Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
89jos	4,062,794	1.93	1.93	934,585	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
89jos	62.0%	1.90	66.0%	91,489	2,011

- 89jos .. `/ngc/projects2/gm/data/archive/2022/variants/snv/89josirem-103814631620-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211215_A01466_BH2GHYDSX3-RHGM_LABKA_WGS- WGS03516_snv_germline_raw.haplotype_caller.vcf.gz`
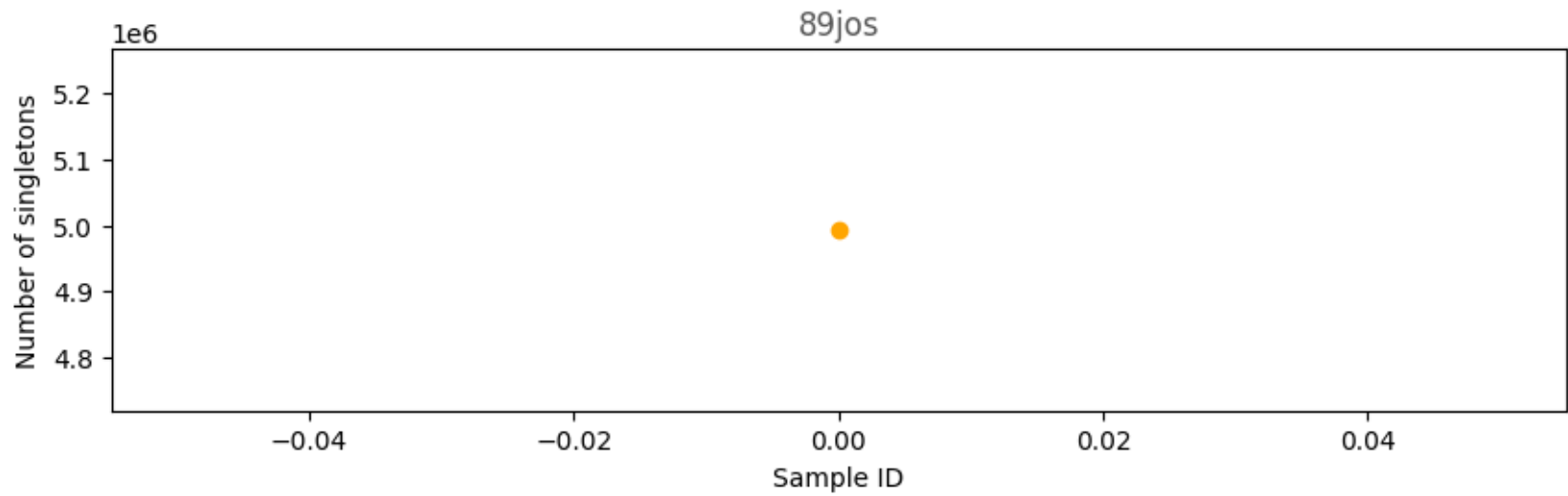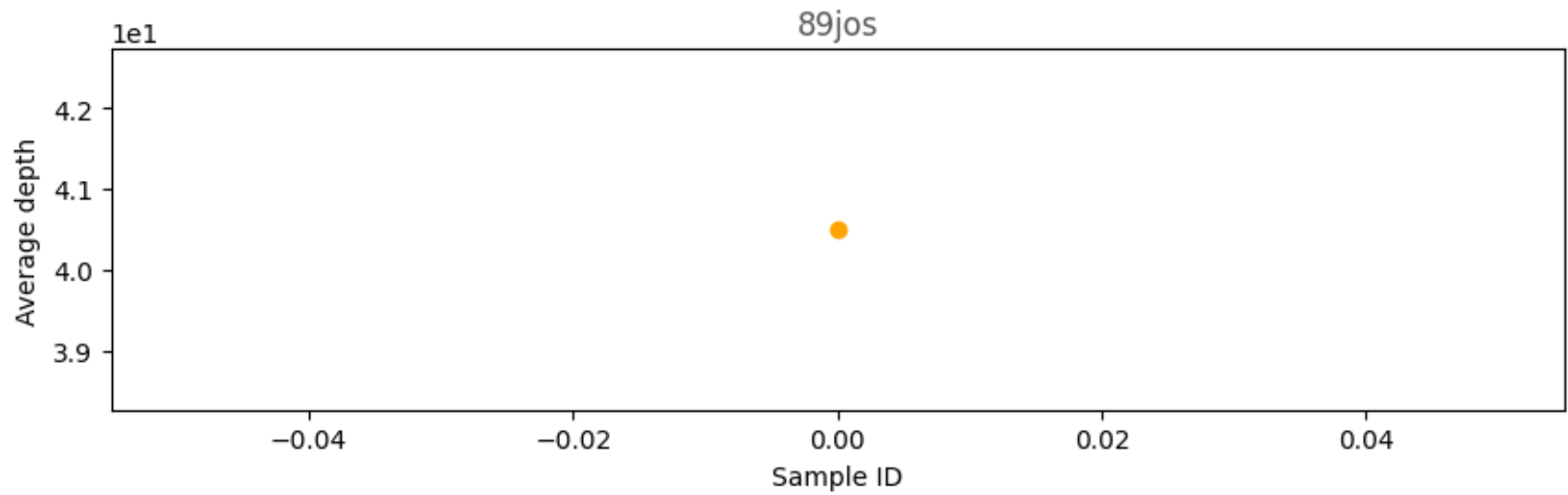
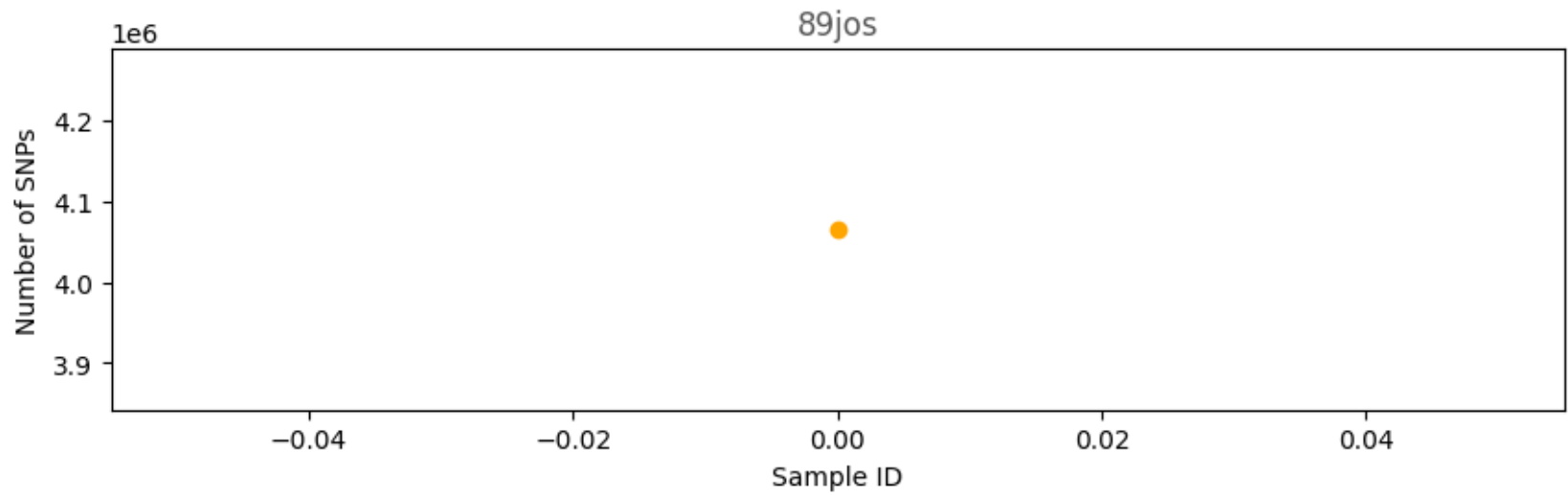# Ts/Tv by sample
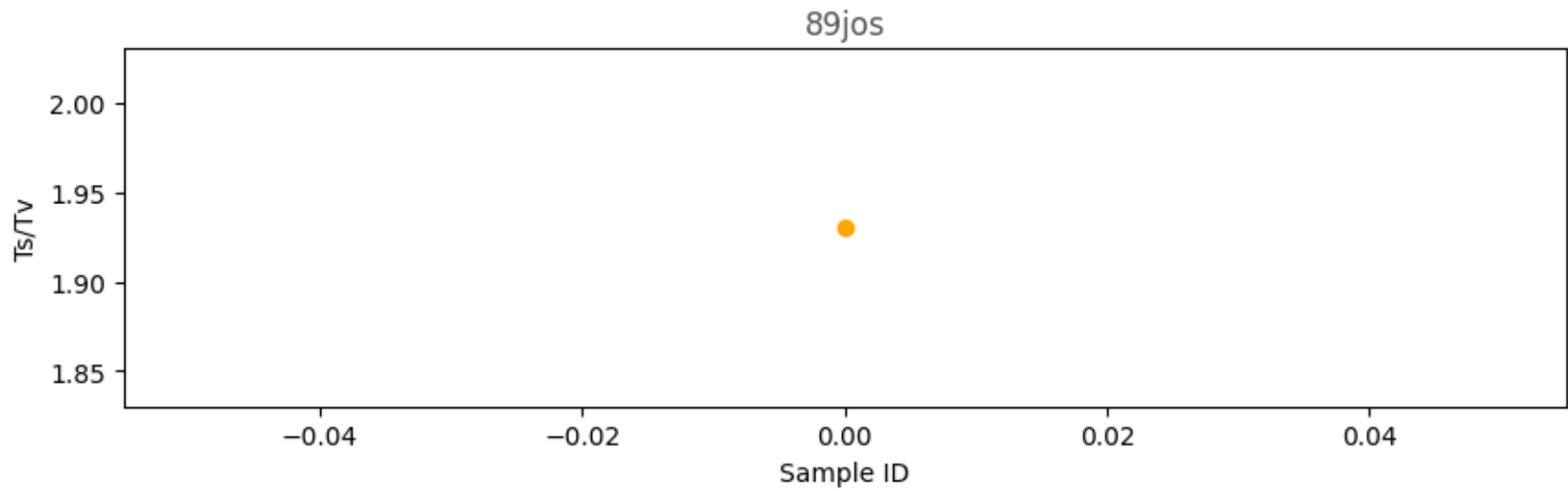
Hets vs non-ref Homs by sample
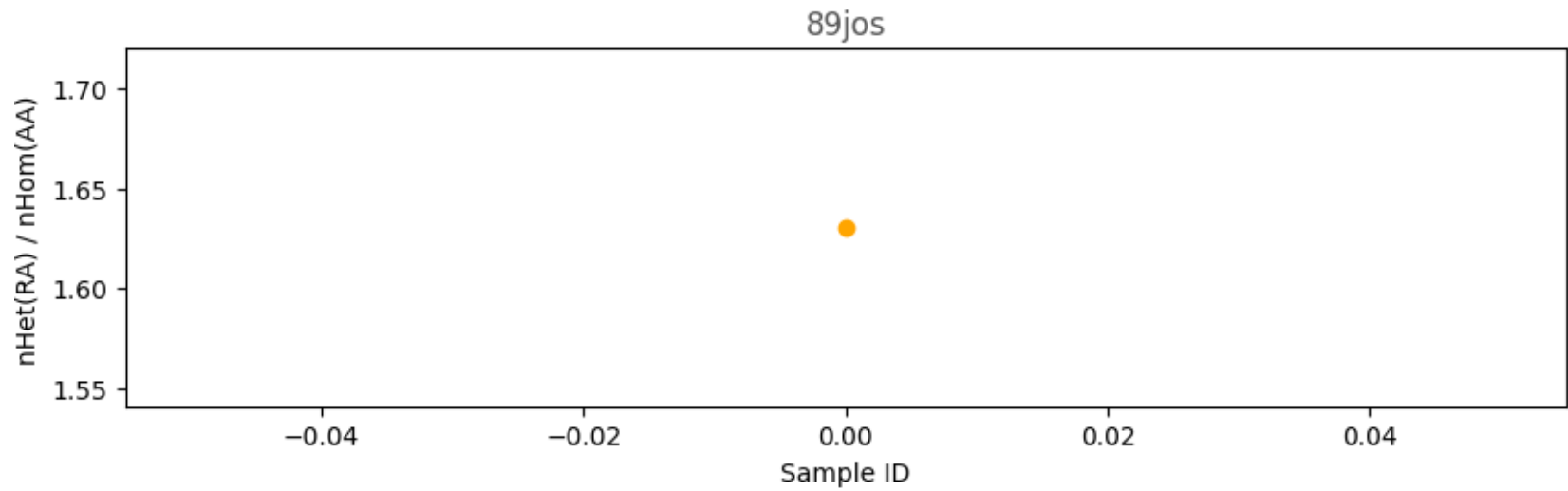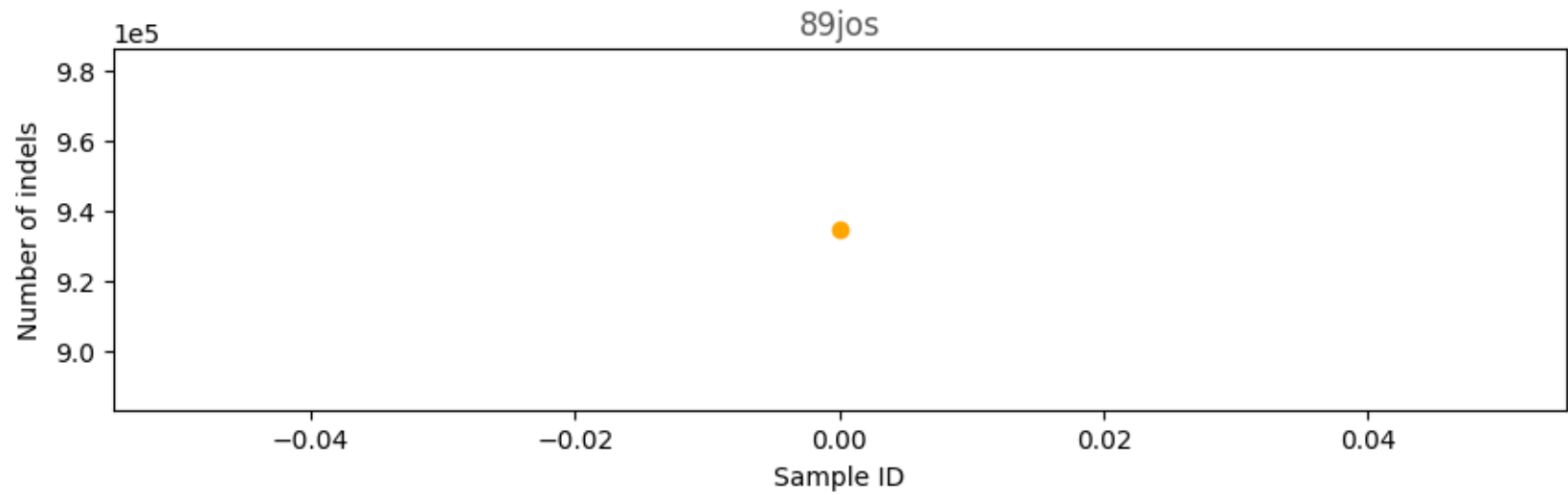
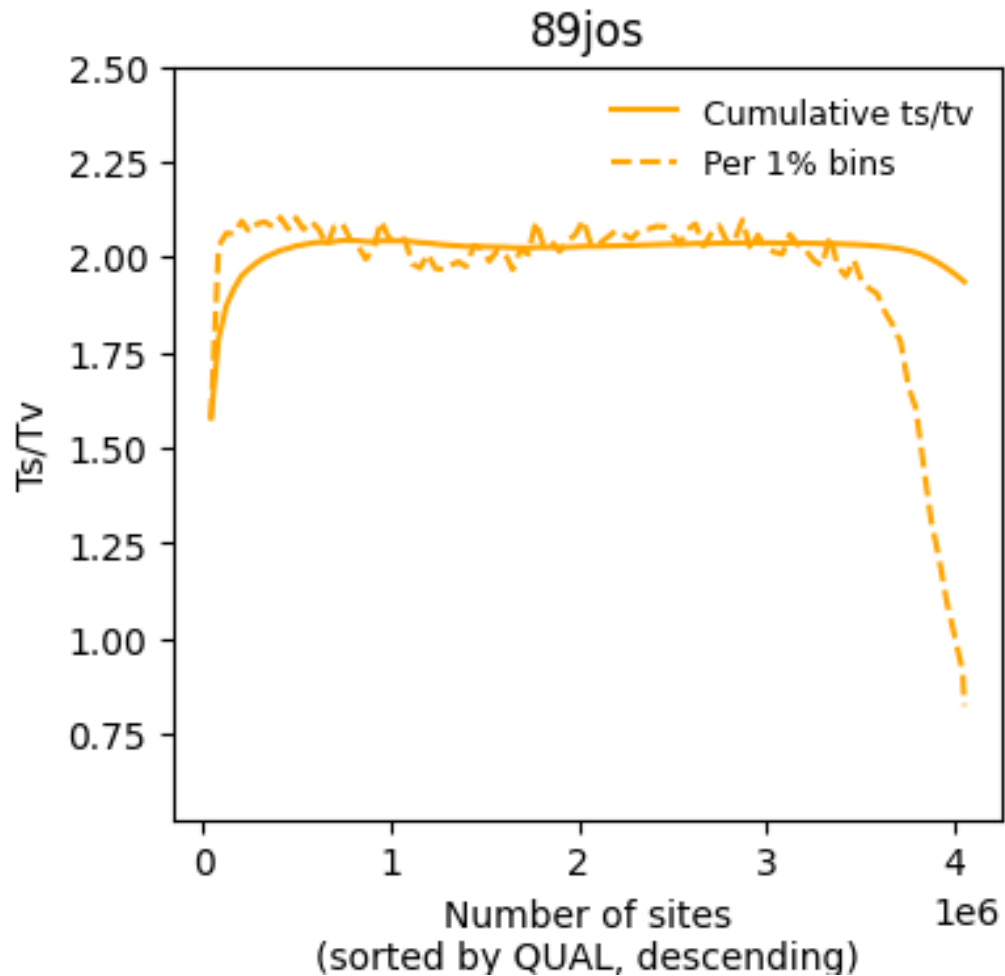# Singletons by sample (hets and homs)
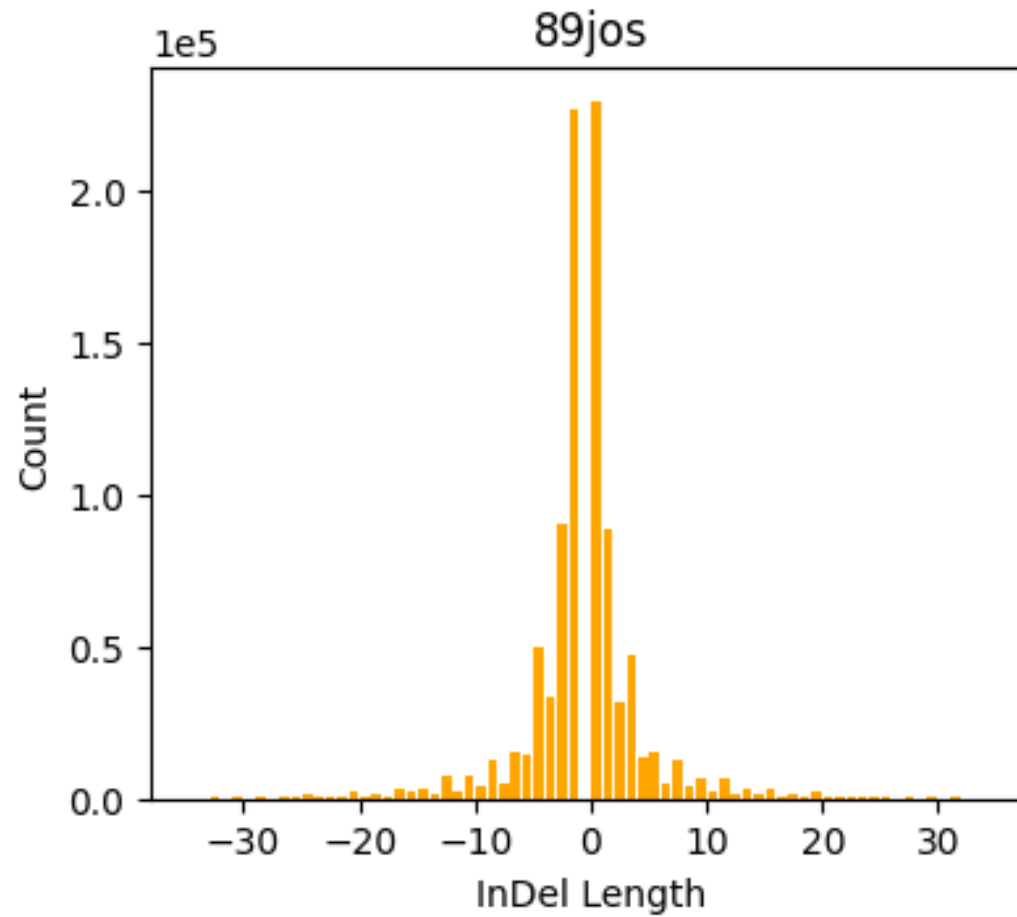
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

89jos

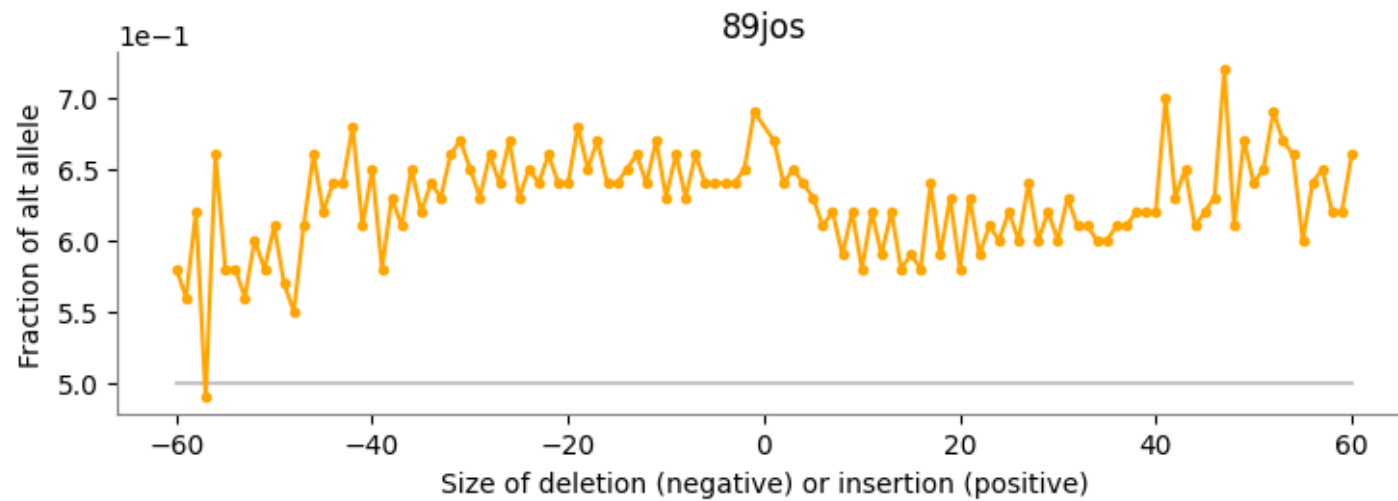# Indel distribution

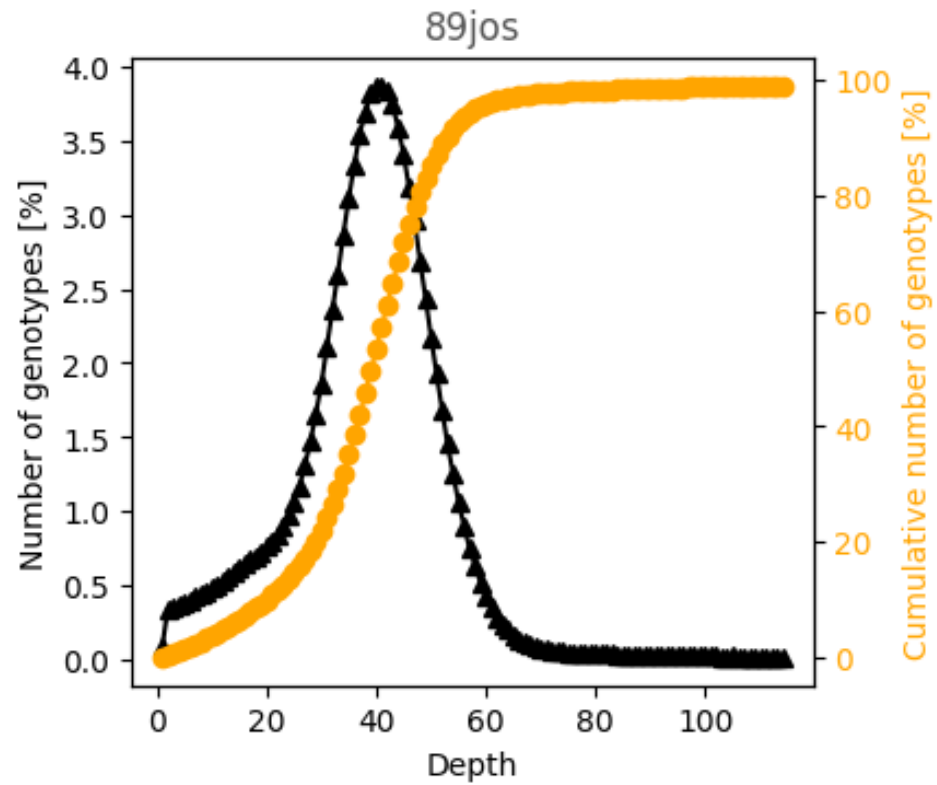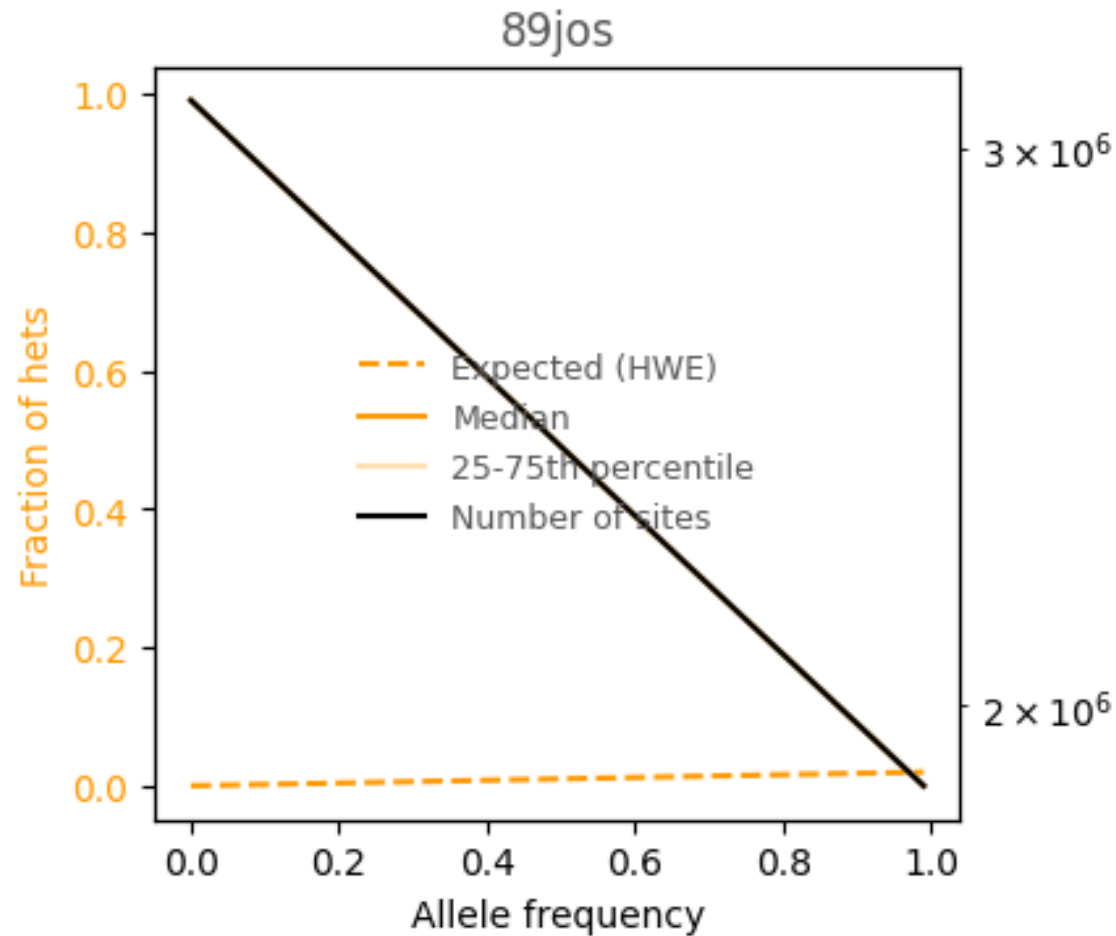# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types