Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
43lie	4,086,710	1.93	1.93	952,746	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
43lie	62.9%	1.90	68.2%	103,120	2,183

- 43lie .. `/ngc/projects2/gm/data/archive/2022/variants/snv/43liefolf-4108499337-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02208-221109_A01176_AH3NY7DSX5-EXT_LABKA_NGCWGS-NGCWGS06090_snv_germline_raw.haplotype_caller.vcf.gz`
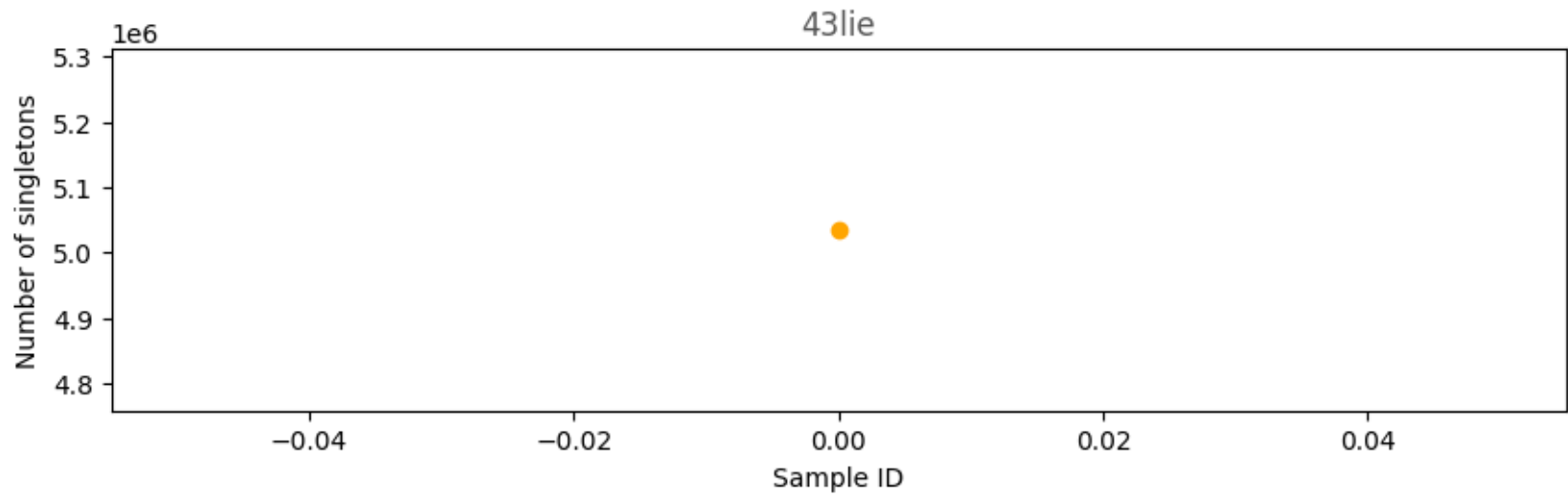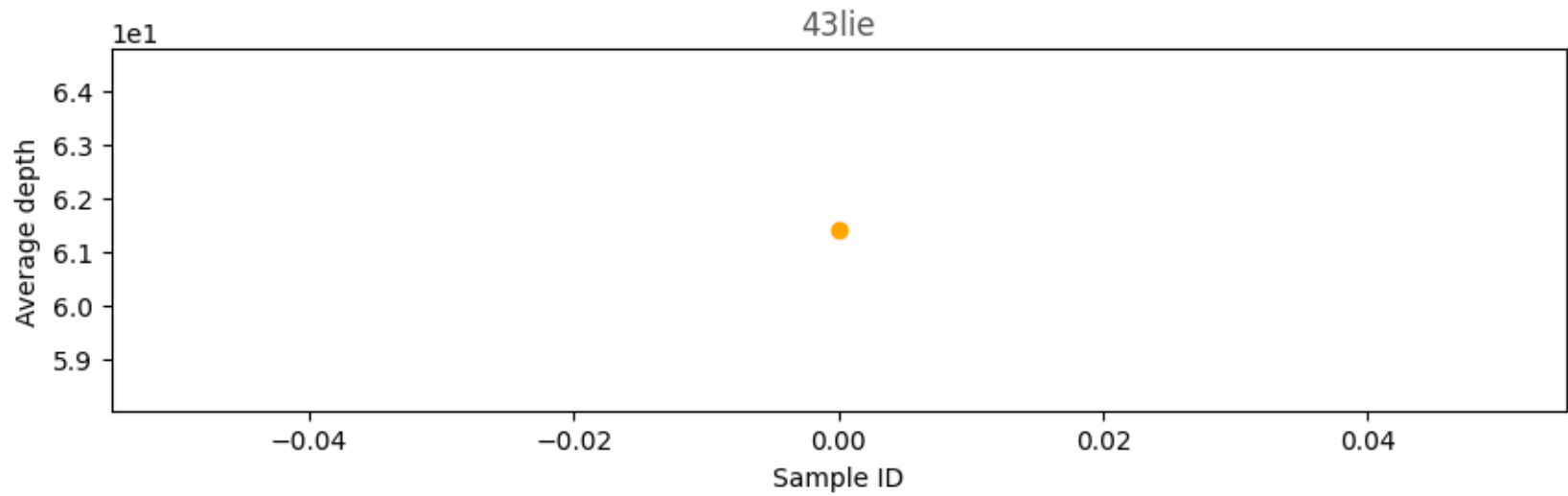
Ts/Tv by sample

43lie
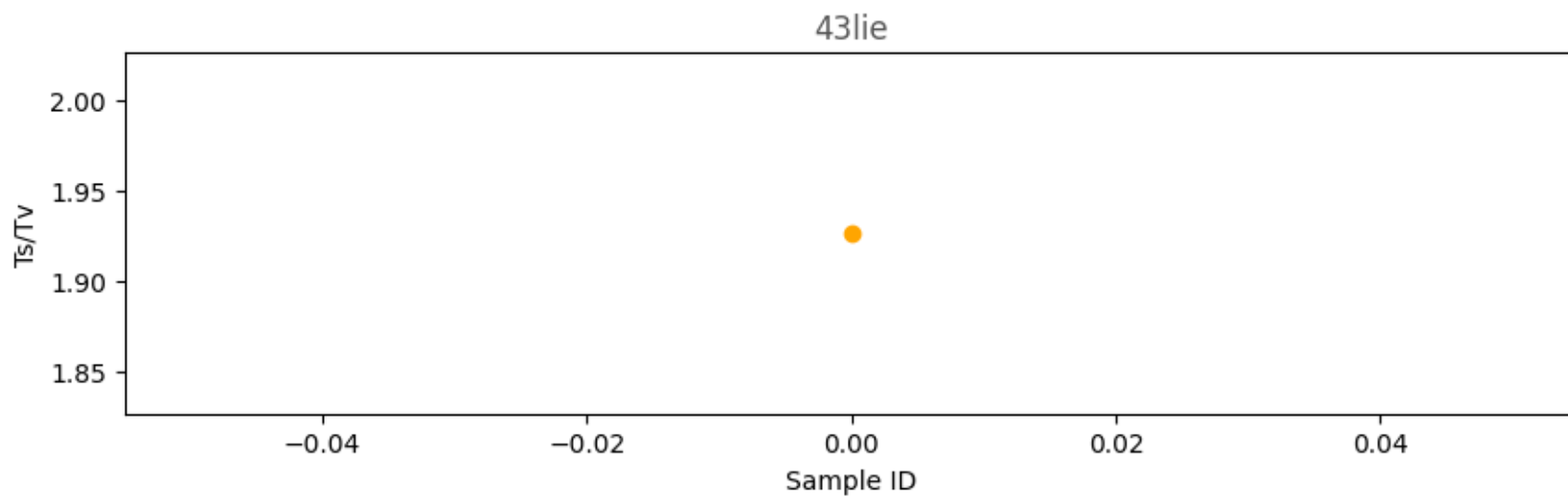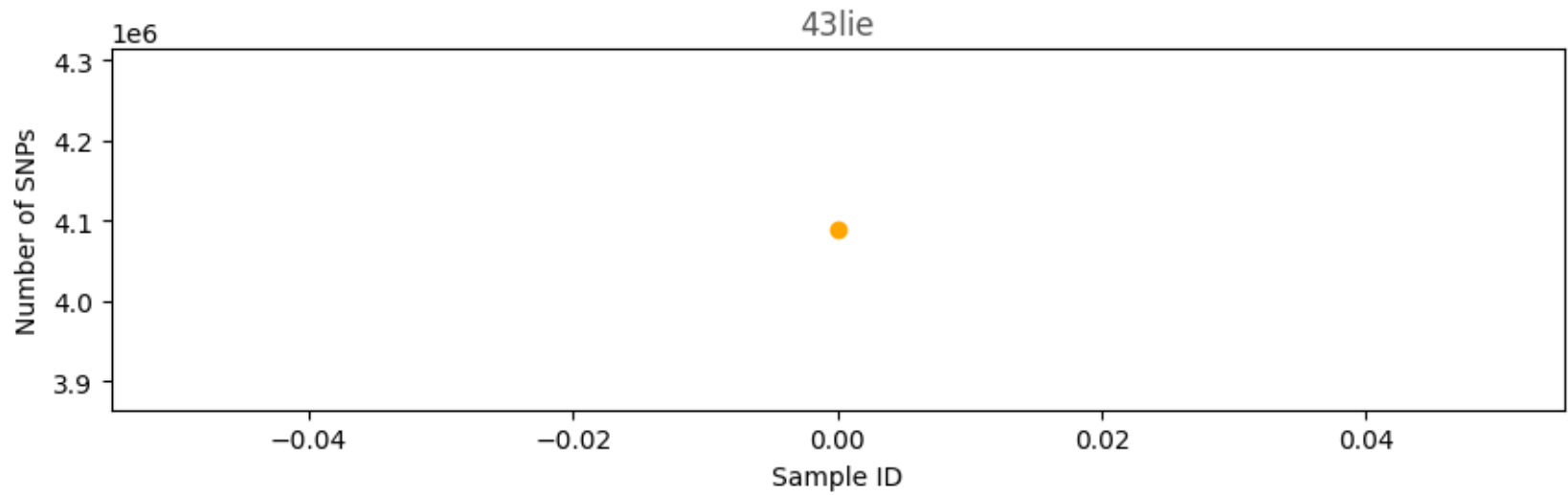
Hets vs non-ref Homs by sample
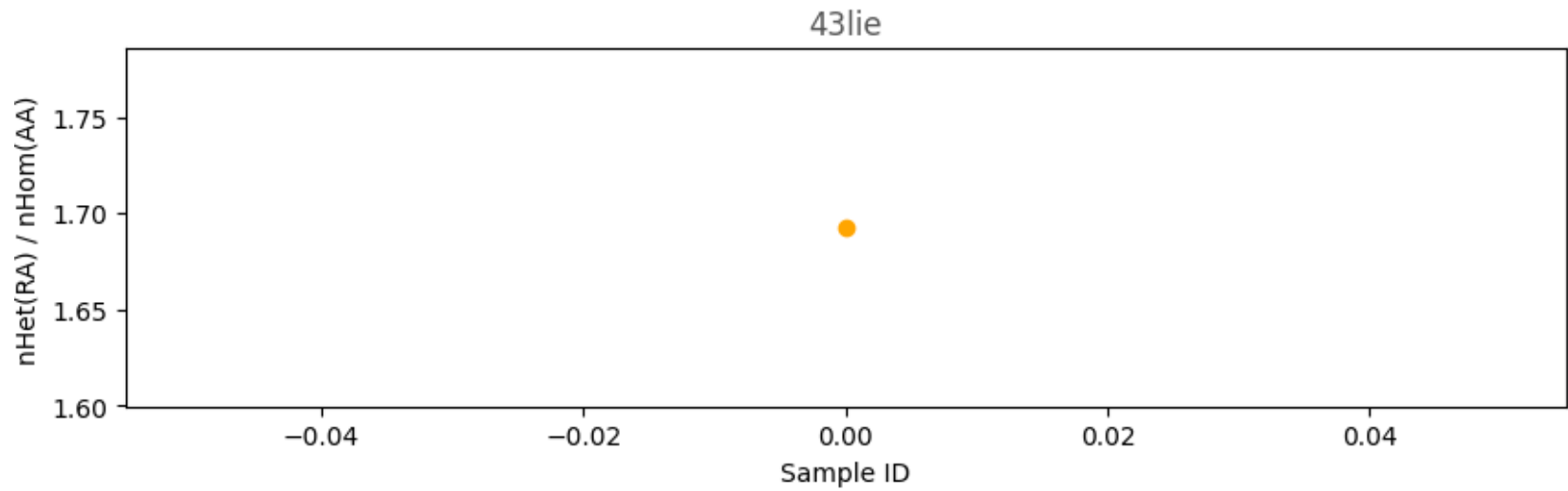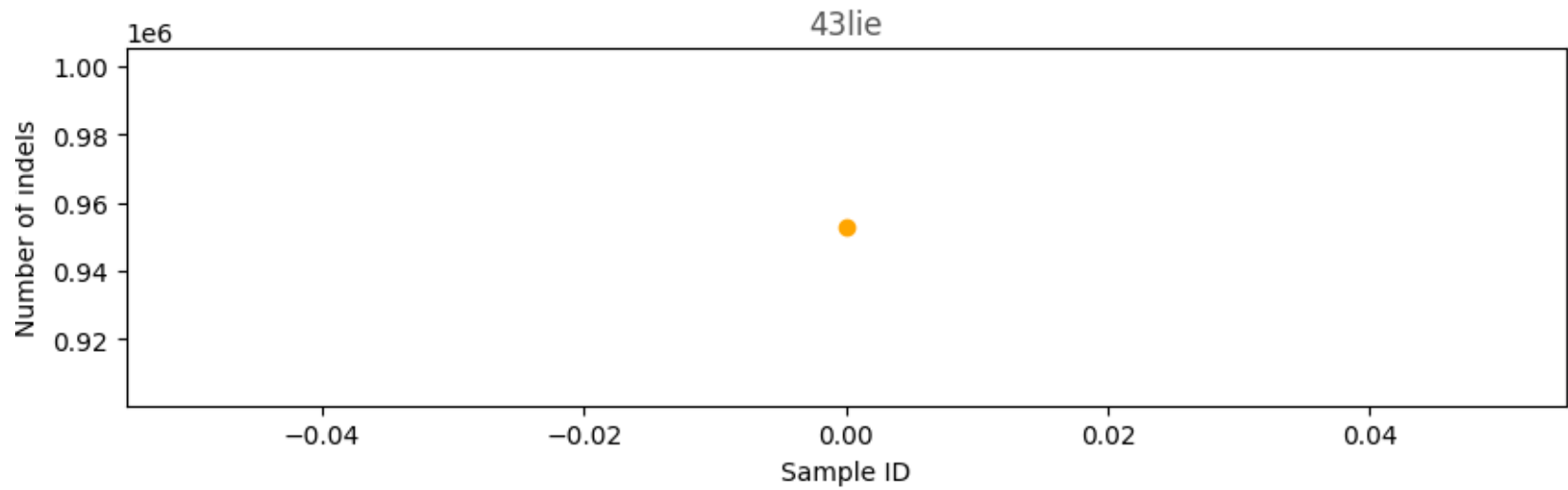
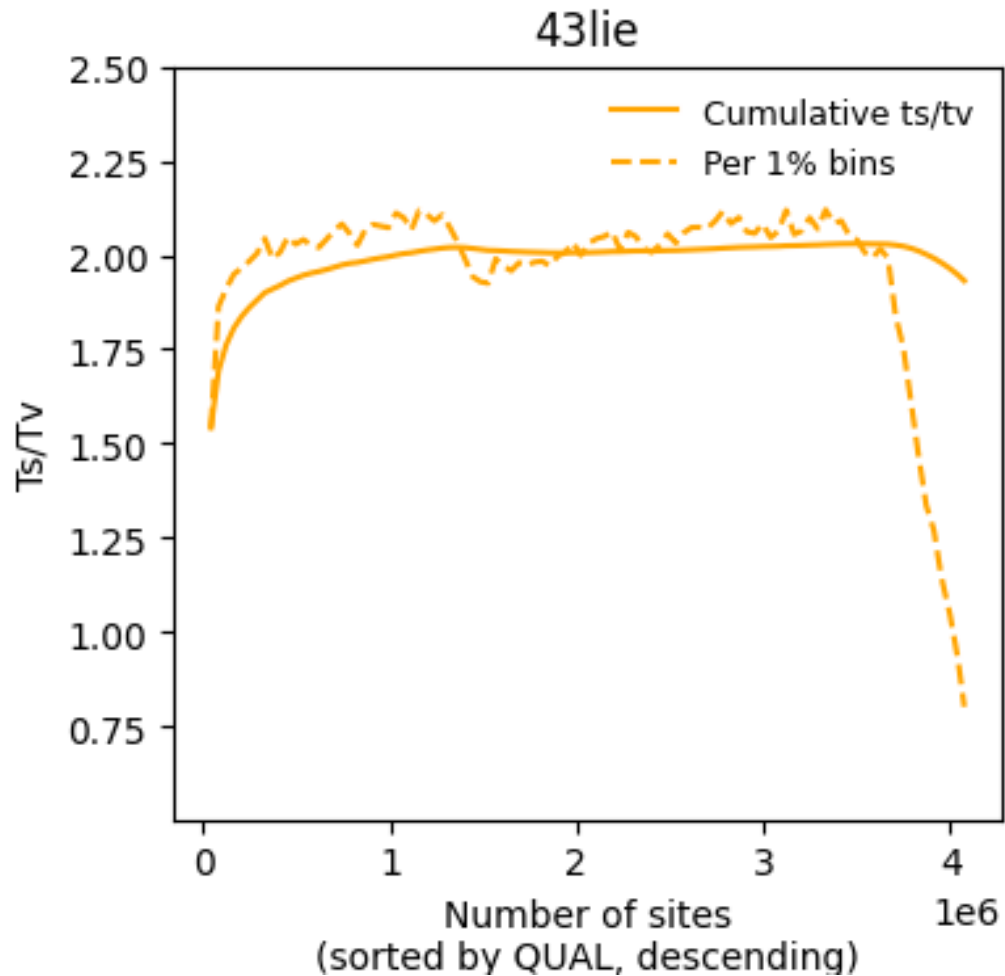# Singletons by sample (hets and homs)

Average depth by sample

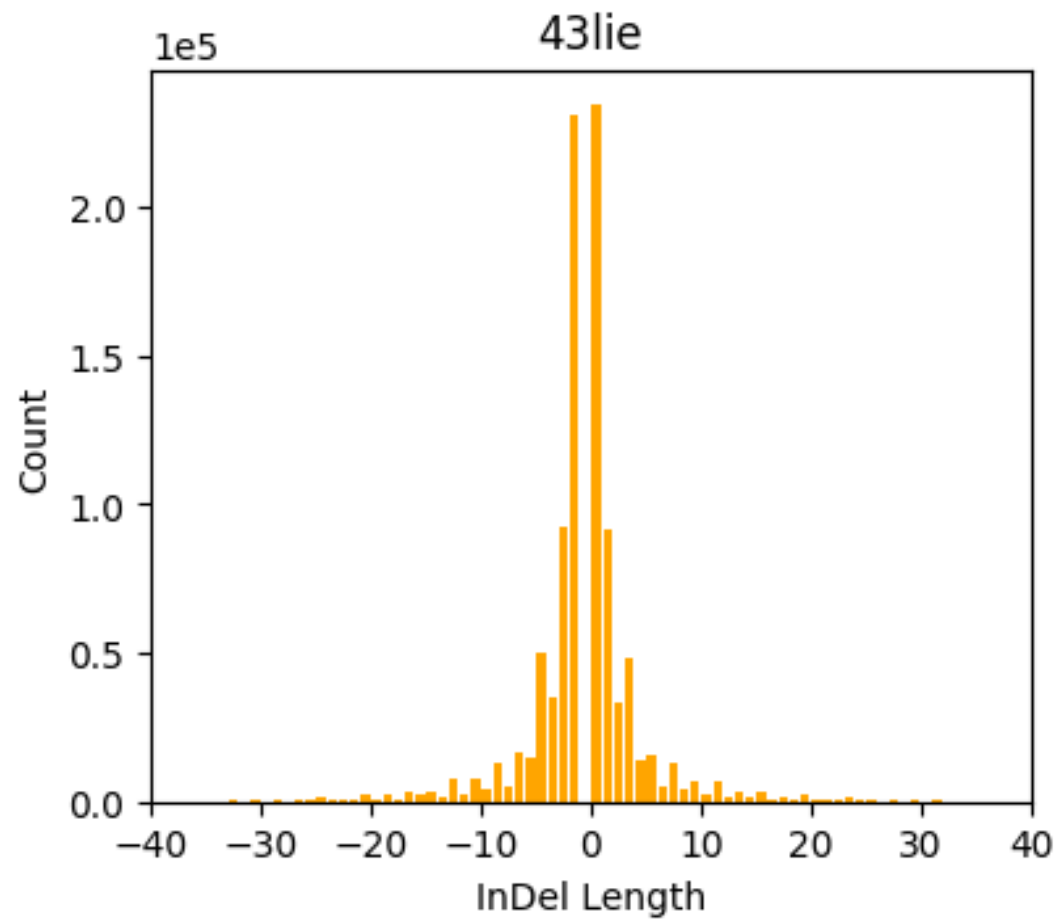# Number of SNPs by sample

Number of indels by sample
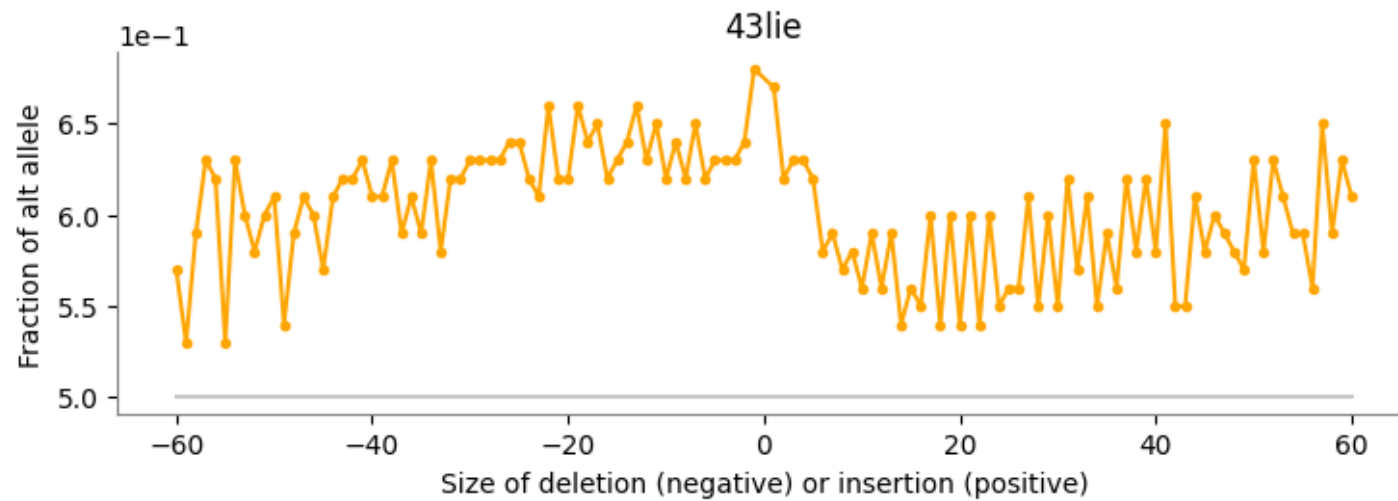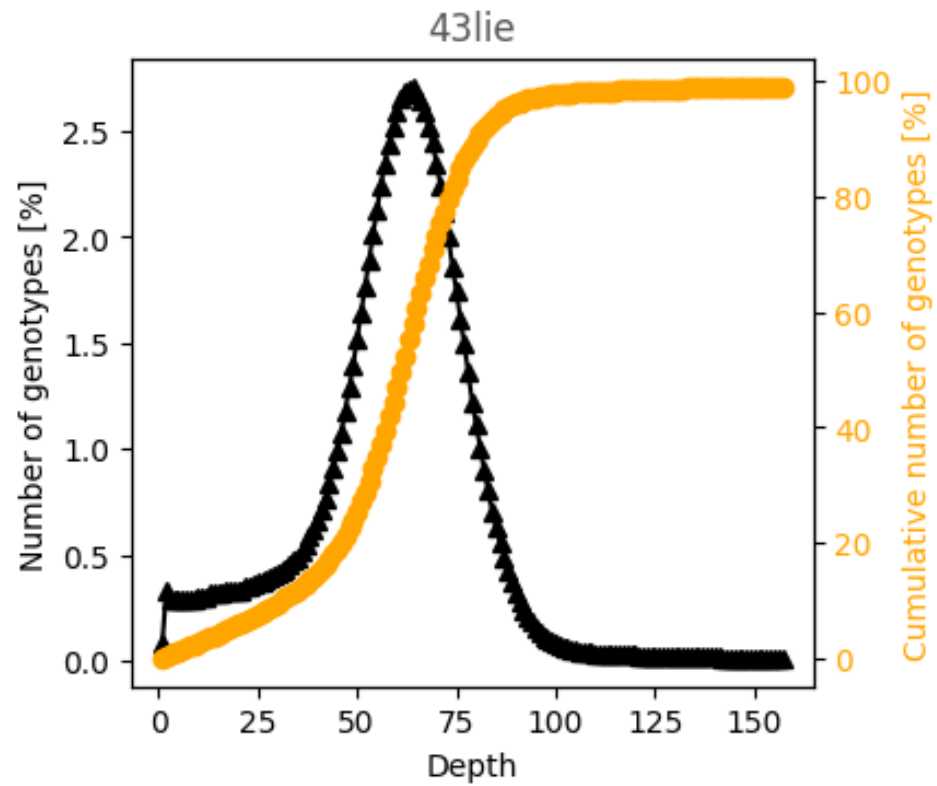
Ts/Tv stratified by QUAL
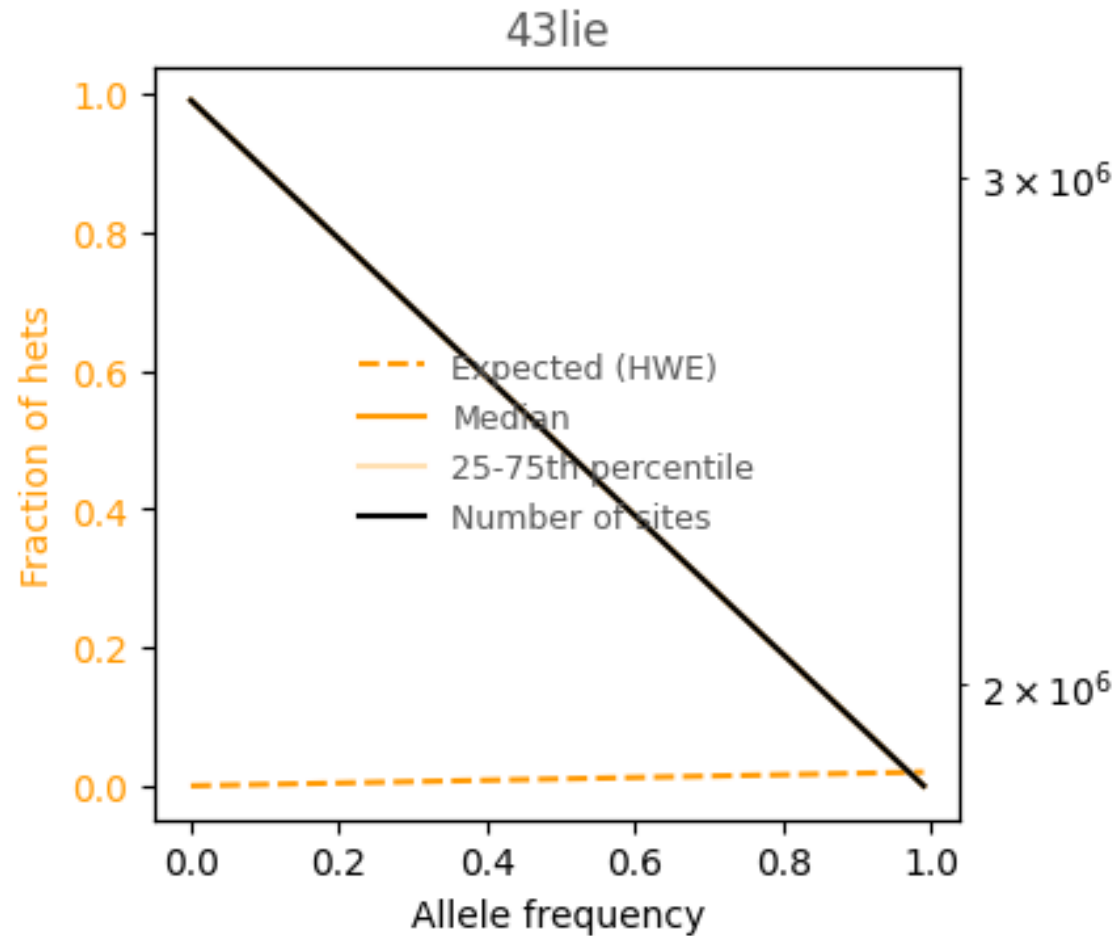

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Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types