Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
52iln	4,065,701	1.94	1.94	943,551	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
52iln	62.0%	1.91	66.3%	95,337	1,985

- 52iln .. `/ngc/projects2/gm/data/archive/2022/variants/snv/52ilngstm-103917223431-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01907-221005_A01411_AH2YY2DSX5-EXT_LAB`
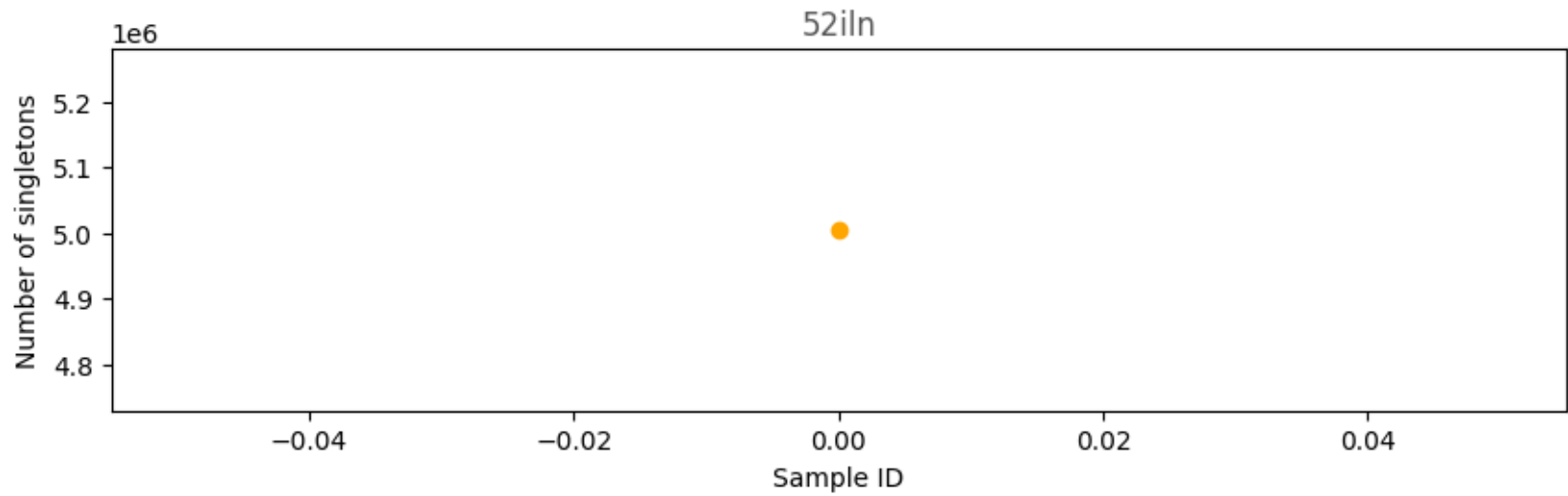 `KA_NGCWGS-NGCWGS05720_snv_germline_raw.haplotype_caller.vcf.gz`
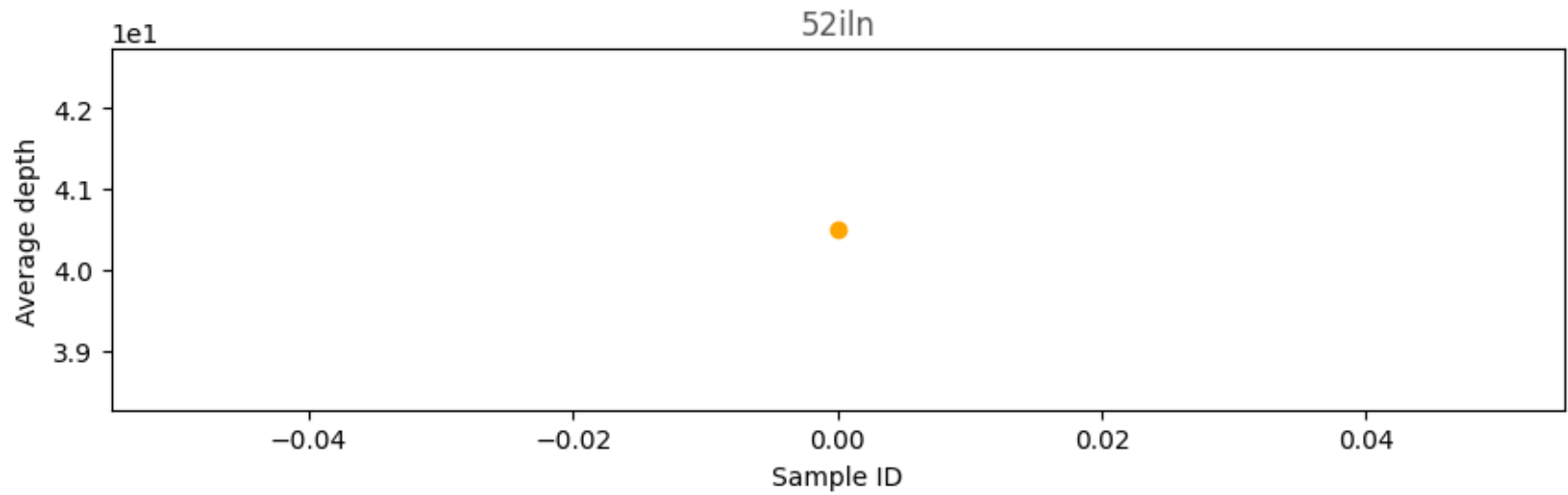
Ts/Tv by sample
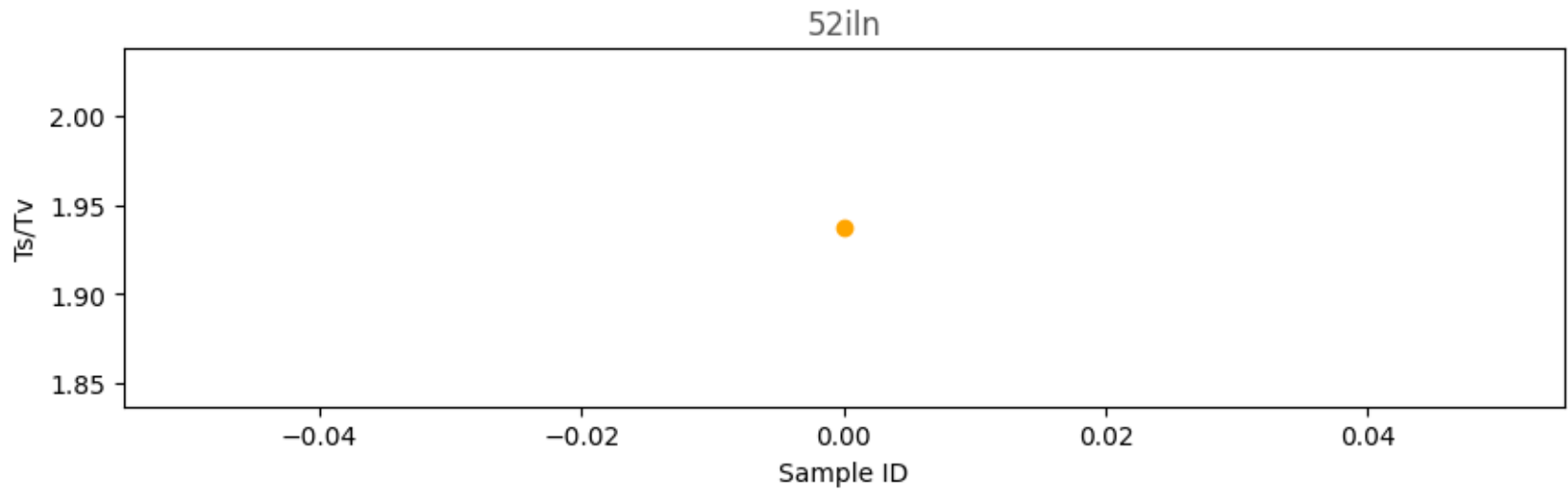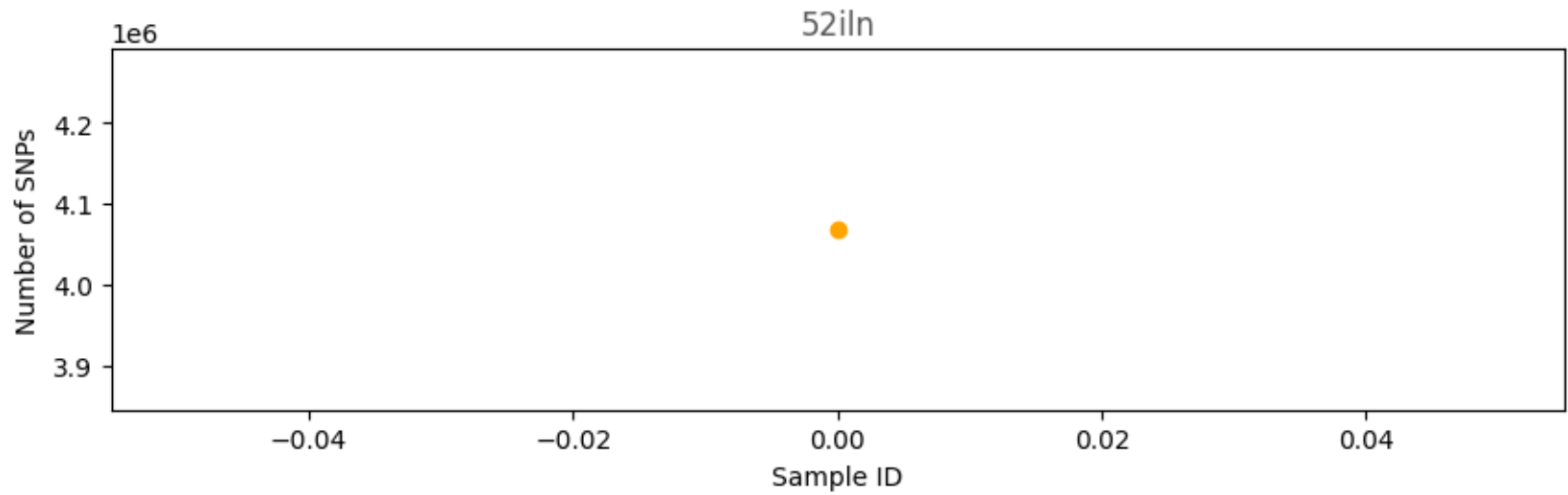
Hets vs non-ref Homs by sample
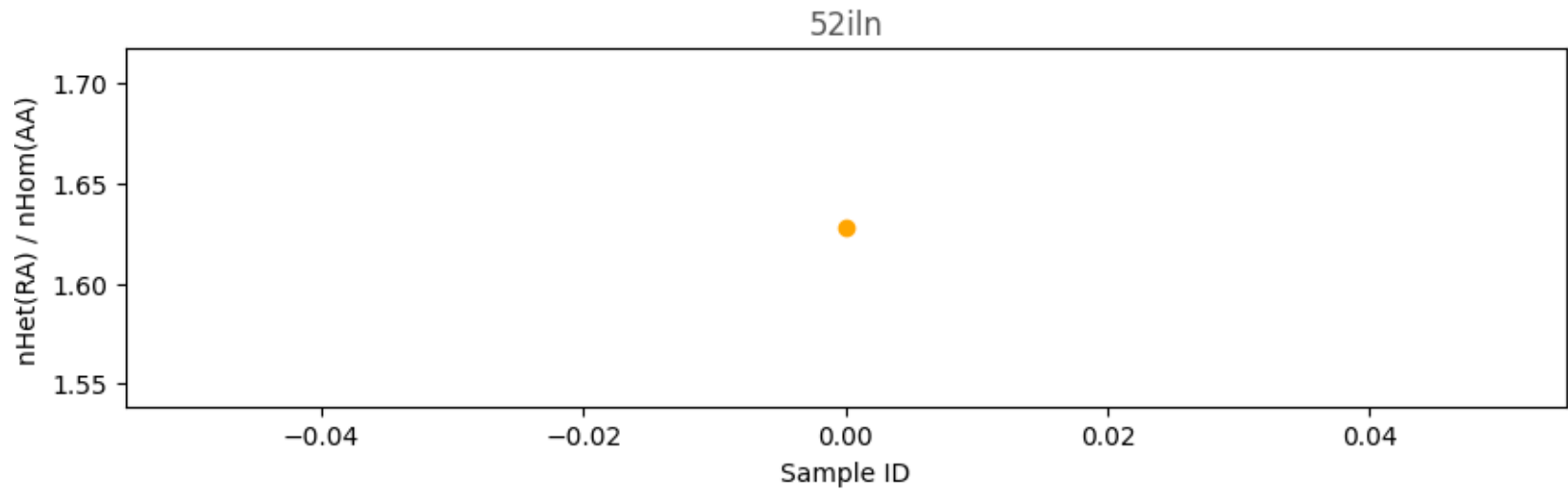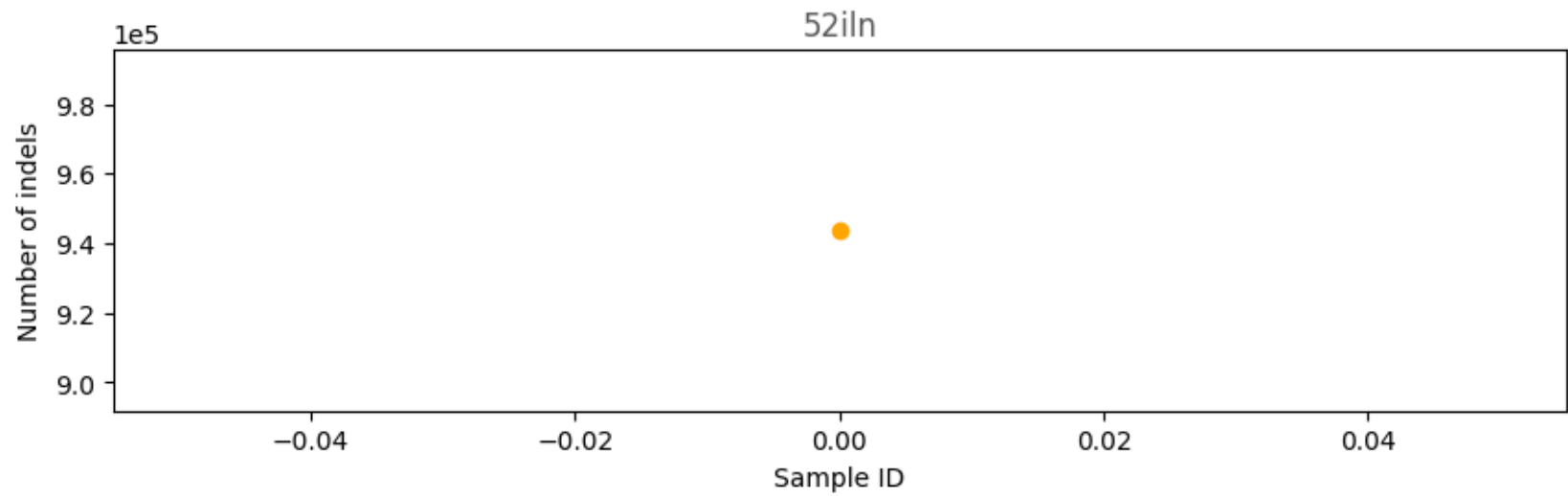
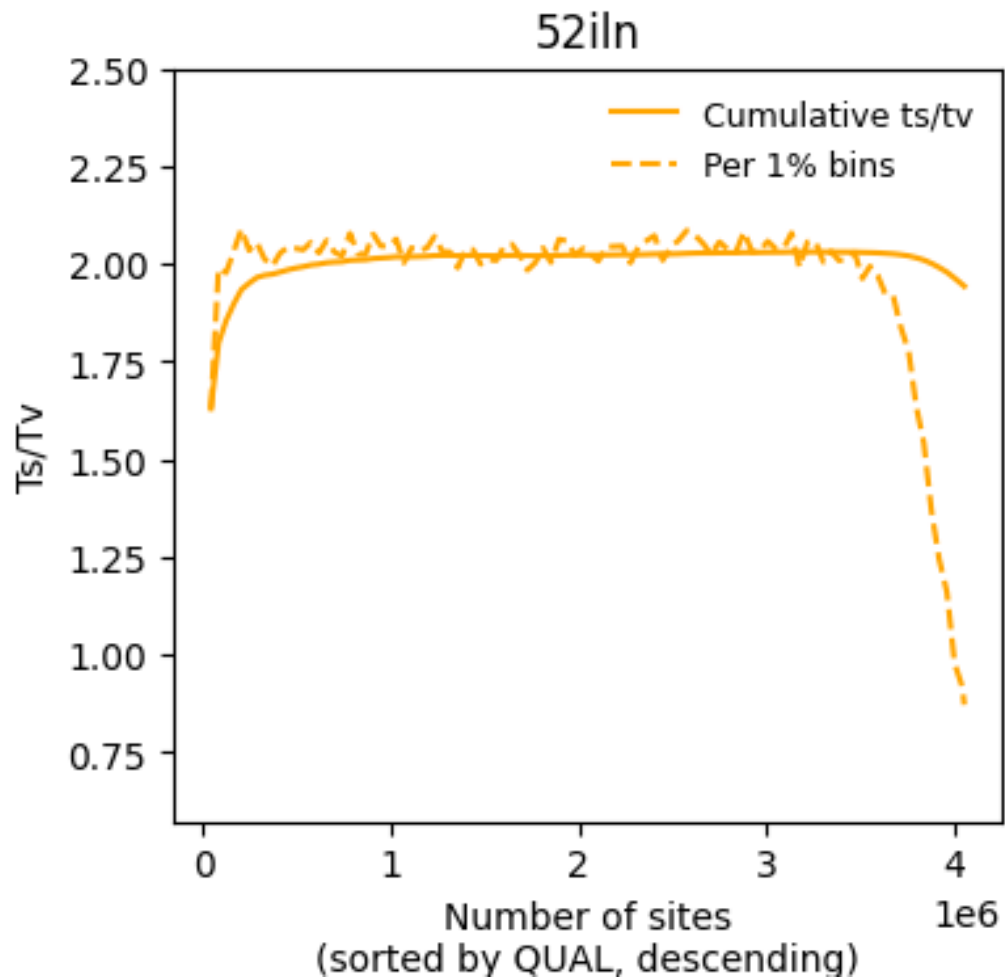# Singletons by sample (hets and homs)

Average depth by sample

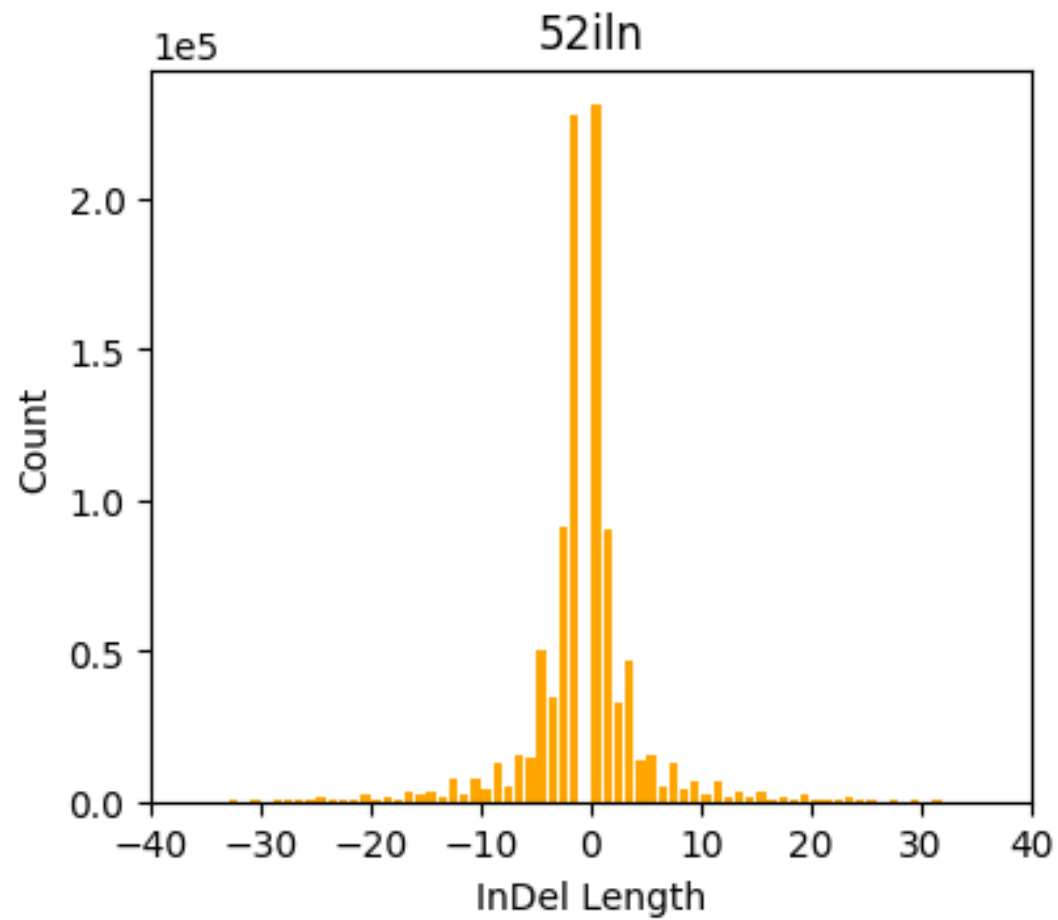# Number of SNPs by sample

Number of indels by sample
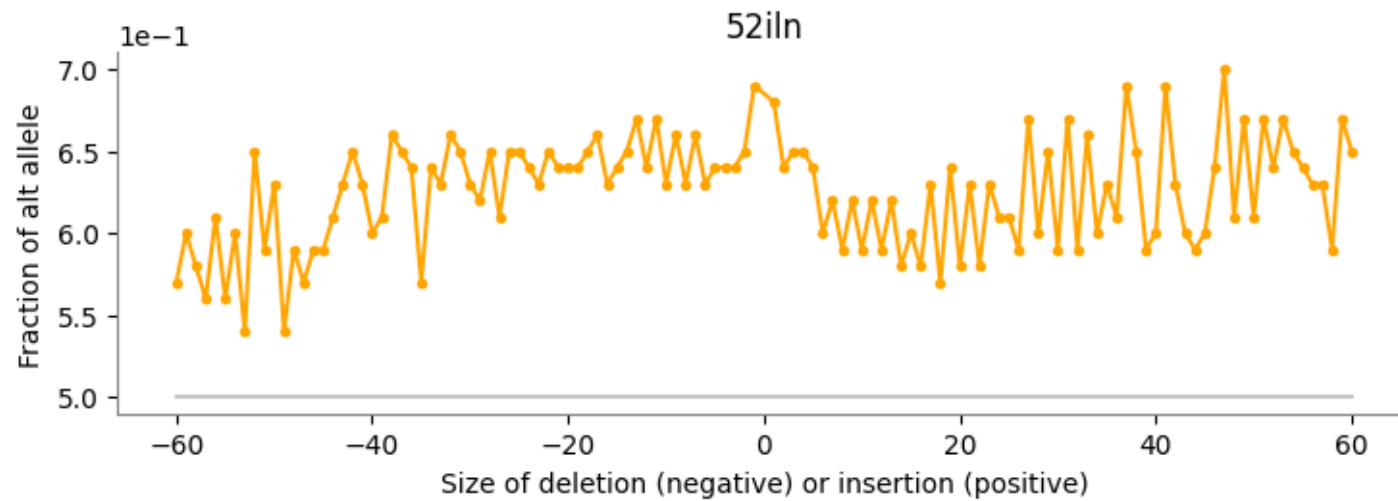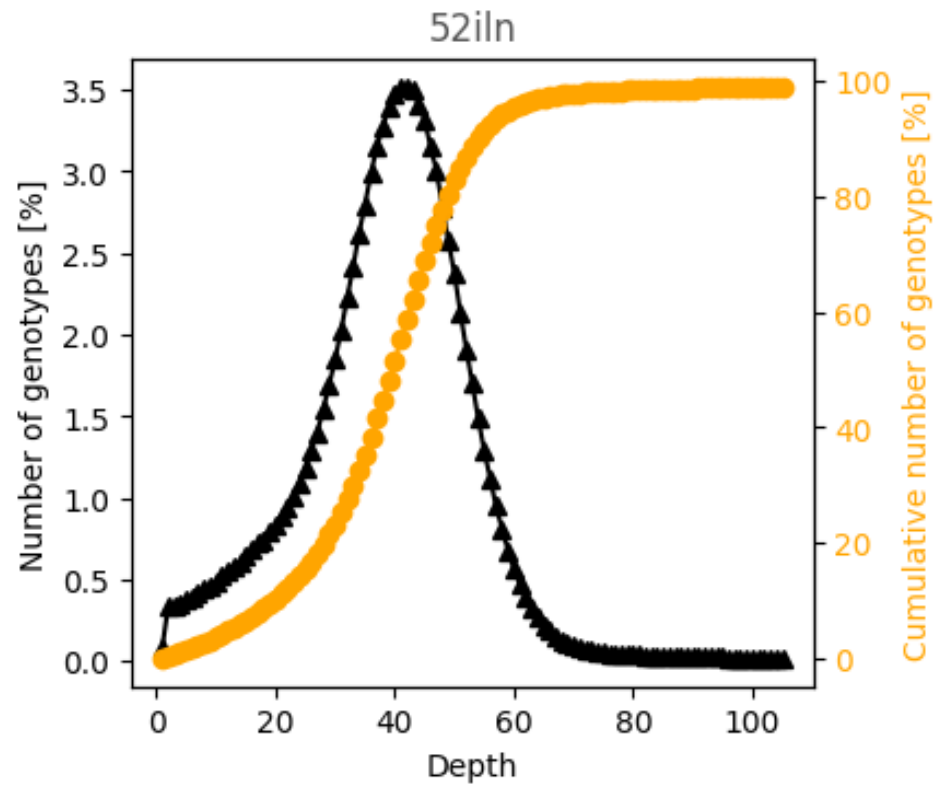
Ts/Tv stratified by QUAL

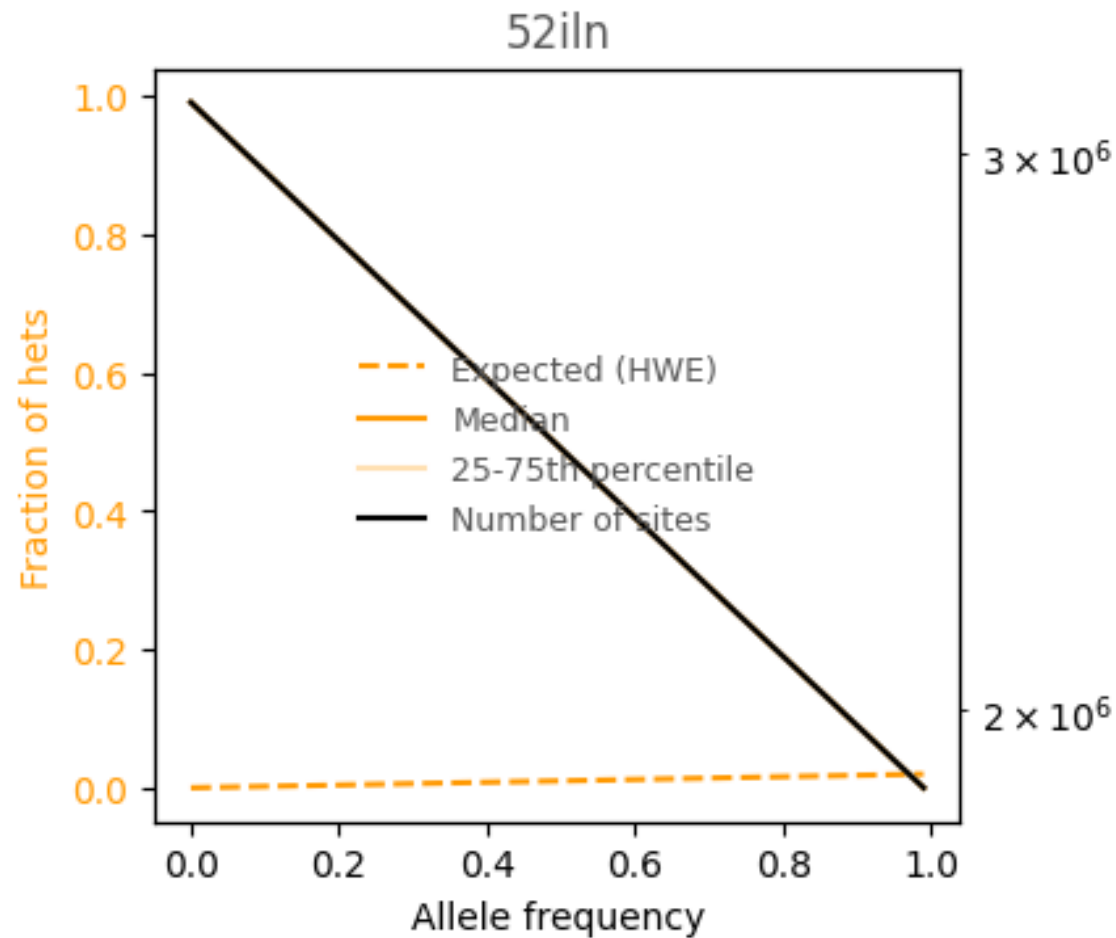

52iln

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types