Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
89olo	4,059,490	1.93	1.94	951,522	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
89olo	62.1%	1.91	67.0%	100,917	1,929

- 89olo .. `/ngc/projects2/gm/data/archive/2022/variants/snv/89oloefum-103917253250-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220928_A01961_AH35JJDSX5-RHGM_LABKA_WGSA KUT-WGSAKUT05635_22RKG023343x01_snv_germline_raw.haplotype_caller.vcf.gz`
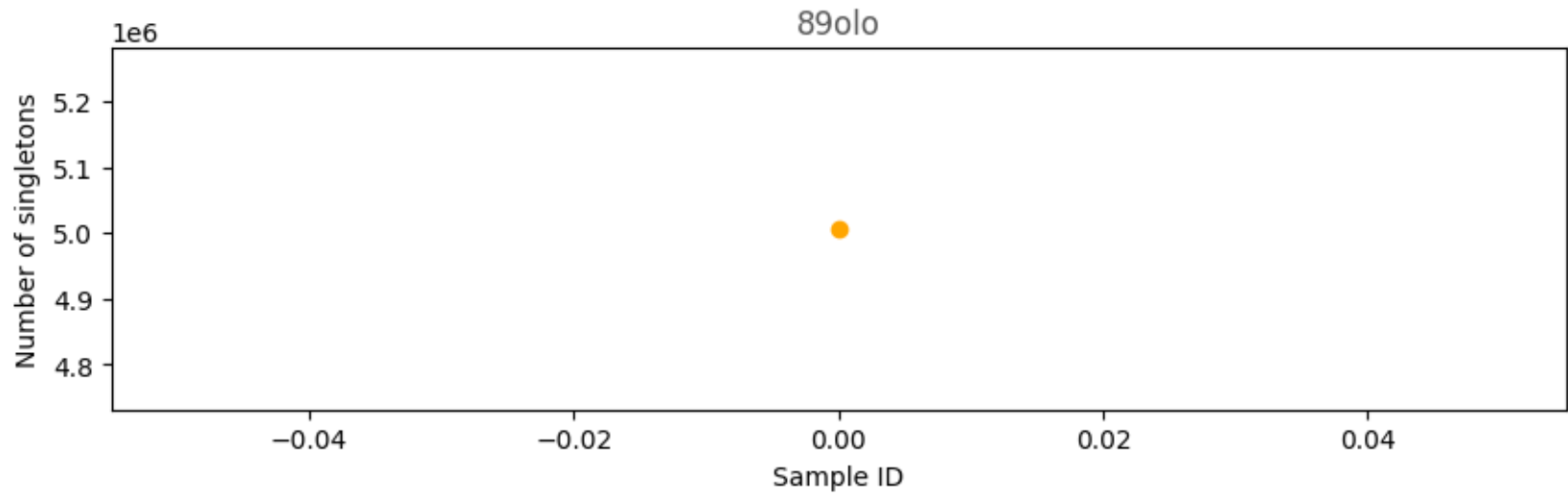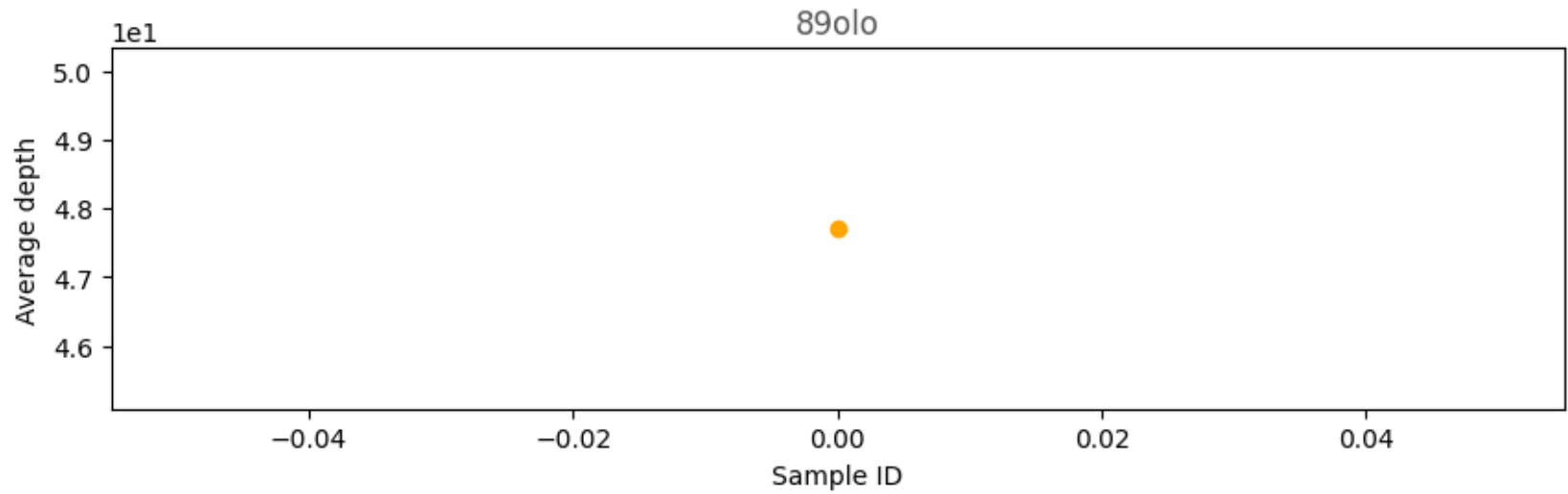
Ts/Tv by sample
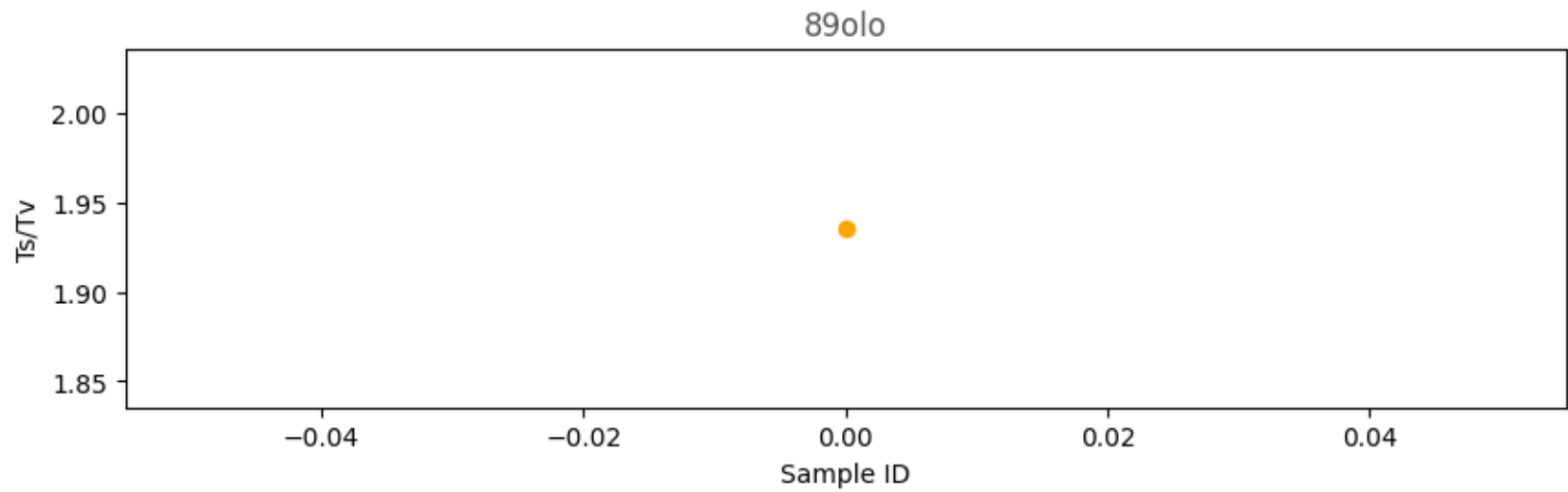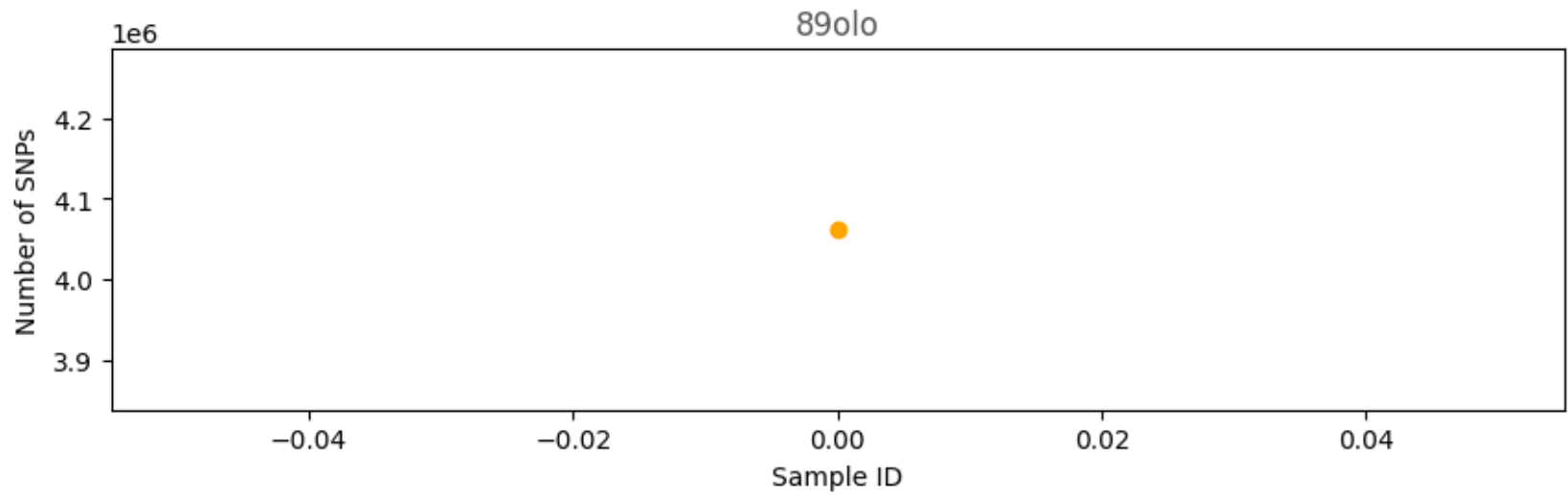
Hets vs non-ref Homs by sample
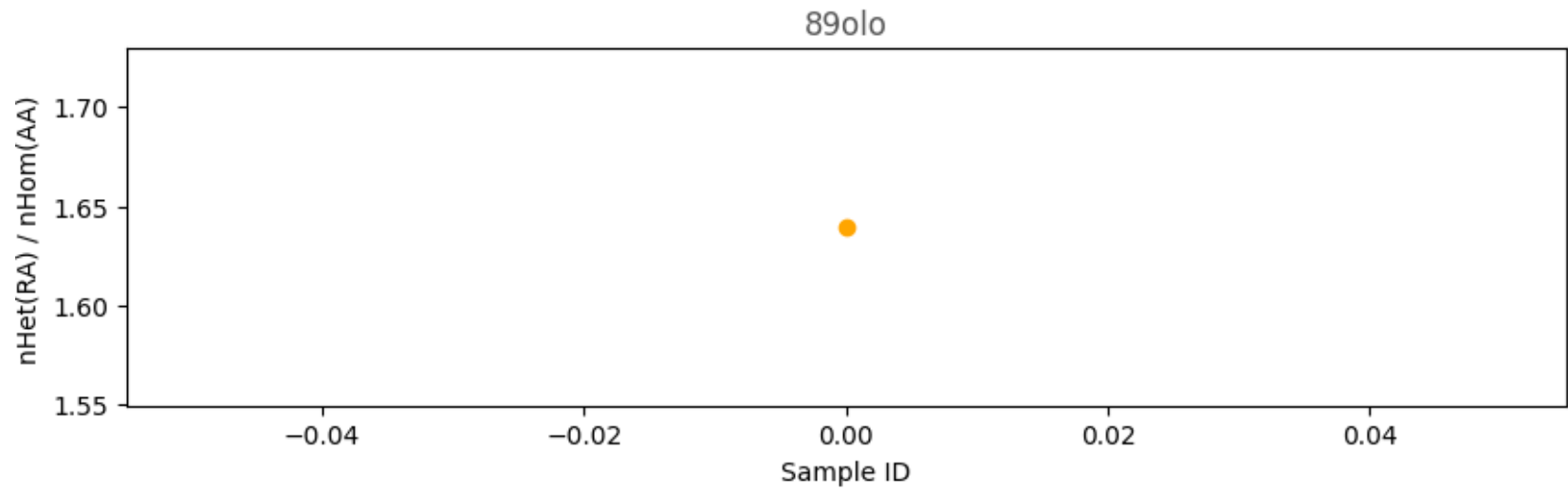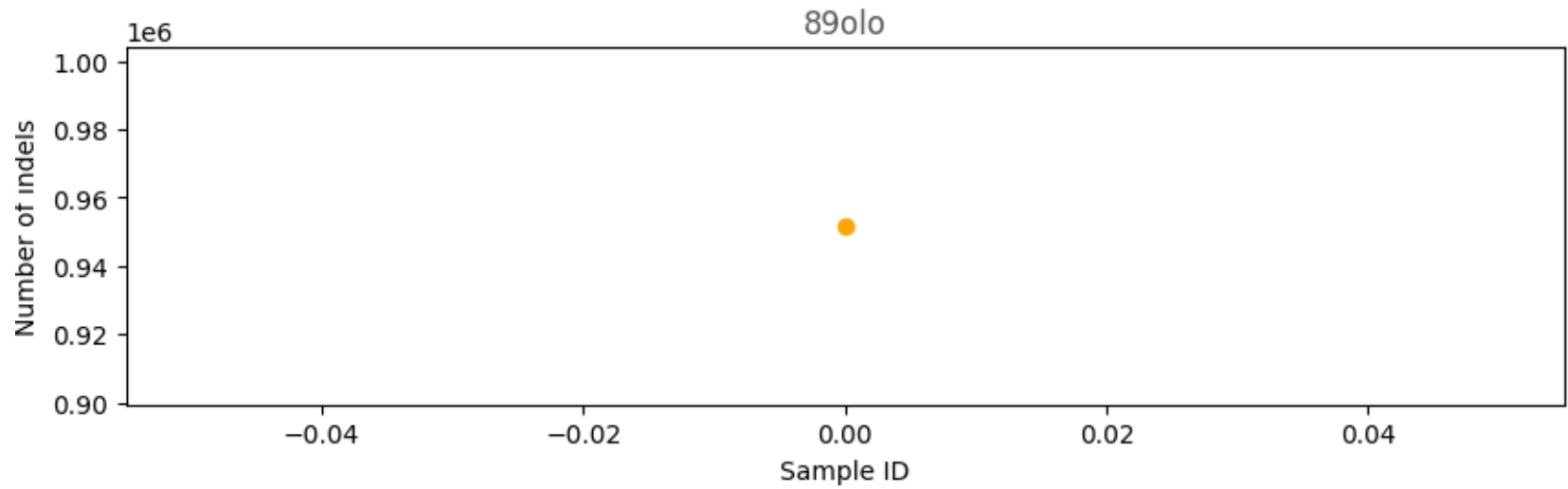
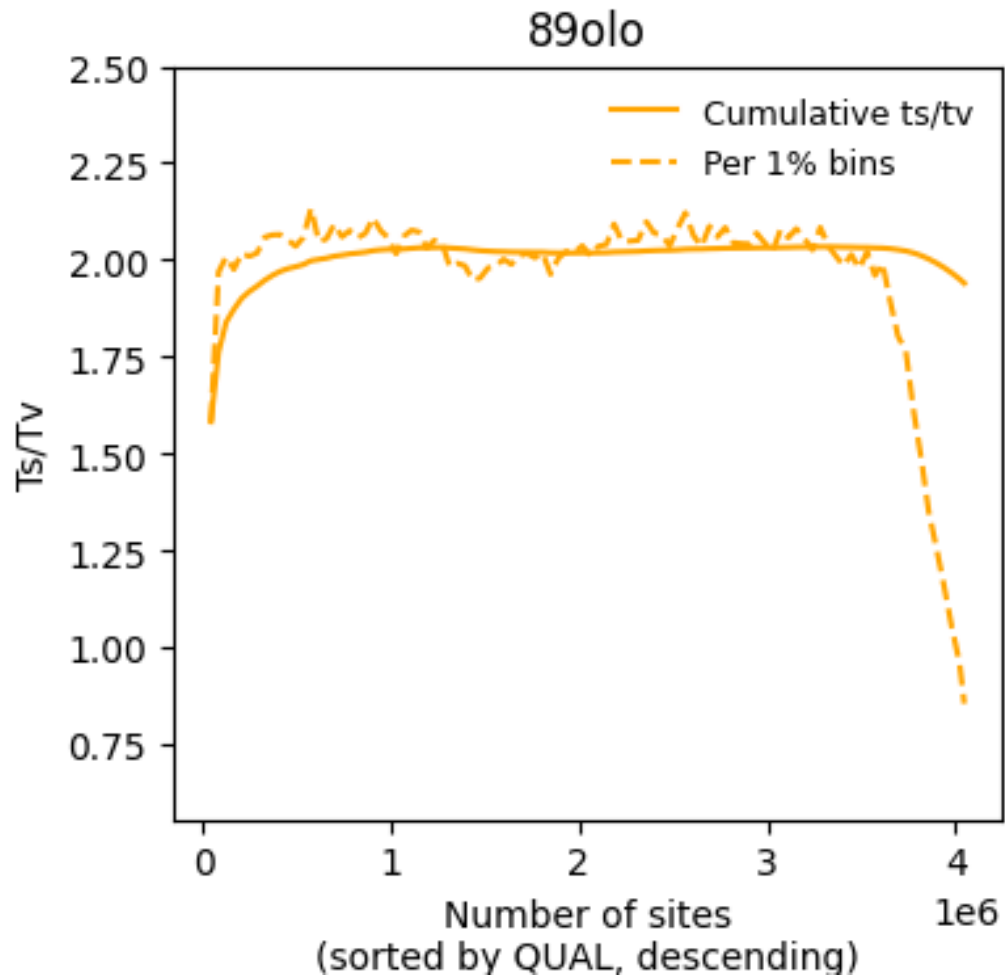# Singletons by sample (hets and homs)

Average depth by sample
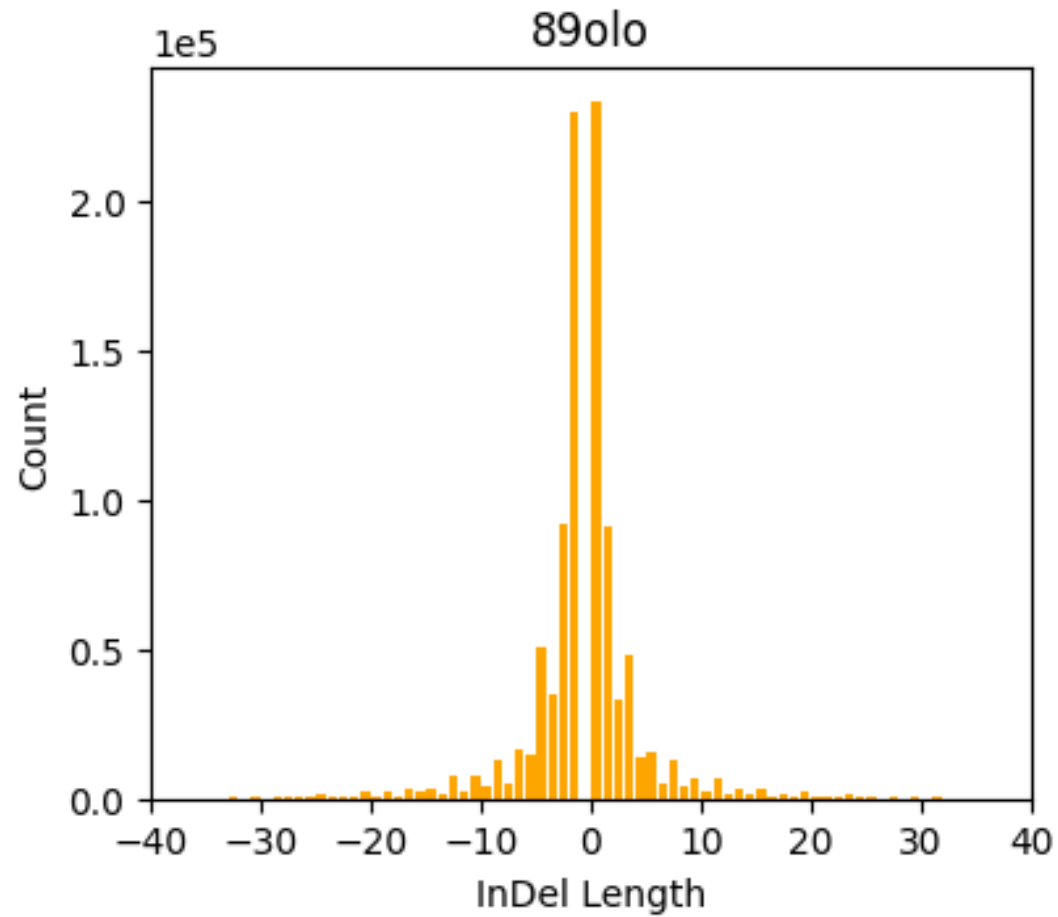
Number of SNPs by sample

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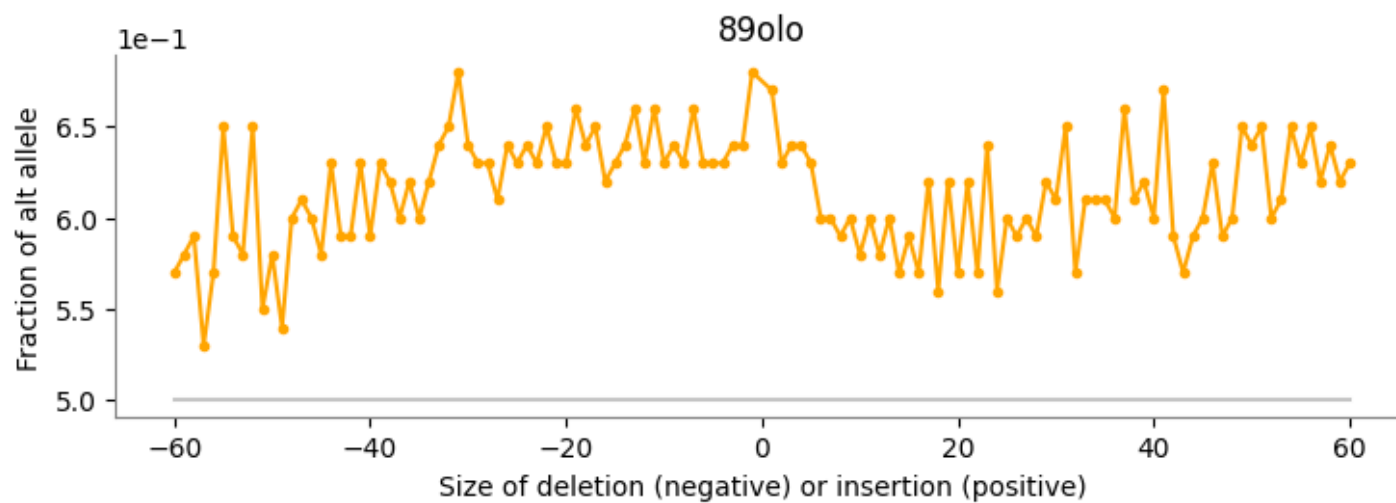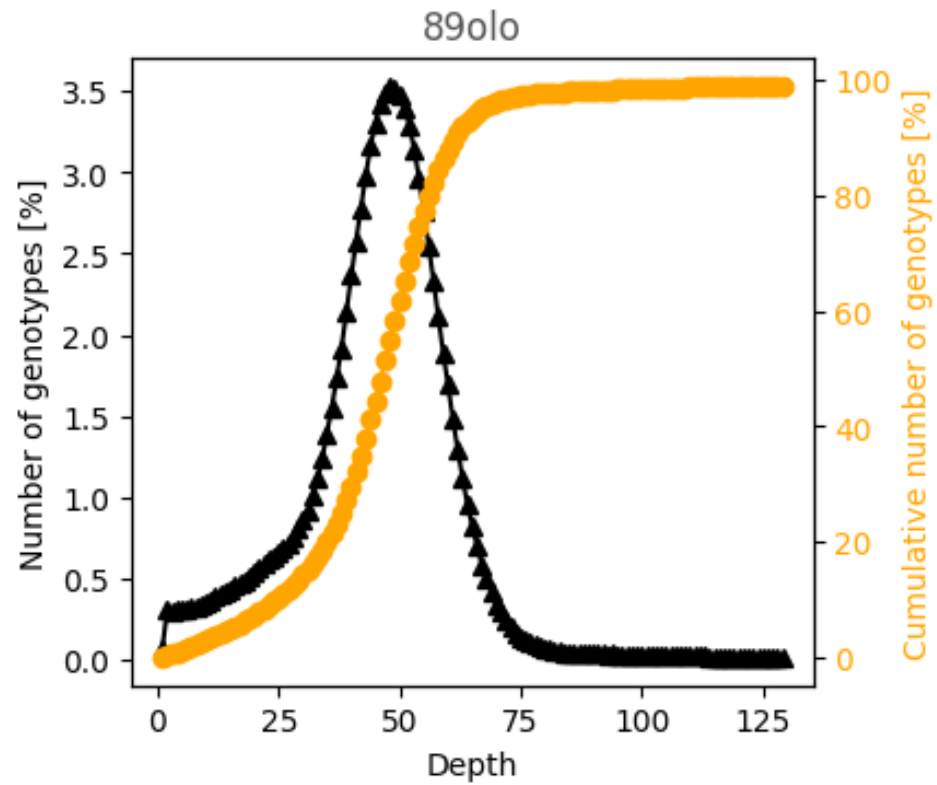
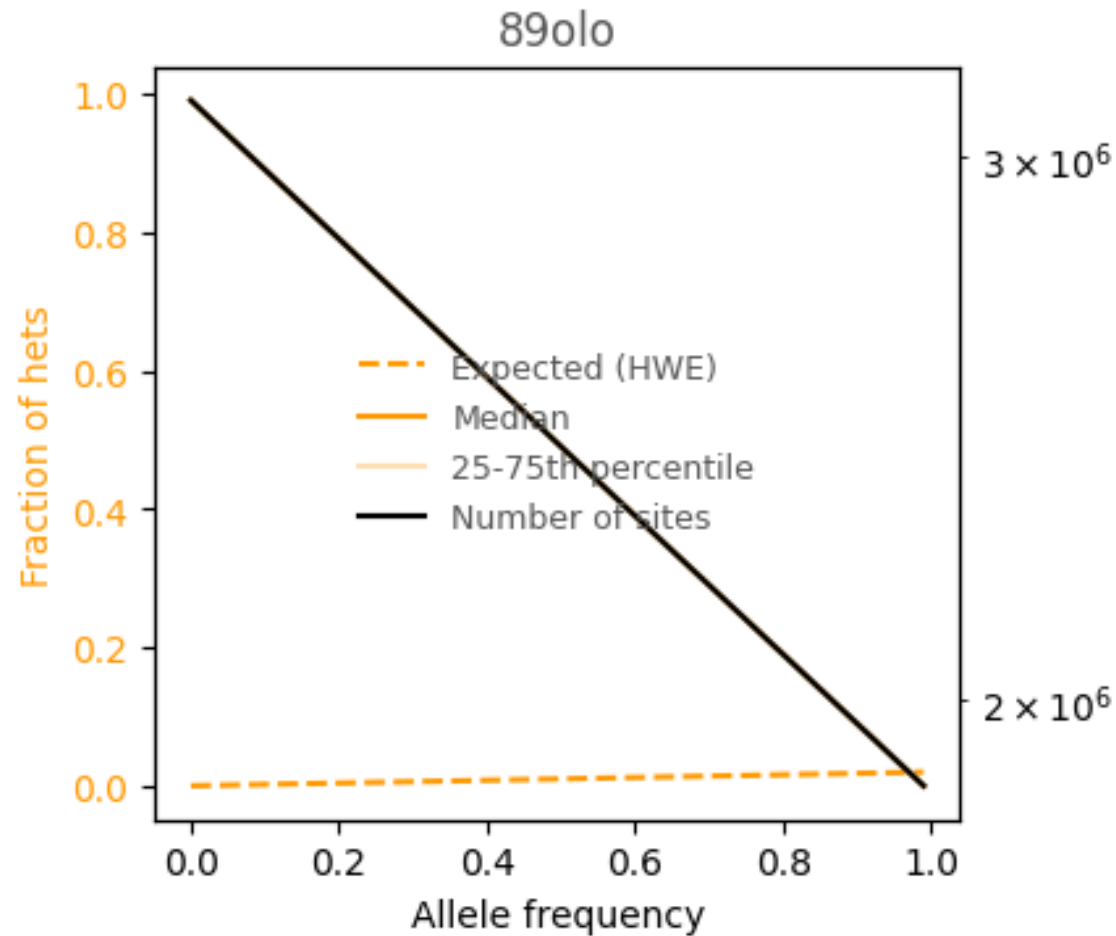# Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types