Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
92cek	4,112,524	1.93	1.94	959,265	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
92cek	63.3%	1.91	68.5%	102,999	2,249

- 92cek .. `/ngc/projects2/gm/data/archive/2022/variants/snv/92cekasif-103901319868-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01561-220916_A01411_BHYNLVDSX3-EXT_LAB KA_NGCWGS-NGCWGS05362_22RKG019631_snv_germline_raw.haplotype_caller.vcf.gz`
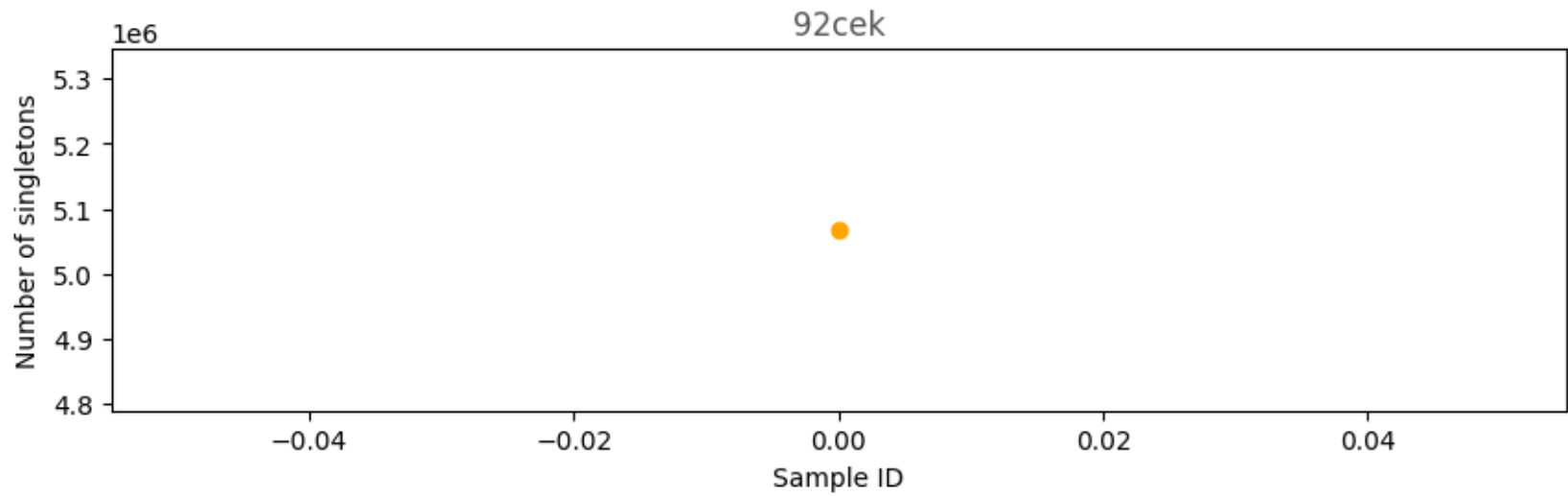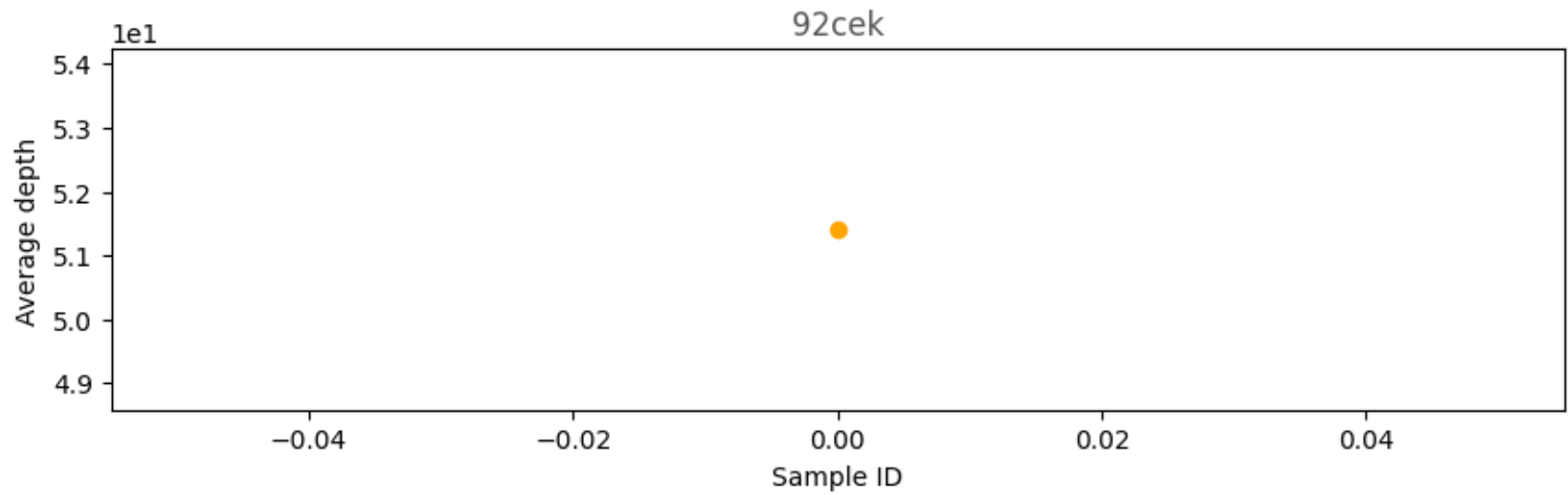
Ts/Tv by sample
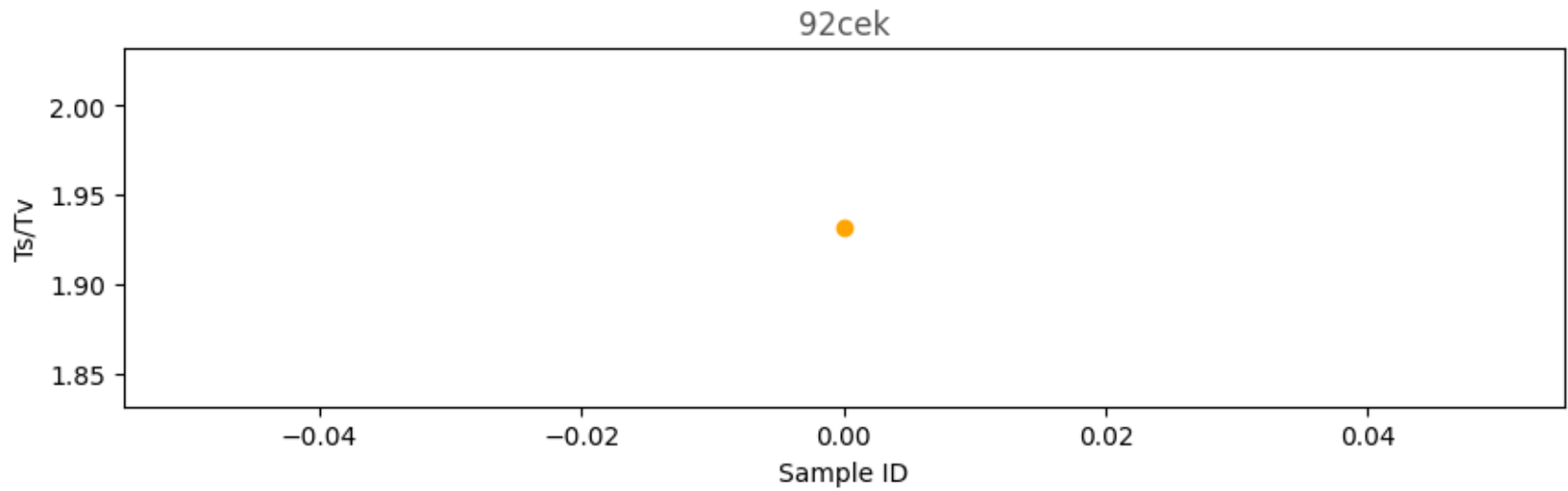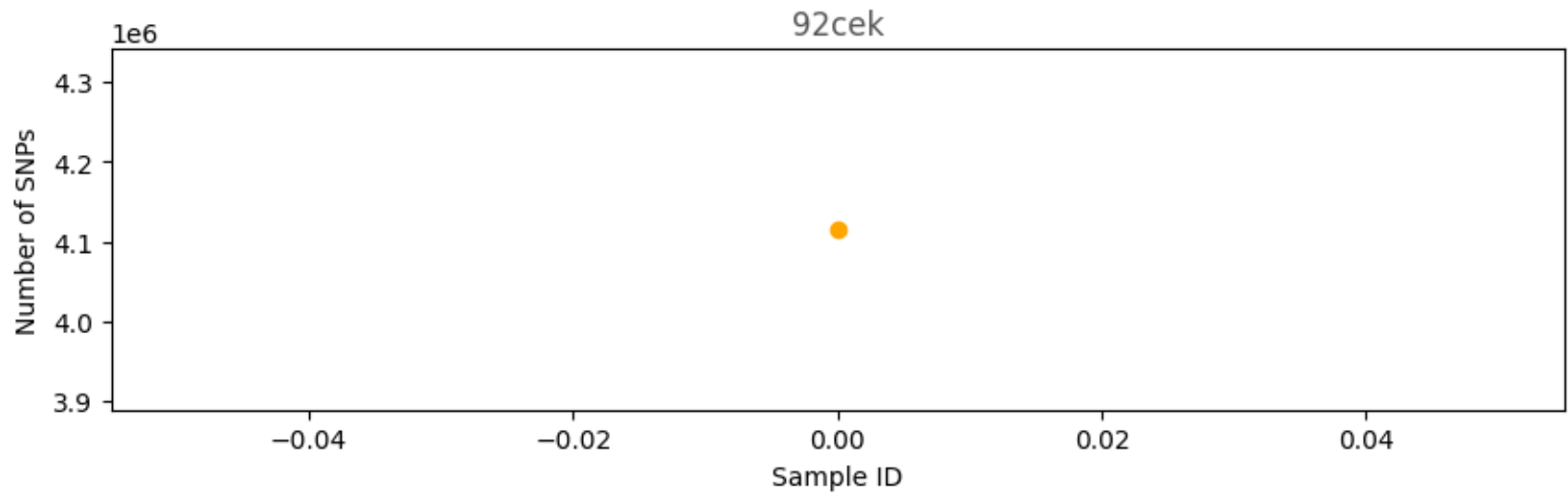
Hets vs non-ref Homs by sample
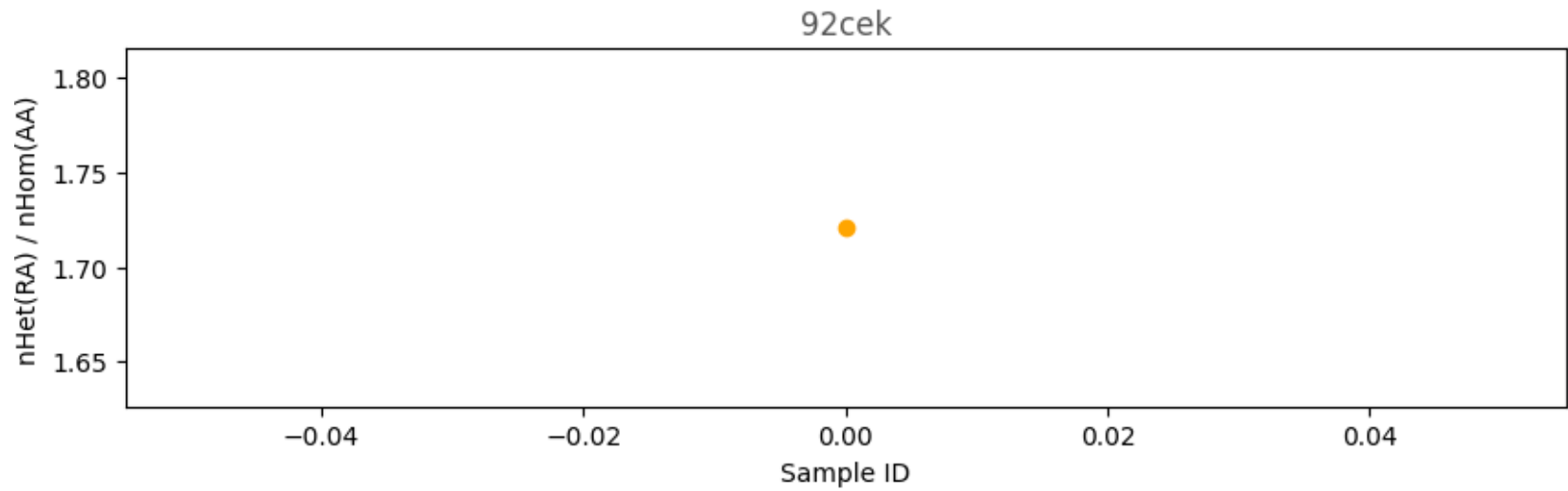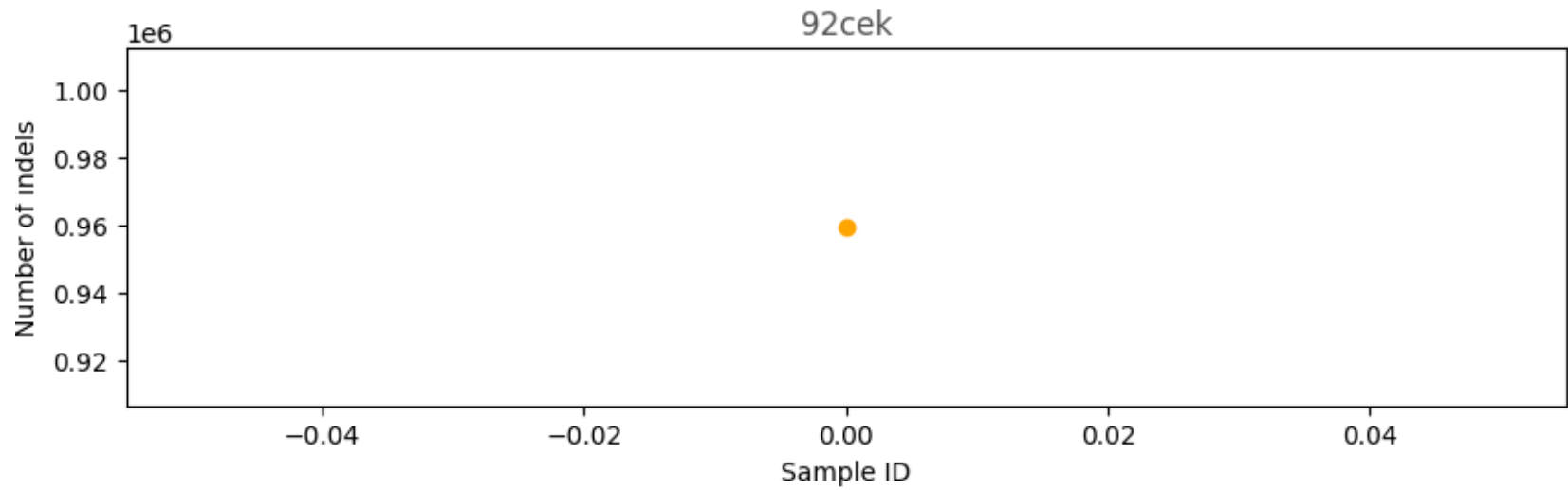
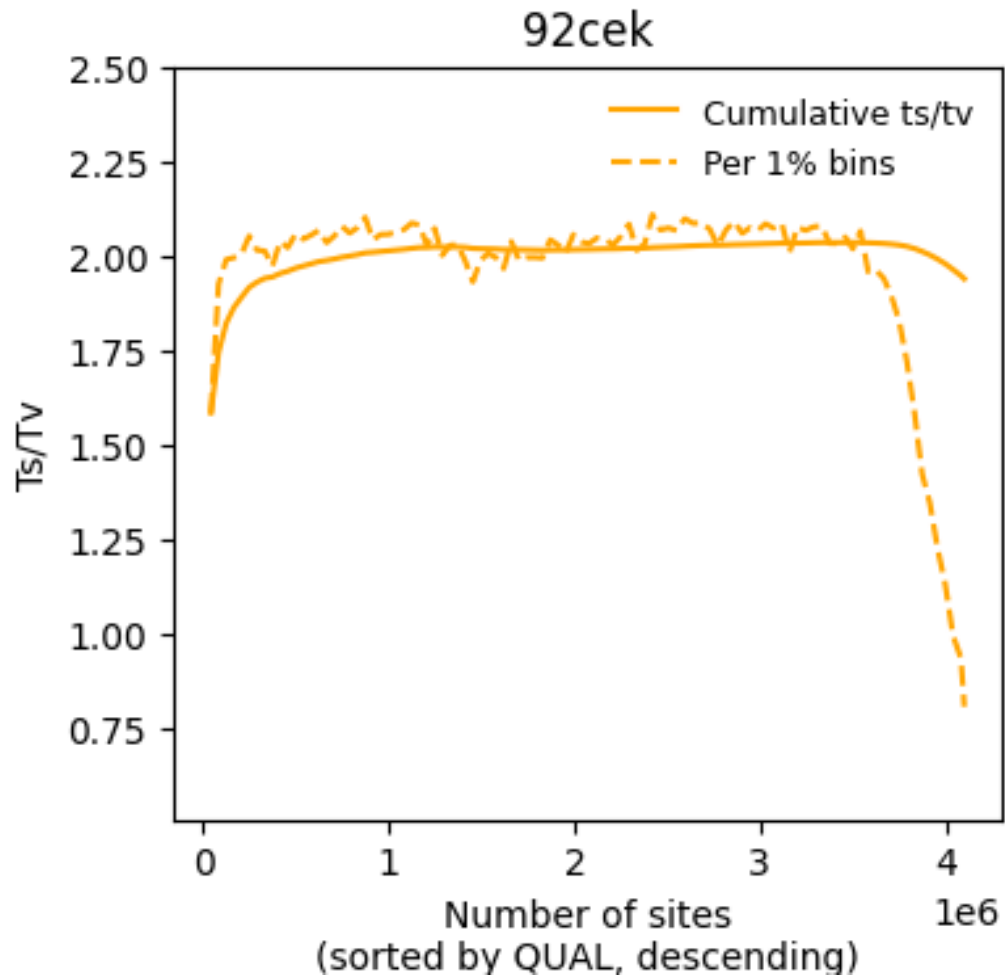# Singletons by sample (hets and homs)
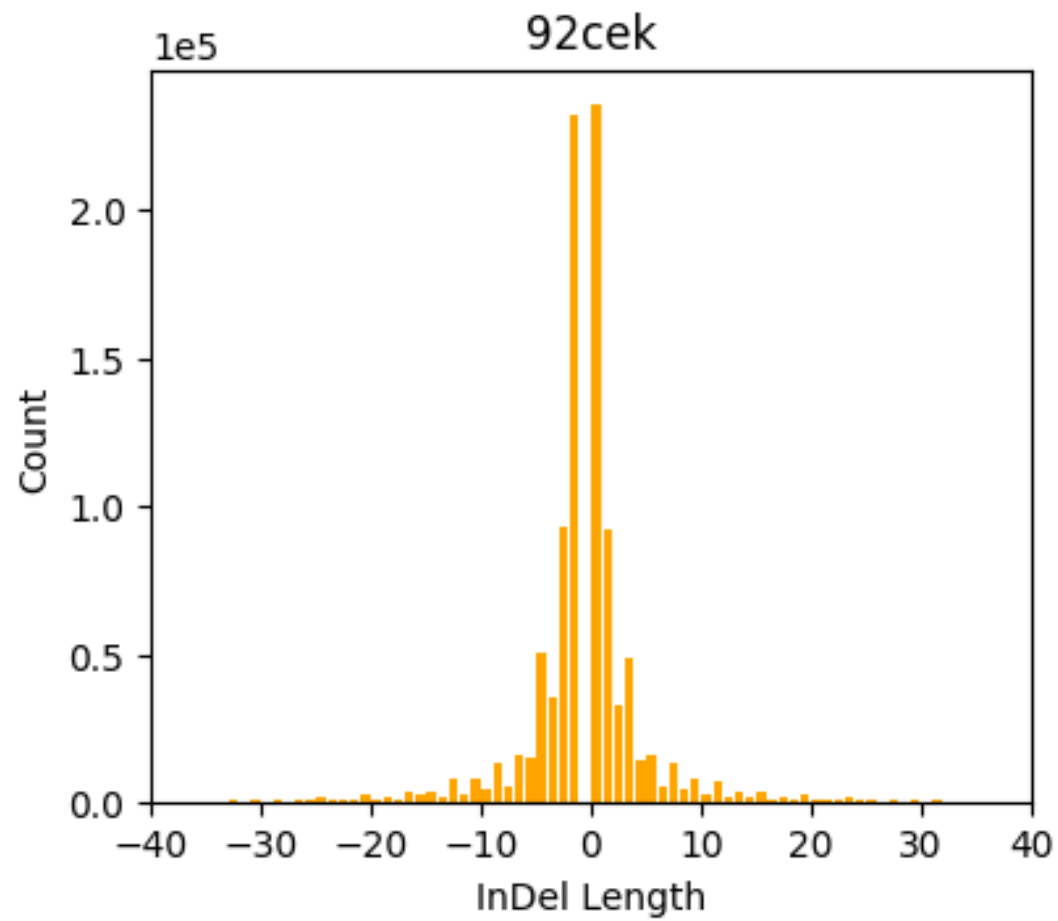
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

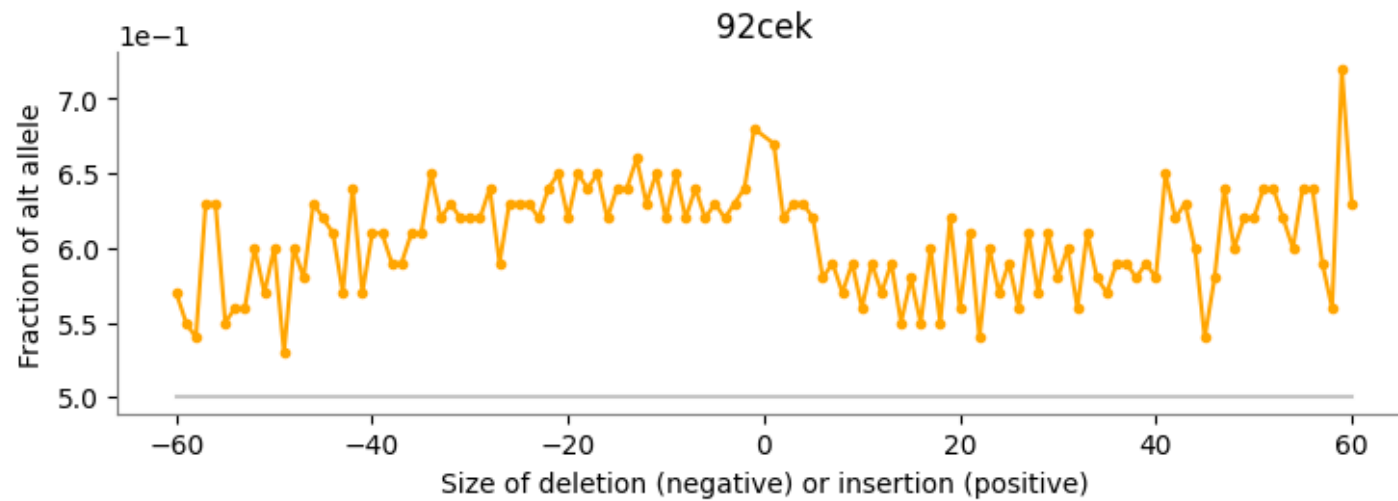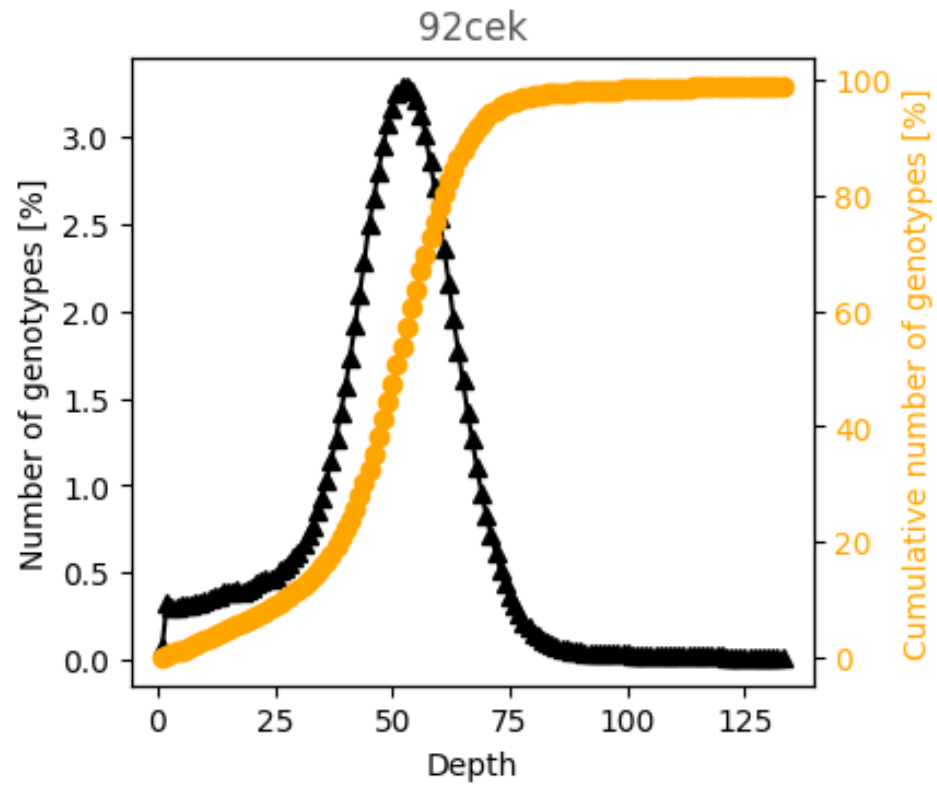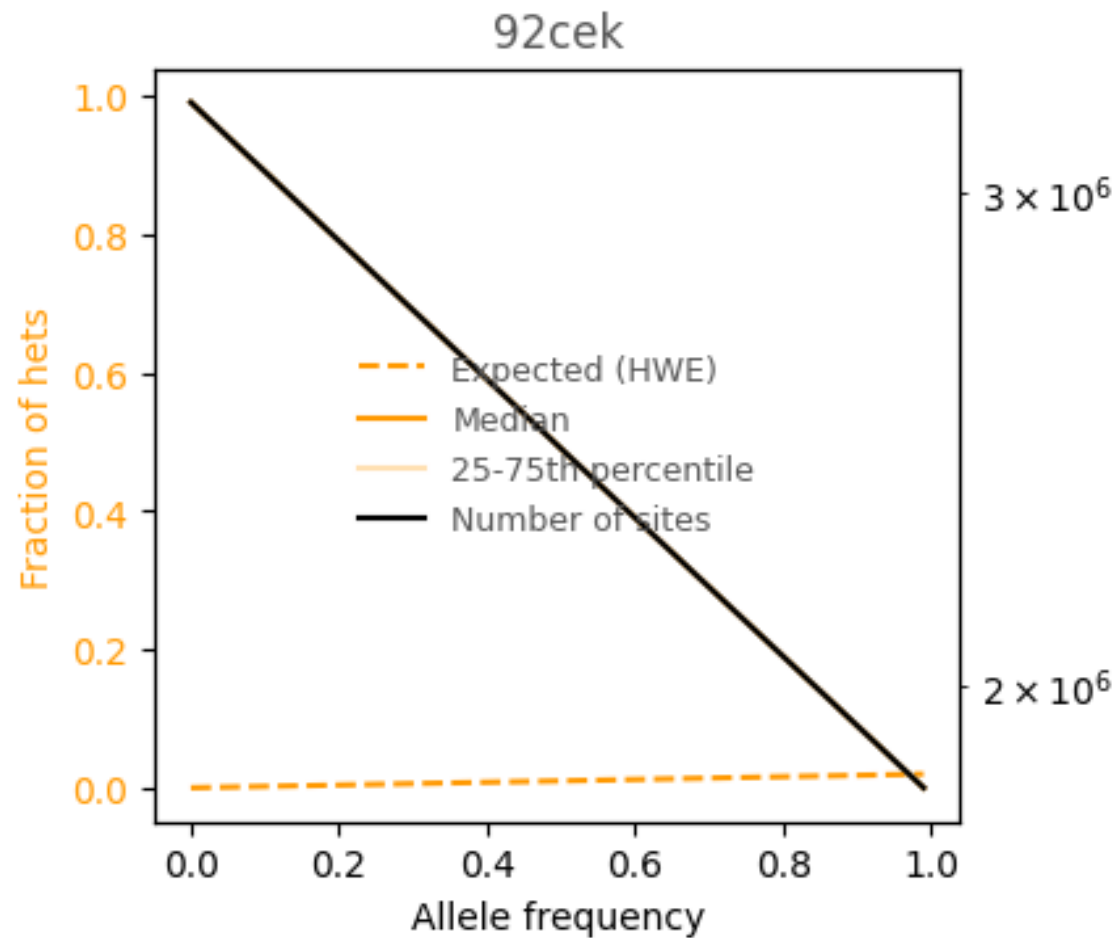# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types