Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
91uli	4,076,767	1.94	1.94	947,682	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
91uli	63.3%	1.91	68.1%	99,515	1,958

- 91uli .. `/ngc/projects2/gm/data/archive/2022/variants/snv/91ulirlyf-103856195917-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220407_A01411_AHGNYMDSX3-RHGM_LABKA_WGSA KUT-WGSAKUT04067_22RKG007984x01_snv_germline_raw.haplotype_caller.vcf.gz`
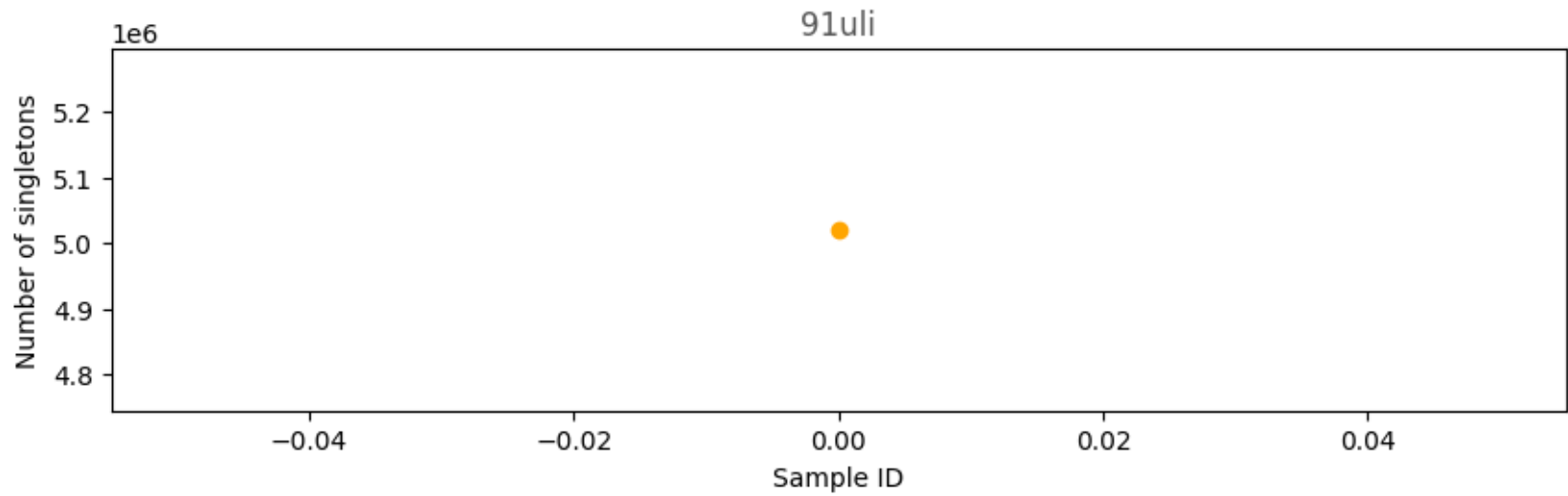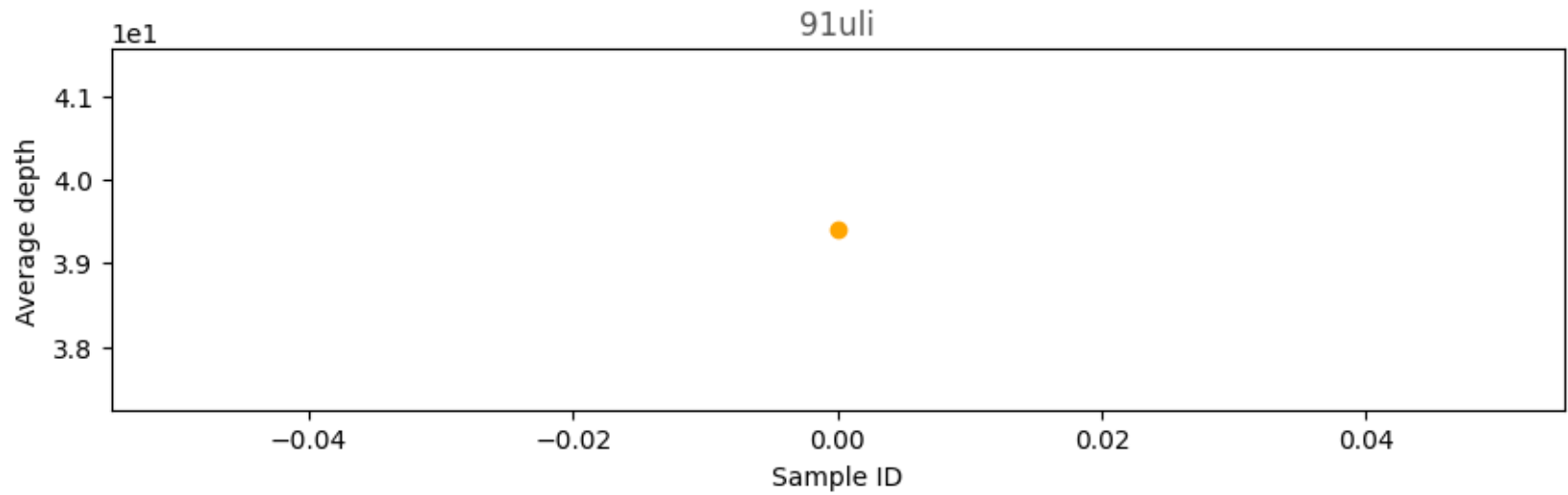
Ts/Tv by sample
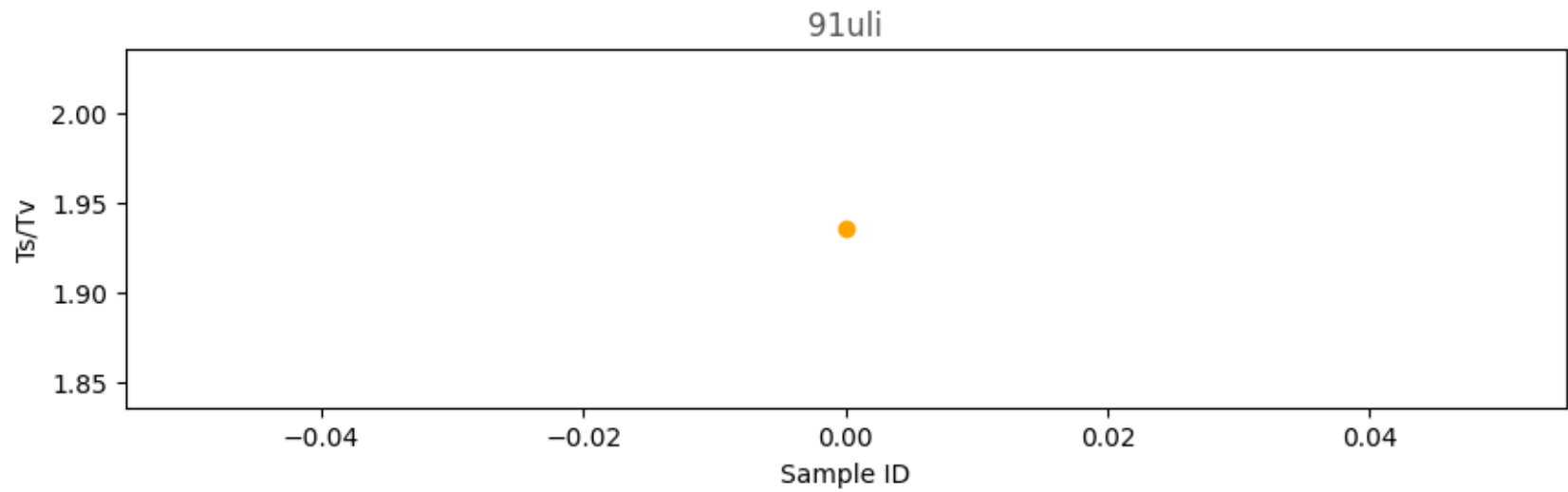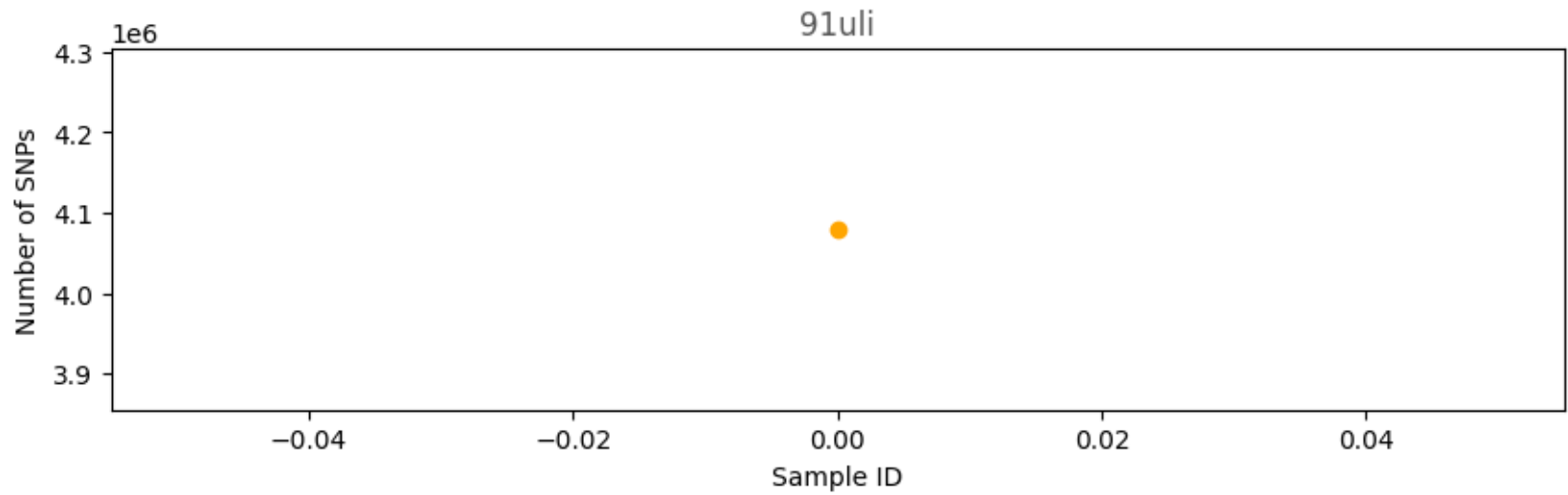
Hets vs non-ref Homs by sample
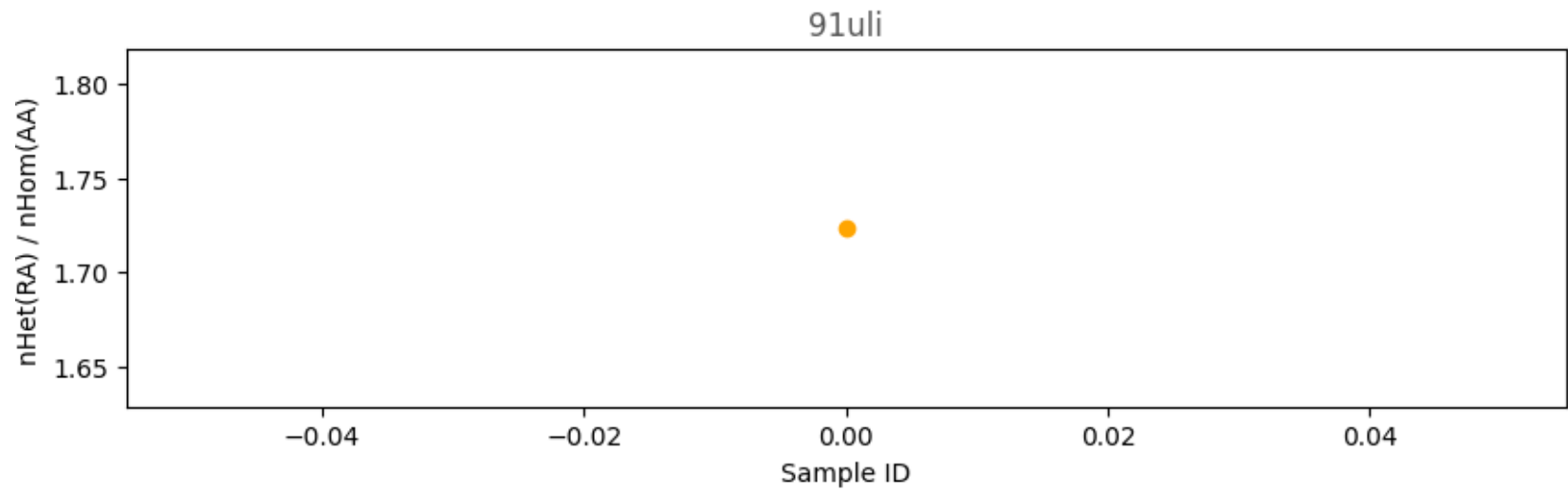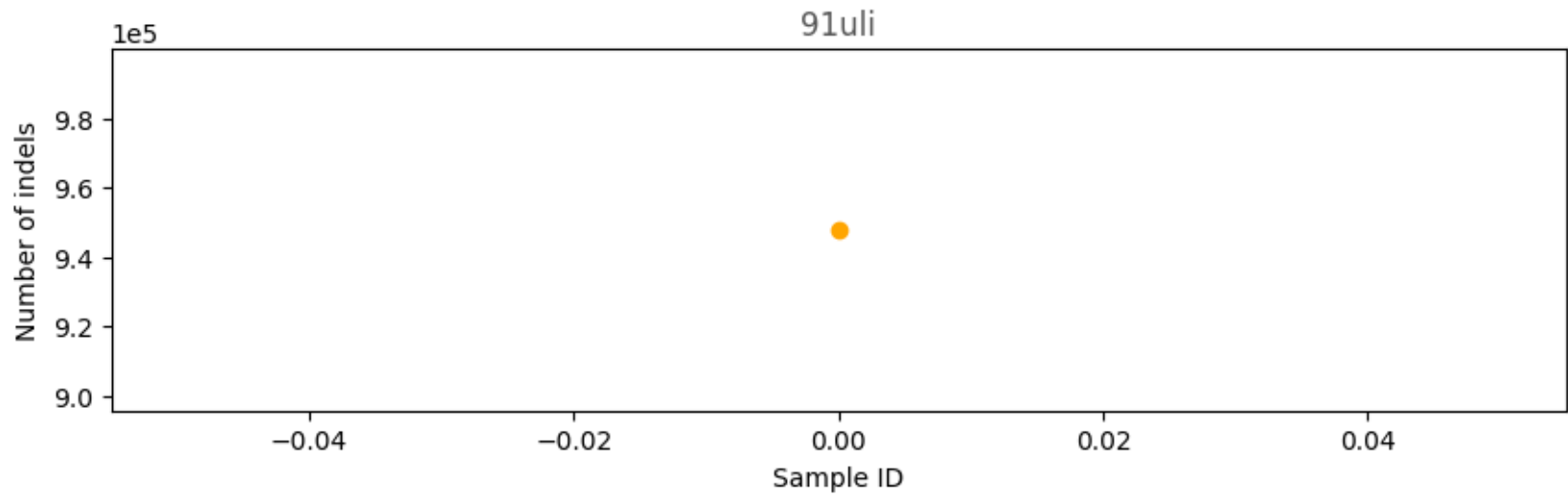
Singletons by sample (hets and homs)
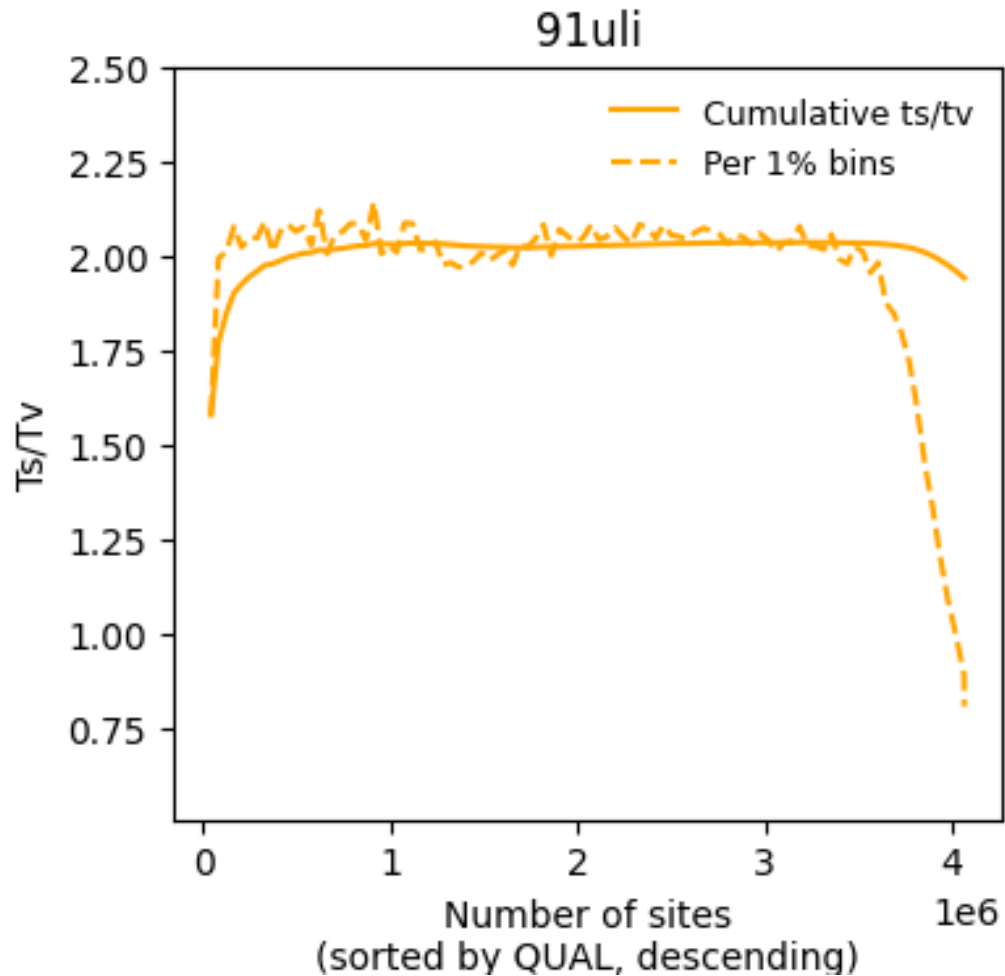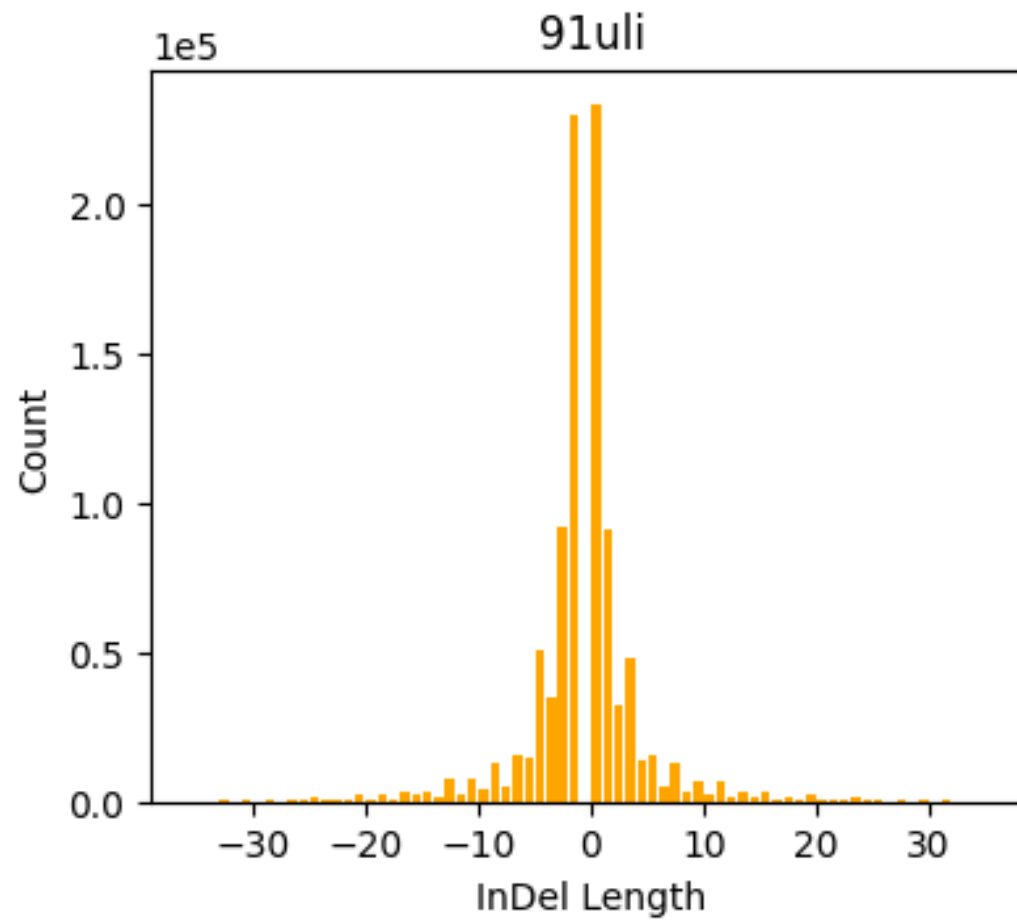
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

8

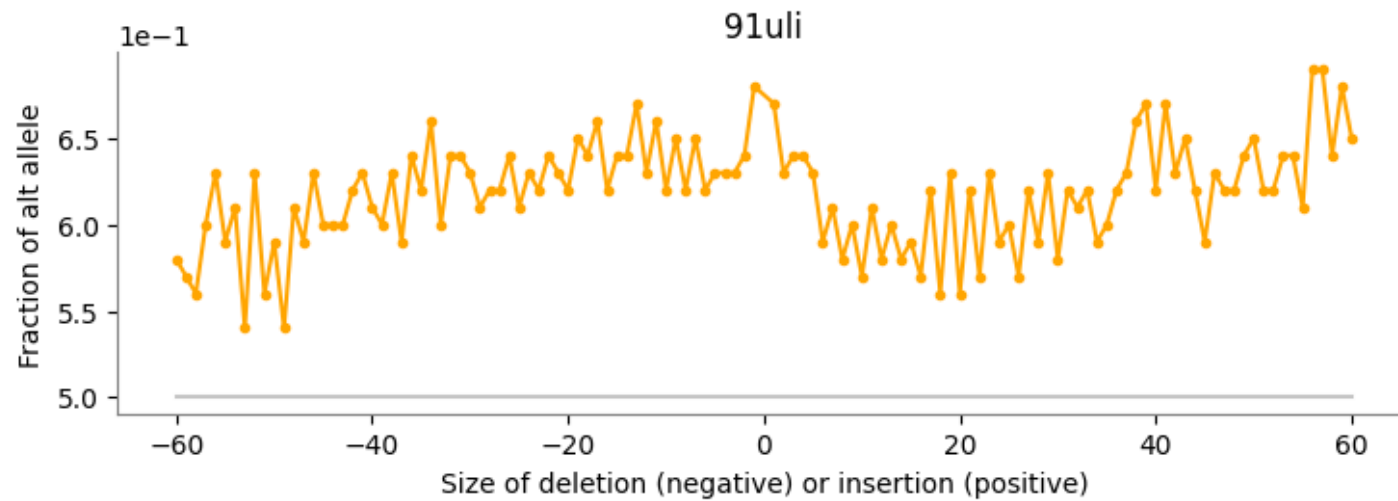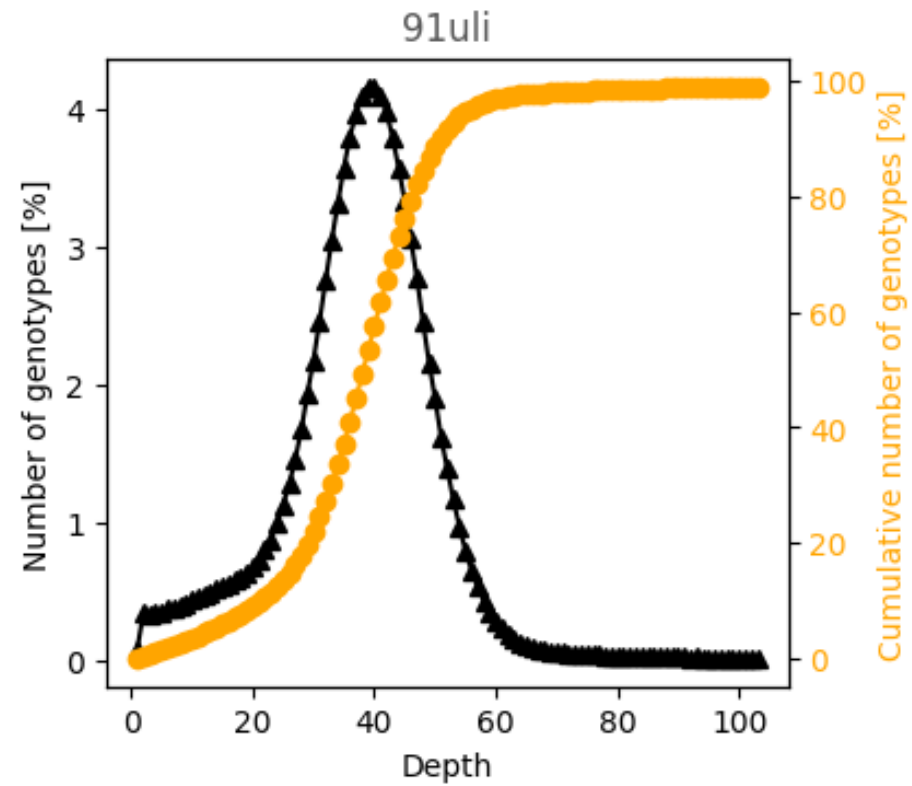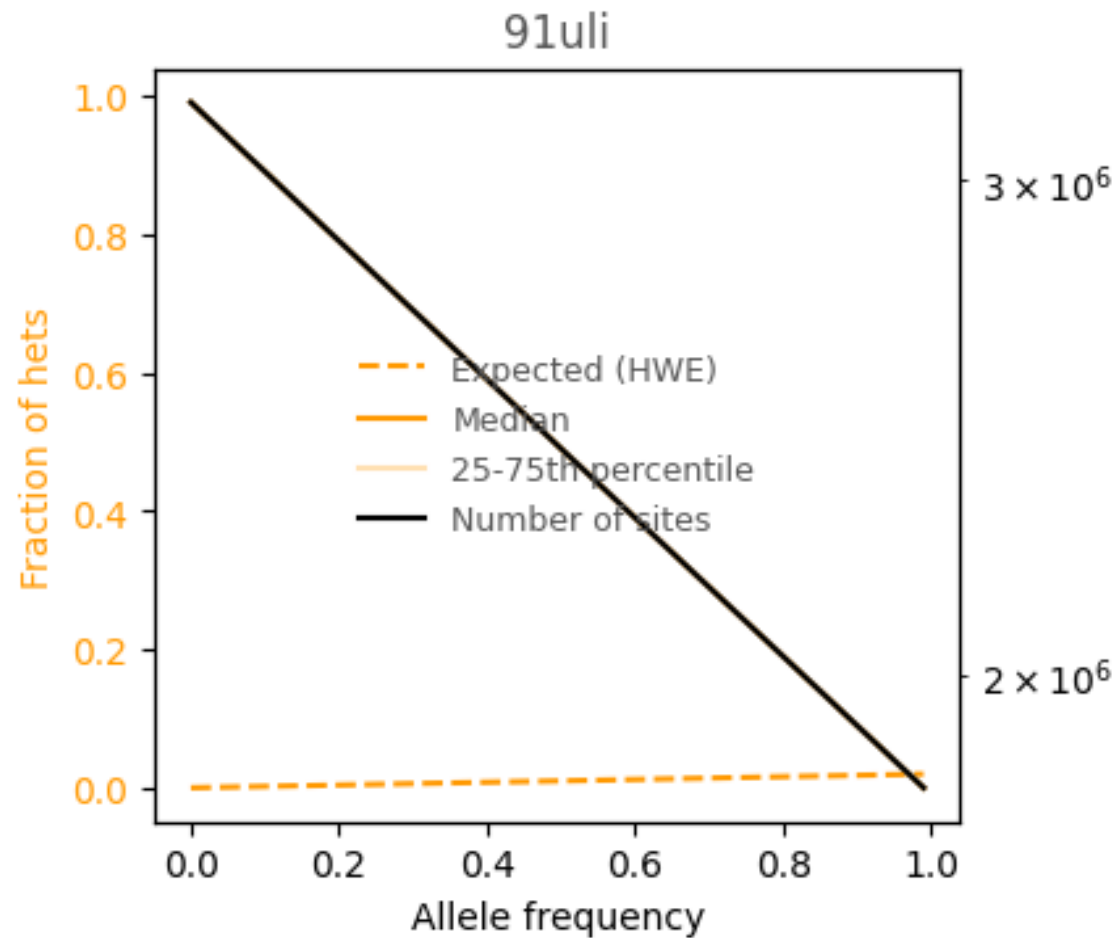# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types