Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
11ite	4,170,760	1.92	1.93	976,429	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
11ite	64.2%	1.89	69.5%	109,505	2,200

- 11ite .. `/ngc/projects2/gm/data/archive/2022/variants/snv/11iterocf-103851571457-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01282-220907_A01411_AHYN3VDSX3-EXT_LAB`
  `KA_NGCWGS-NGCWGS05058_22RKG006946x01_snv_germline_raw.haplotype_caller.vcf.gz`
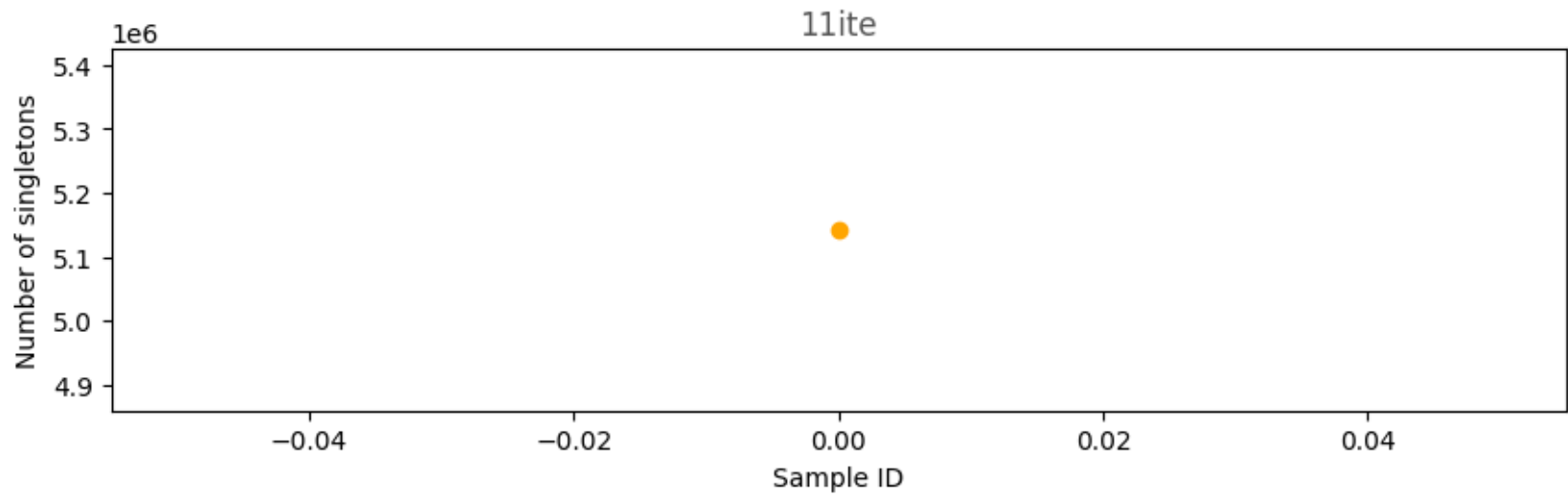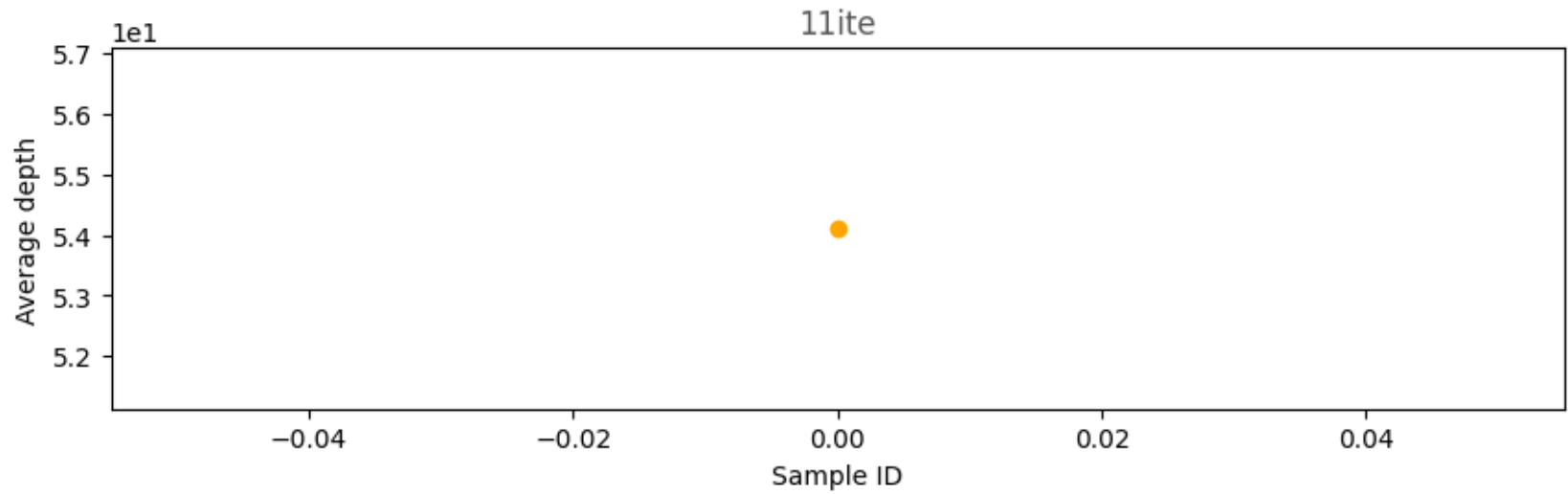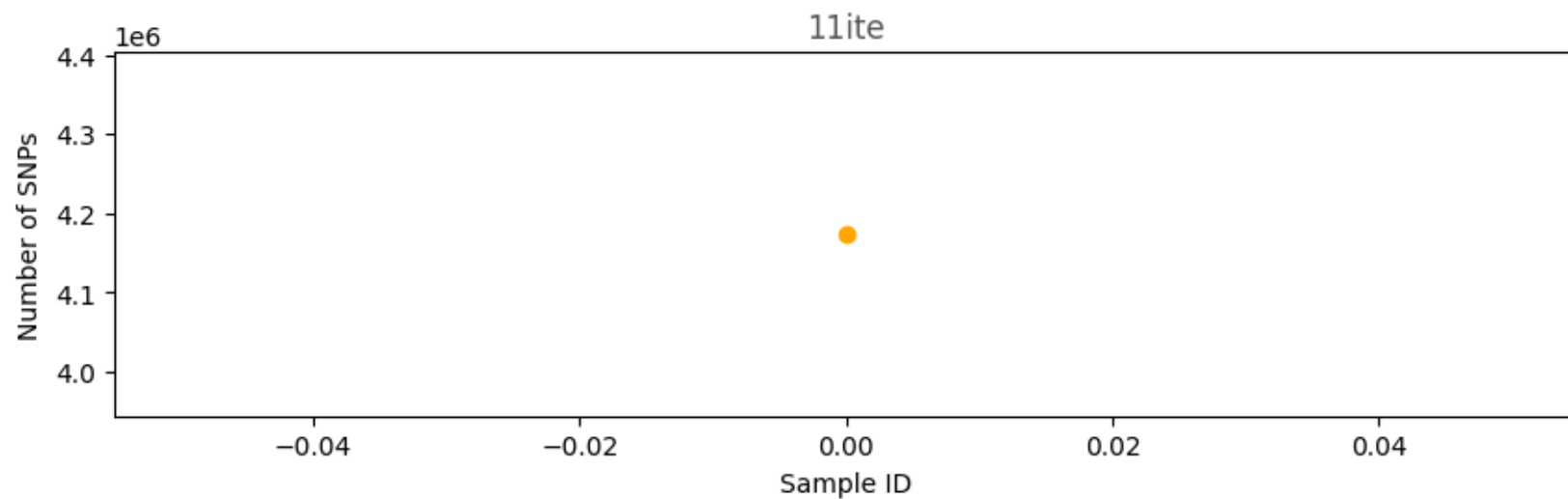
Ts/Tv by sample
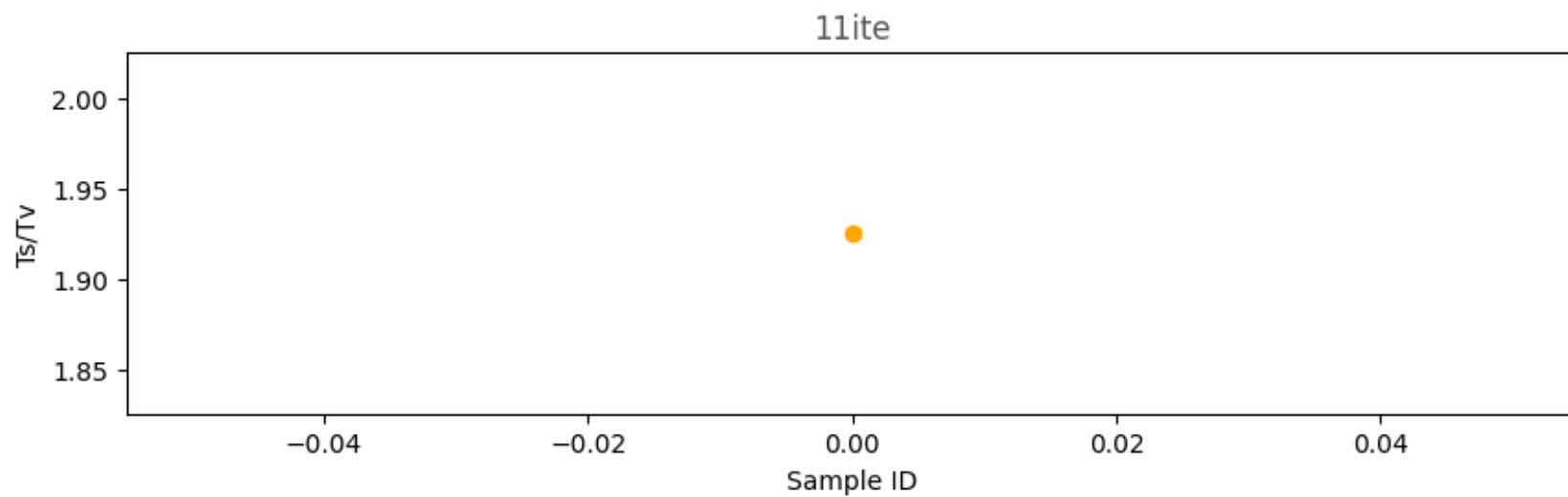
Hets vs non-ref Homs by sample
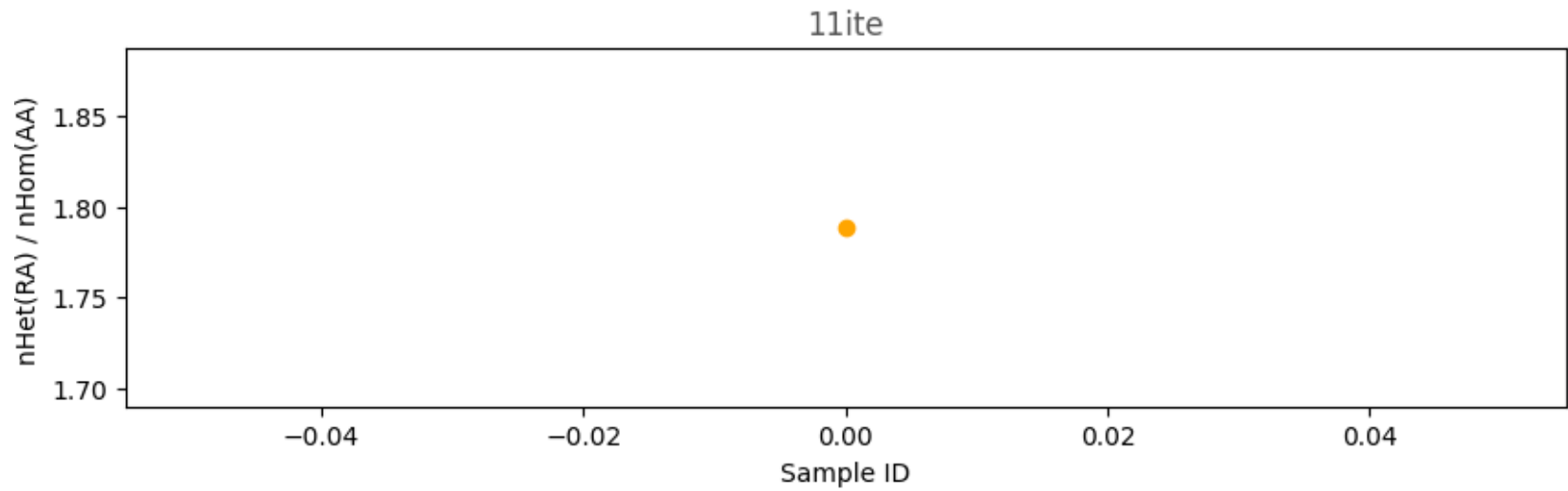
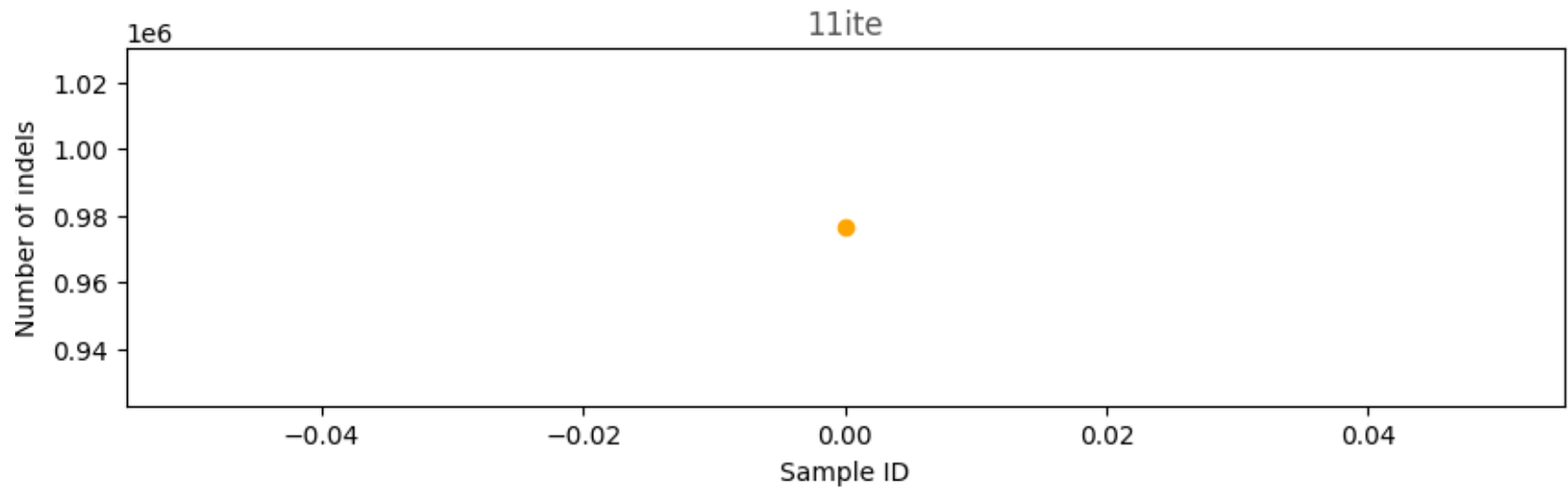# Singletons by sample (hets and homs)

Average depth by sample
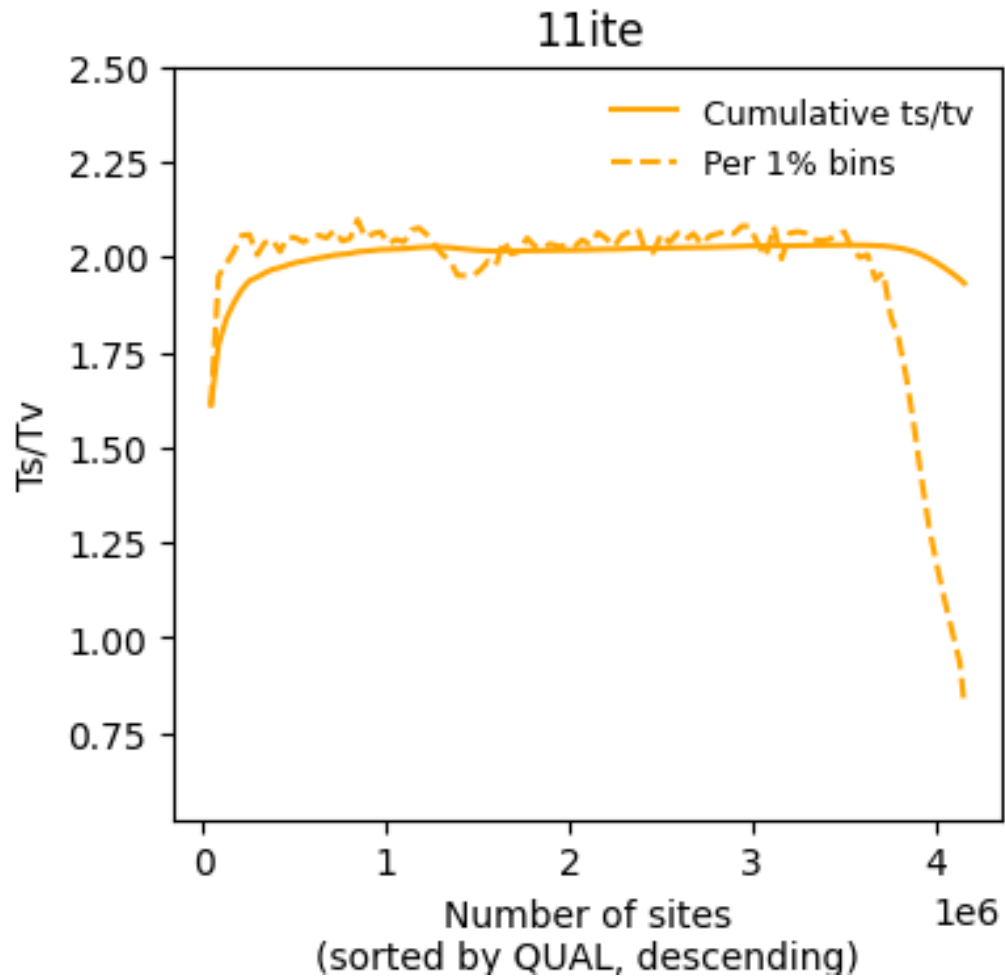
Number of SNPs by sample
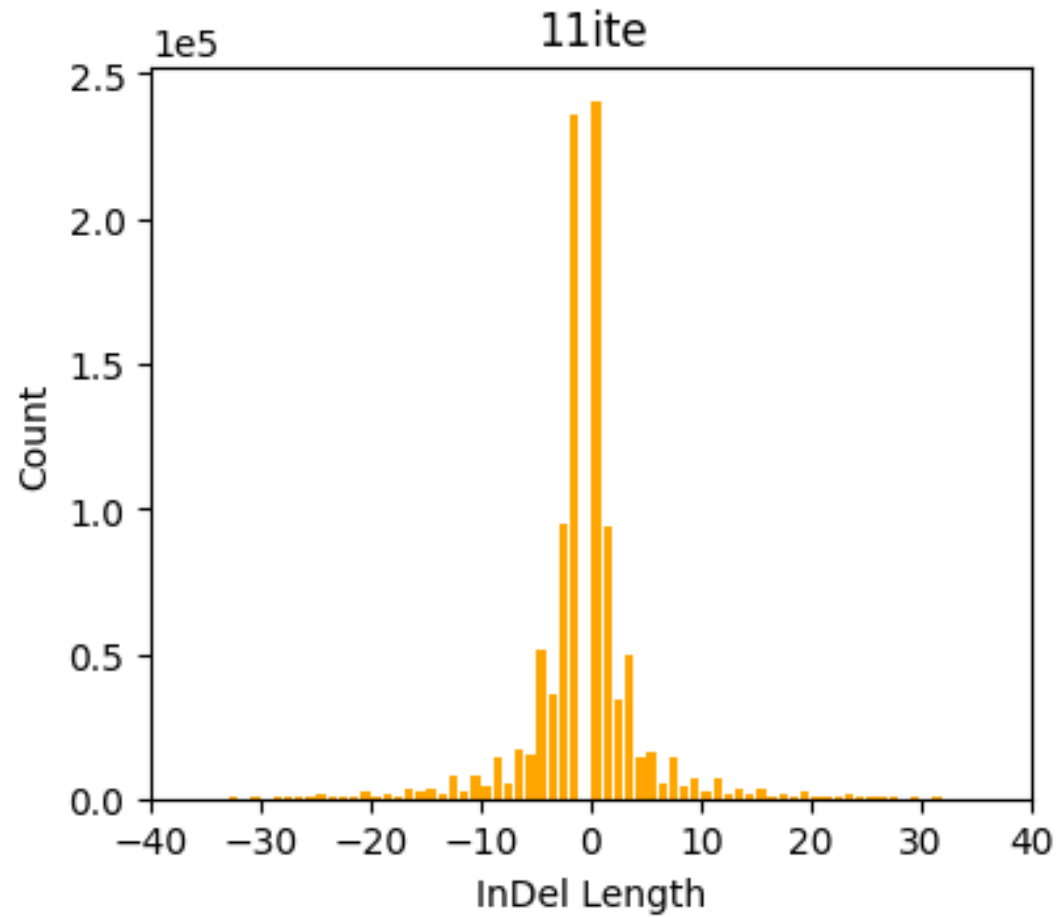
Number of indels by sample
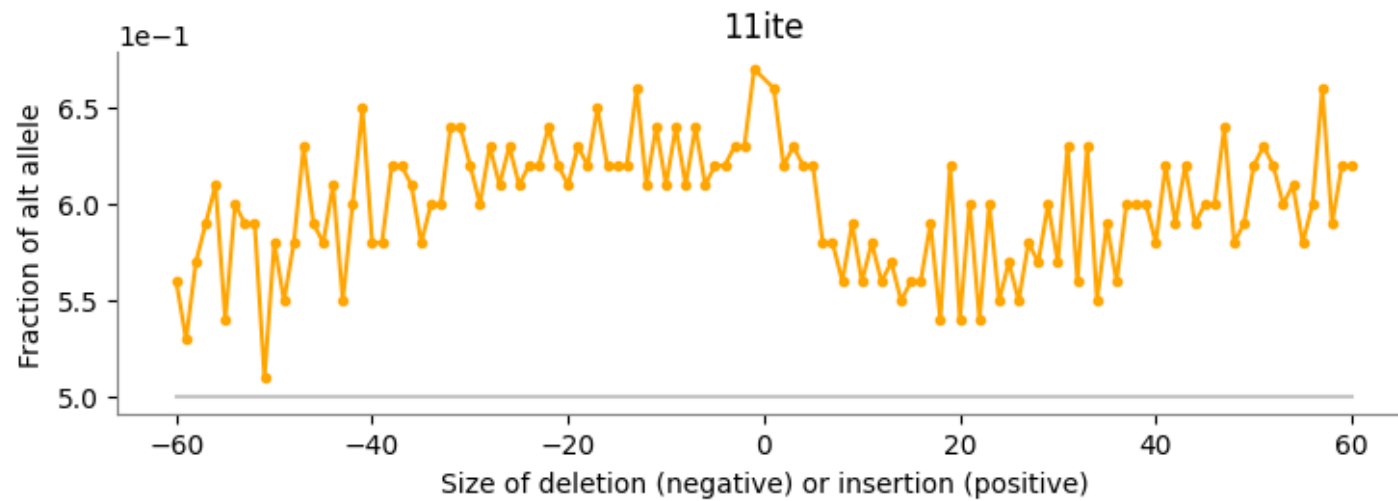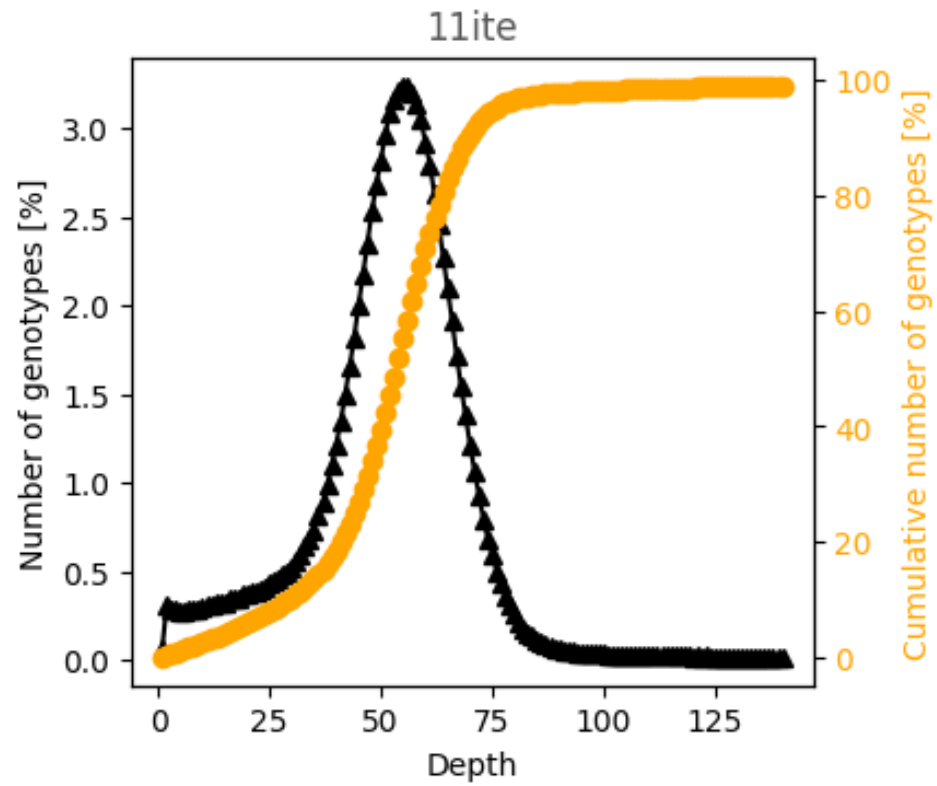
Ts/Tv stratified by QUAL
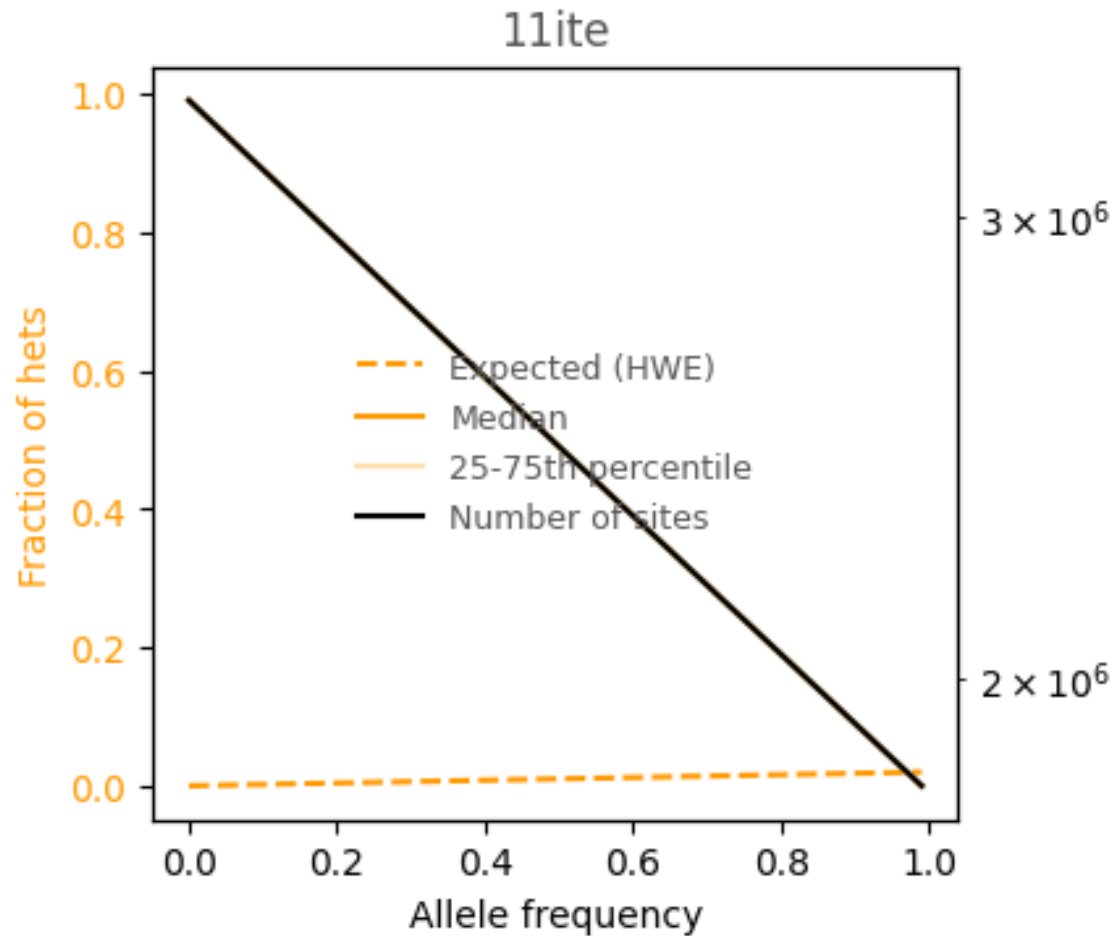
Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types