Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
59ere	4,149,167	1.93	1.93	971,349	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
59ere	63.6%	1.90	69.2%	107,331	2,335

- 59ere .. `/ngc/projects2/gm/data/archive/2022/variants/snv/59ereanof-103870987175-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220518_A01411_AHJ5W2DSX3-RHGM_LABKA_WGS-`
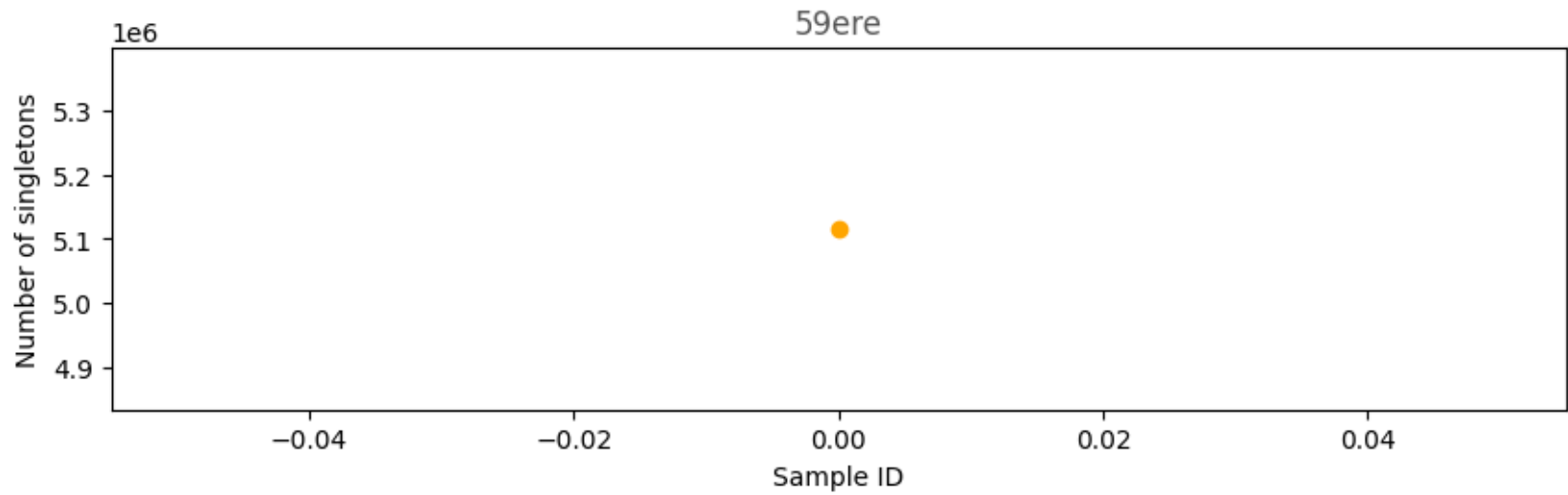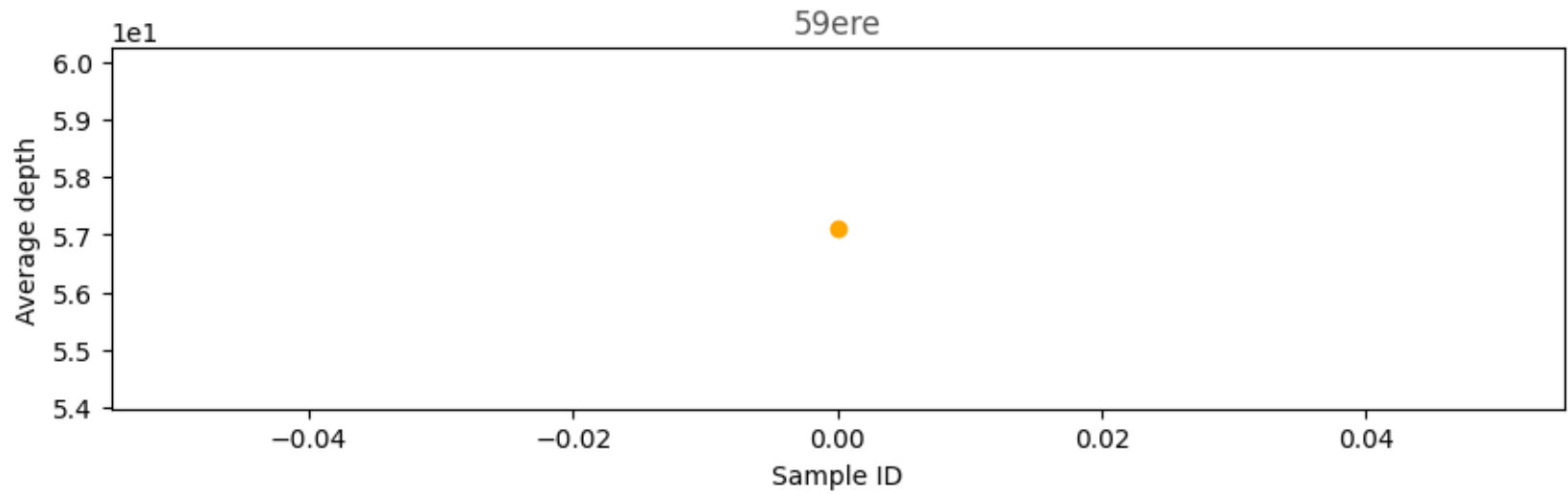 `WGS04288_snv_germline_raw.haplotype_caller.vcf.gz`
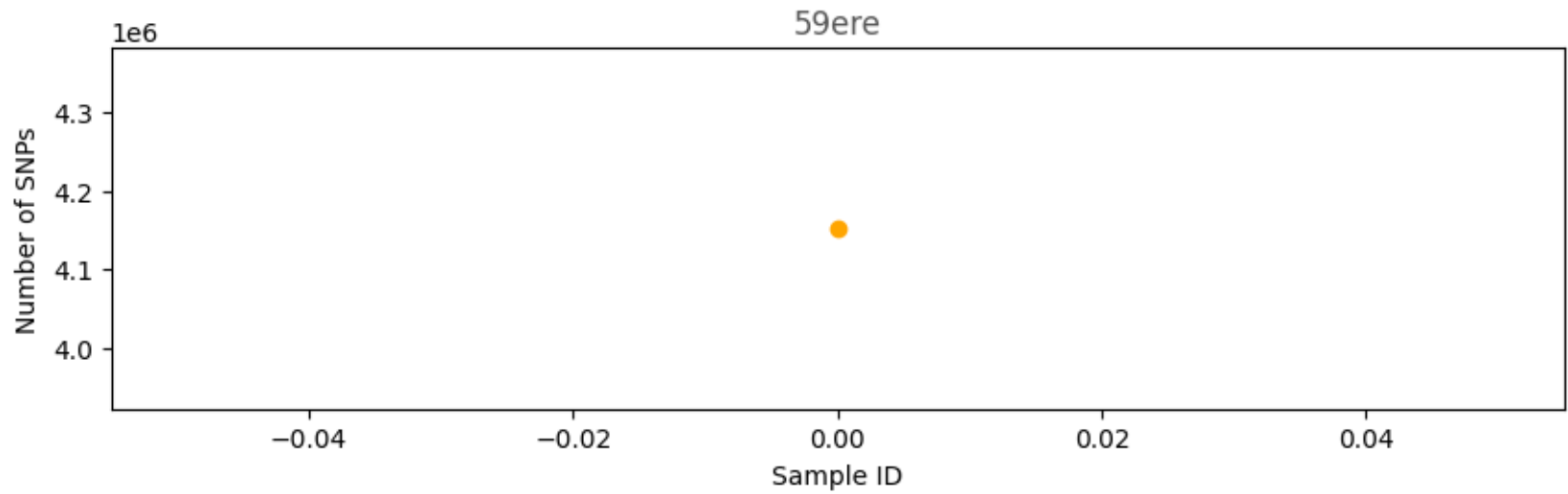
Ts/Tv by sample
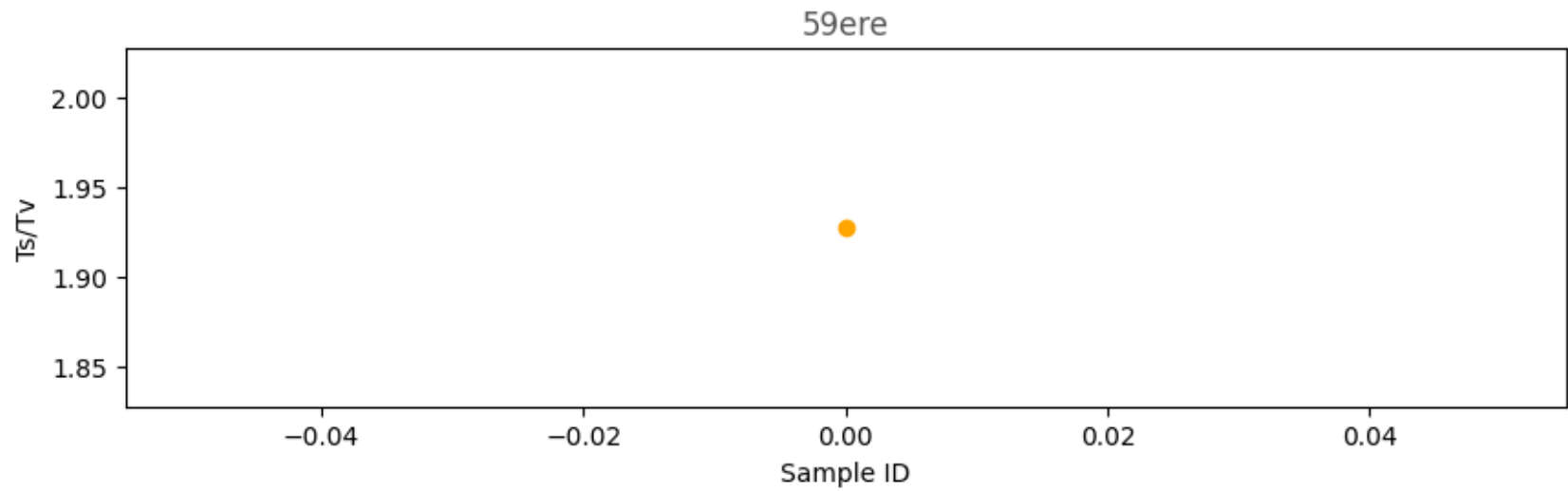
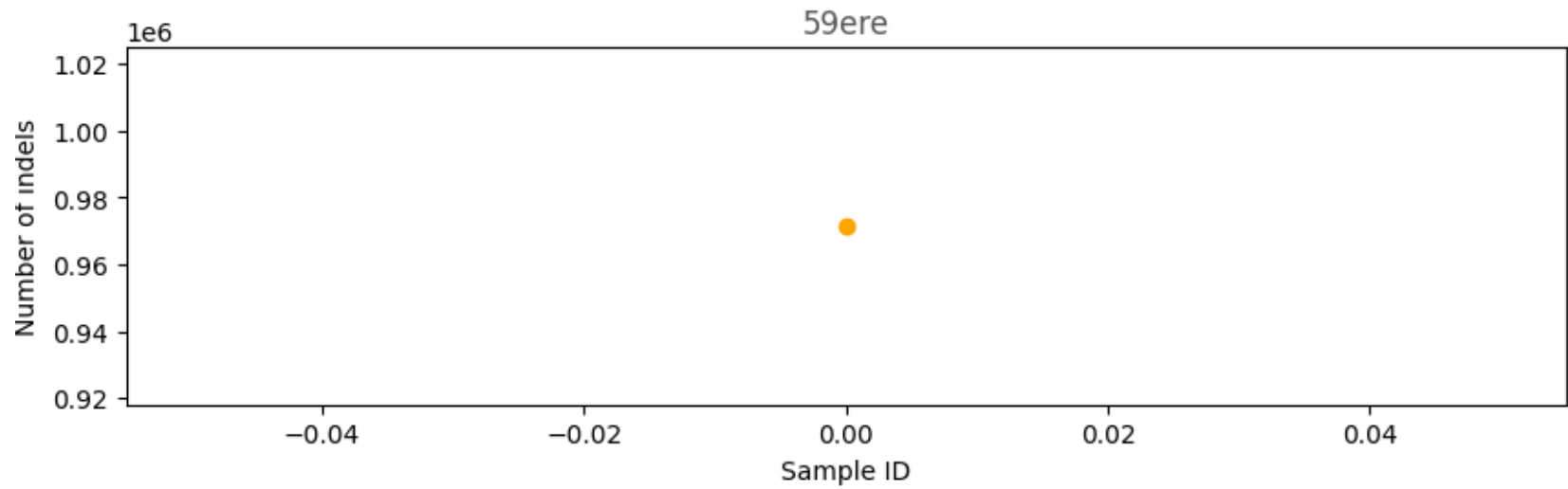# Hets vs non-ref Homs by sample
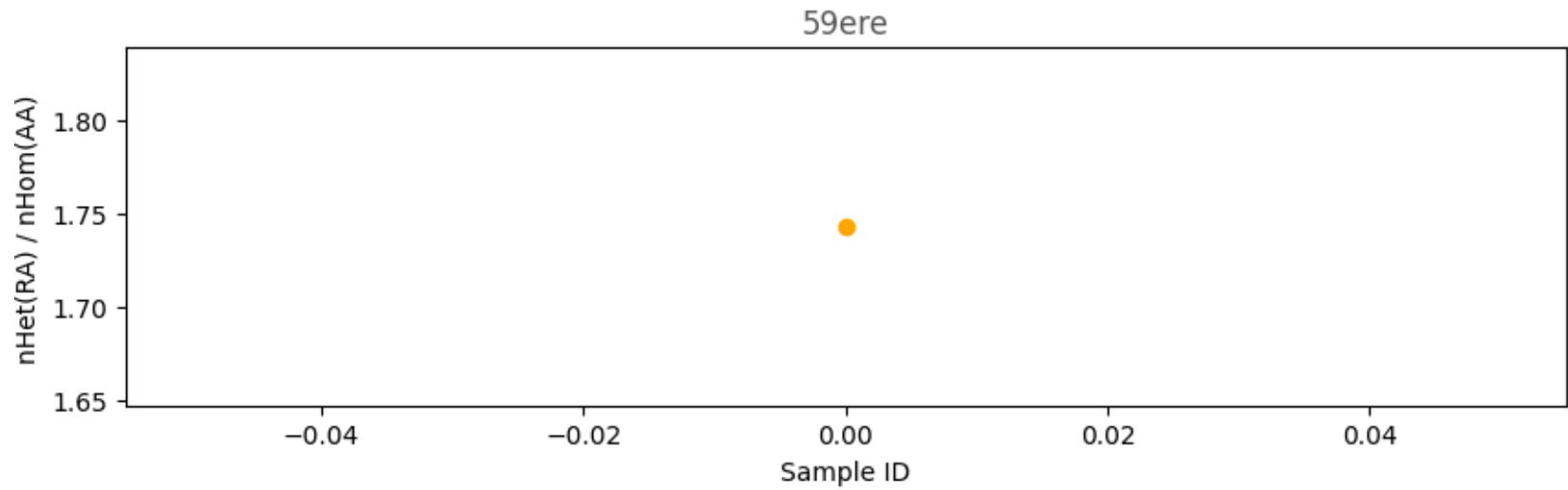
Singletons by sample (hets and homs)
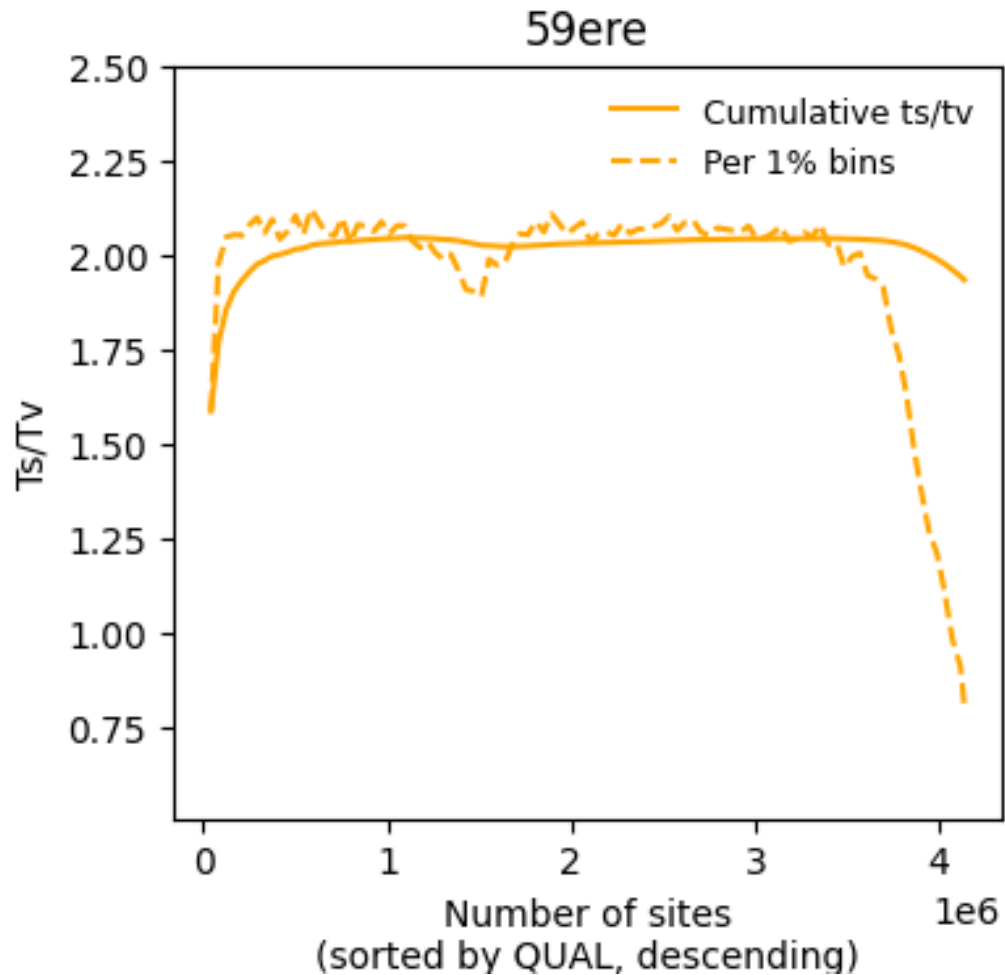
Average depth by sample

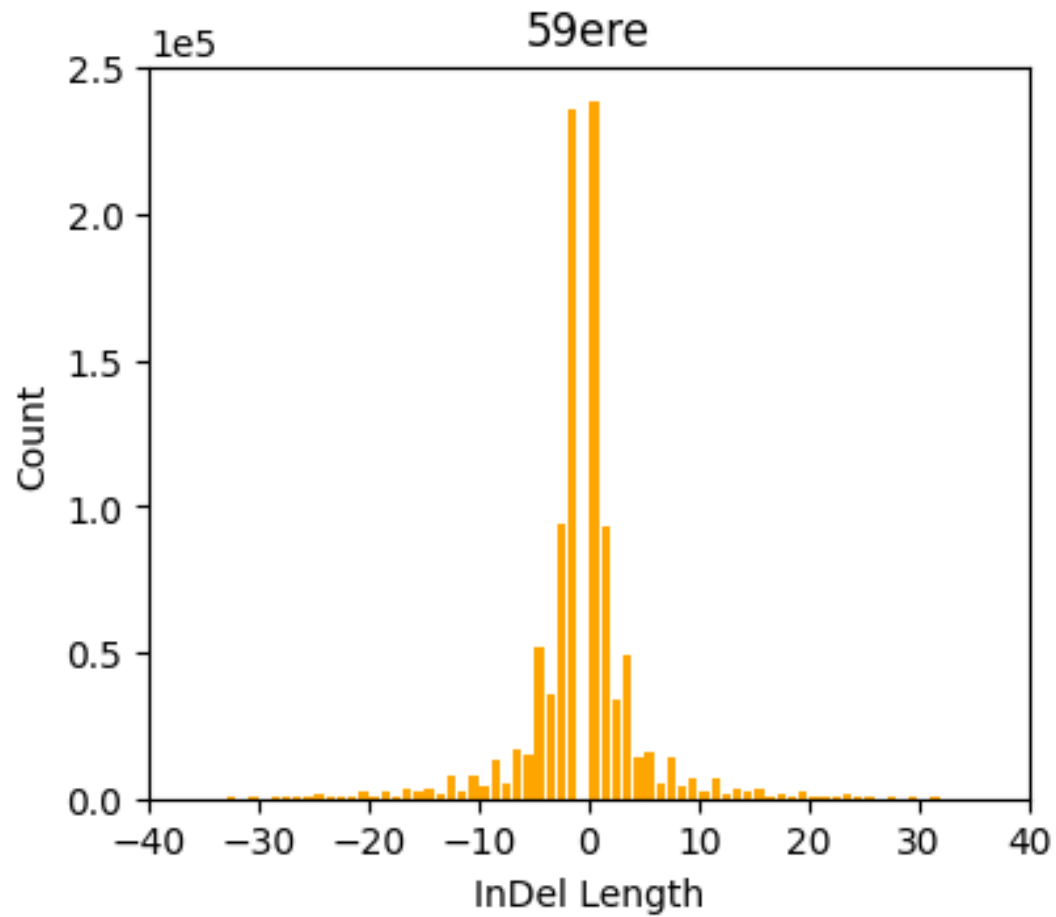# Number of SNPs by sample

59ere

Number of indels by sample
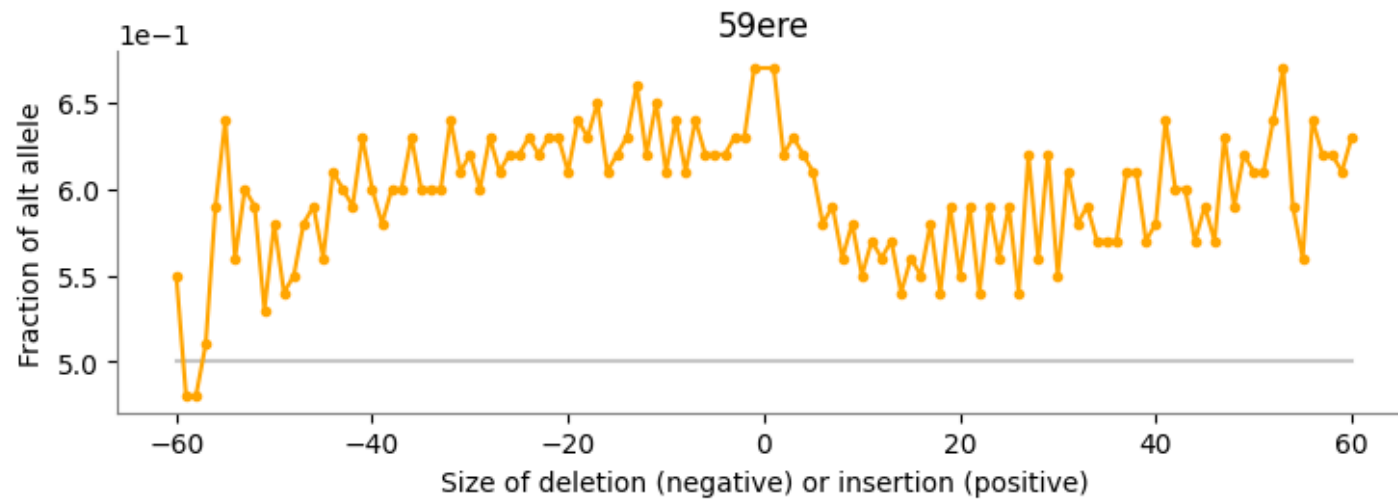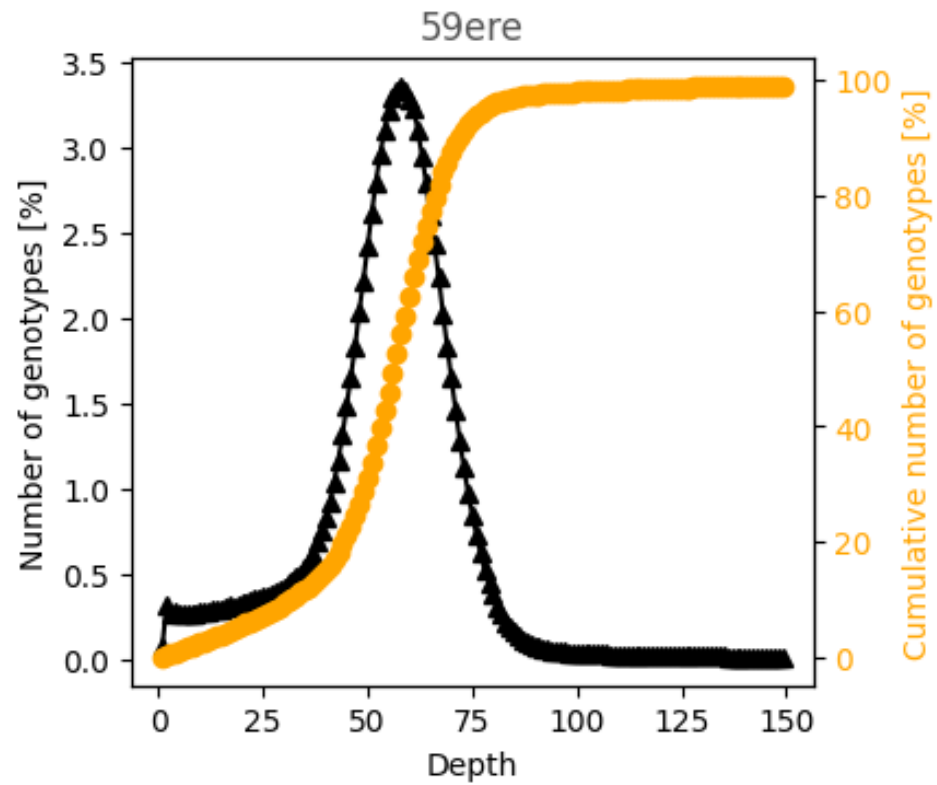
Ts/Tv stratified by QUAL

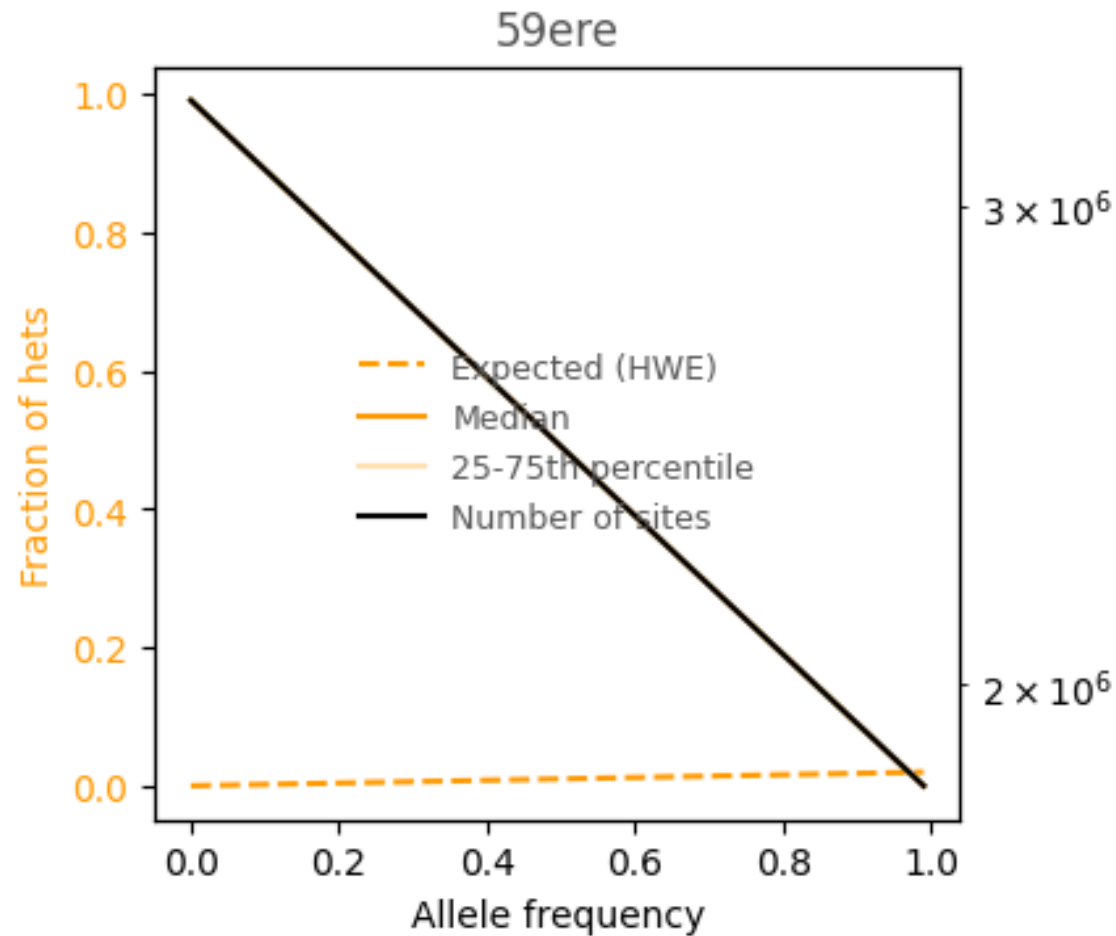

59ere

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types