Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
47ste	4,028,153	1.94	1.95	920,494	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
47ste	62.0%	1.91	65.5%	86,910	1,857

- 47ste .. `/ngc/projects2/gm/data/archive/2022/variants/snv/47stendim-103843971107-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220302_A01411_AH5T5WDSX3-RHGM_LABKA_WGS-WGS03933_snv_germline_raw.haplotype_caller.vcf.gz`

Ts/Tv by sample
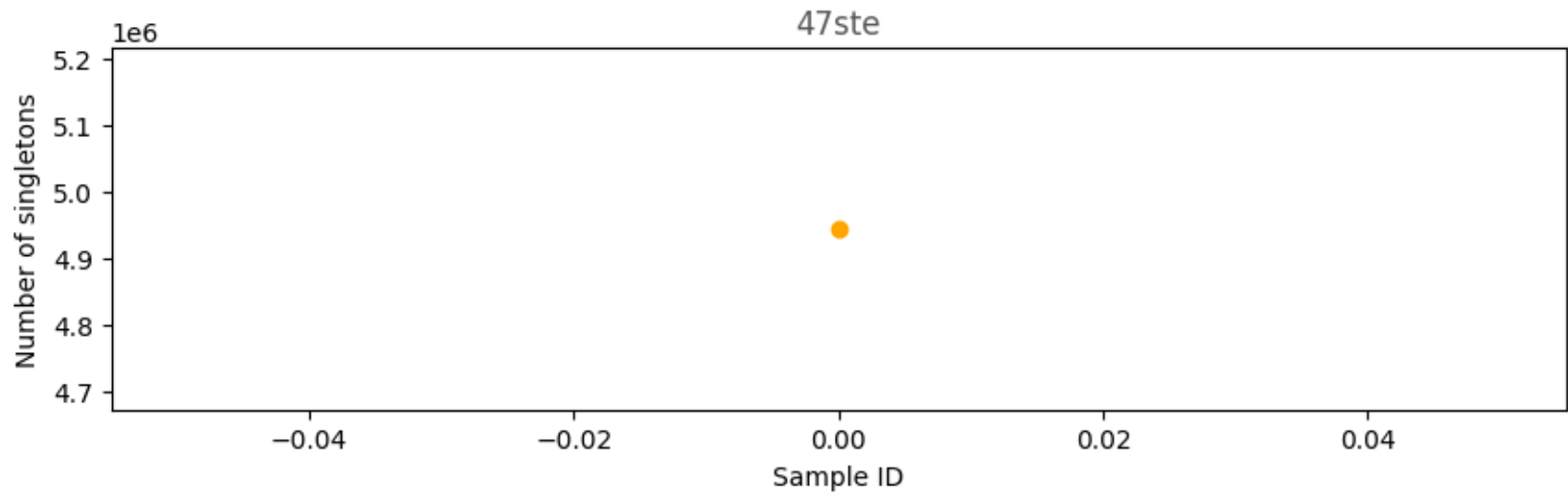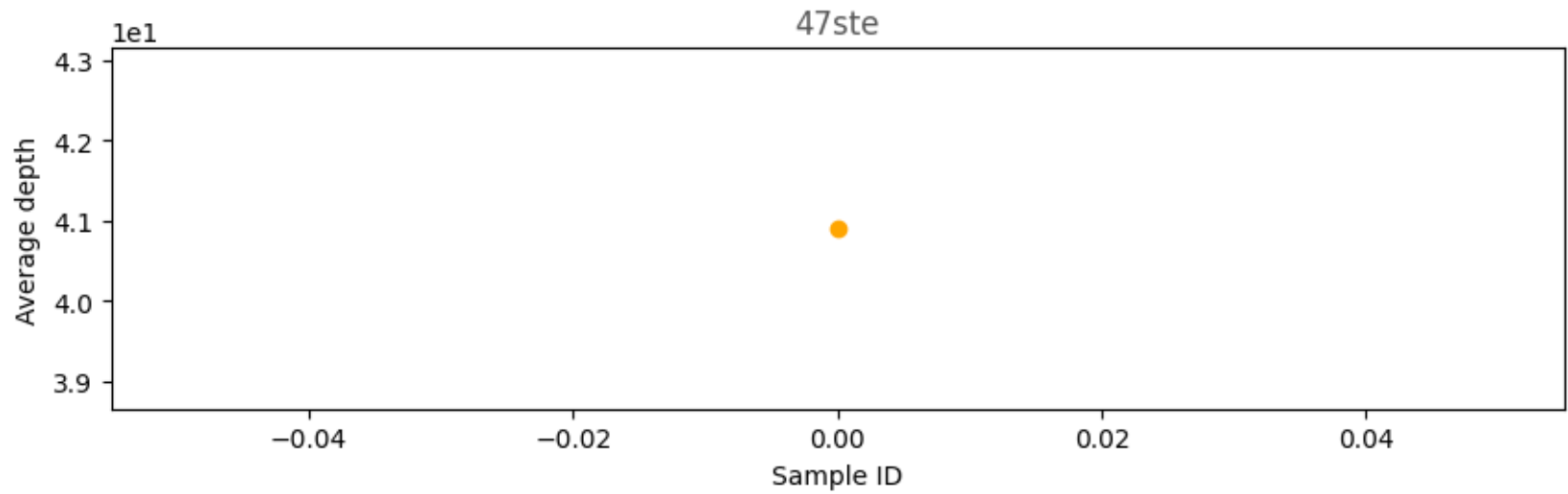

47ste
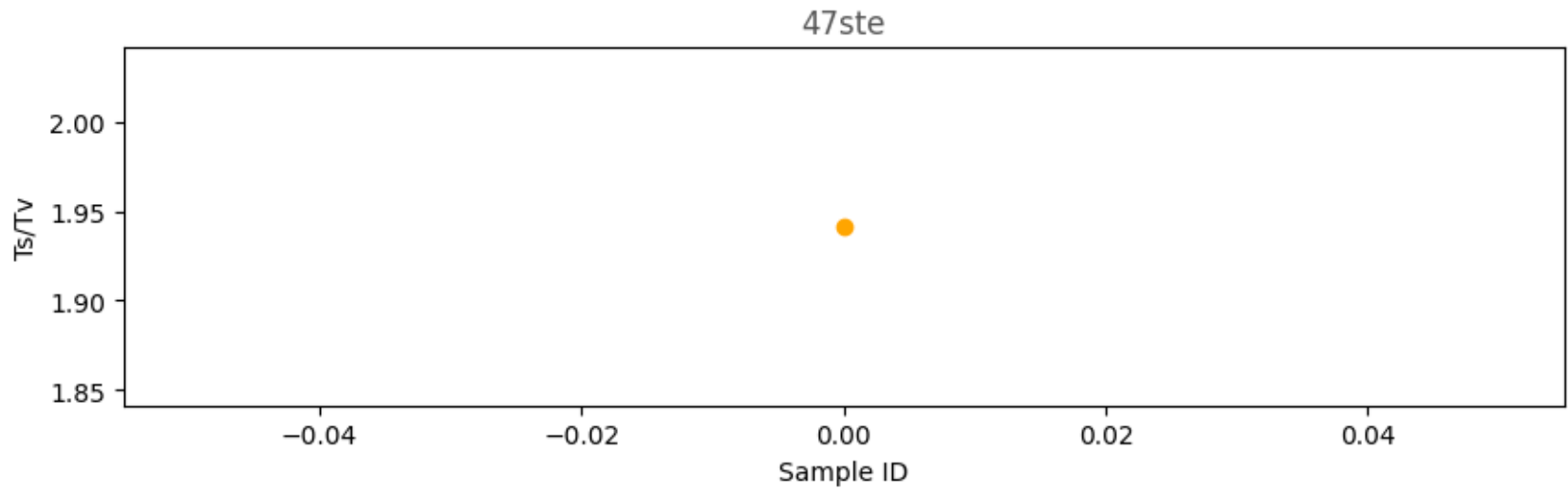
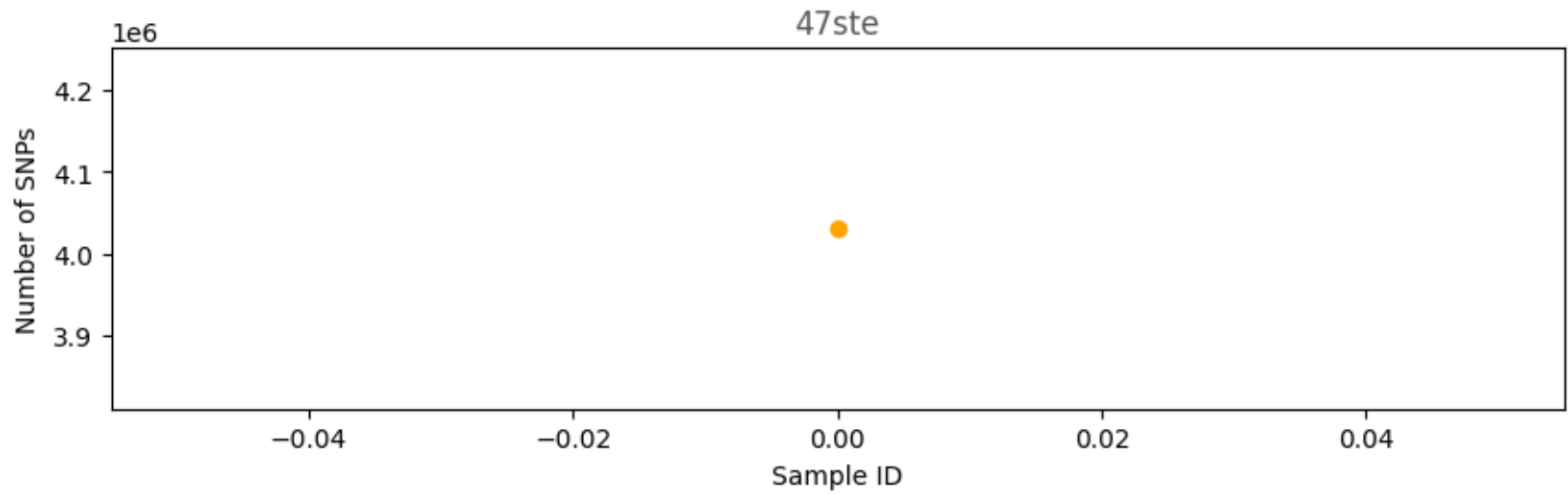# Hets vs non-ref Homs by sample
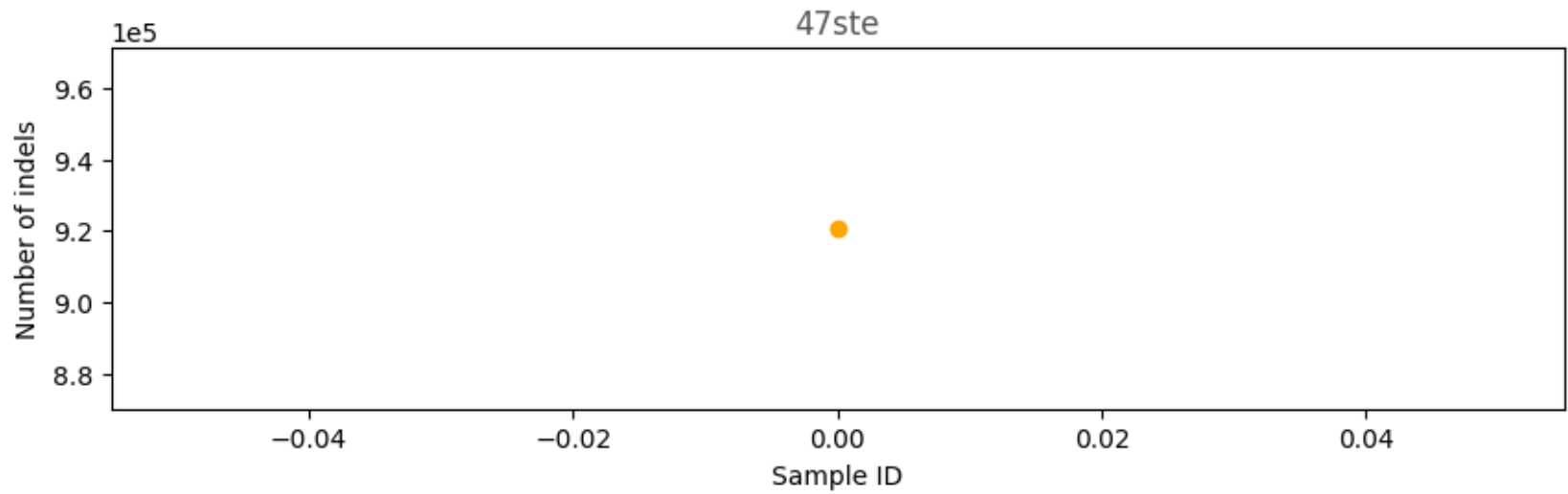

47ste
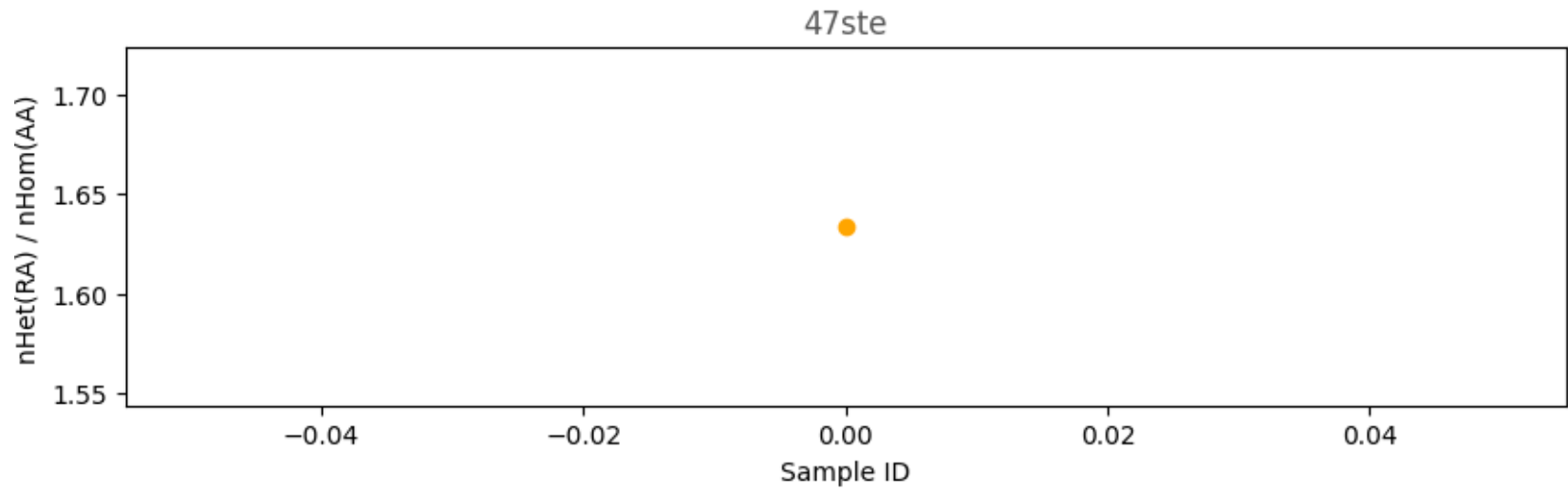
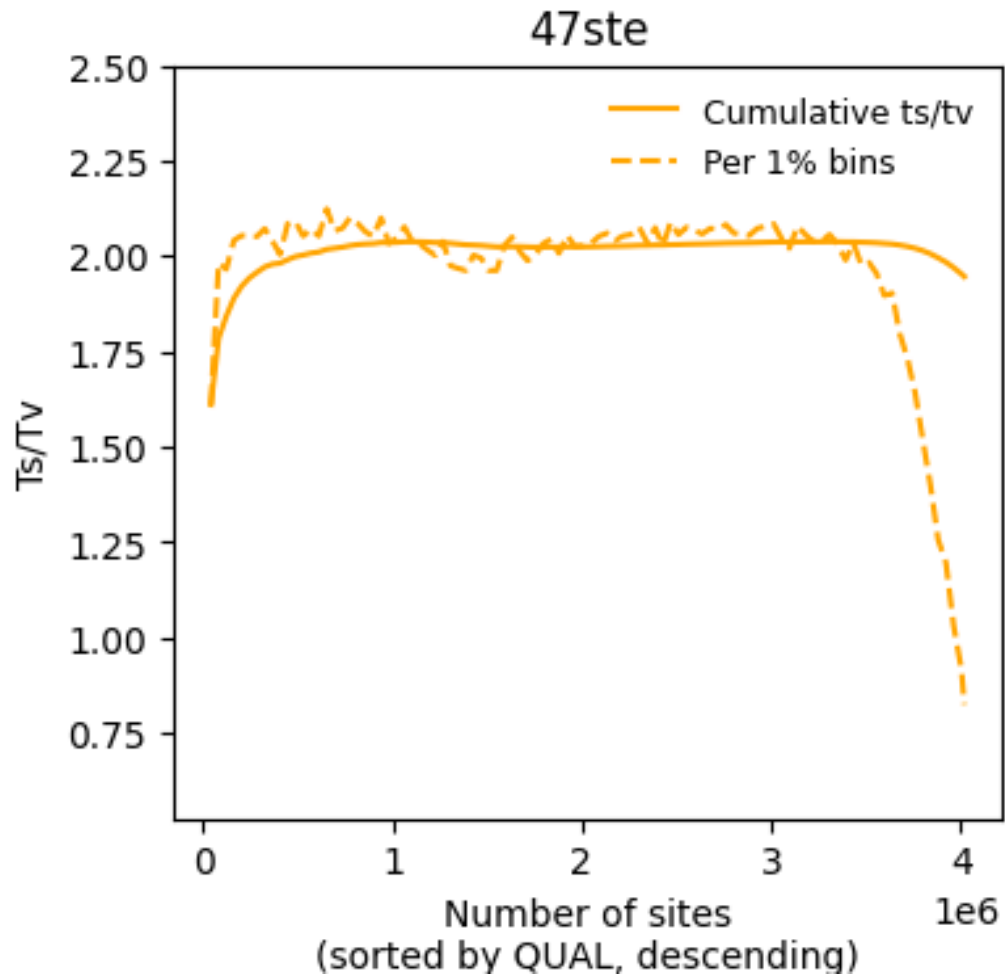# Singletons by sample (hets and homs)

Average depth by sample
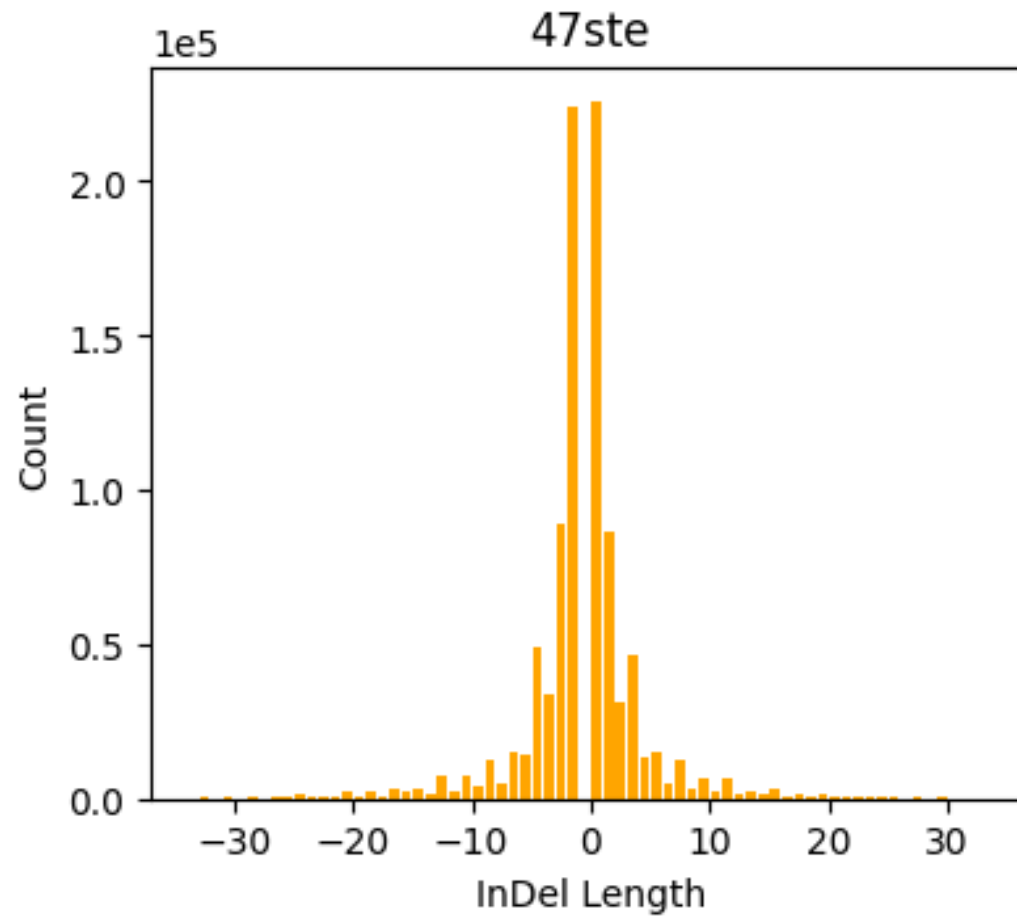
Number of SNPs by sample

Number of indels by sample
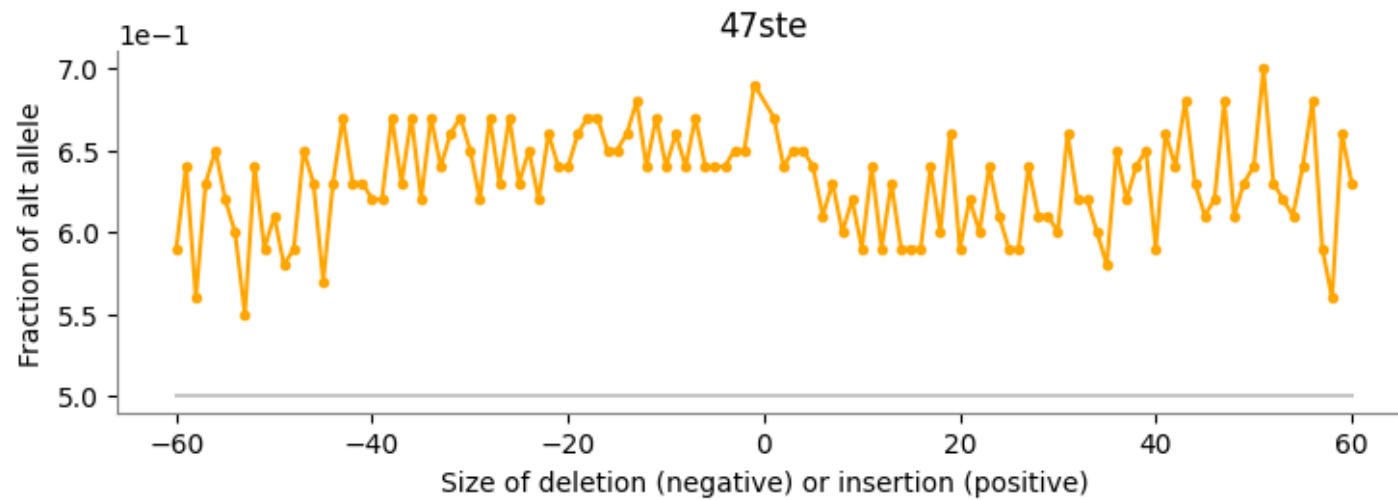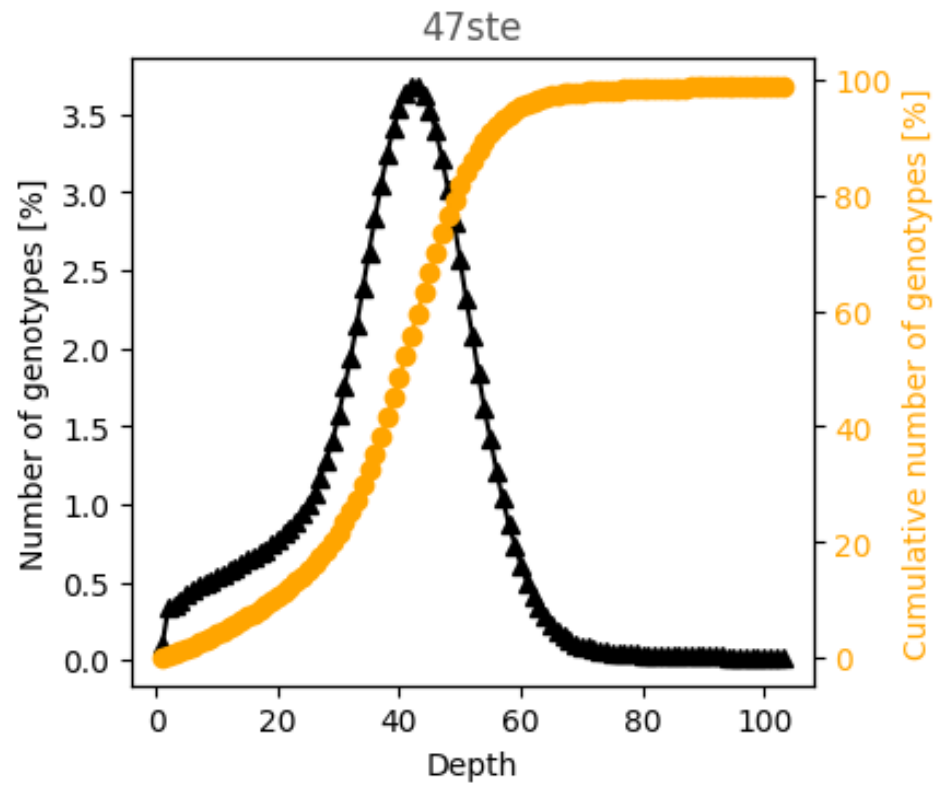
Ts/Tv stratified by QUAL

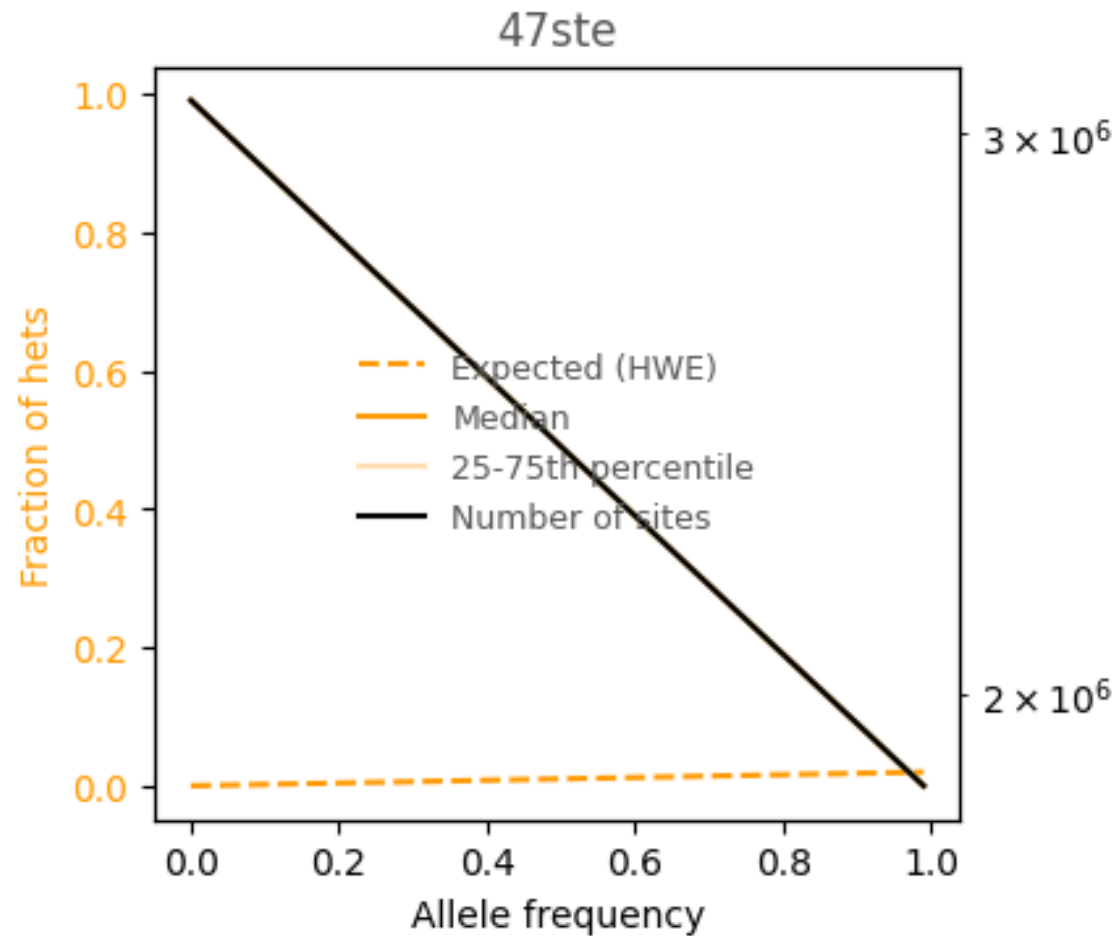47ste

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Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types