Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
59ive	4,069,710	1.94	1.95	936,745	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
59ive	63.1%	1.91	67.3%	94,494	2,077

- 59ive .. `/ngc/projects2/gm/data/archive/2022/variants/snv/59iveieaf-103870542181-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00817-220525_A01176_AHJ5HGDSX3-EXT_LAB`
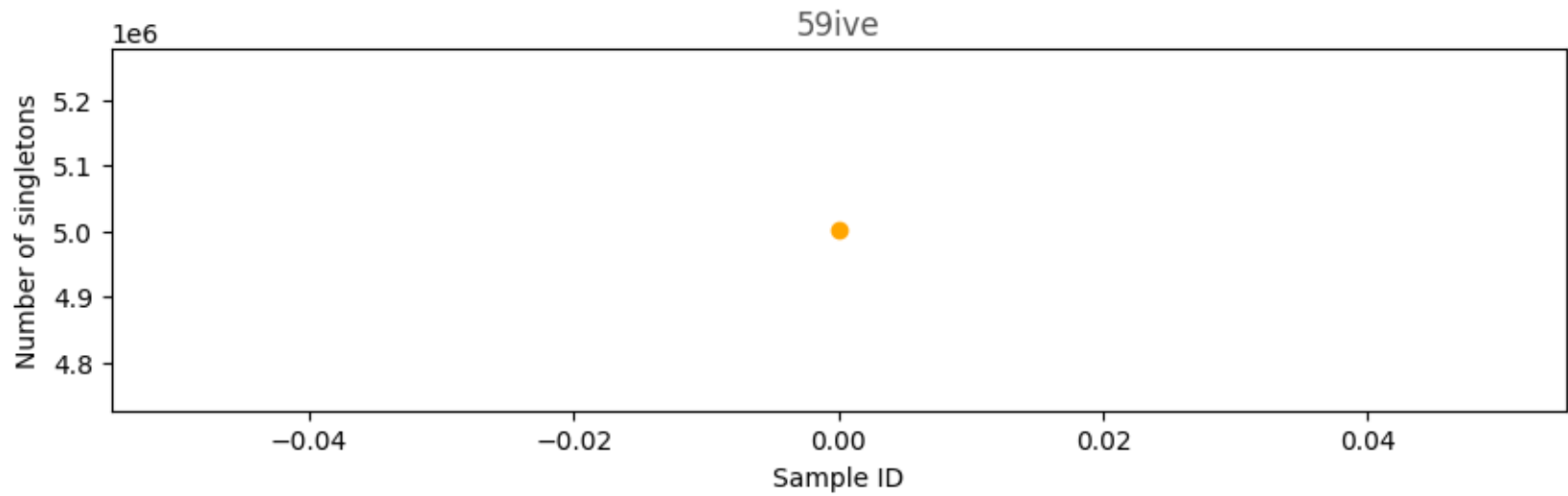 `KA_NGCWGS-NGCWGS04323_snv_germline_raw.haplotype_caller.vcf.gz`
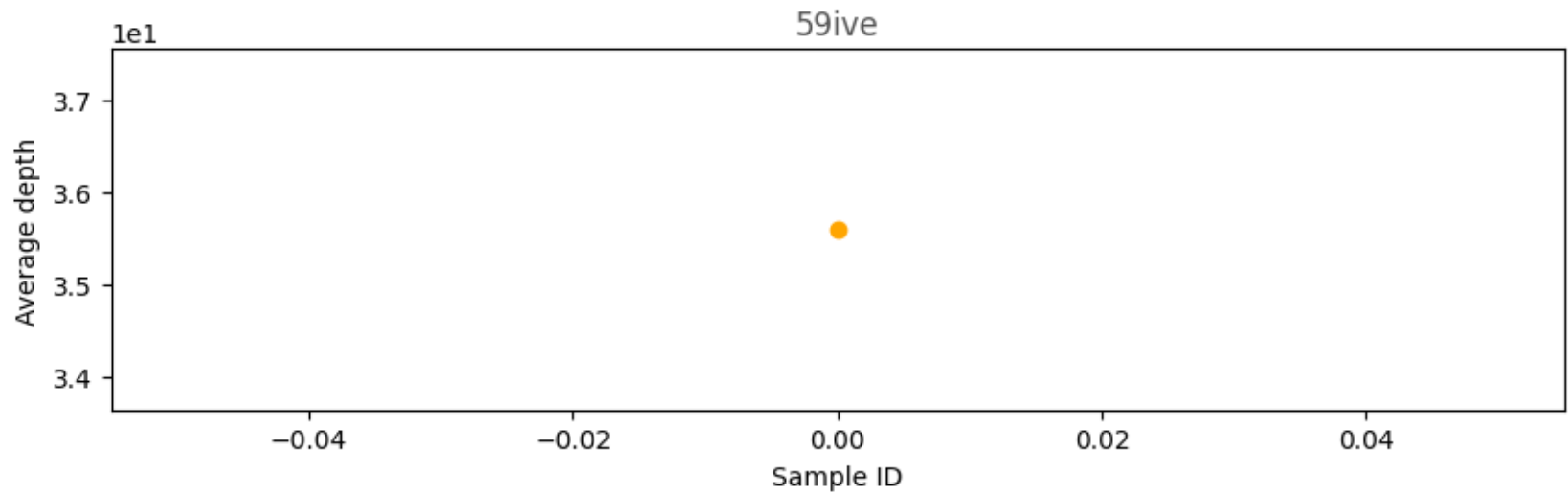
Ts/Tv by sample
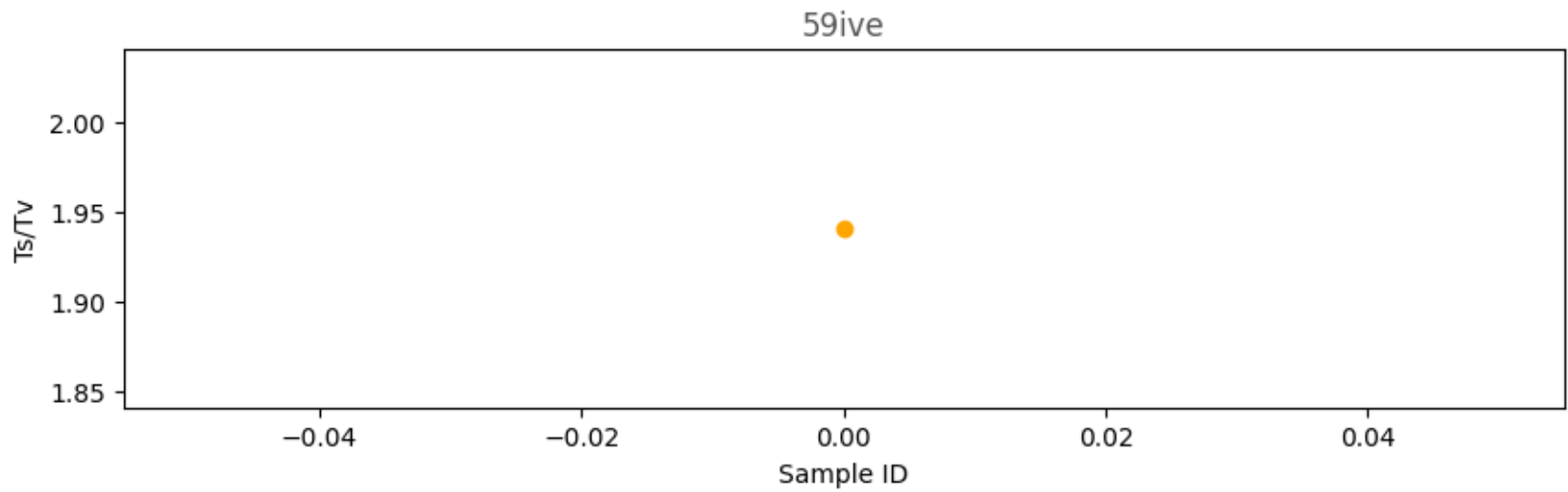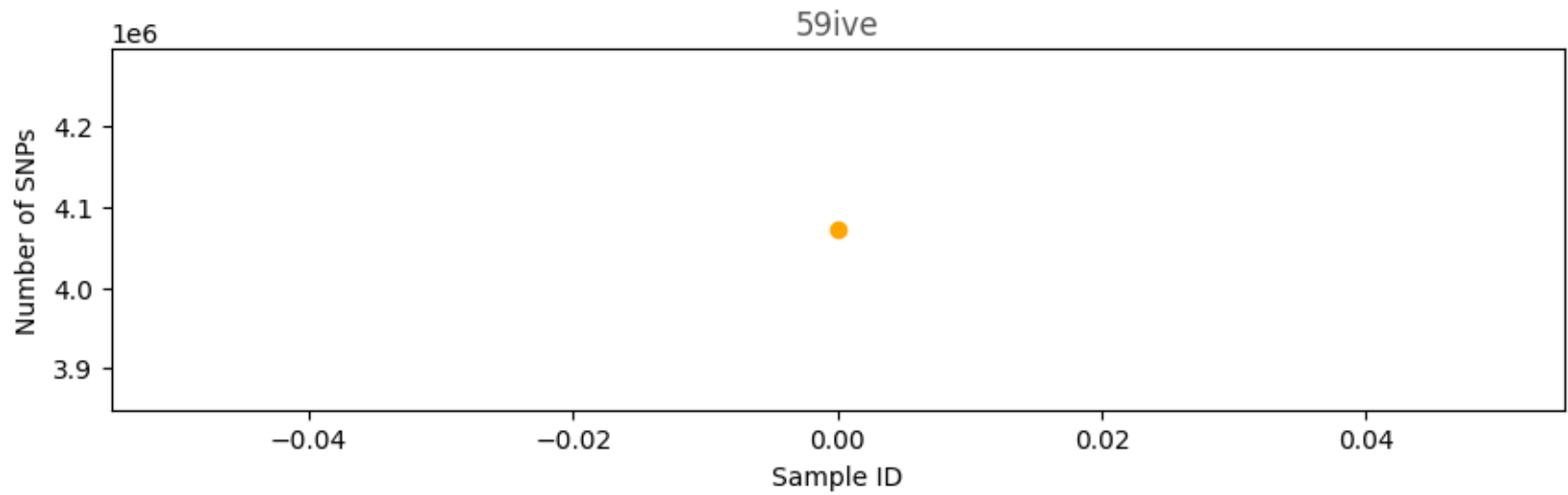
Hets vs non-ref Homs by sample
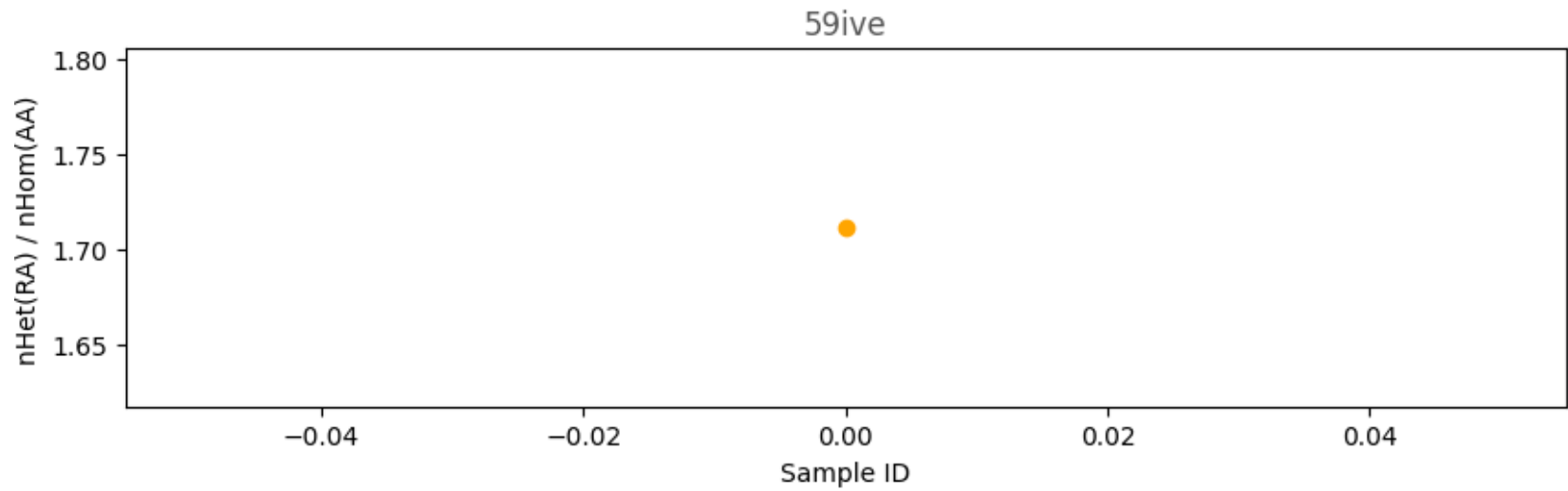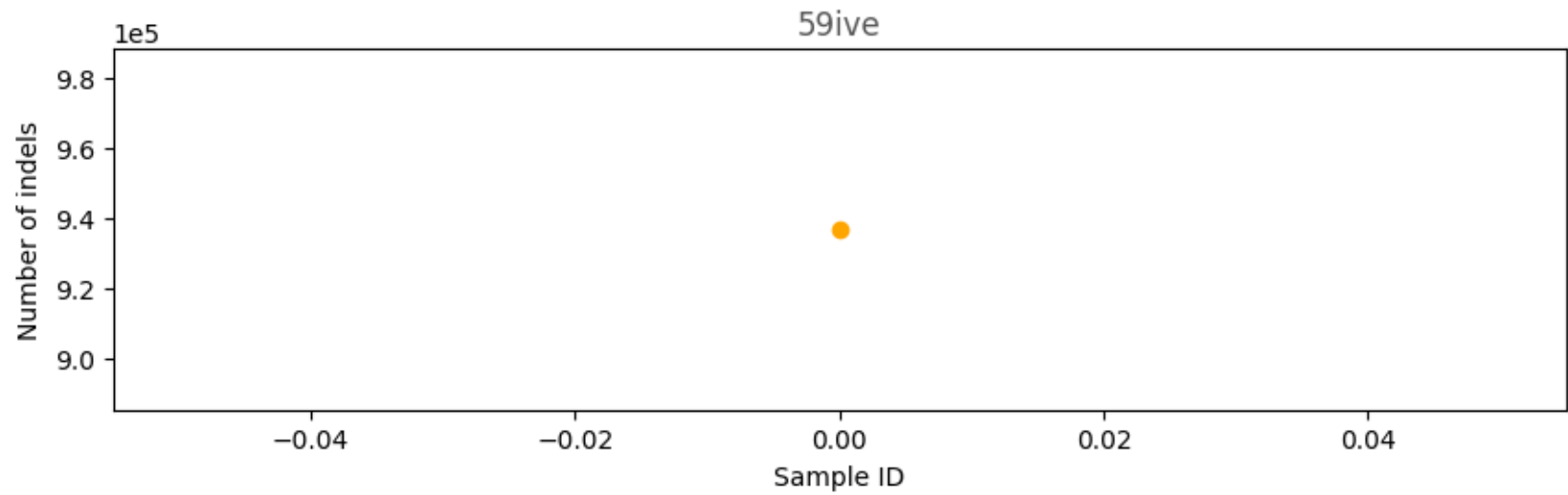
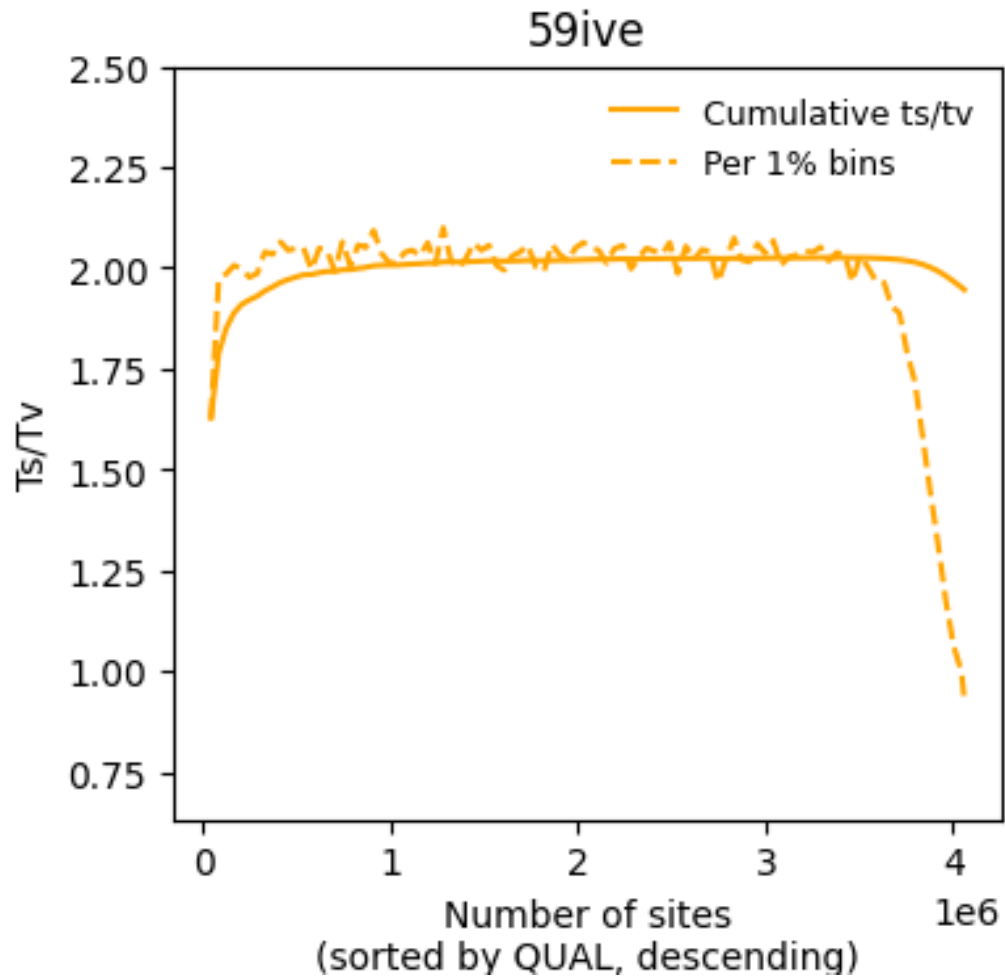# Singletons by sample (hets and homs)

Average depth by sample
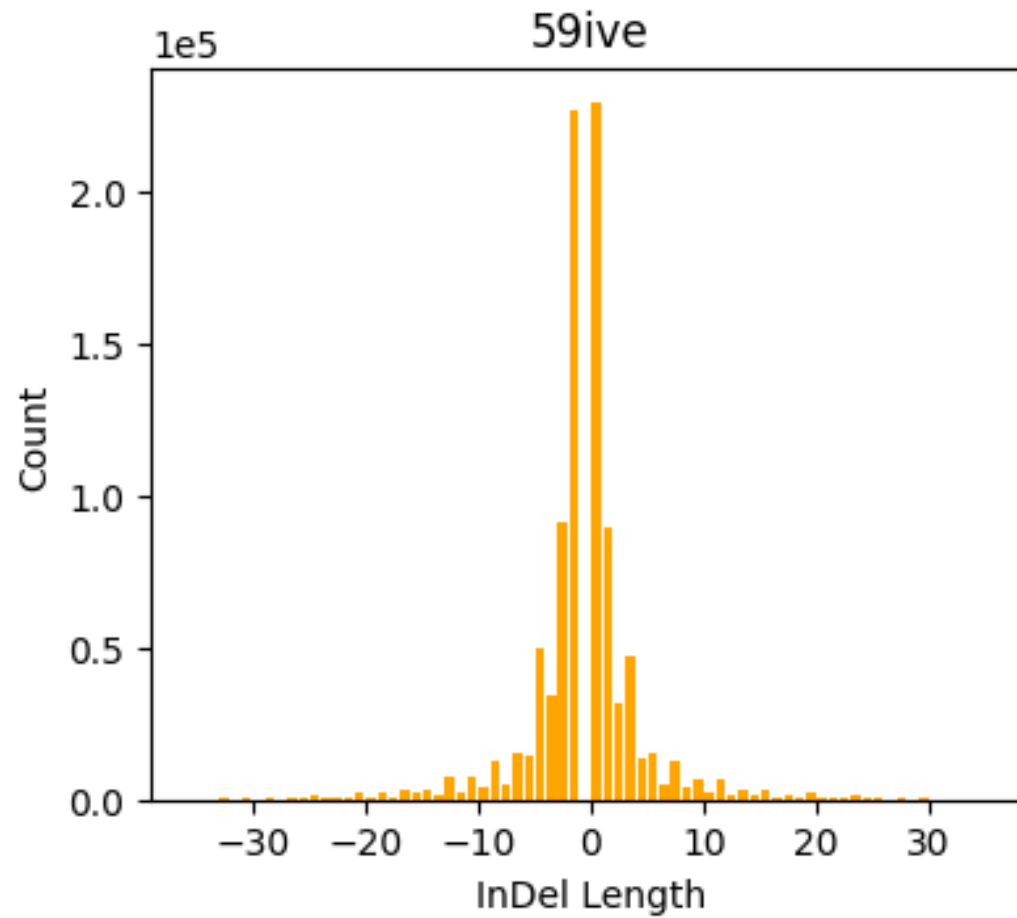
Number of SNPs by sample

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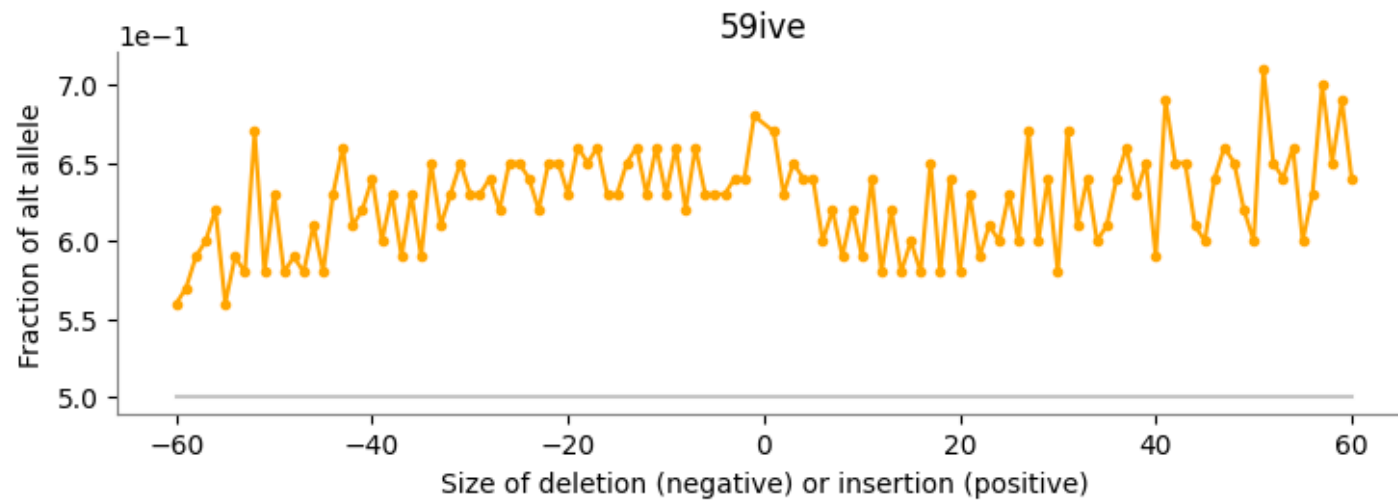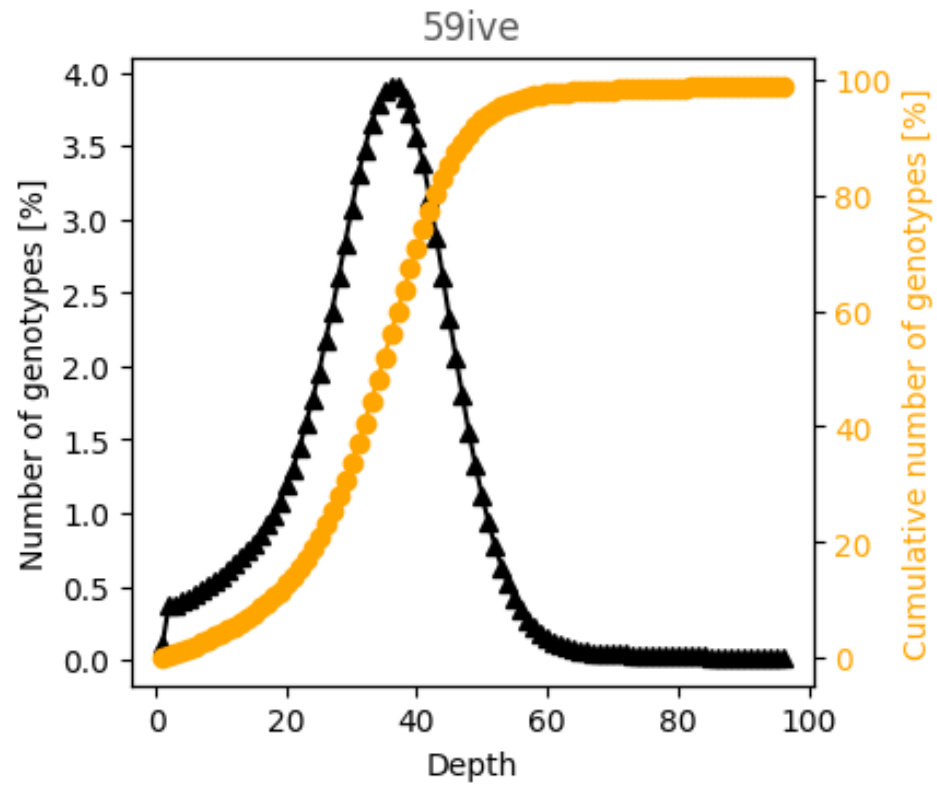
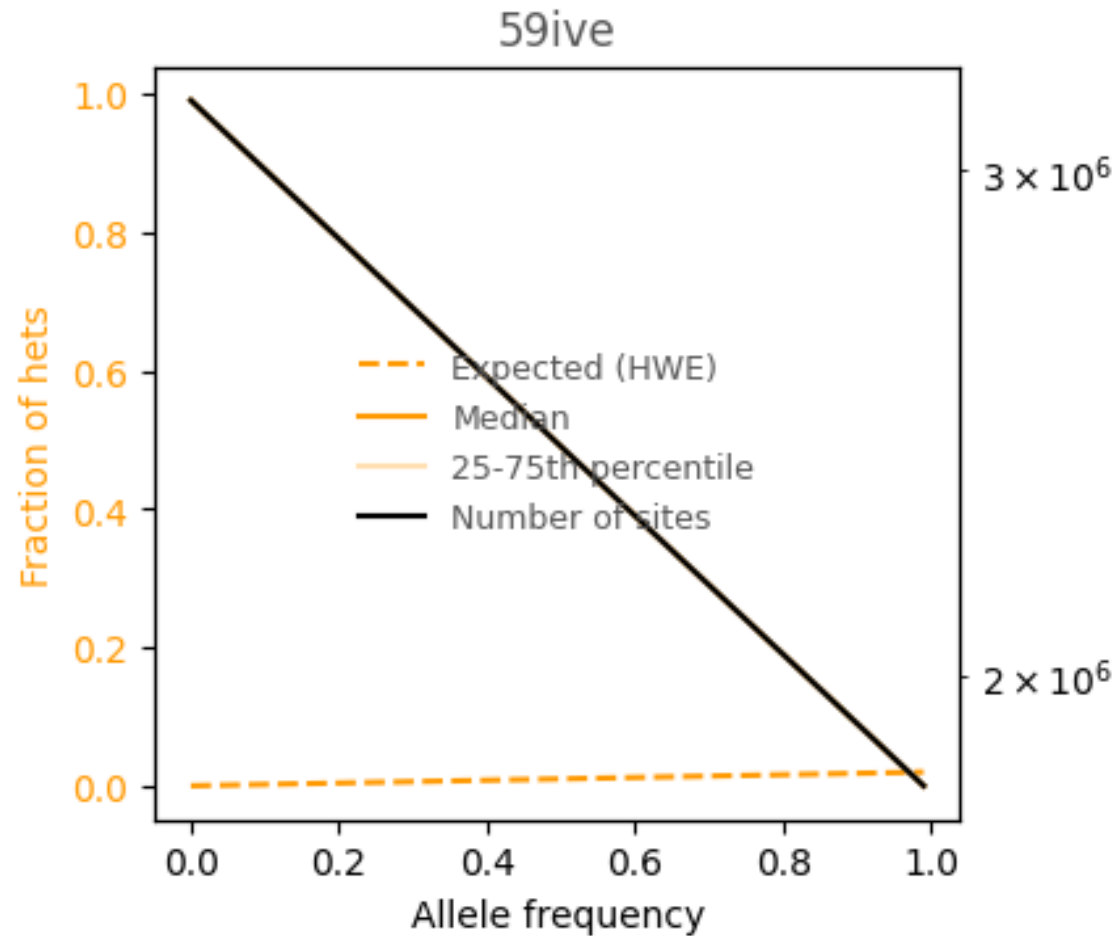# Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types