Summary Numbers

| Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
52inf	4,102,852	1.92	1.93	965,879	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
52inf	63.0%	1.89	68.7%	108,143	2,134

- 52inf .. `/ngc/projects2/gm/data/archive/2022/variants/snv/52infthof-103814429219-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211215_A01466_AHVN2CDSX2-RHGM_LABKA_WGS-WGS03492_snv_germline_raw.haplotype_caller.vcf.gz`
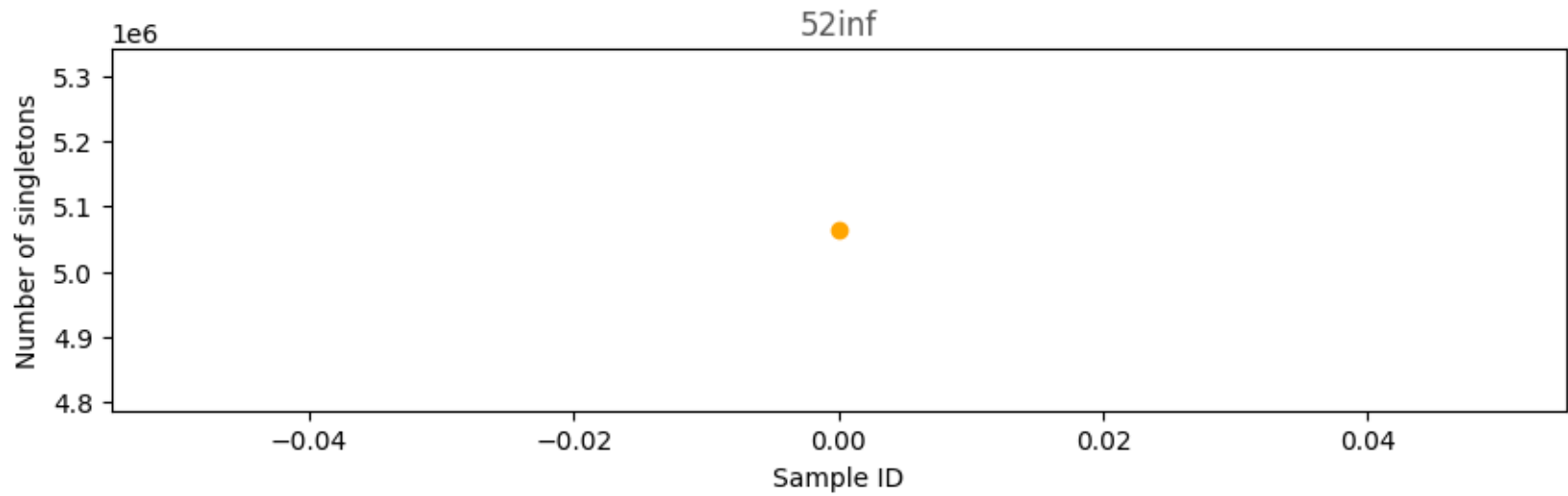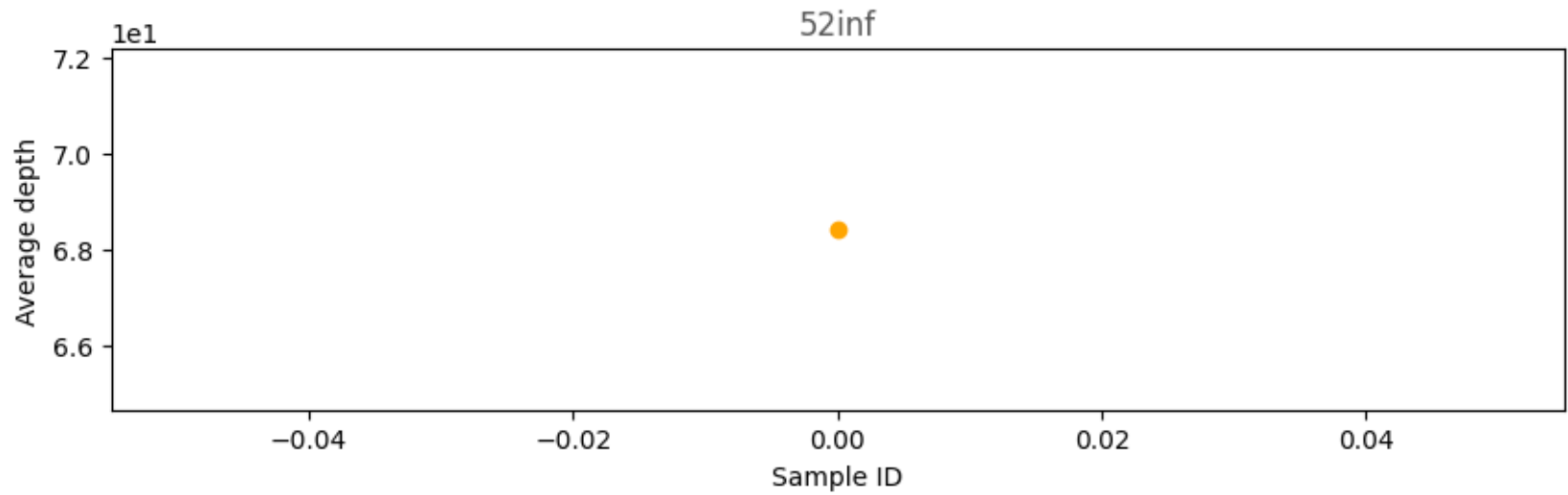
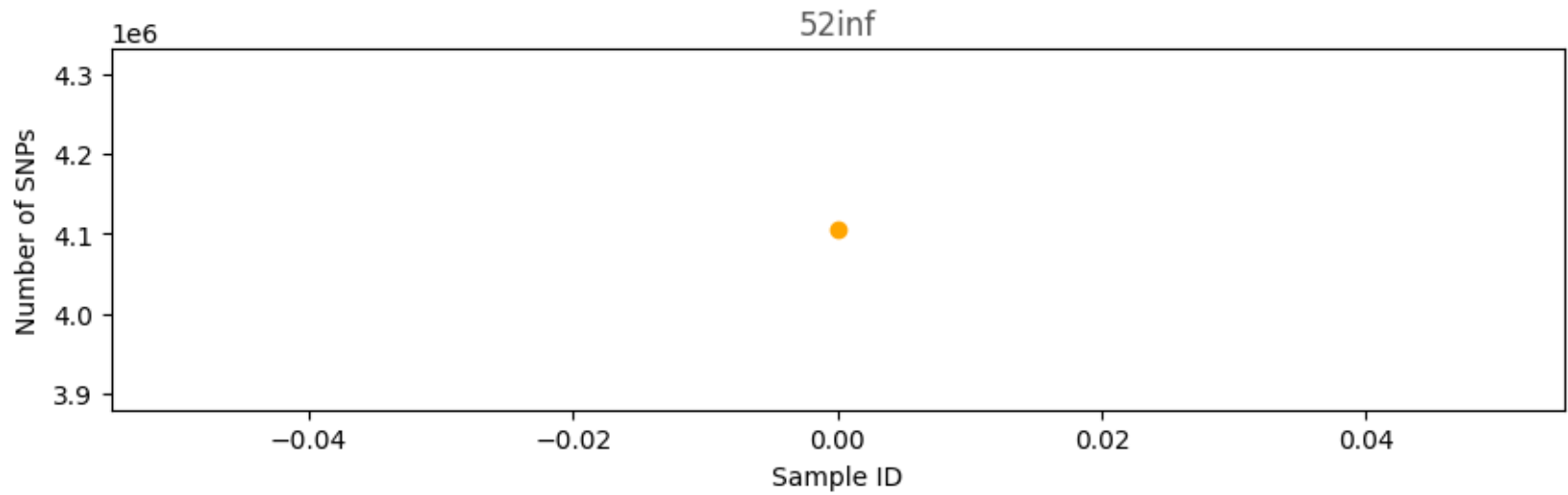# Ts/Tv by sample

52inf
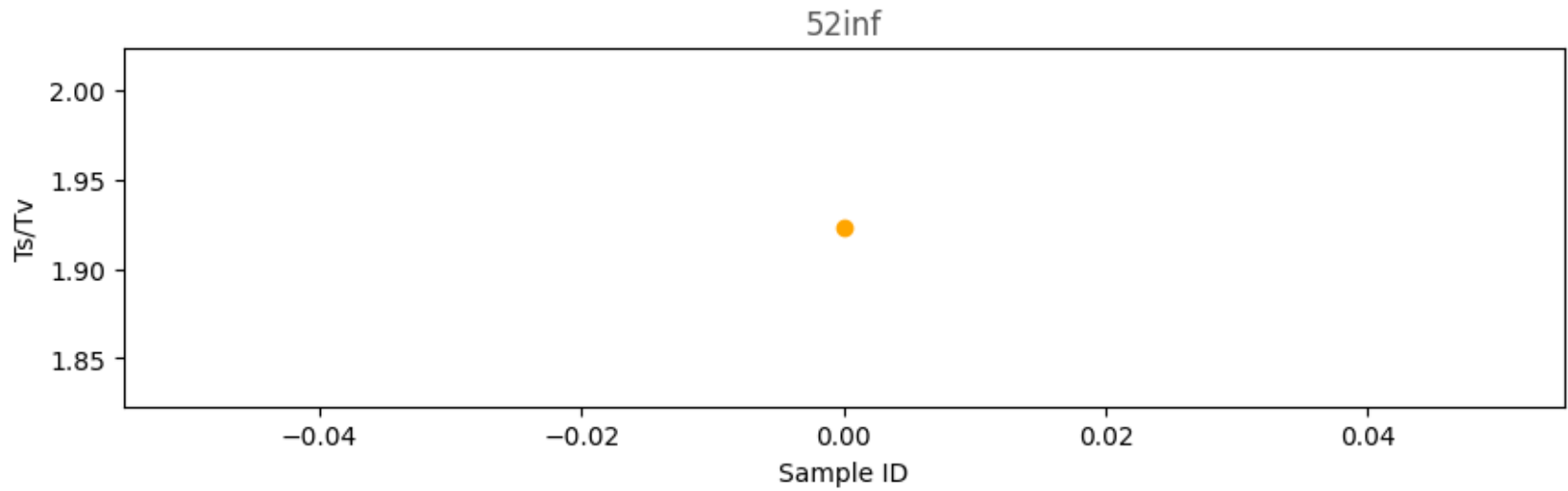
Hets vs non-ref Homs by sample
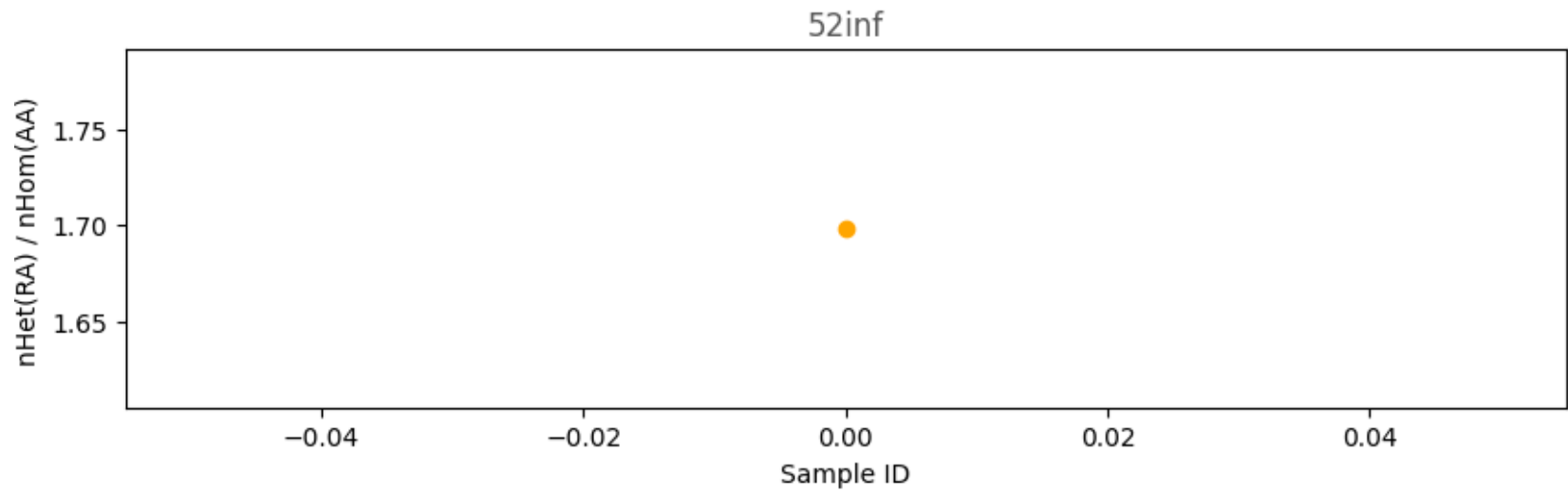
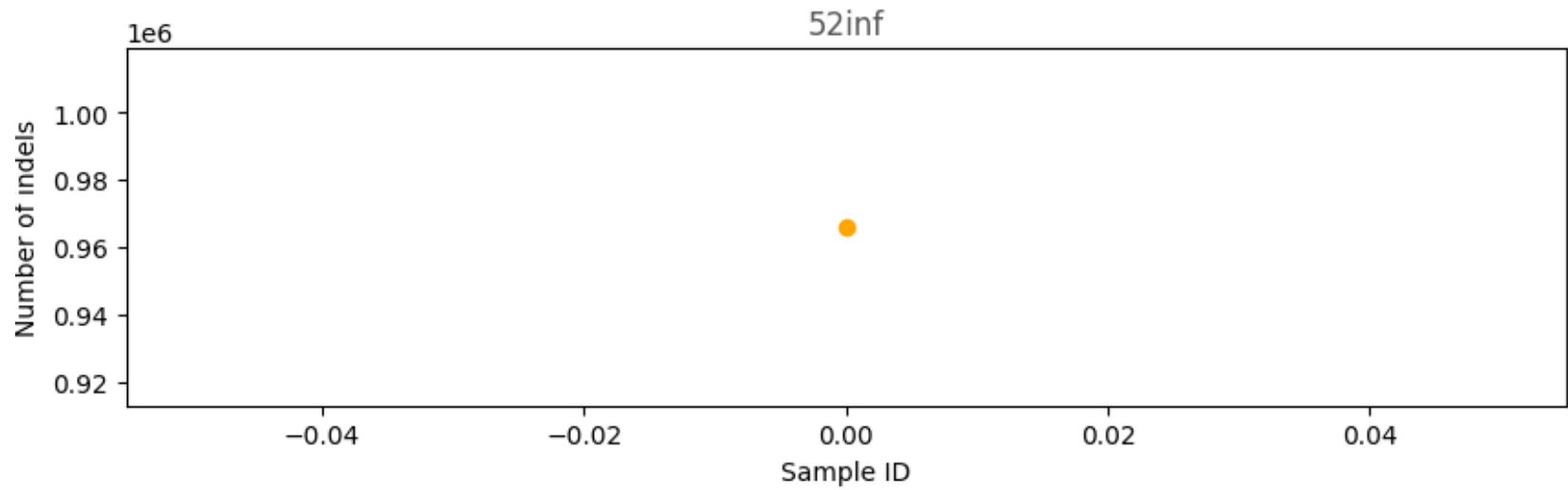# Singletons by sample (hets and homs)
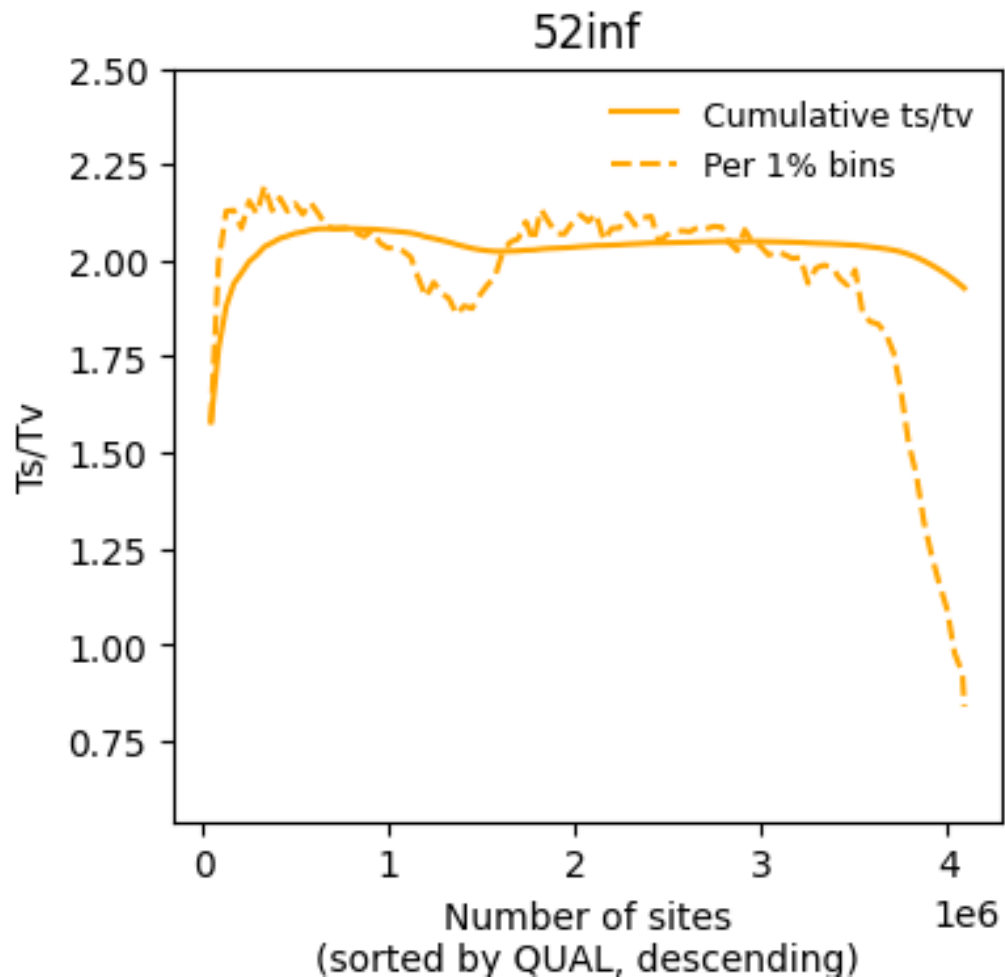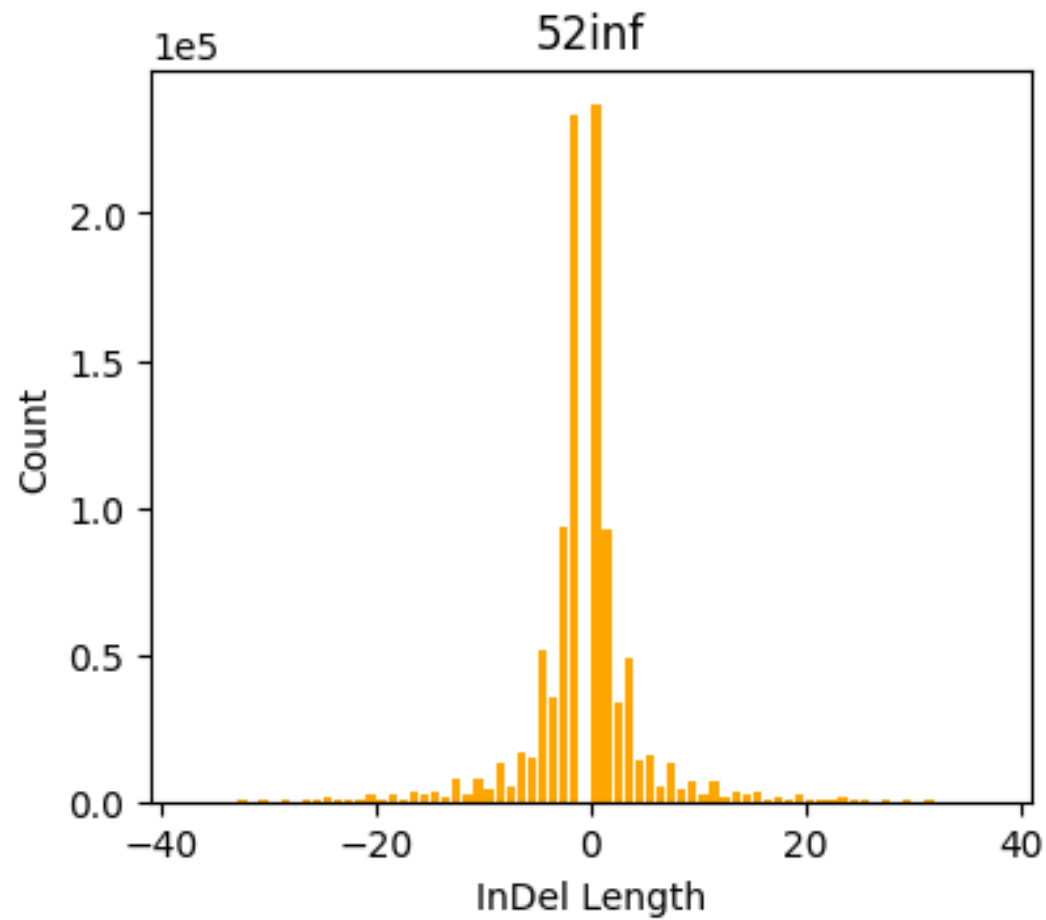
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

52inf

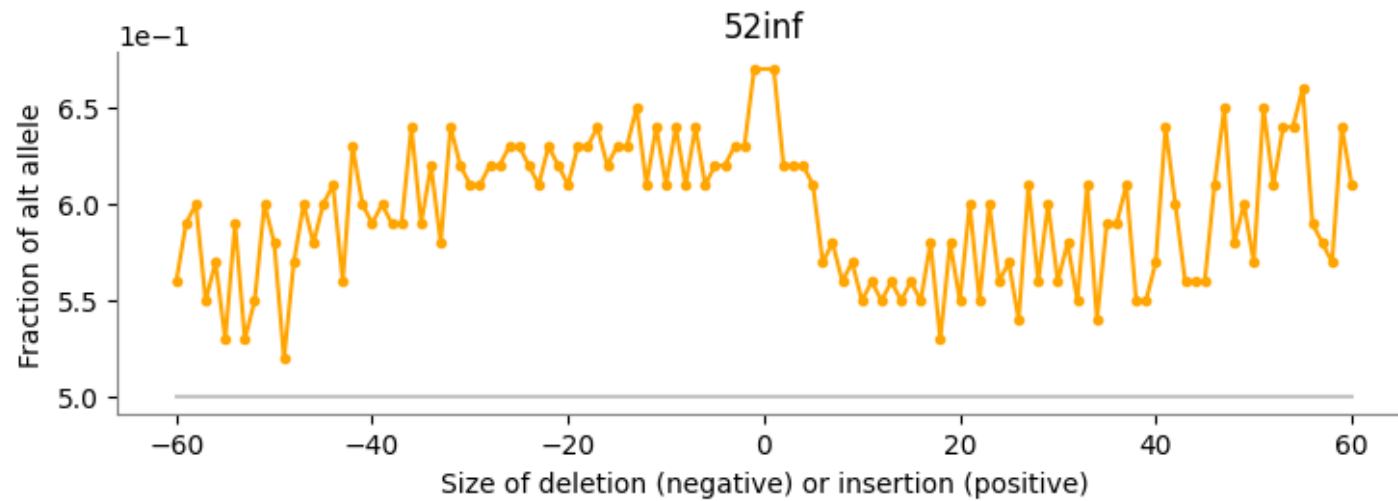# Indel distribution

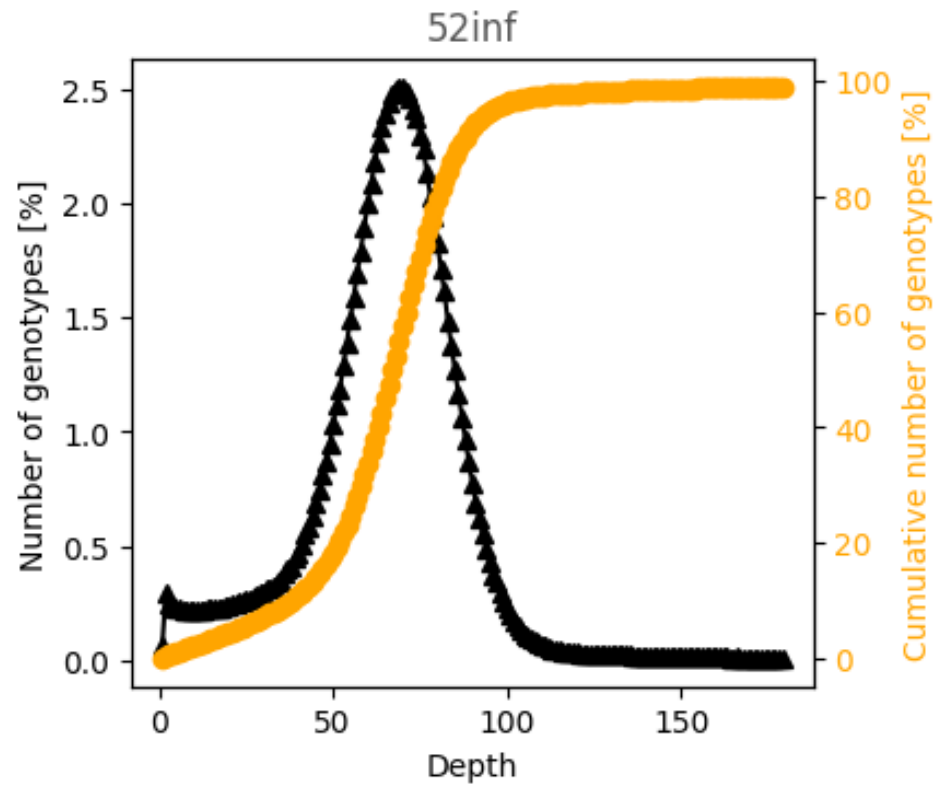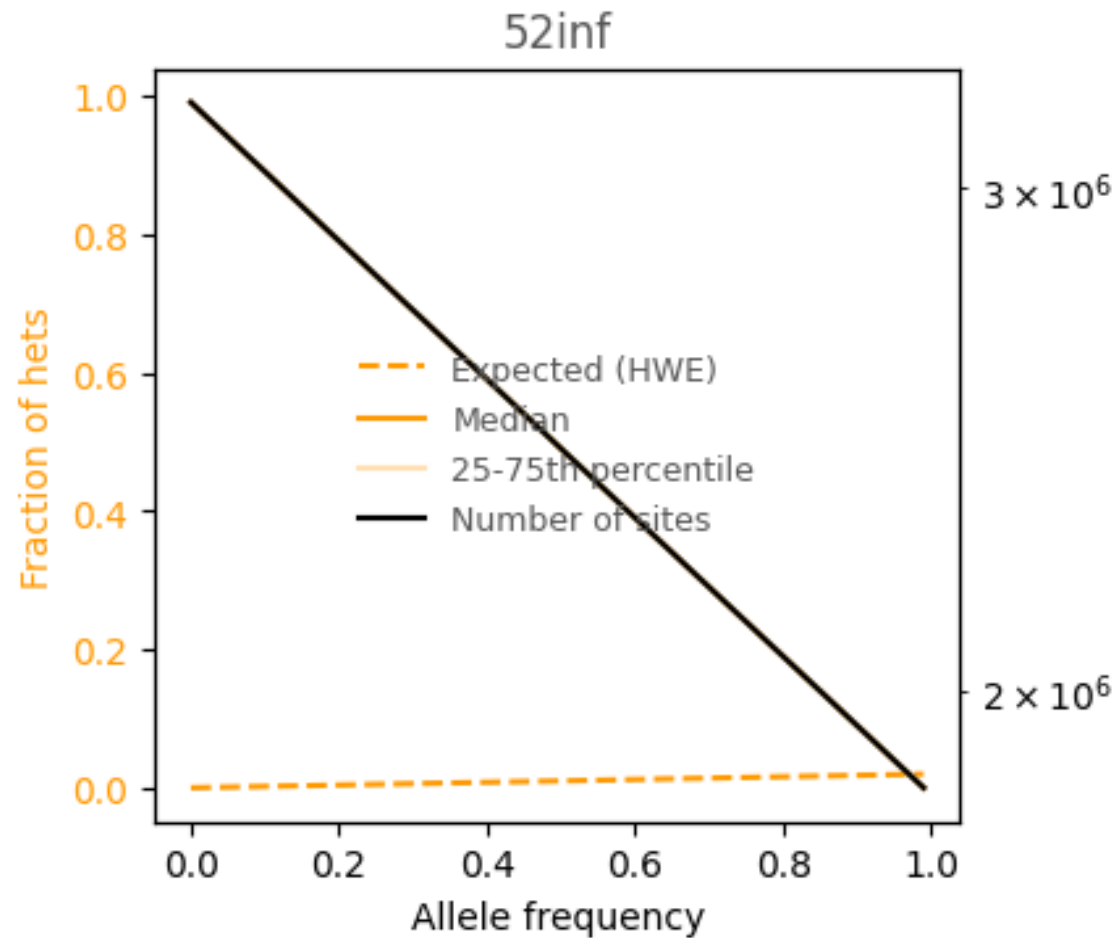# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types