Summary Numbers

| Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90rar	4,074,632	1.93	1.93	950,970	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90rar	61.7%	1.90	66.6%	100,204	2,047

- 90rar .. `/ngc/projects2/gm/data/archive/2022/variants/snv/90rarnhem-110297313740-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-221014_A00559_BH3FK3DSX5-RHGM_LABKA_WGSA` `KUT-WGSAKUT05830_22RKG024520x01_snv_germline_raw.haplotype_caller.vcf.gz`
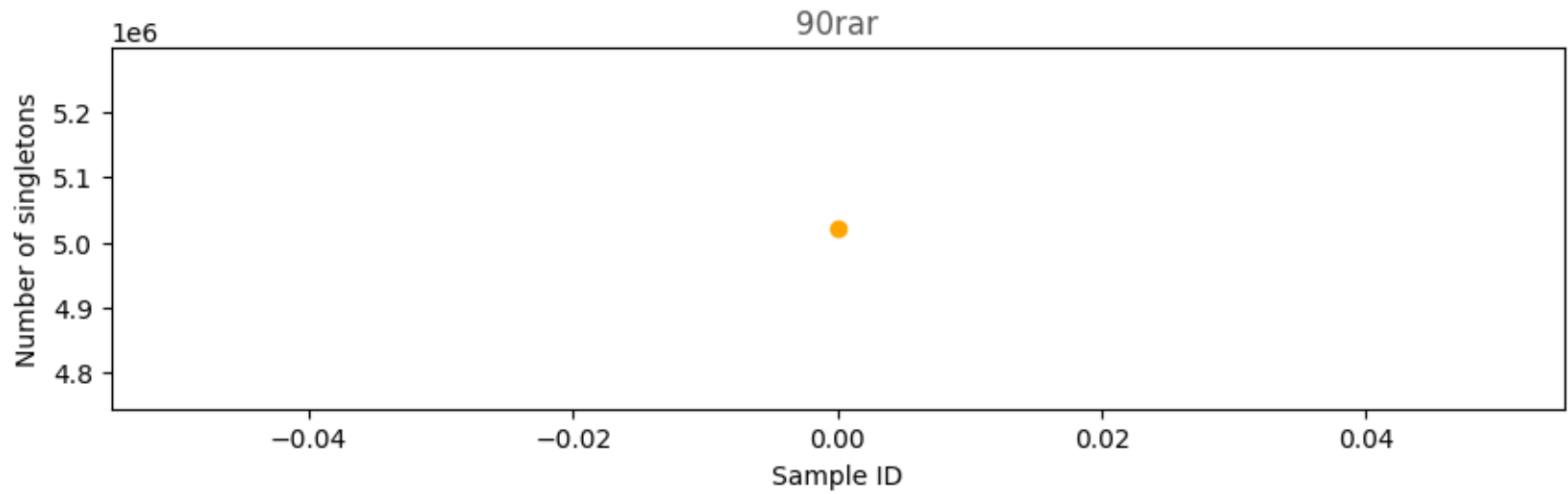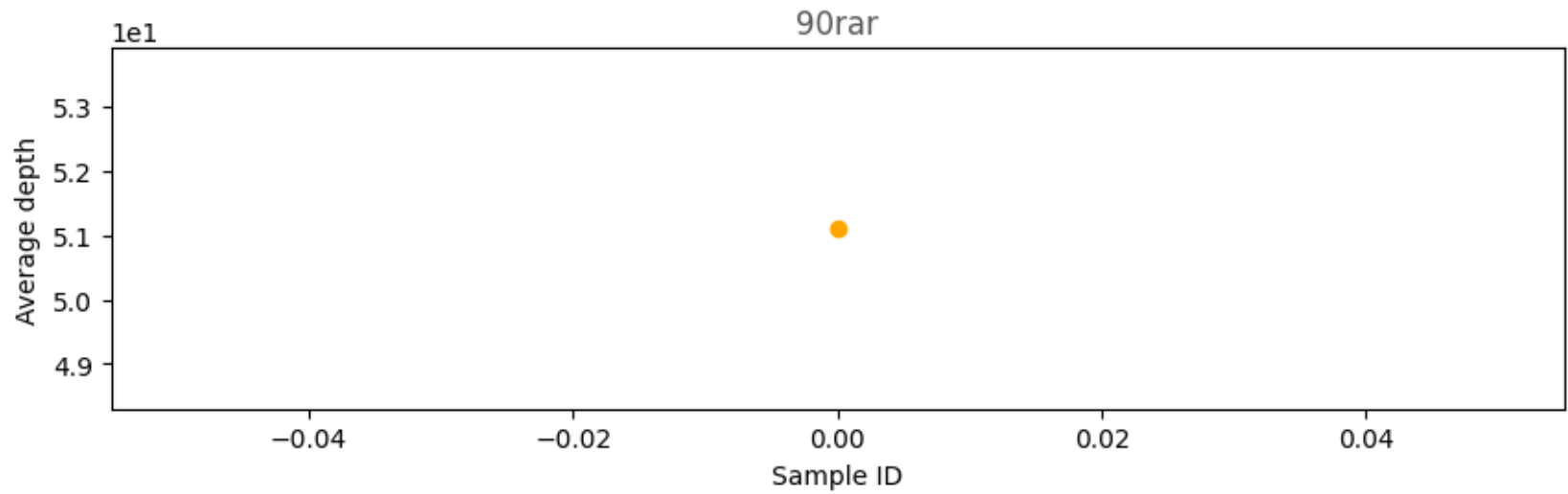
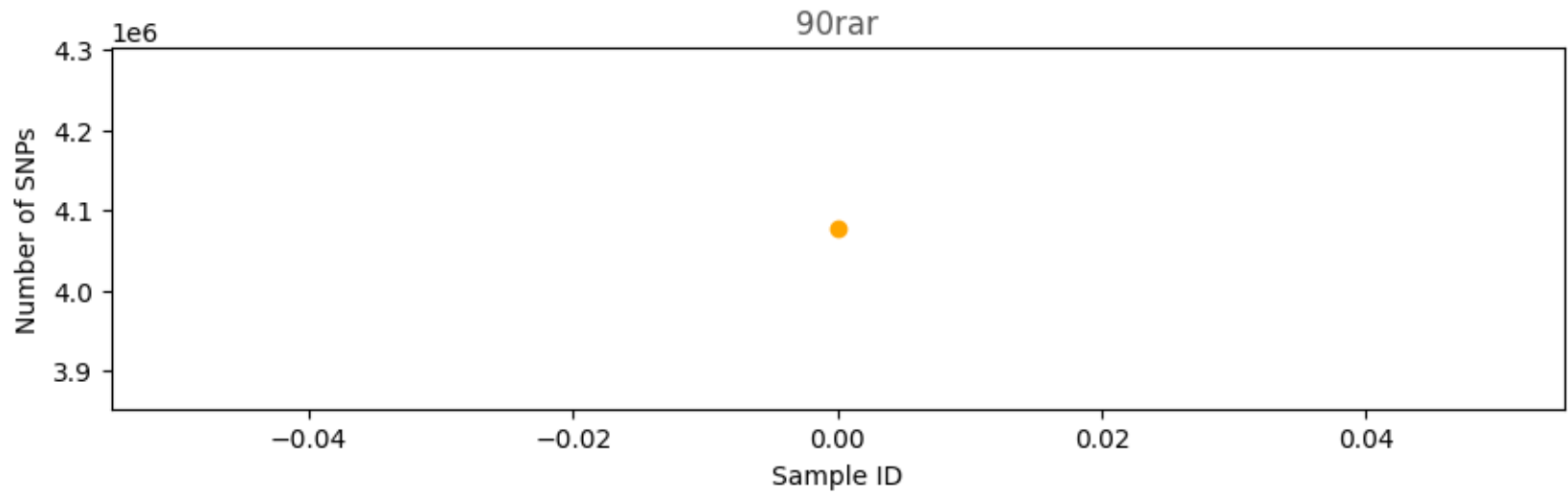# Ts/Tv by sample
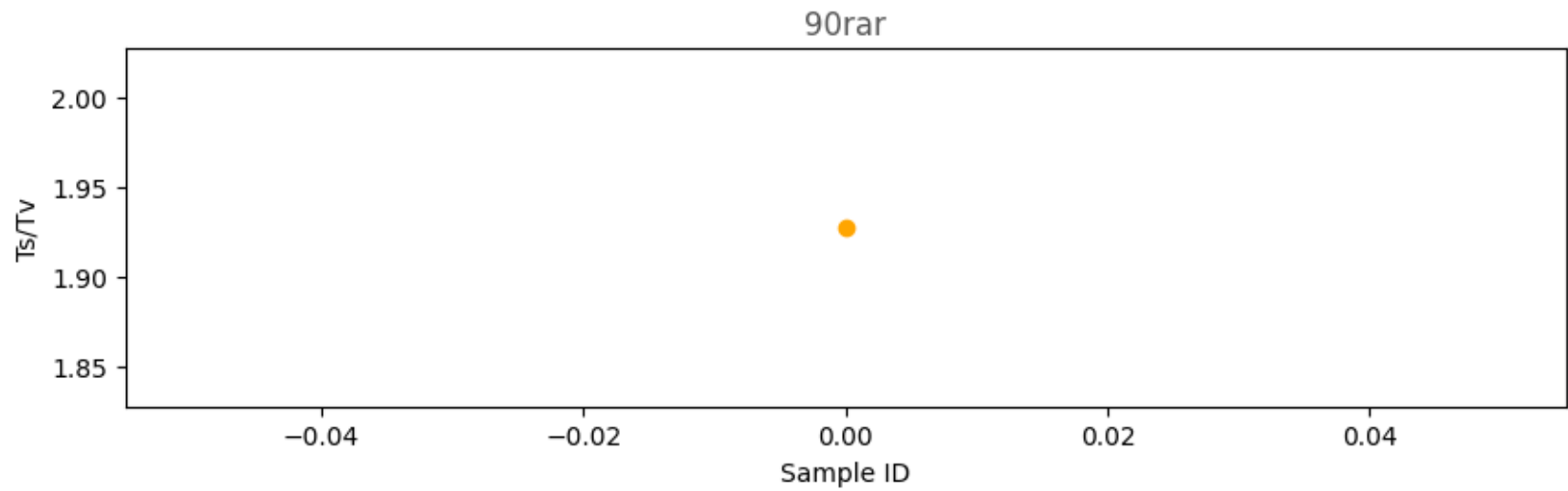
Hets vs non-ref Homs by sample
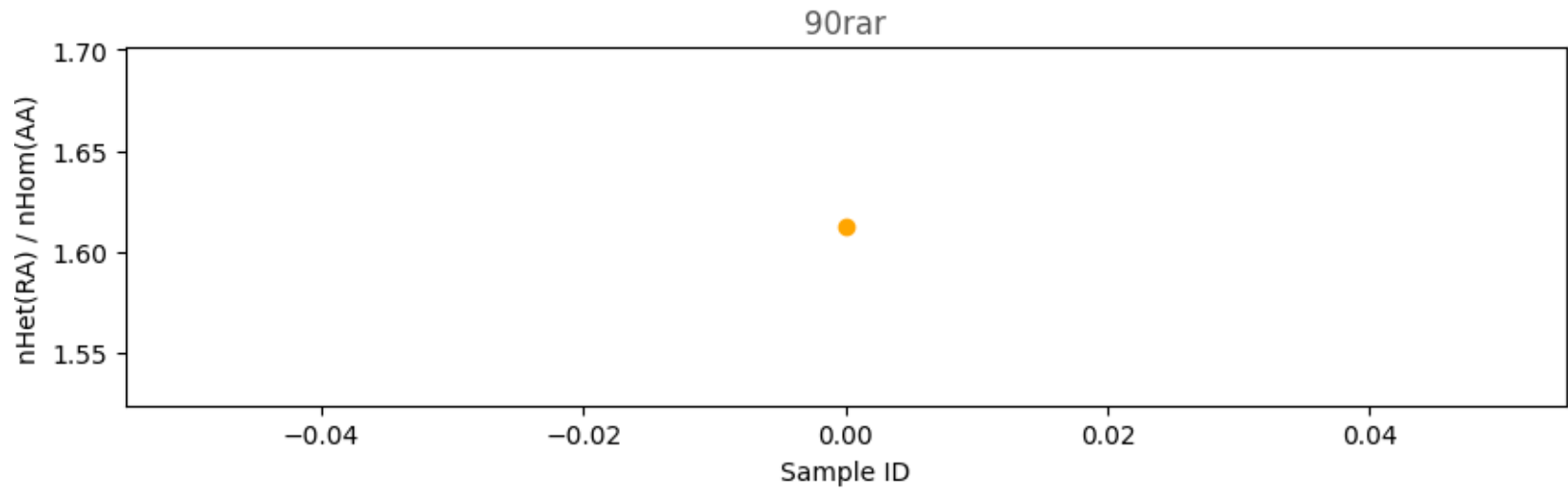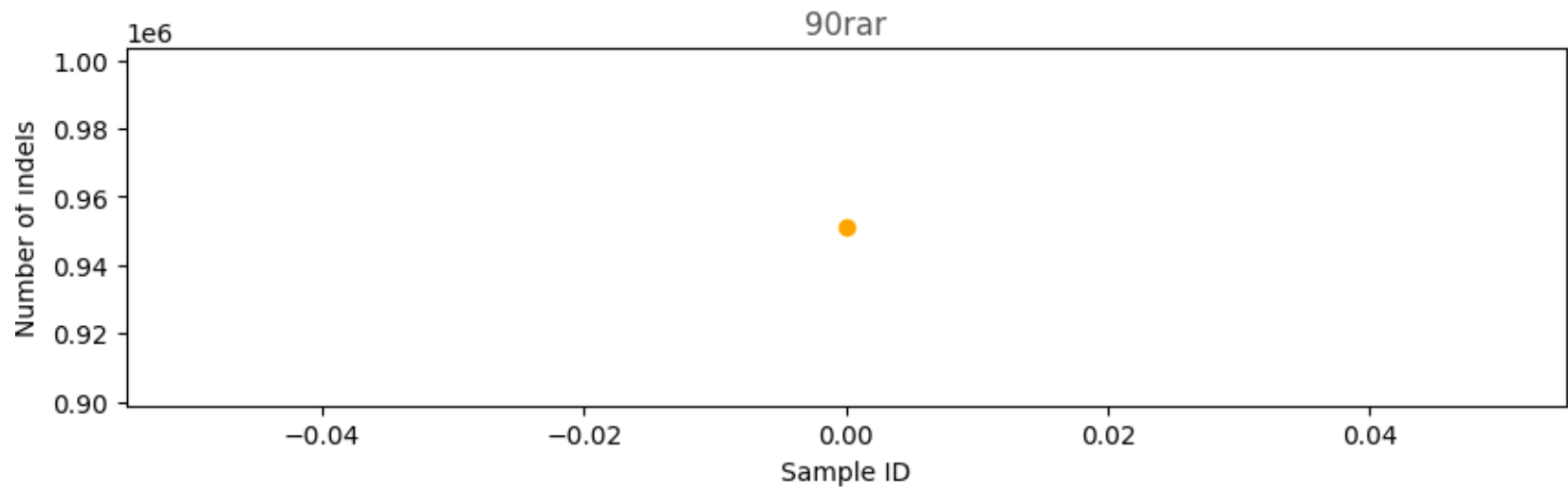
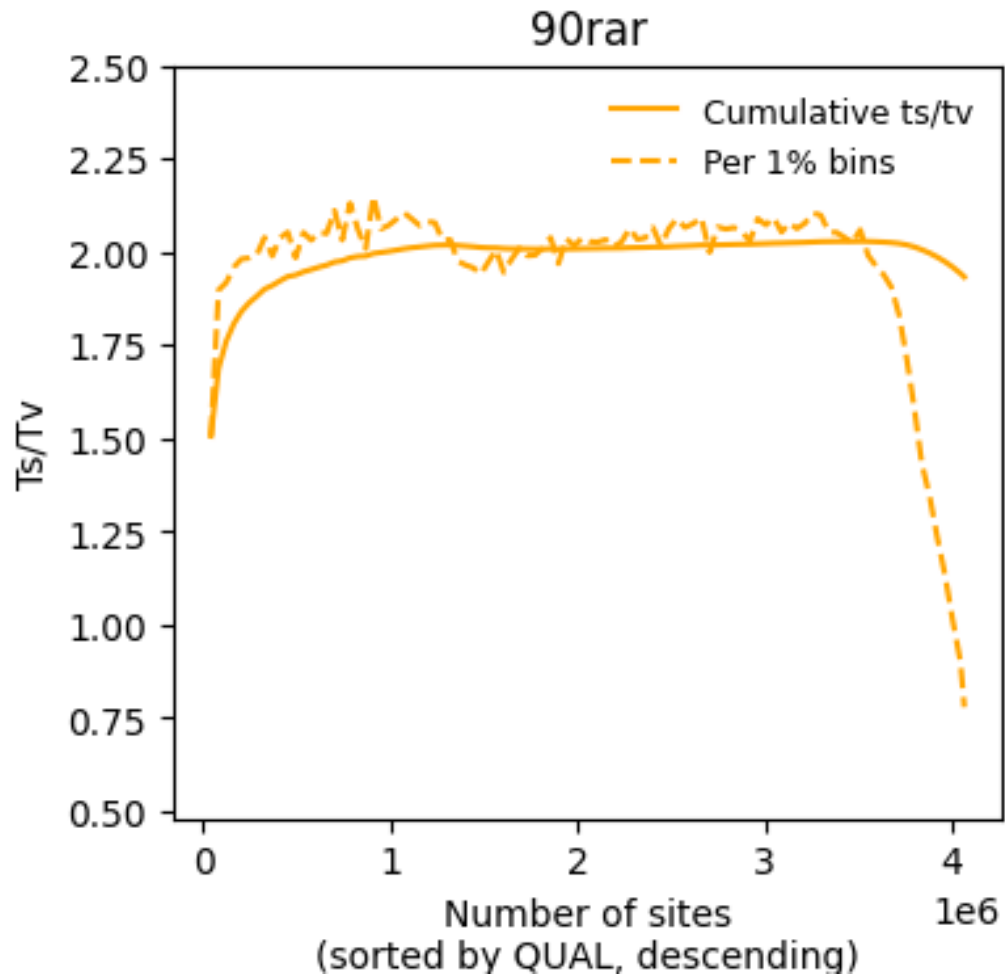# Singletons by sample (hets and homs)
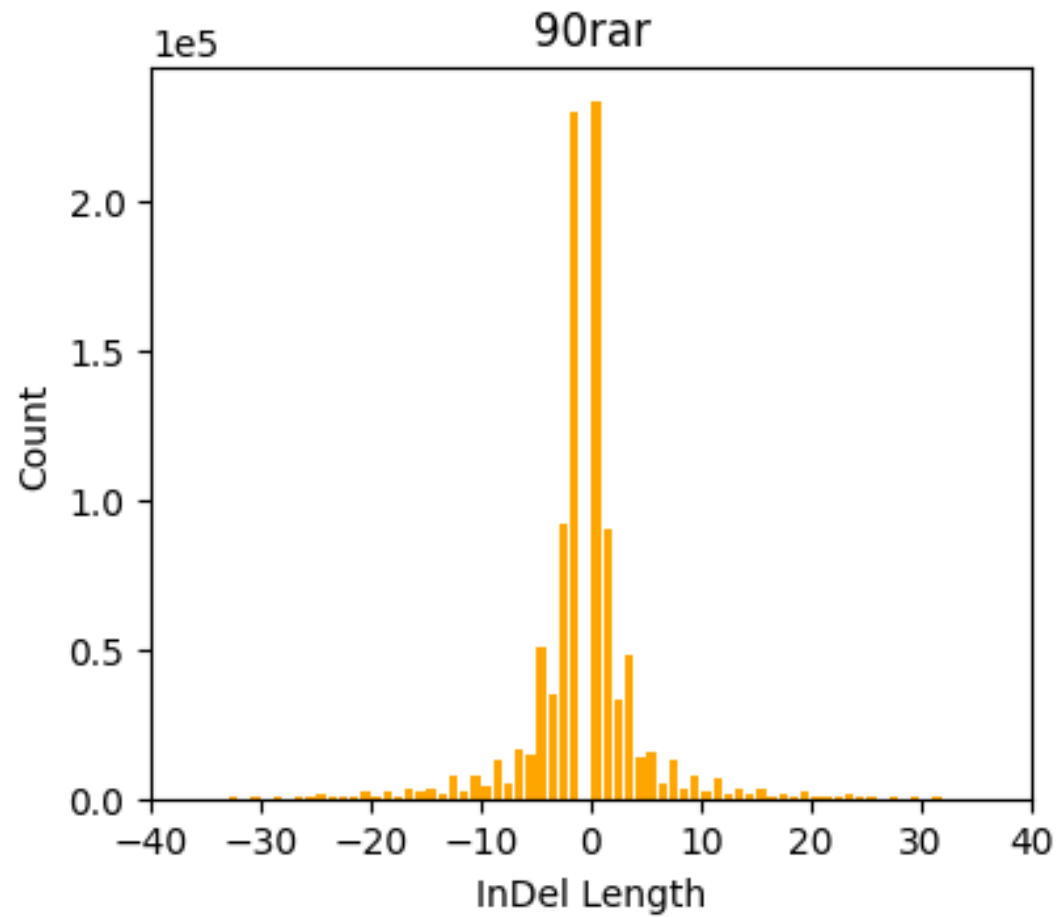
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

90rar

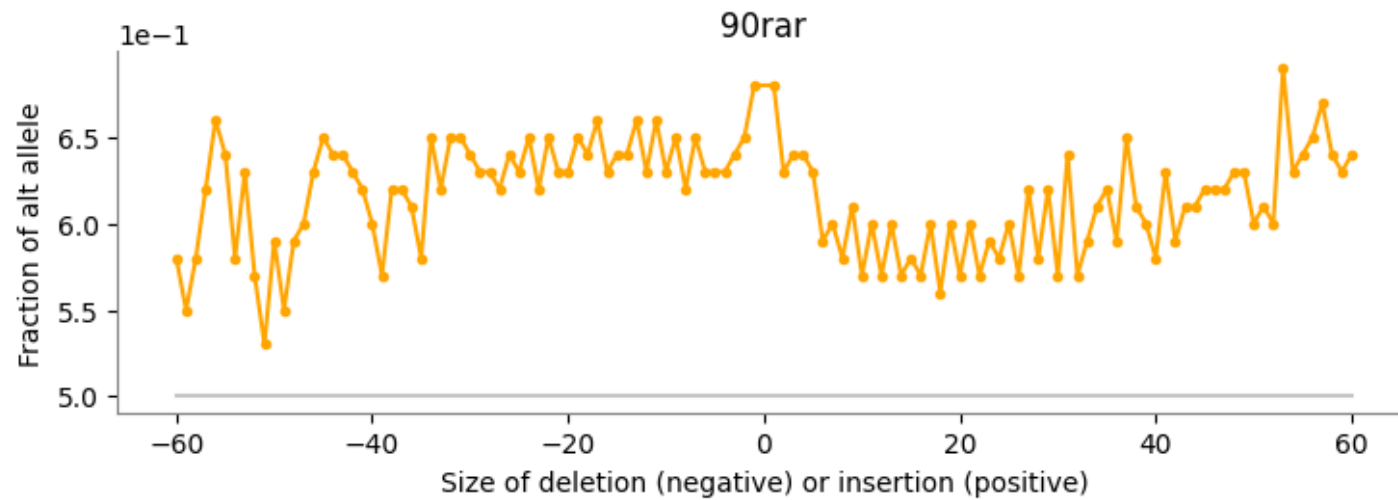# Indel distribution

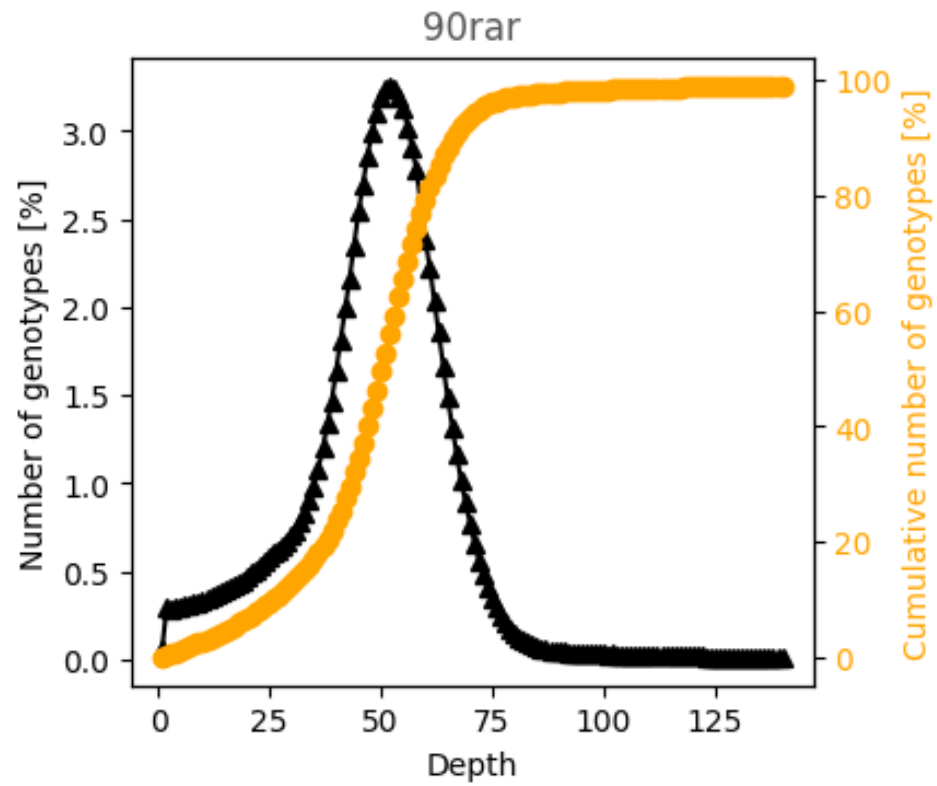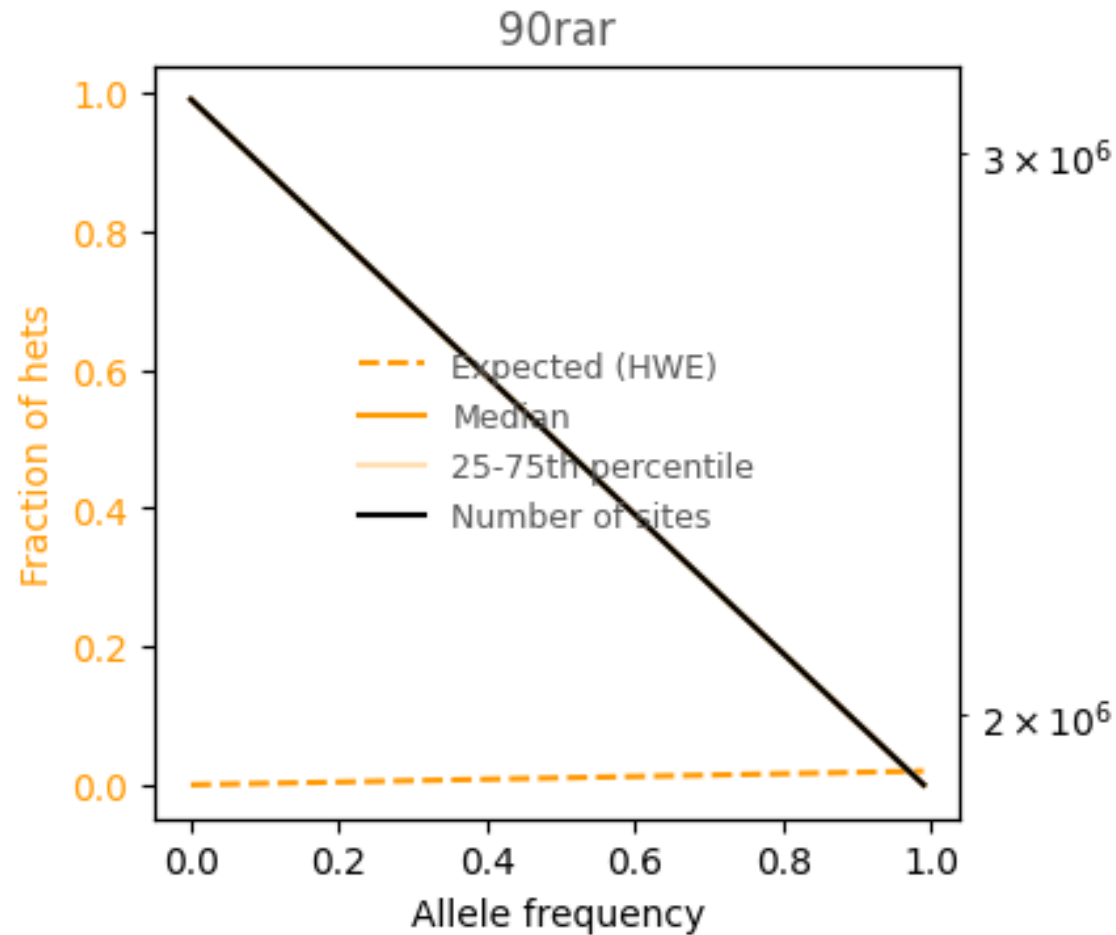# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types