Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90ceo	4,068,438	1.94	1.95	940,026	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90ceo	63.2%	1.92	67.5%	93,678	1,893

- 90ceo .. `/ngc/projects2/gm/data/archive/2022/variants/snv/90ceoetof-103804479949-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211117_A01411_BHT773DSX2-RHGM_LABKA_WGS- WGS03333_snv_germline_raw.haplotype_caller.vcf.gz`
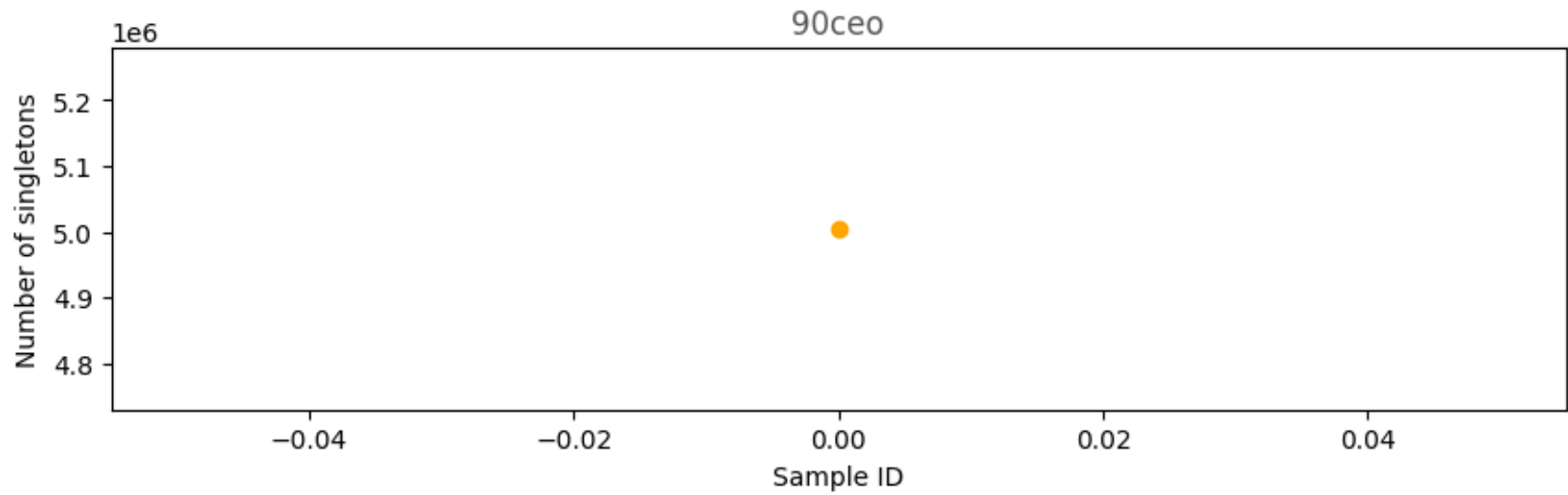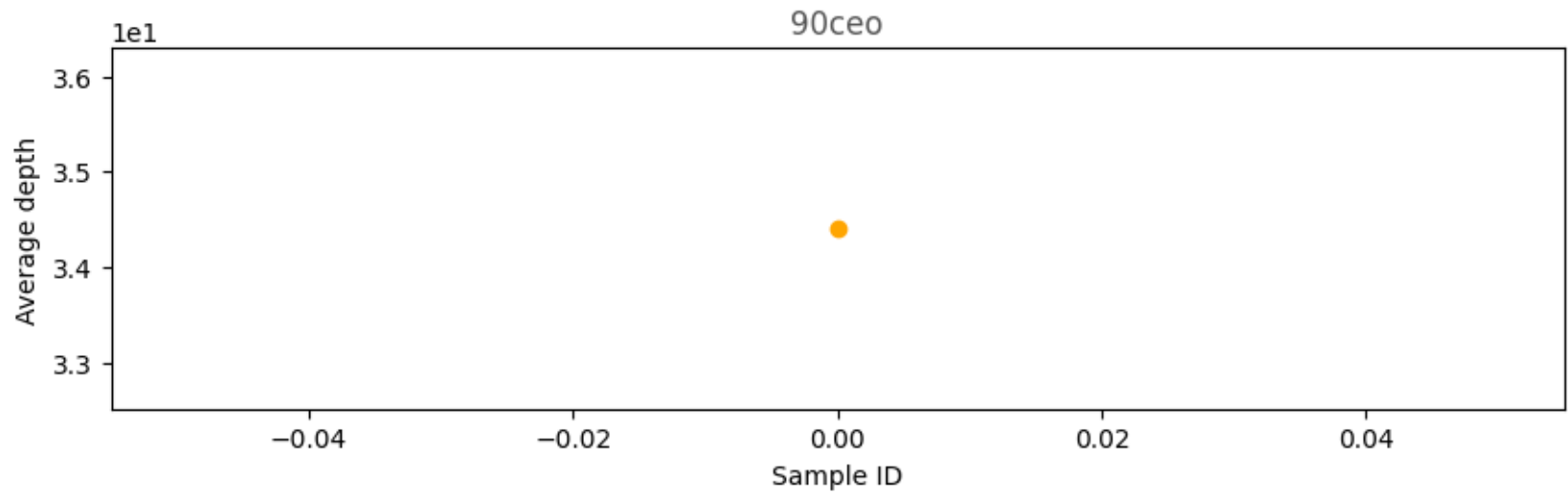
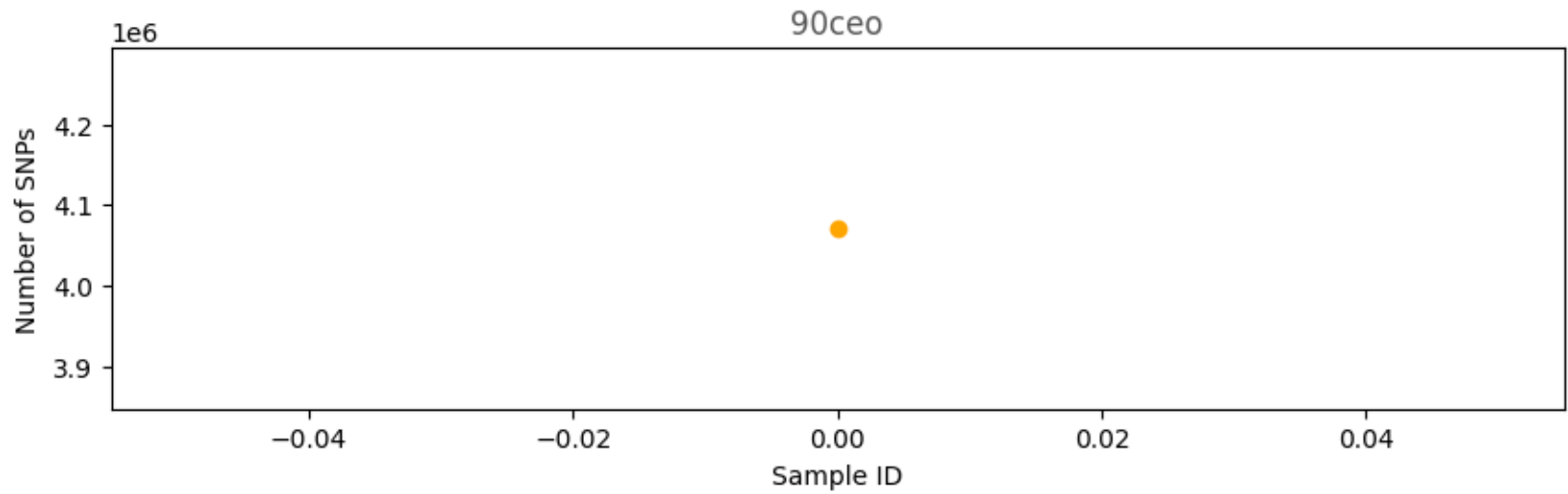# Ts/Tv by sample
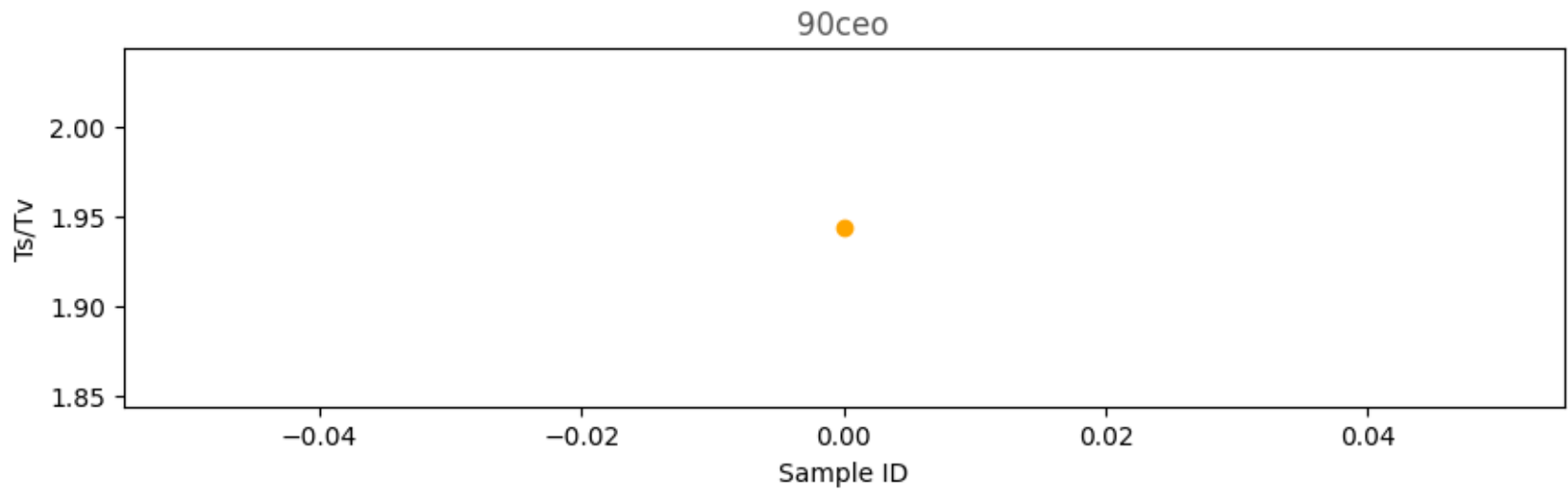
Hets vs non-ref Homs by sample
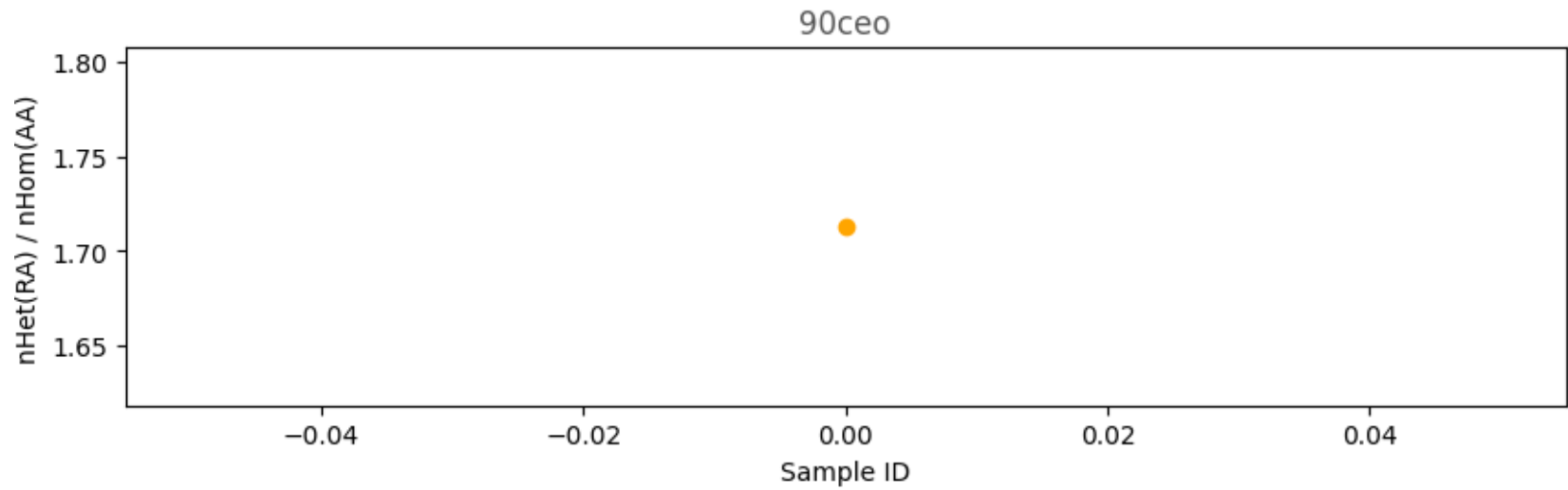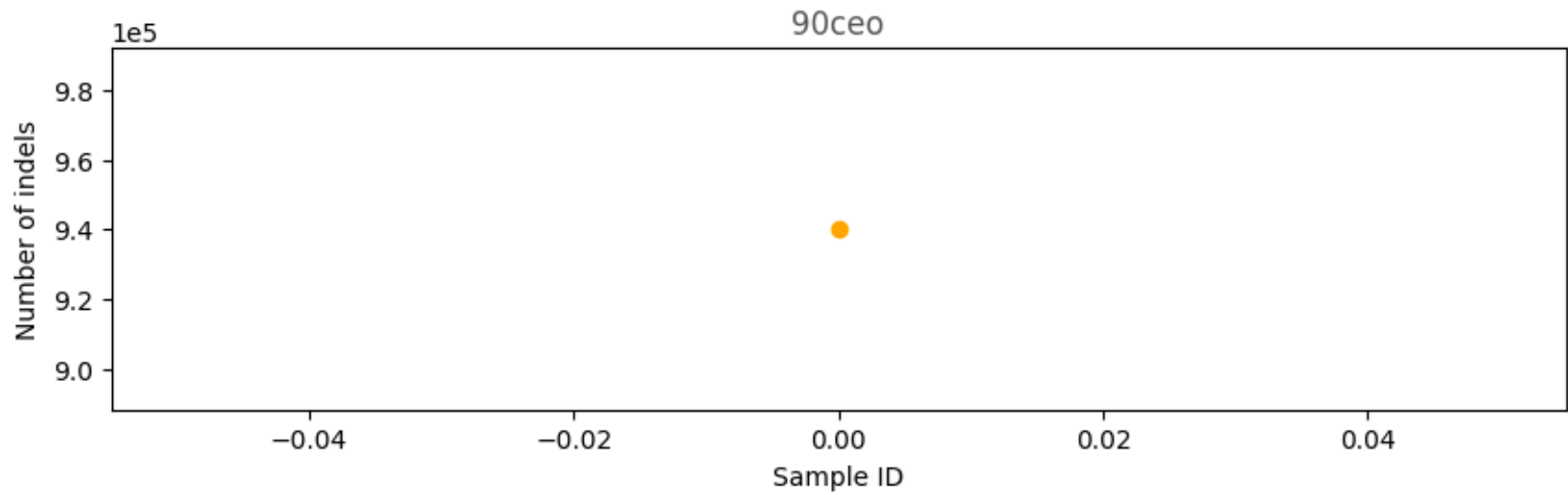
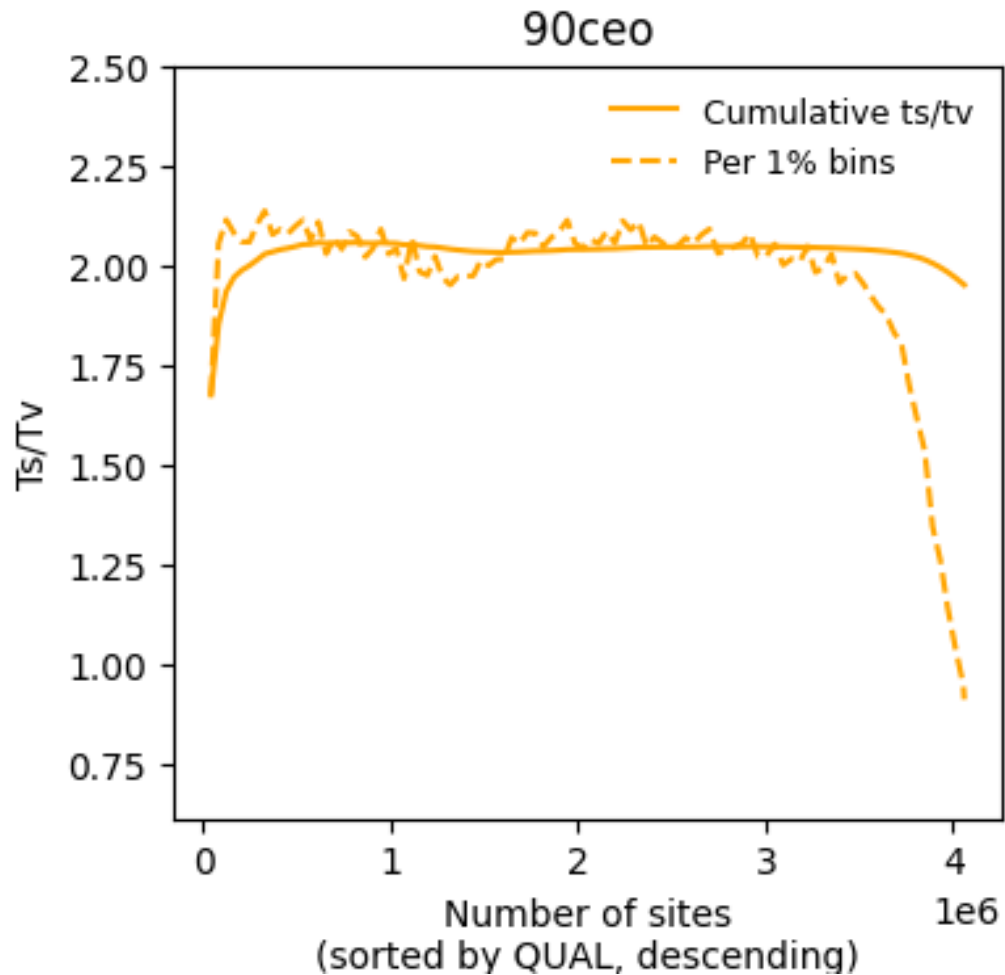# Singletons by sample (hets and homs)
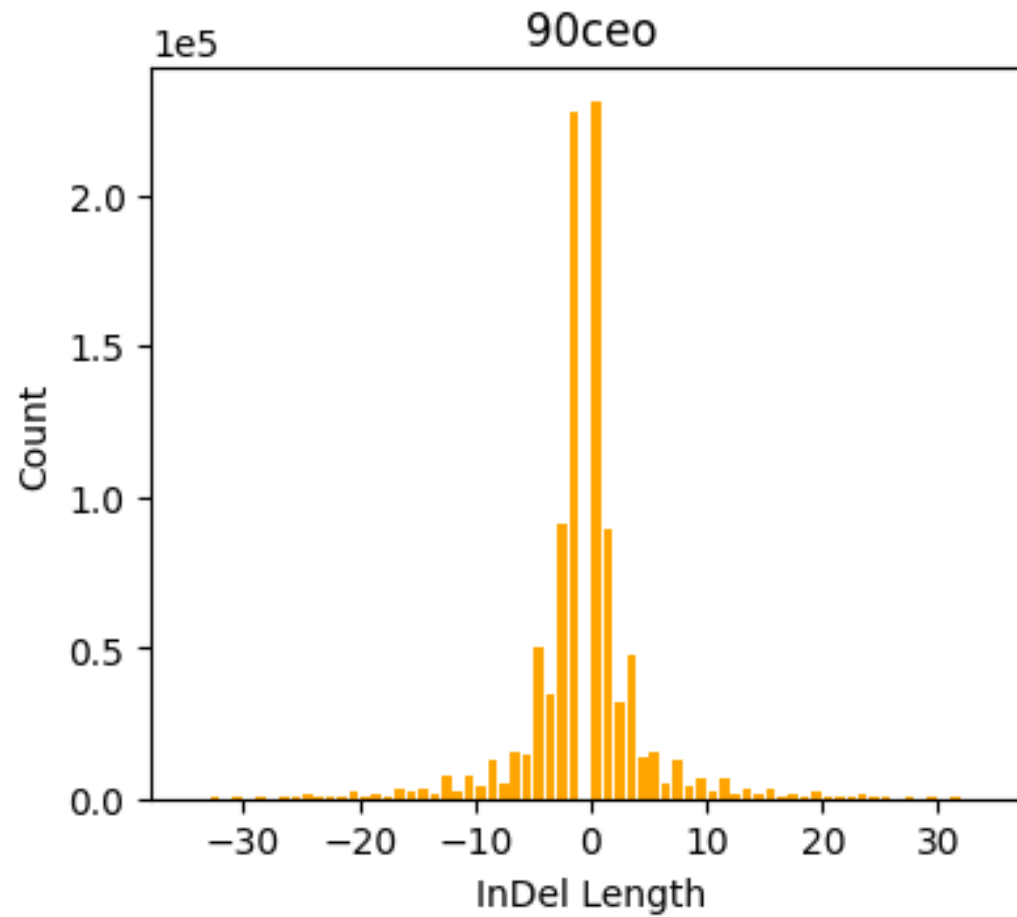
Average depth by sample

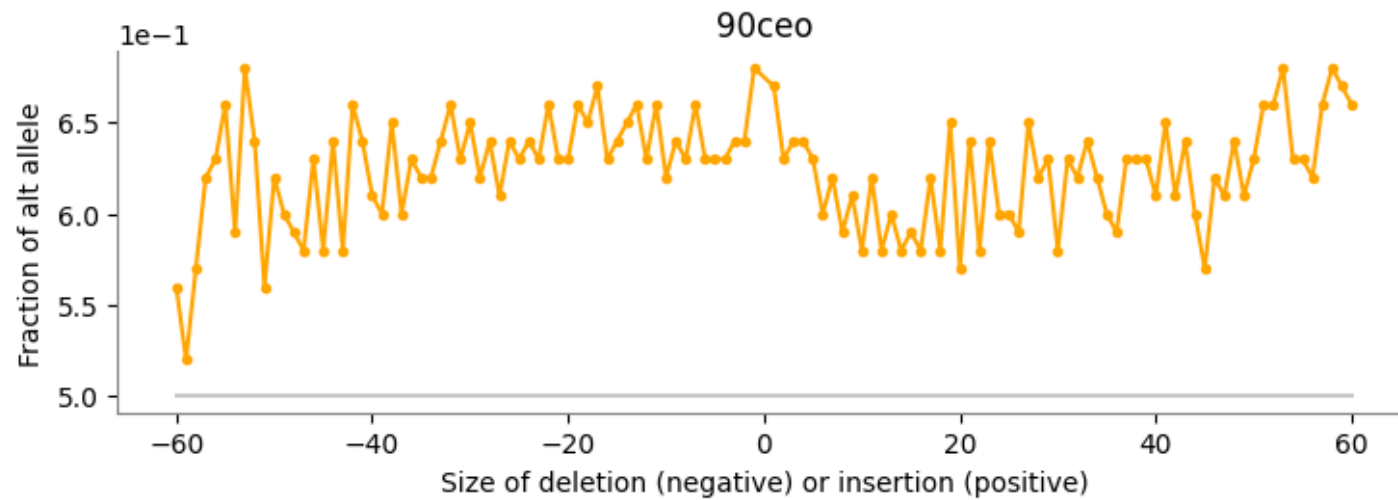# Number of SNPs by sample

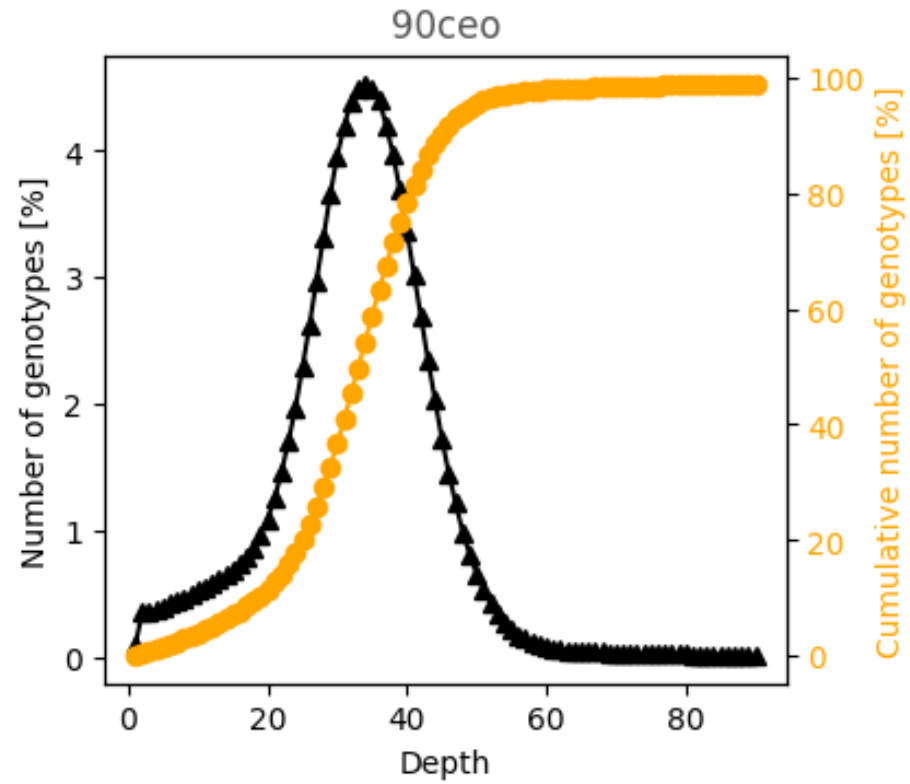# Number of indels by sample
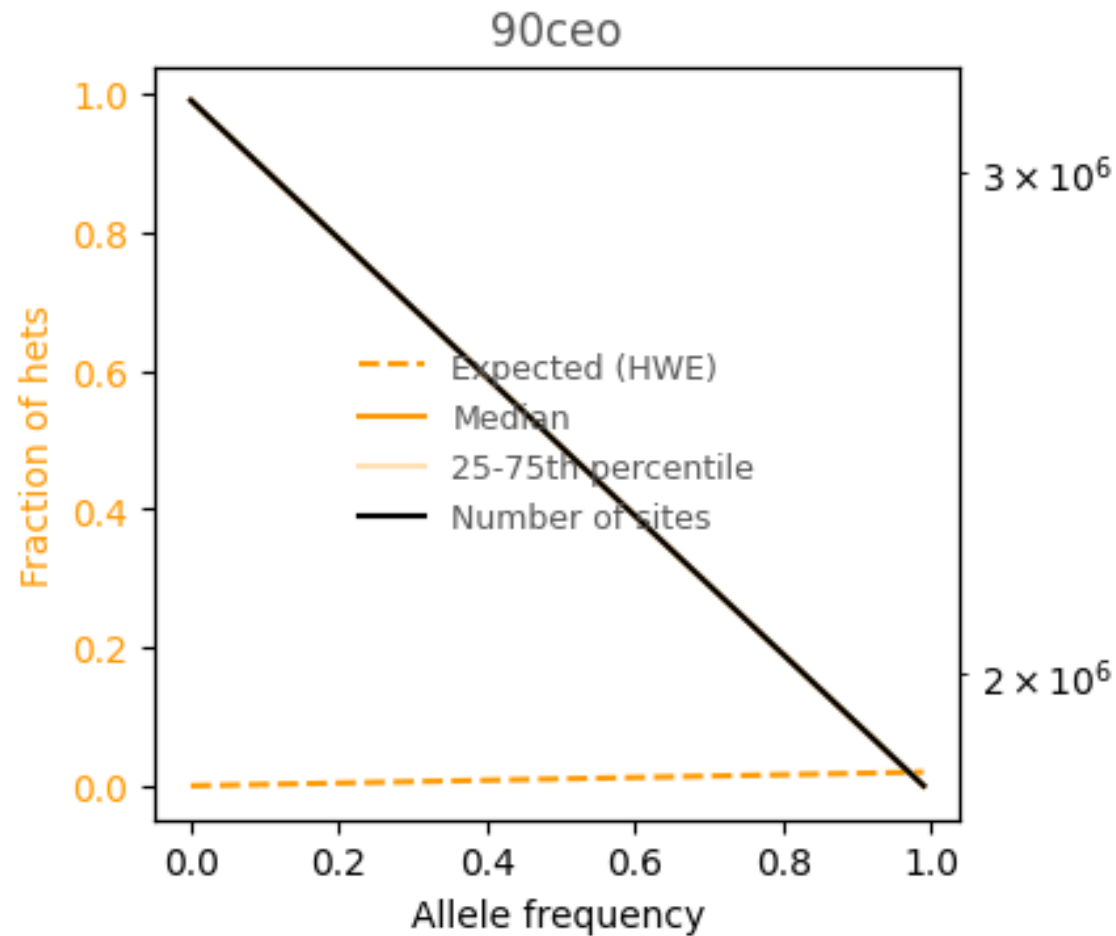
Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

90ceo

Substitution types