Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
77rar	4,589,966	1.95	1.95	1,057,432	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
77rar	65.8%	1.93	70.8%	122,182	2,393

- 77rar .. `/ngc/projects2/gm/data/archive/2022/variants/snv/77raratim-103886312350-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01567-220916_A01411_BHYNLVDSX3-EXT_LAB KA_NGCWGS-NGCWGS05368_22RKG015825_snv_germline_raw.haplotype_caller.vcf.gz`
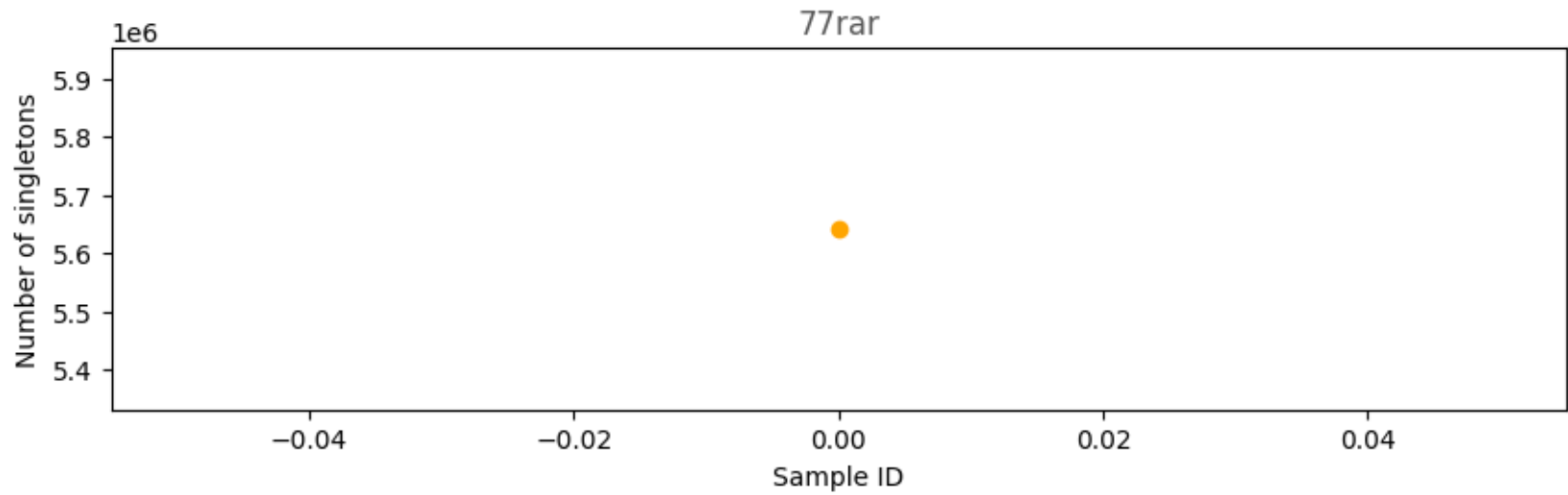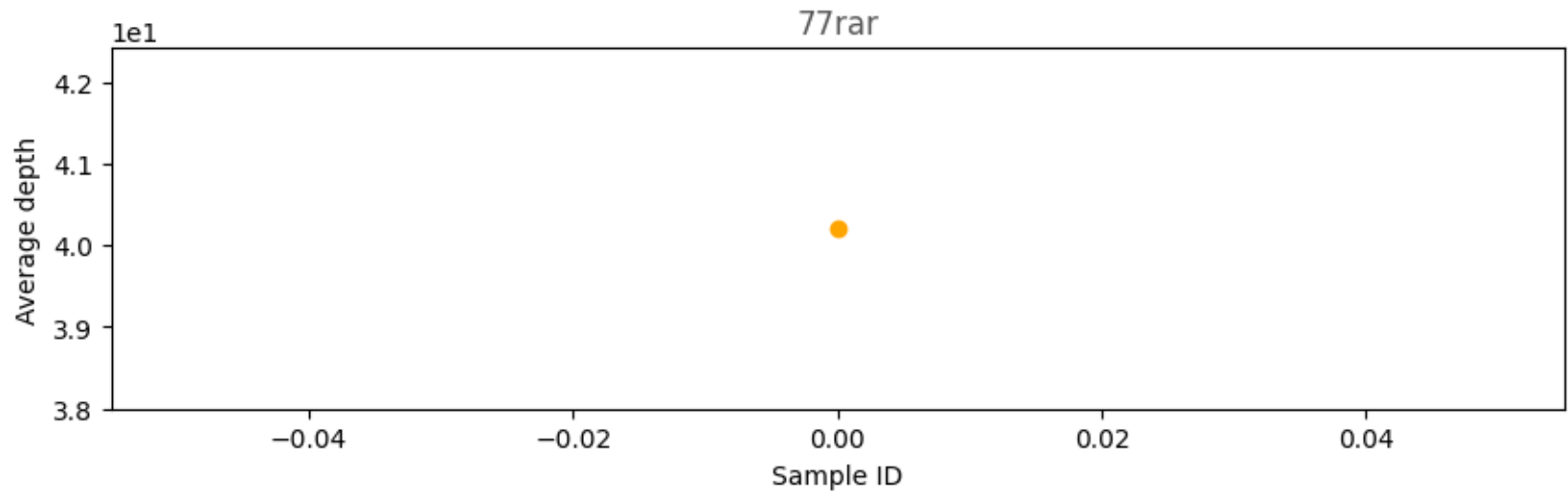
Ts/Tv by sample
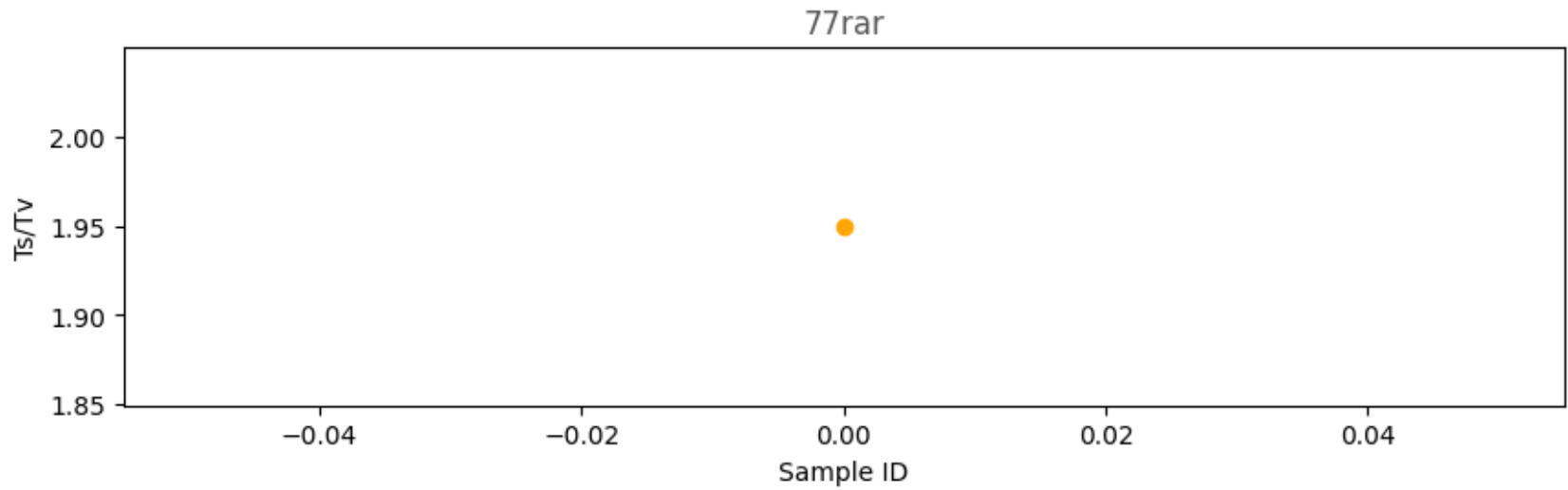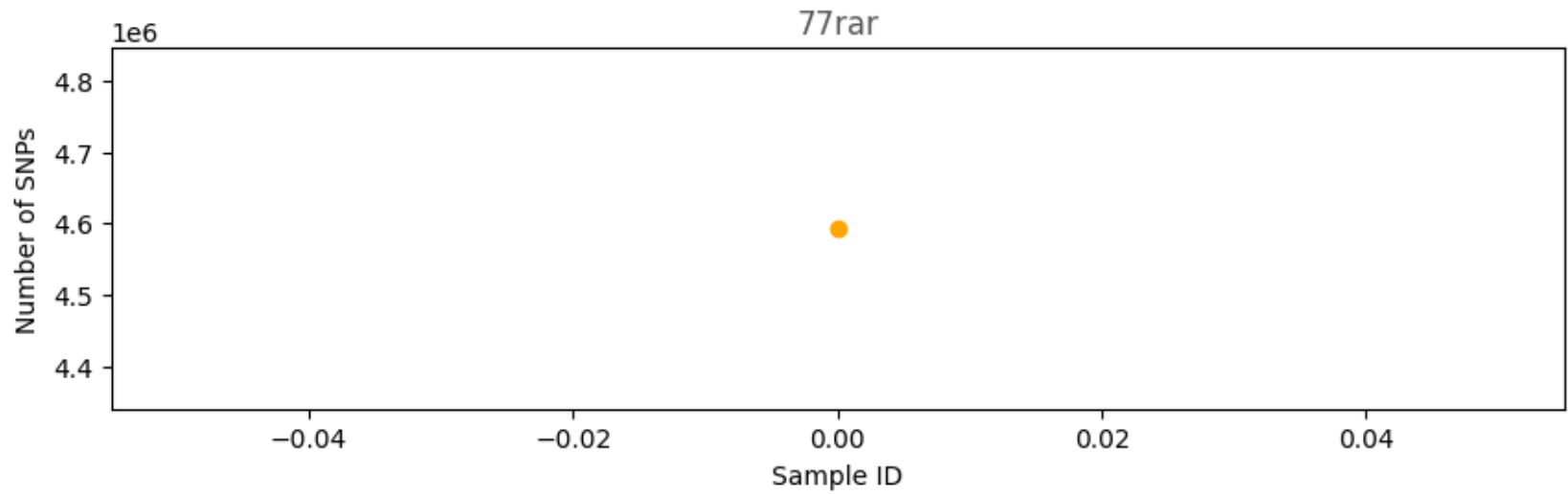
Hets vs non-ref Homs by sample
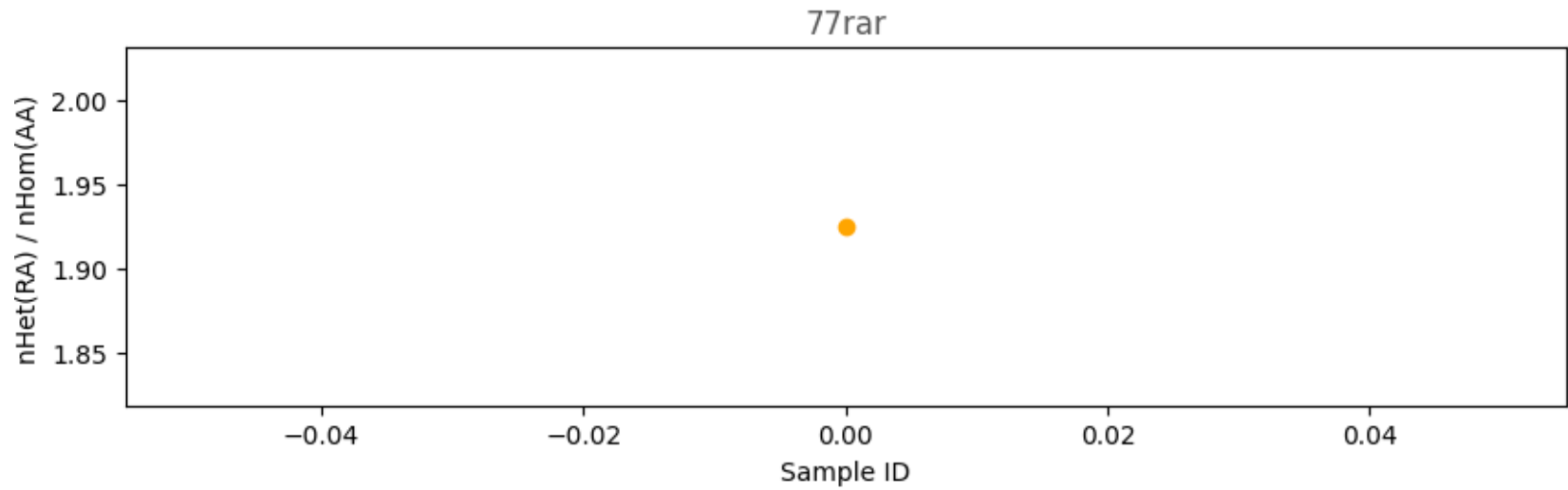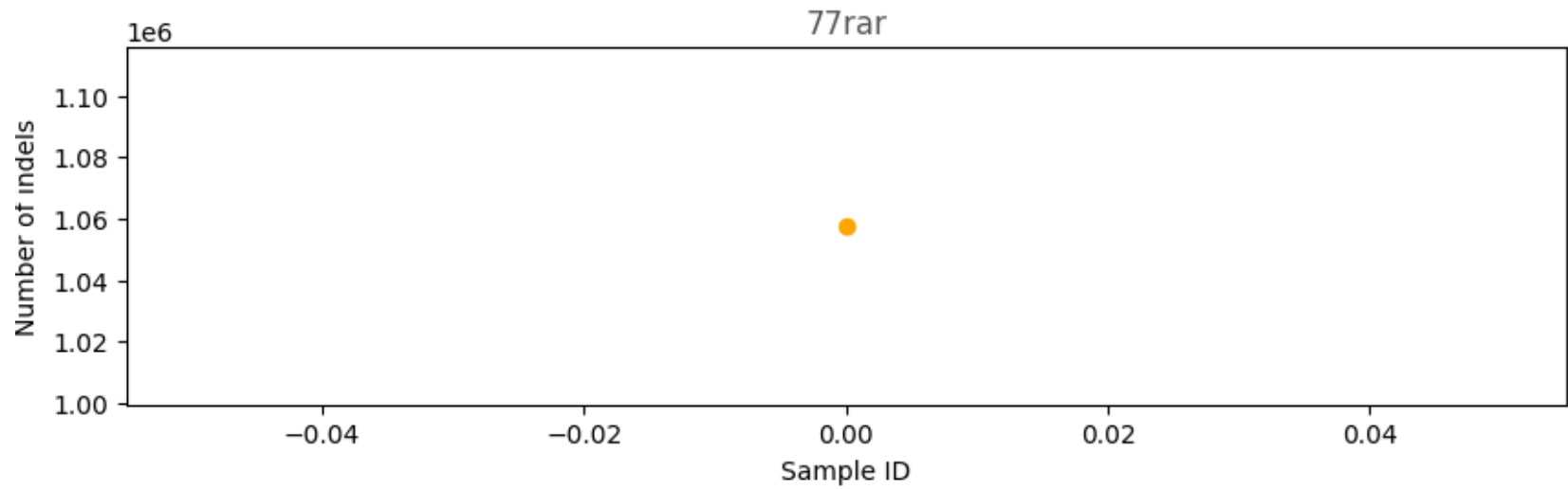
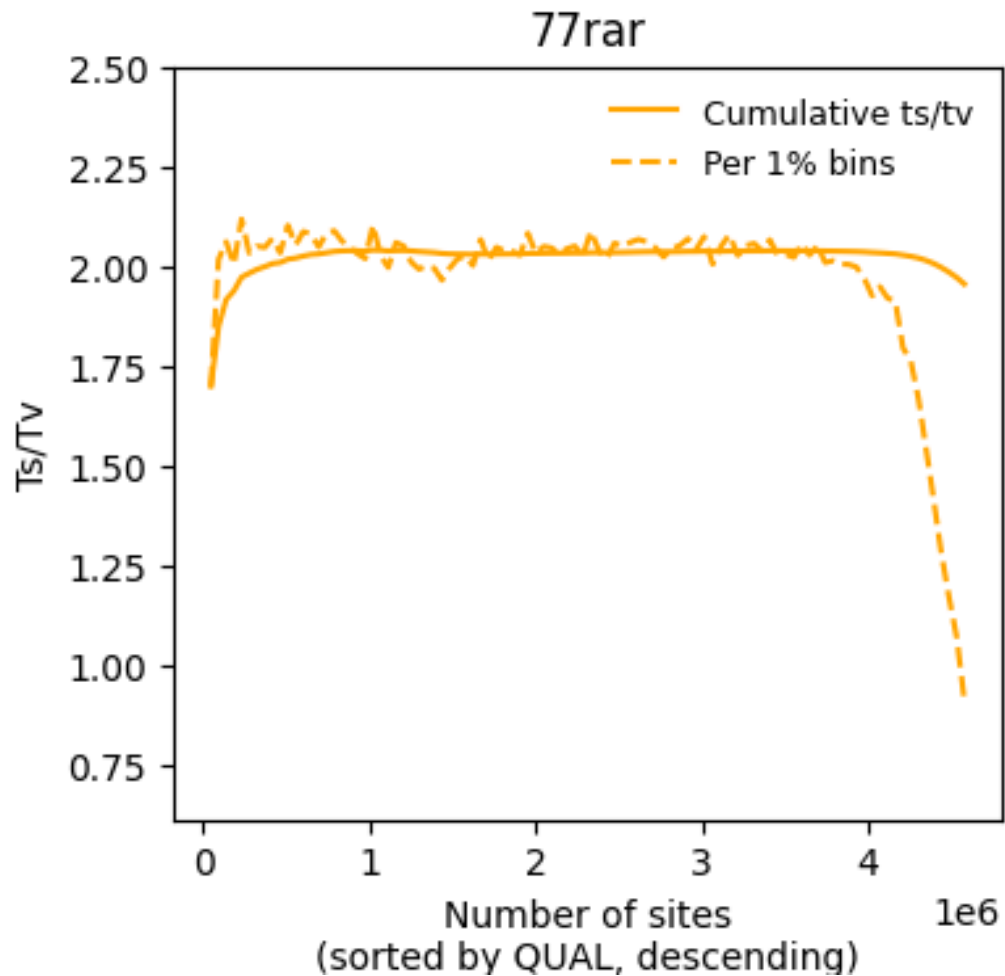# Singletons by sample (hets and homs)

Average depth by sample

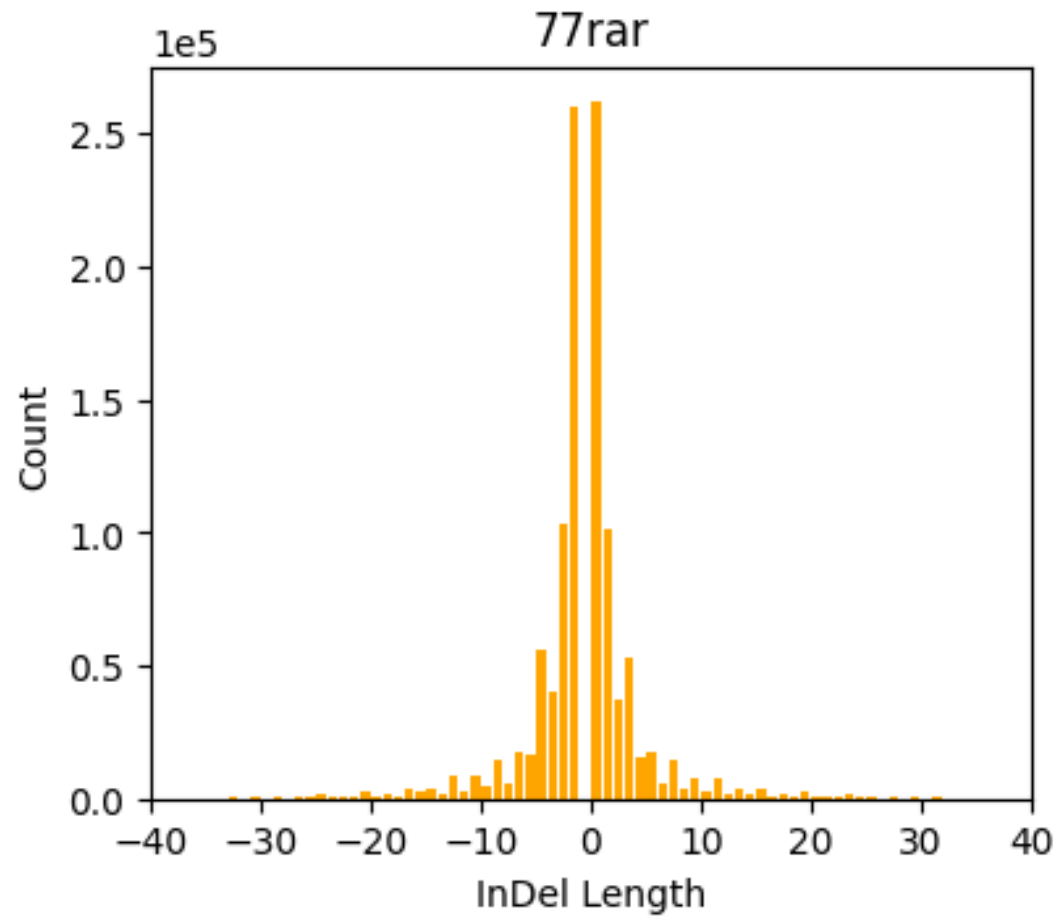# Number of SNPs by sample

Number of indels by sample
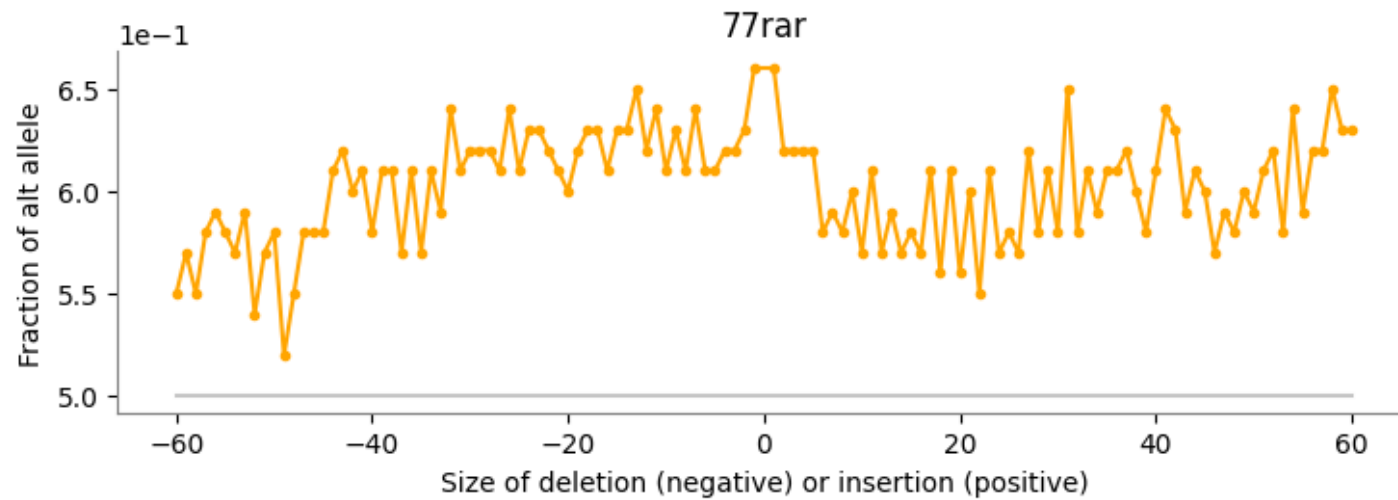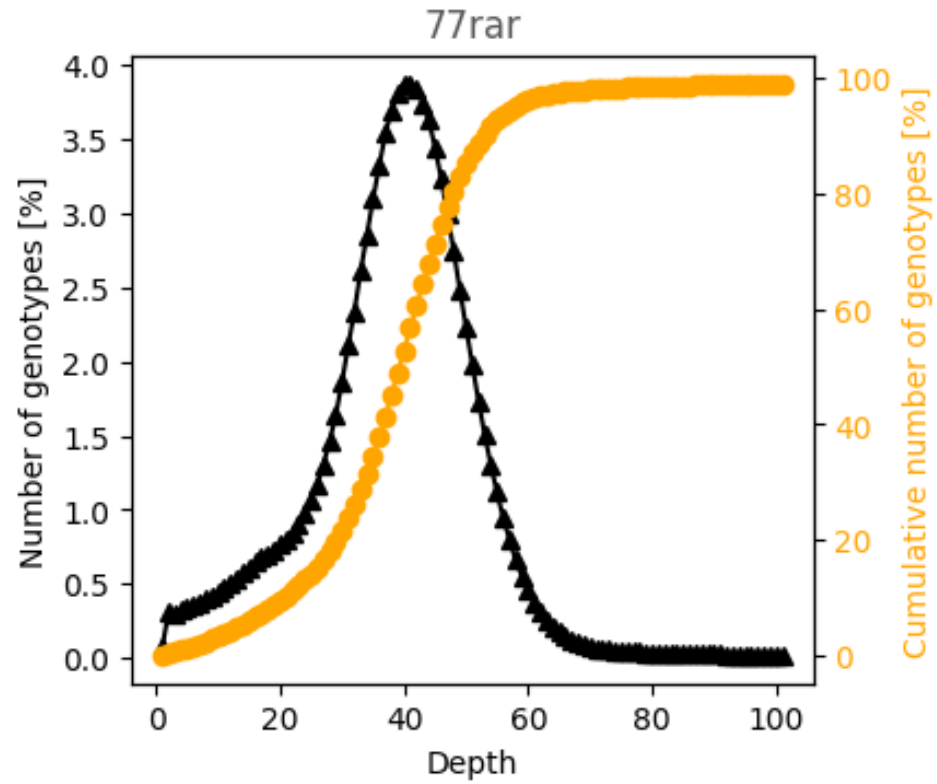
Ts/Tv stratified by QUAL

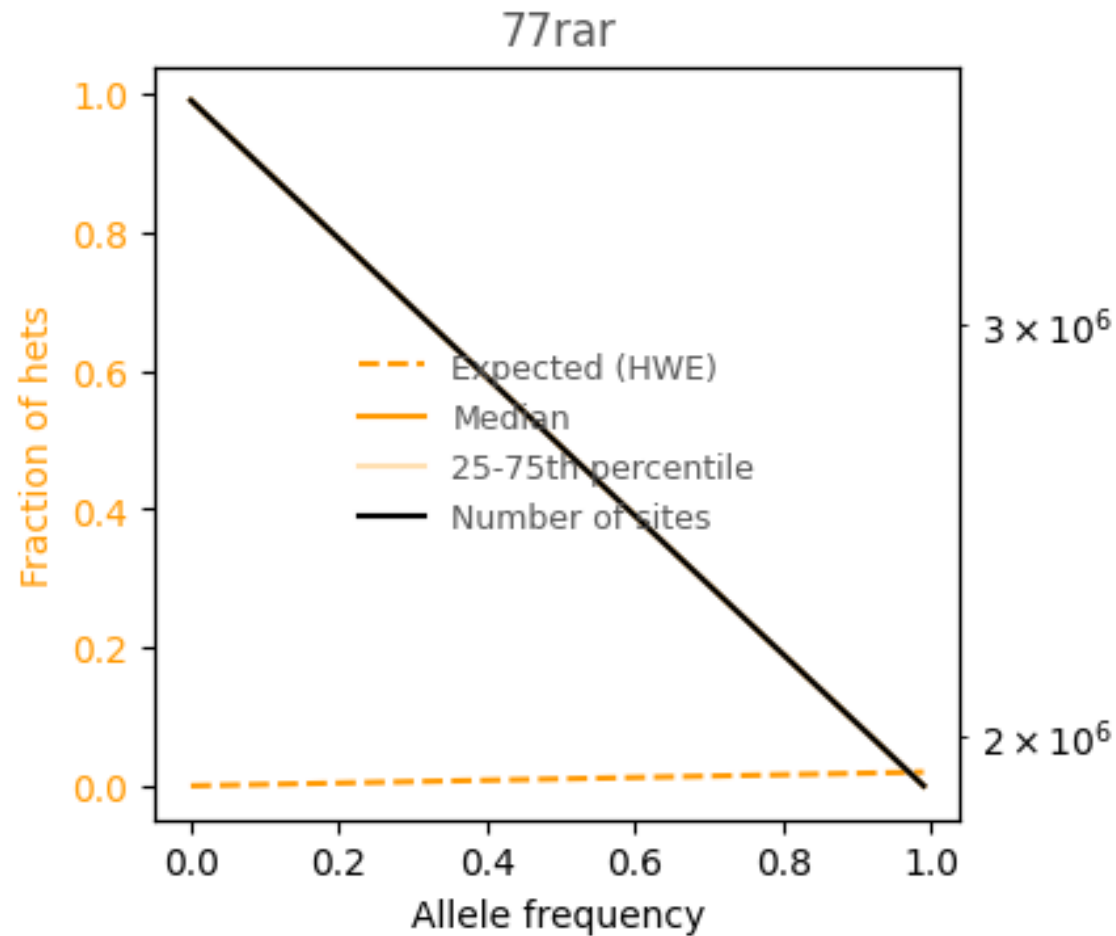

77rar

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types