Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
91rck	4,118,112	1.94	1.94	967,580	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
91rck	63.5%	1.91	69.1%	106,881	2,151

- 91rck .. `/ngc/projects2/gm/data/archive/2022/variants/snv/91rcknonf-103825128827-Normal_B`
`lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220107_A01176_BH5TGWDSX3-RHGM_LABKA_WGSA`
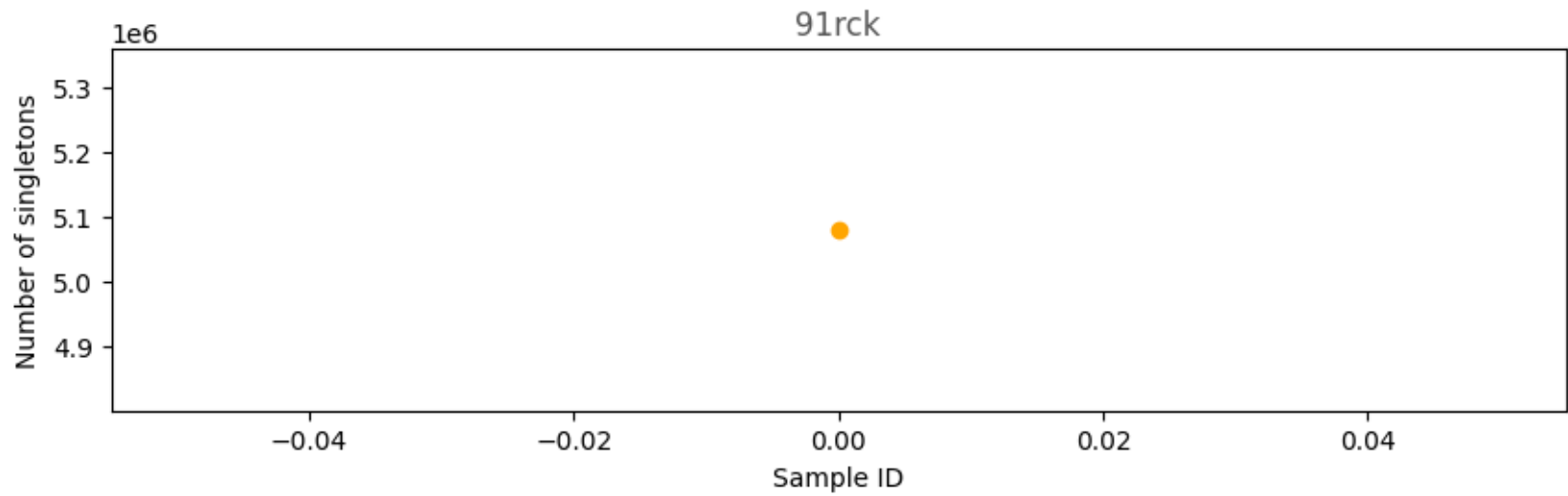`KUT-WGSAKUT03635_22RKG000104x01_snv_germline_raw.haplotype_caller.vcf.gz`
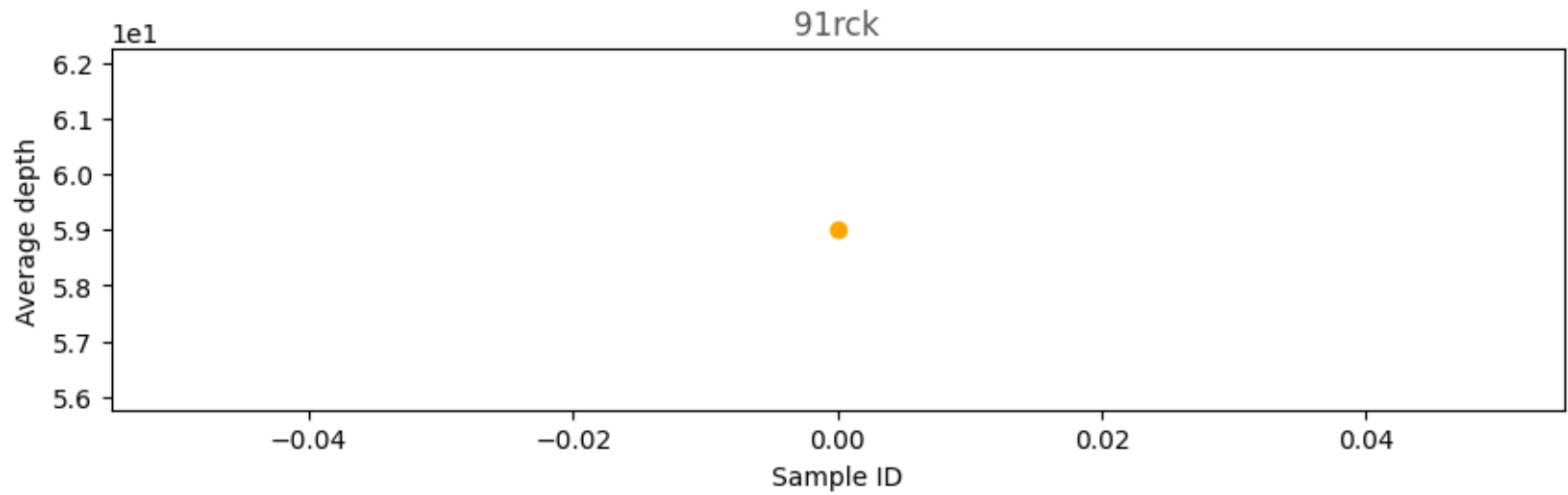
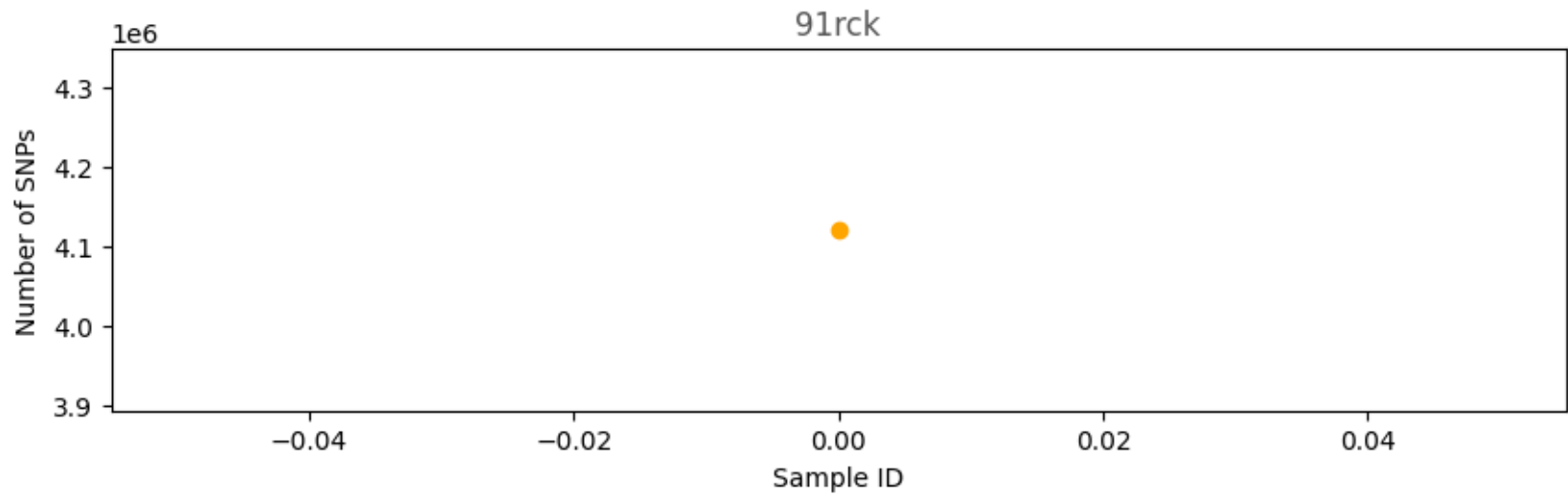# Ts/Tv by sample
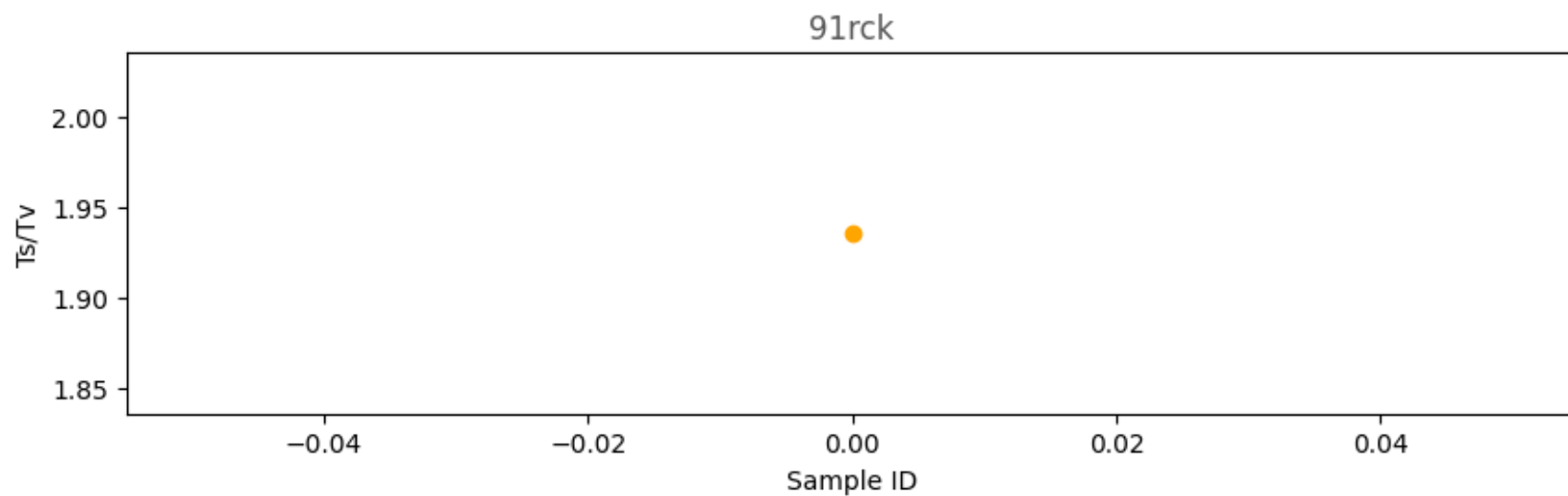
Hets vs non-ref Homs by sample
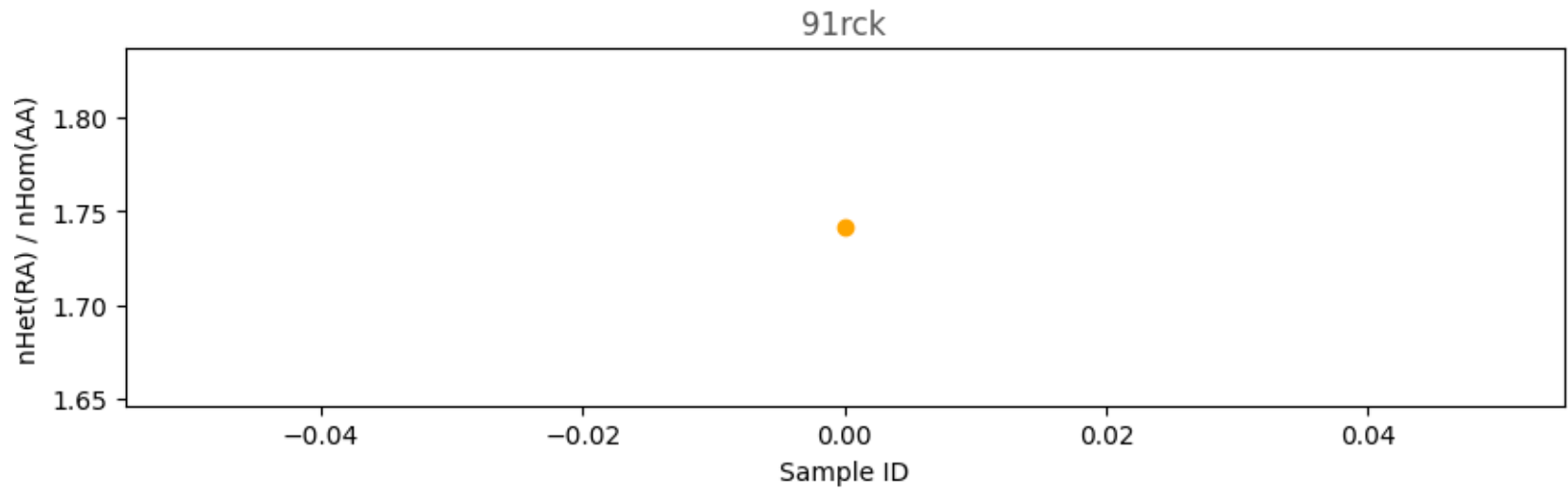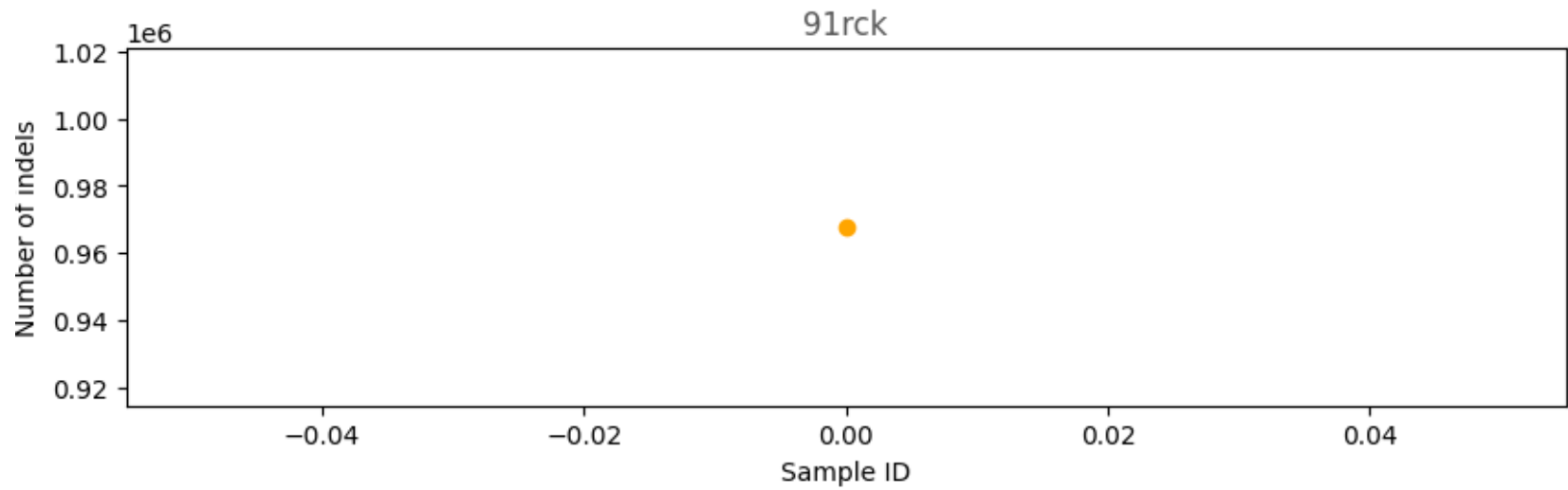
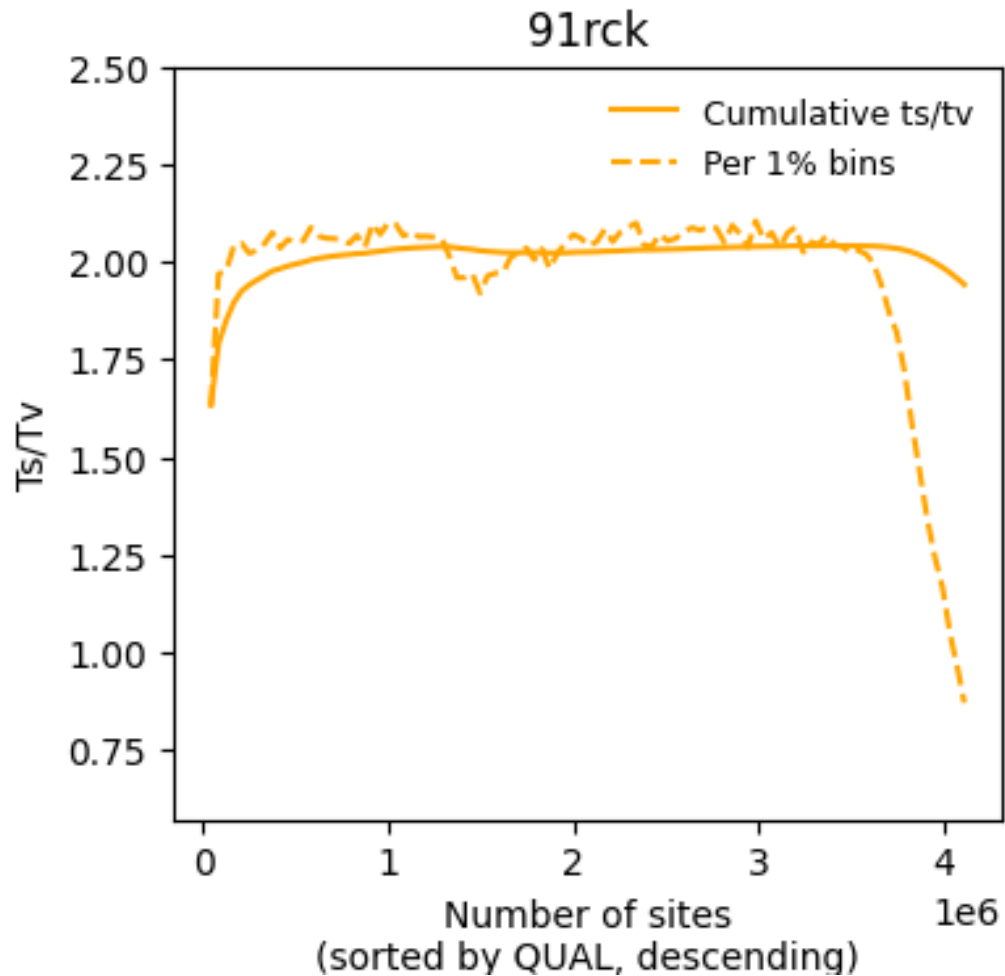# Singletons by sample (hets and homs)
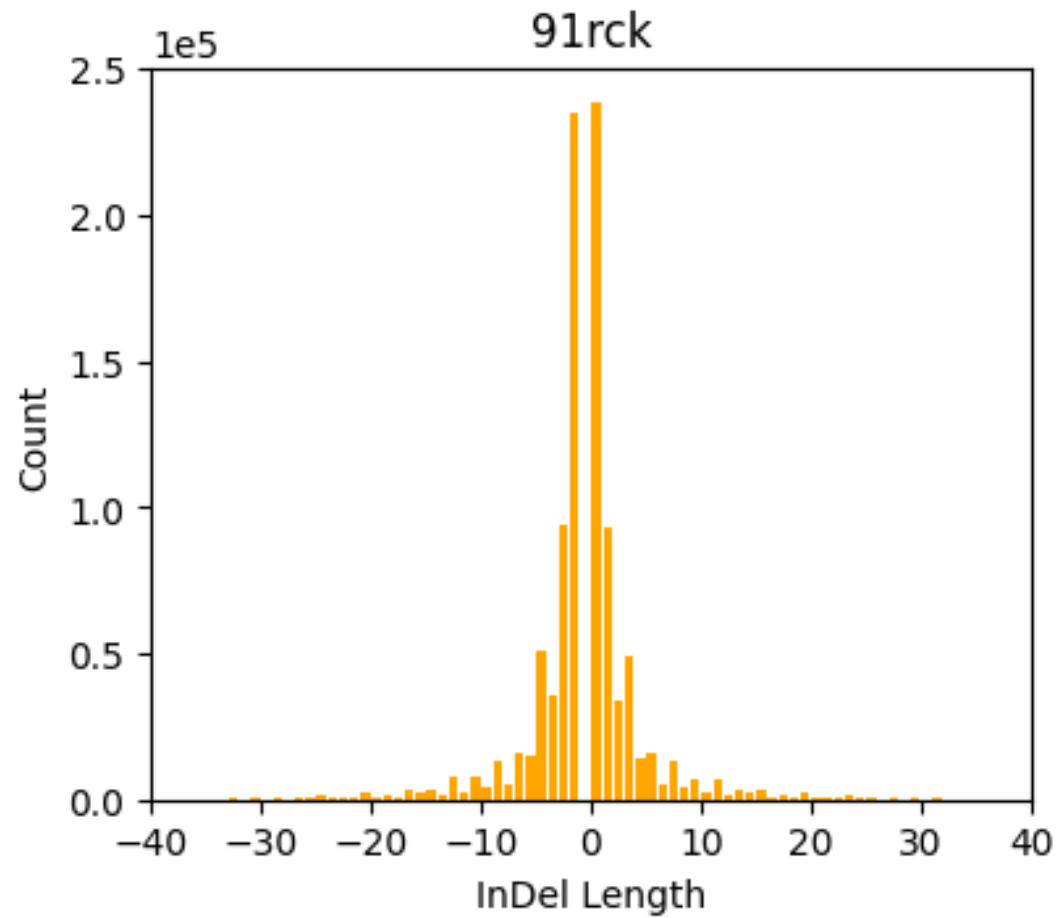
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

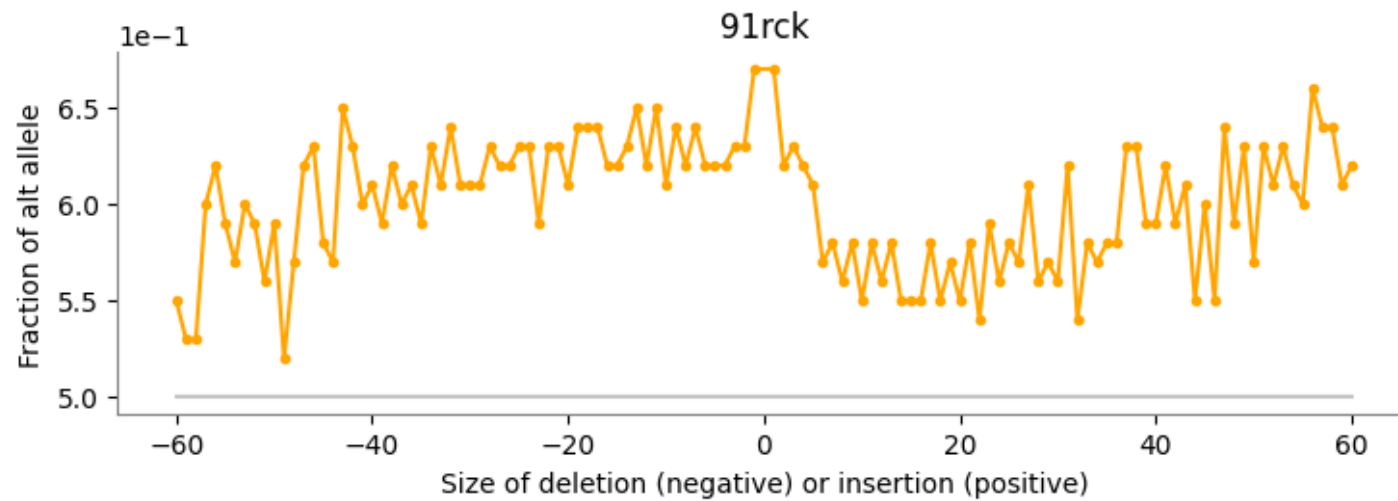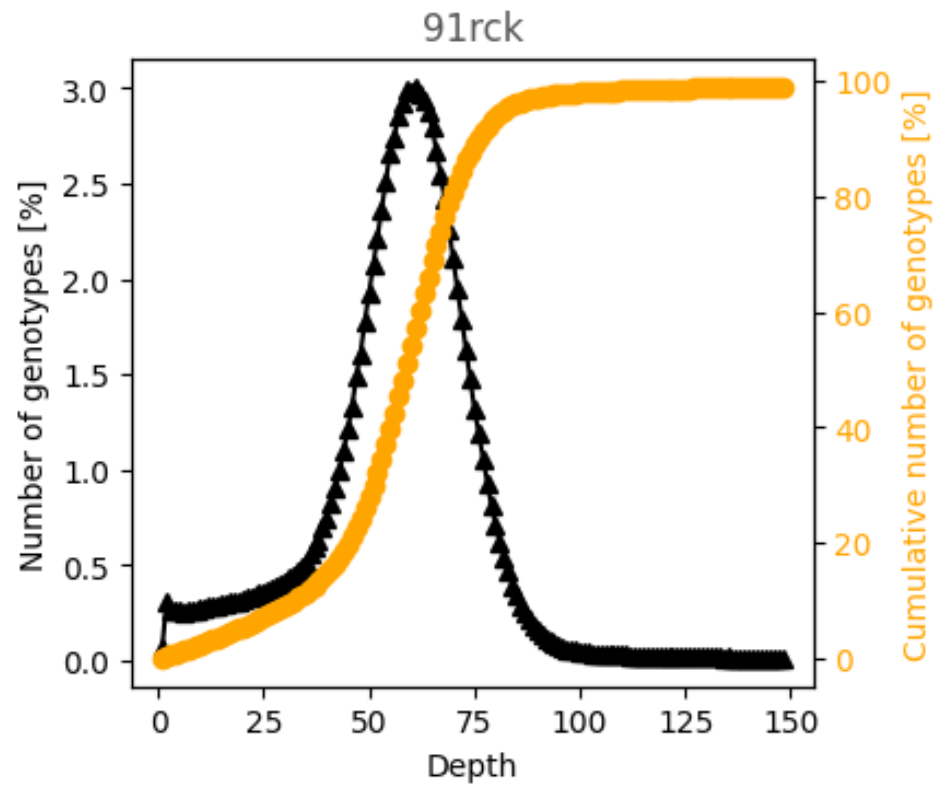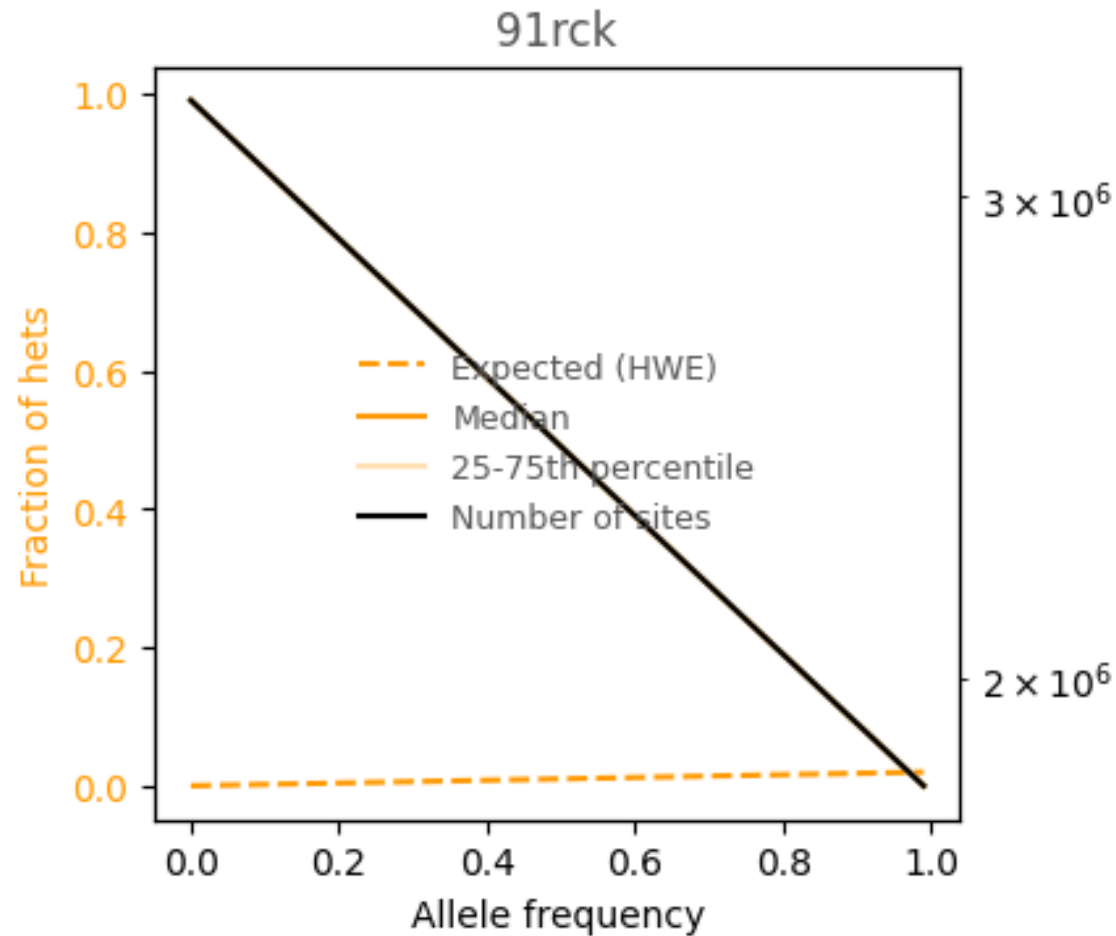# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types