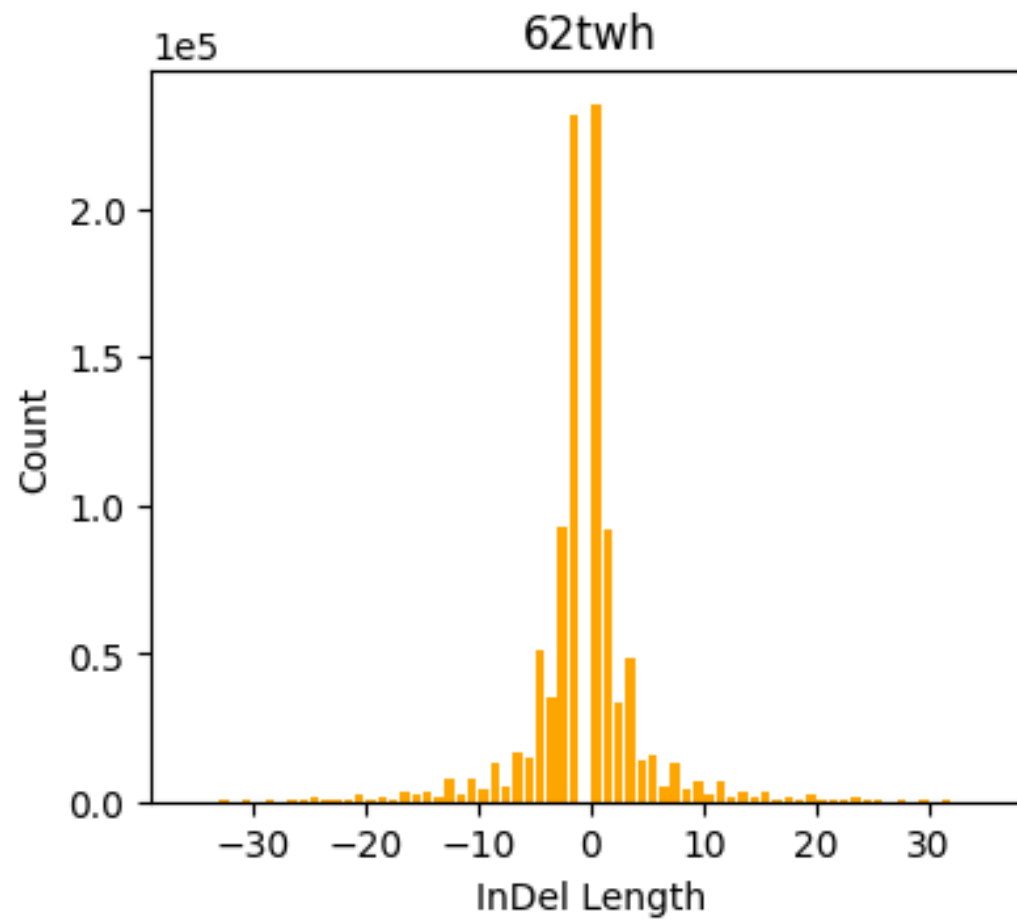
Number of SNPs by sample

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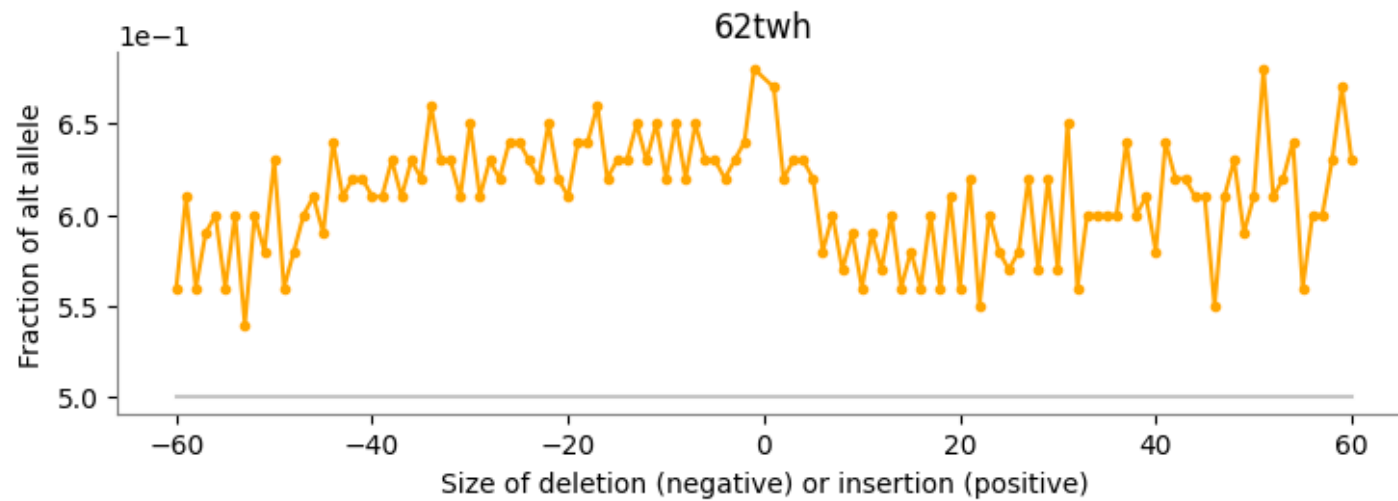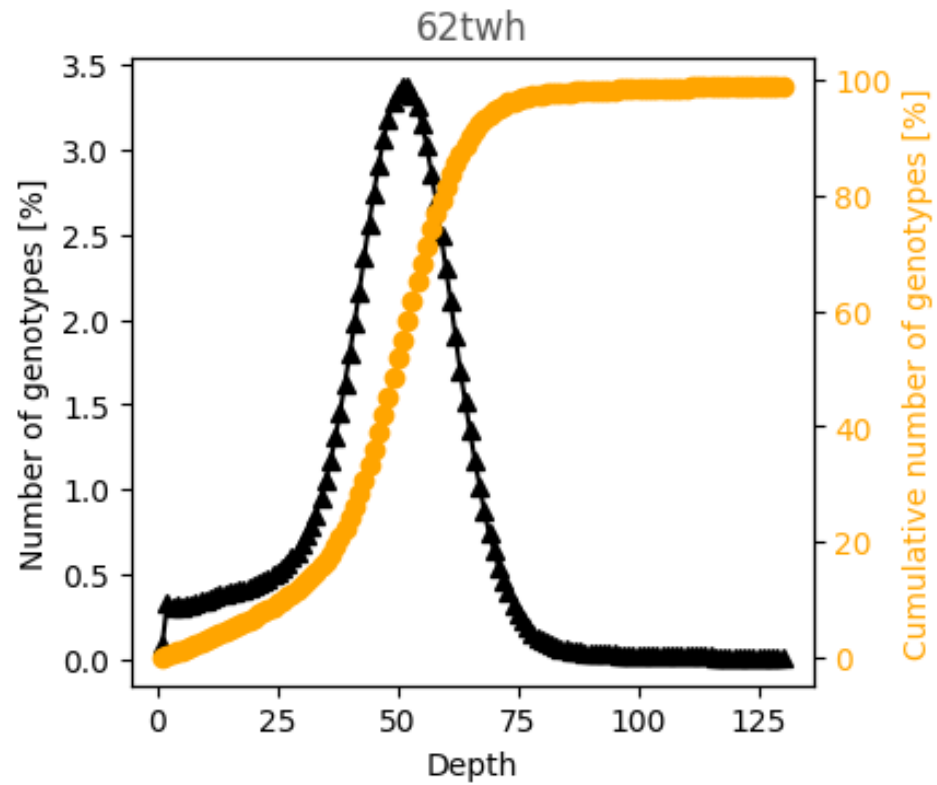
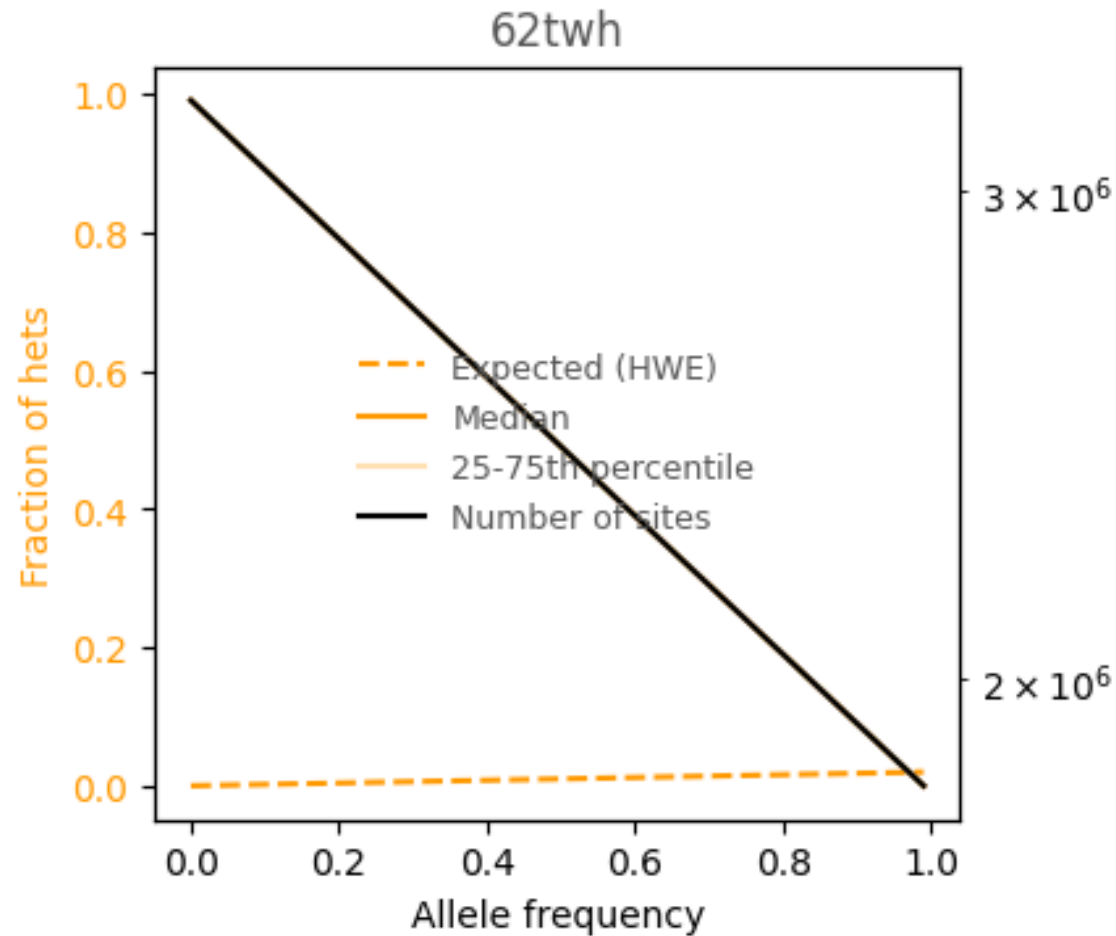# Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types