Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
58iss	4,005,208	1.97	1.97	903,763	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
58iss	62.5%	1.95	65.0%	72,824	1,721

- 58iss .. `/ngc/projects2/gm/data/archive/2022/variants/snv/58issancf-103840959157-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220316_A01411_BHCF77DSX3-RHGM_LABKA_WGS-`
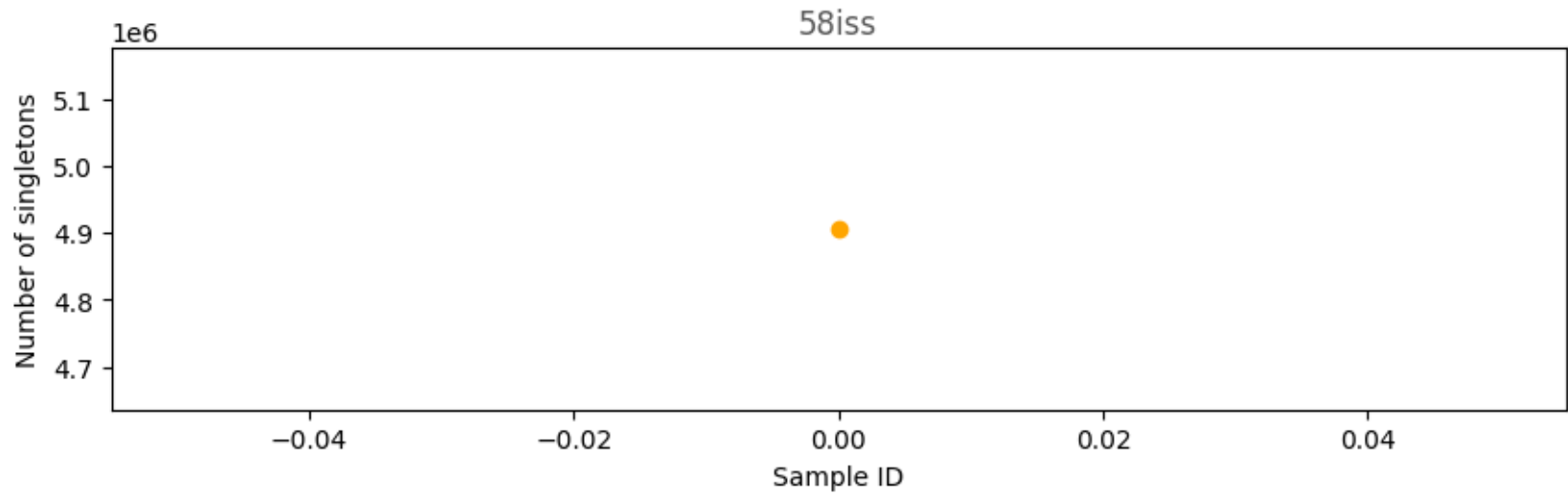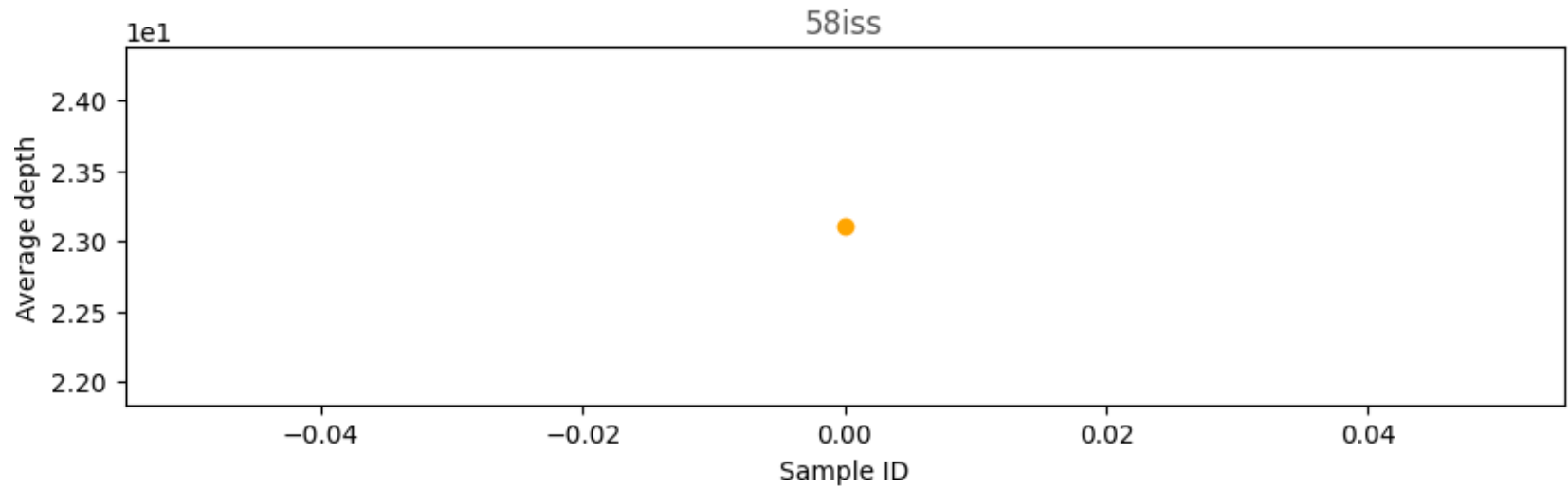 `WGS03920NY_snv_germline_raw.haplotype_caller.vcf.gz`
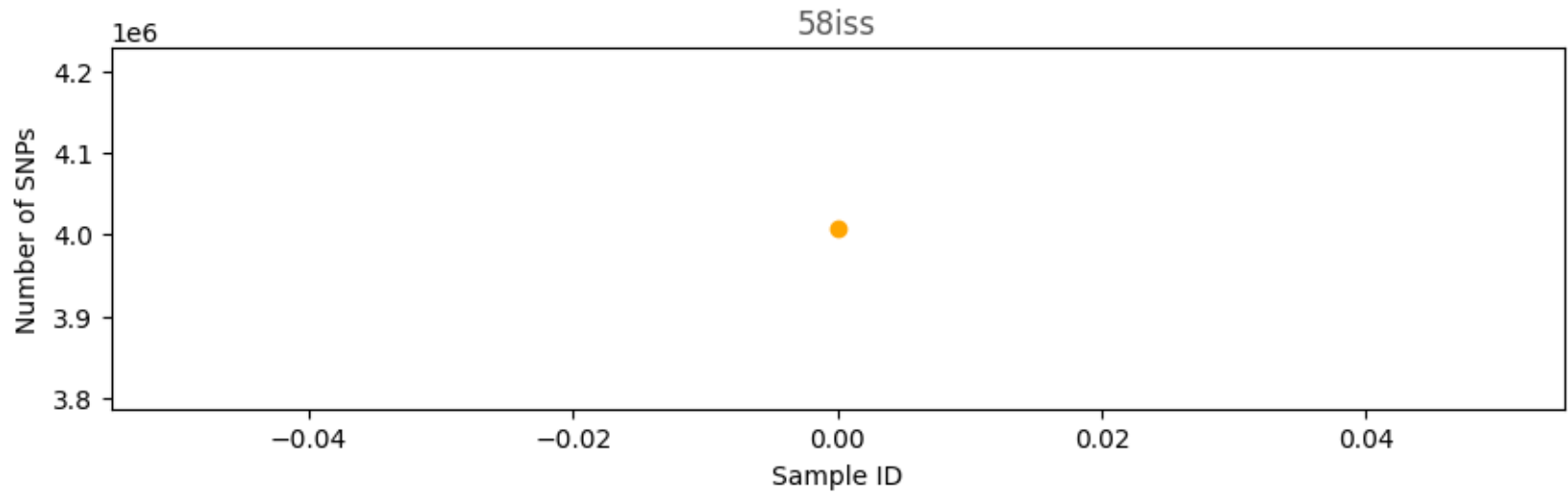
Ts/Tv by sample
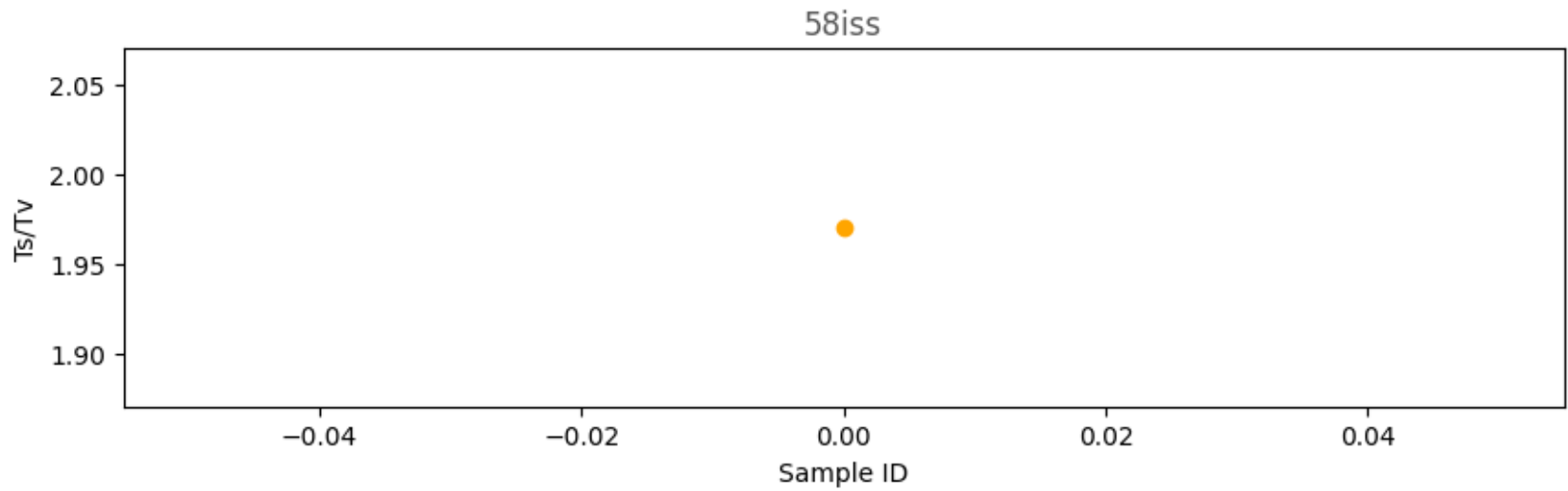
Hets vs non-ref Homs by sample
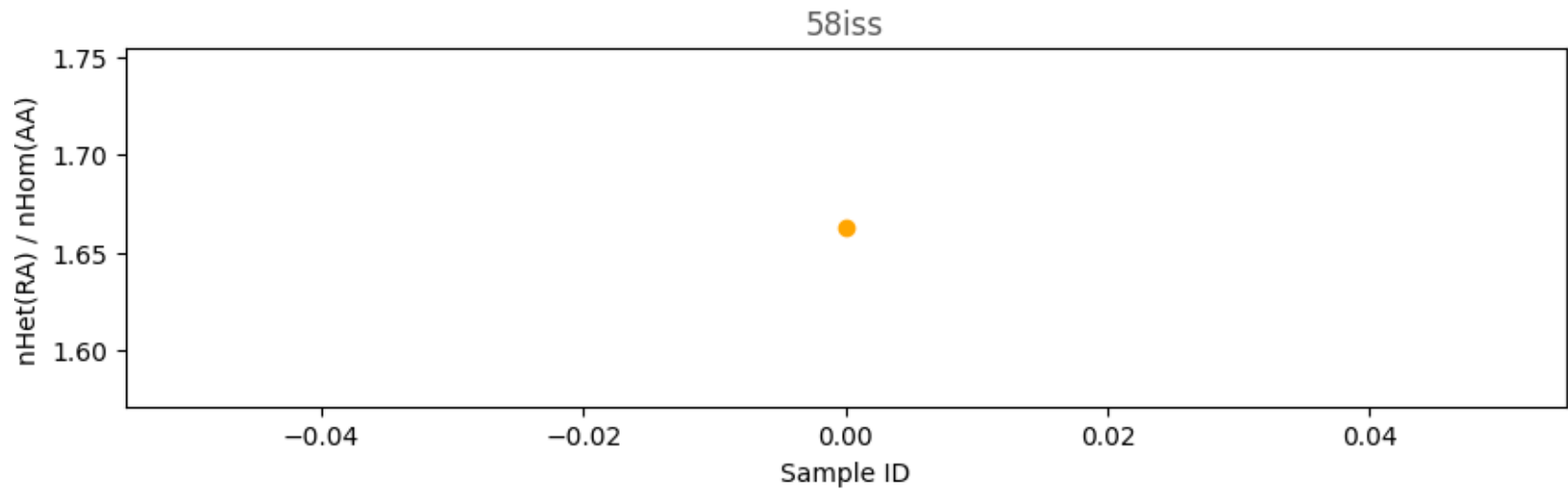
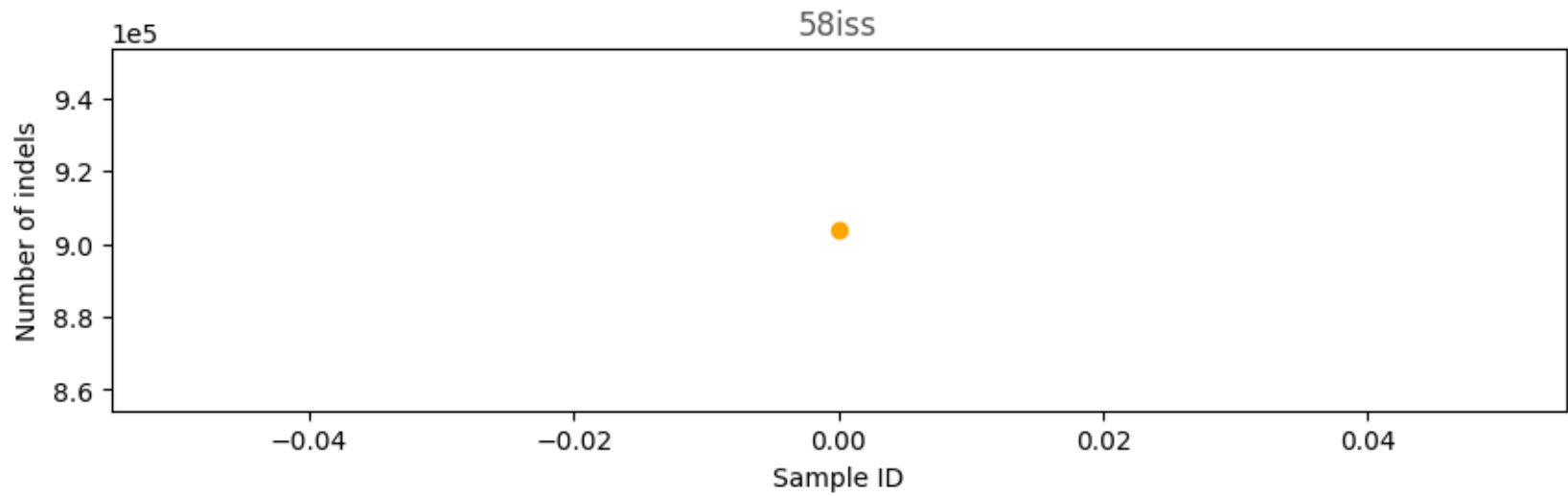# Singletons by sample (hets and homs)
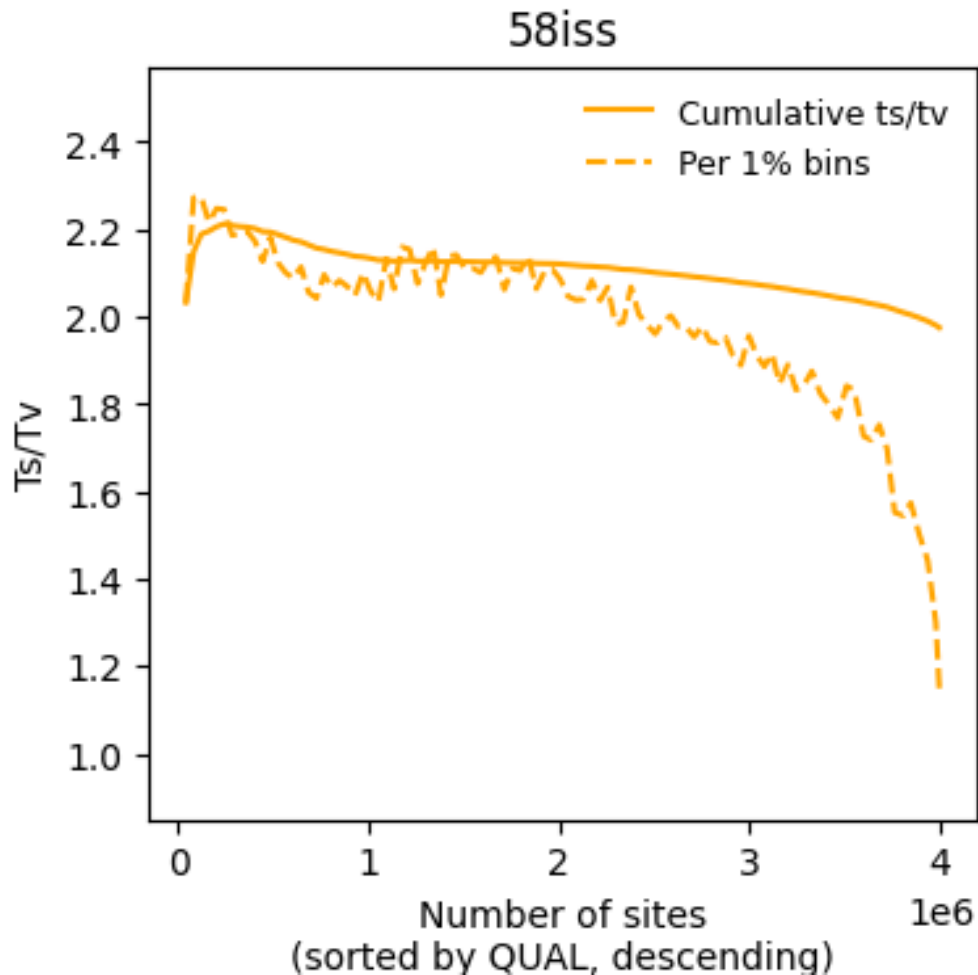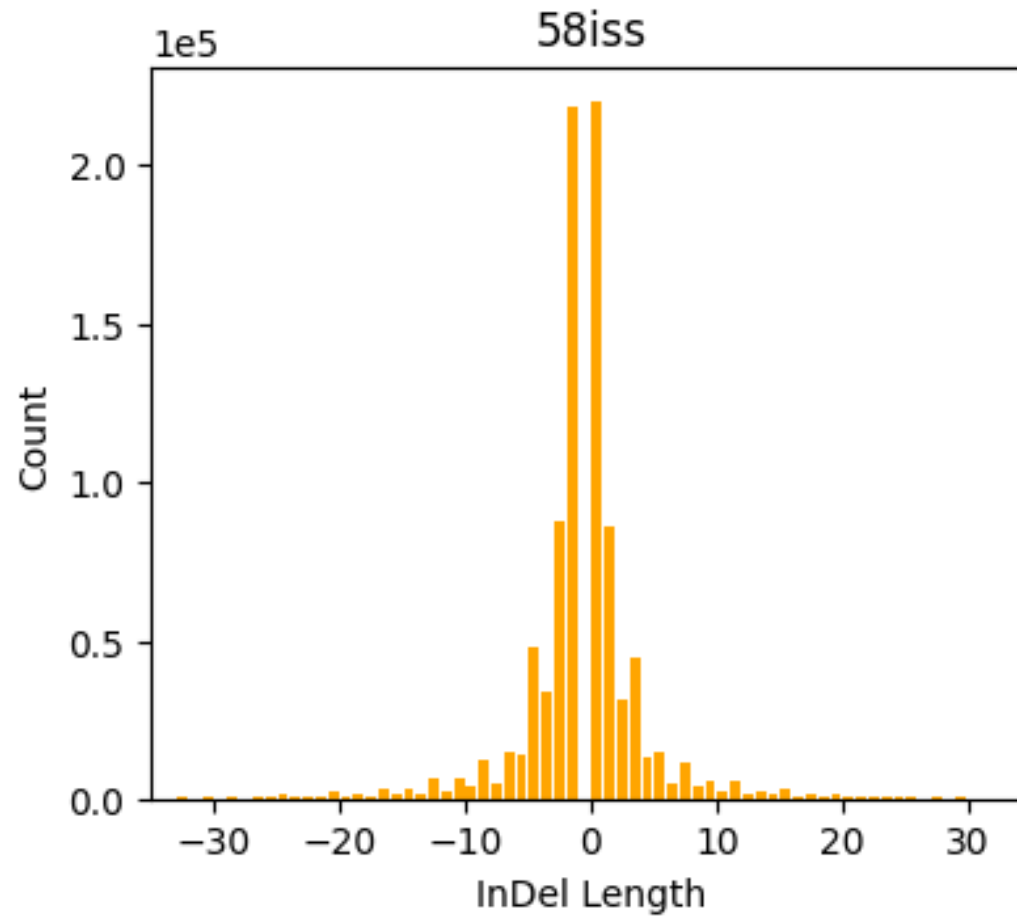
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

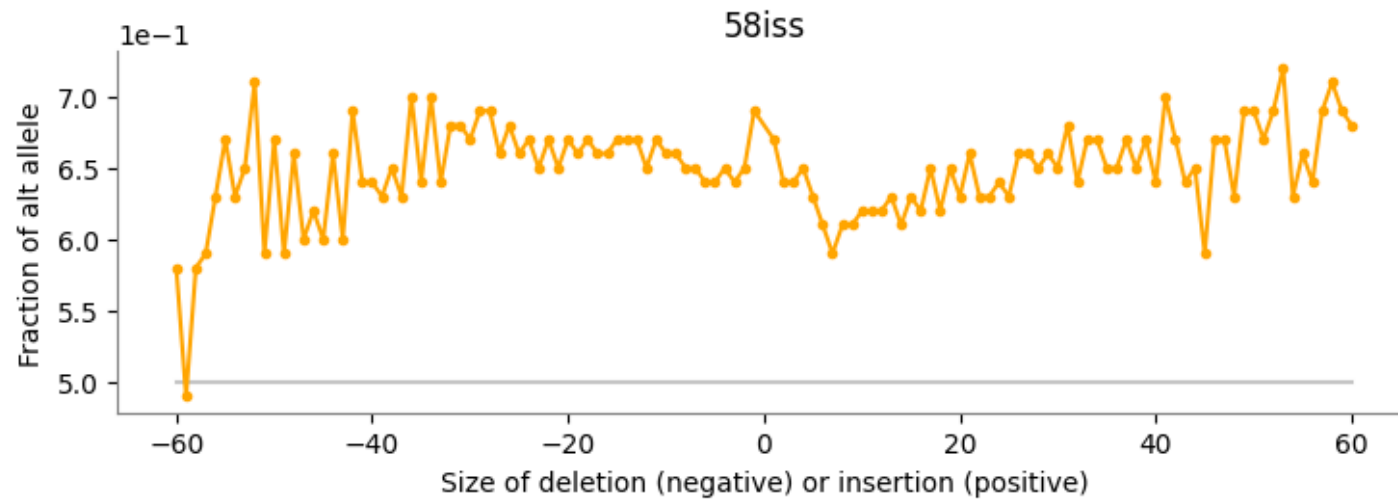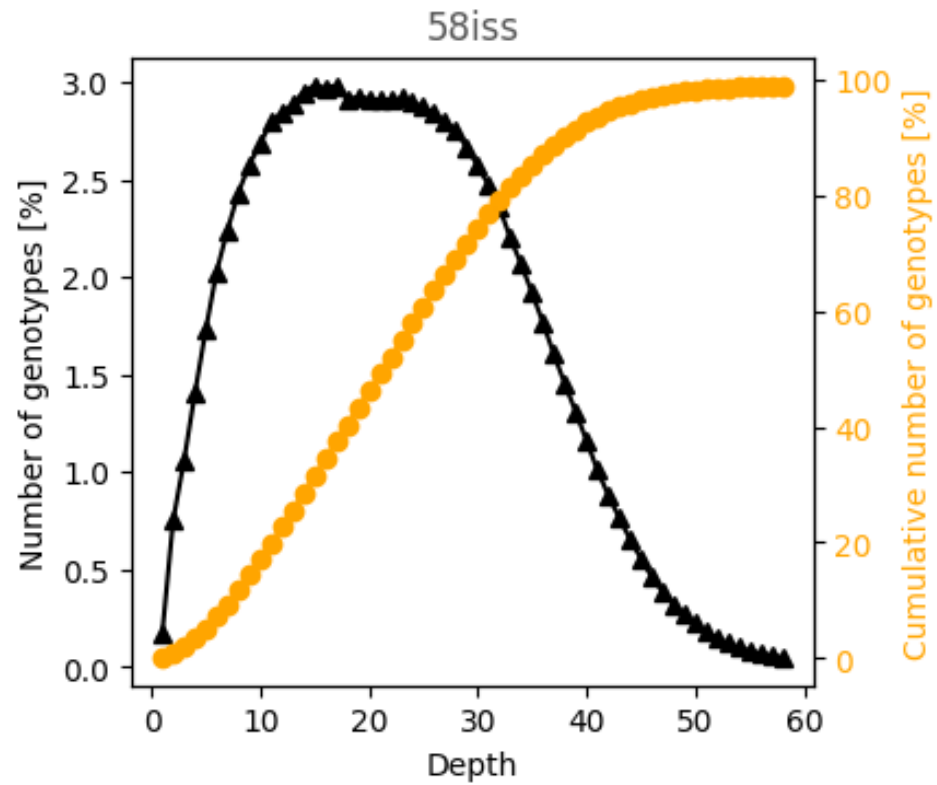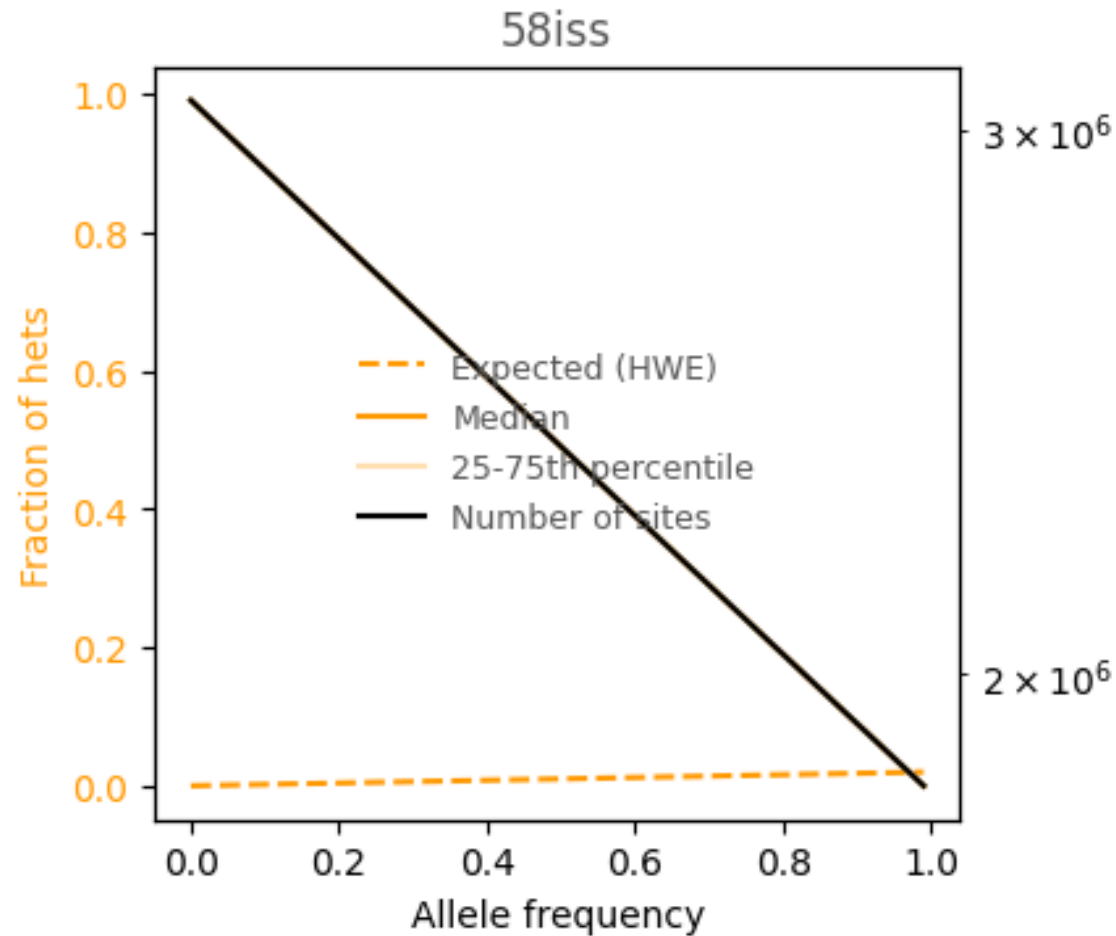# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types