Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
72uom	4,133,264	1.91	1.91	968,327	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
72uom	62.8%	1.87	68.2%	109,592	2,213

- 72uom .. `/ngc/projects2/gm/data/archive/2022/variants/snv/72uomesam-103893787377-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01039-220727_A01176_AHNW3KDSX3-EXT_LAB`
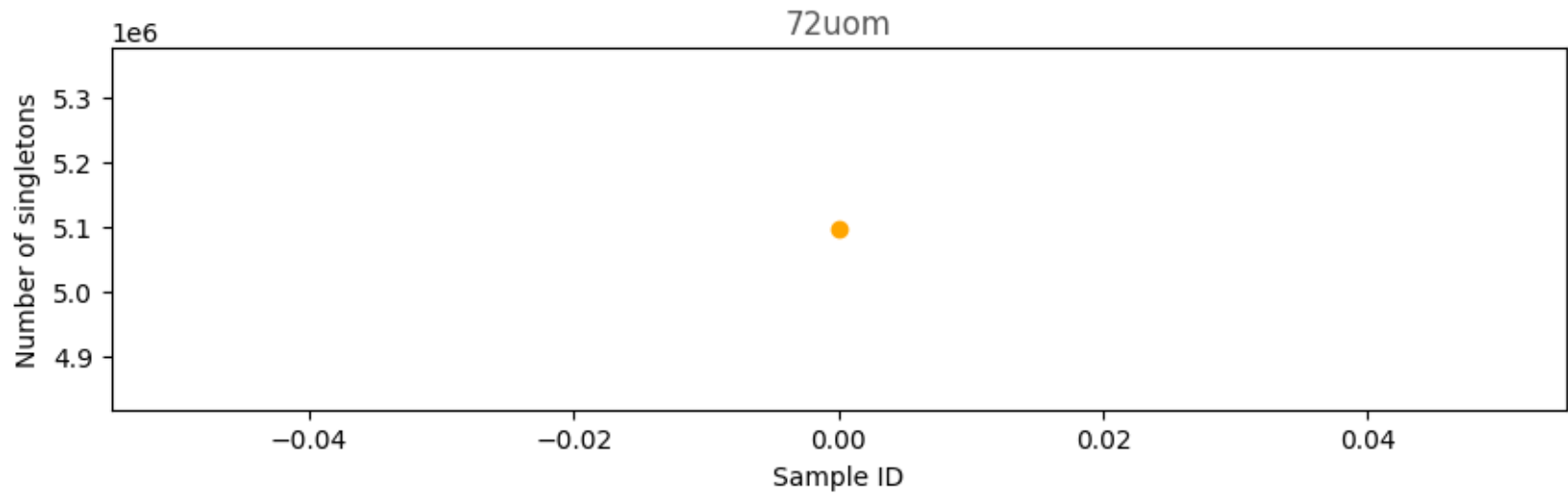 `KA_NGCWGS-NGCWGS04714_22RKG016216x01_snv_germline_raw.haplotype_caller.vcf.gz`
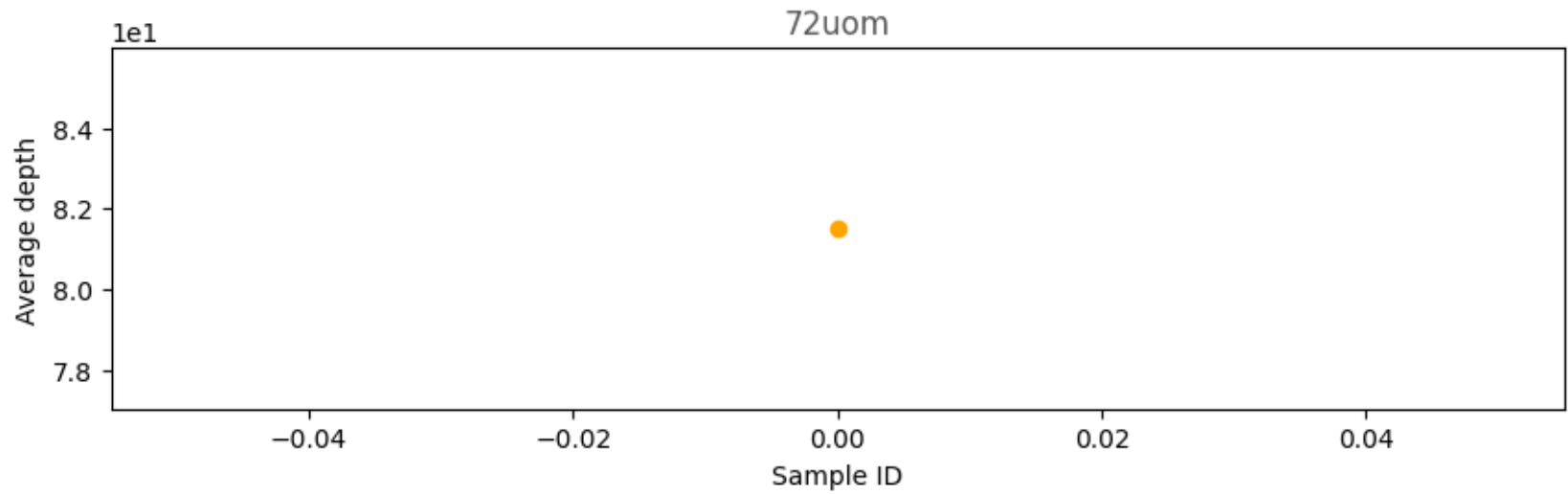
Ts/Tv by sample

72uom
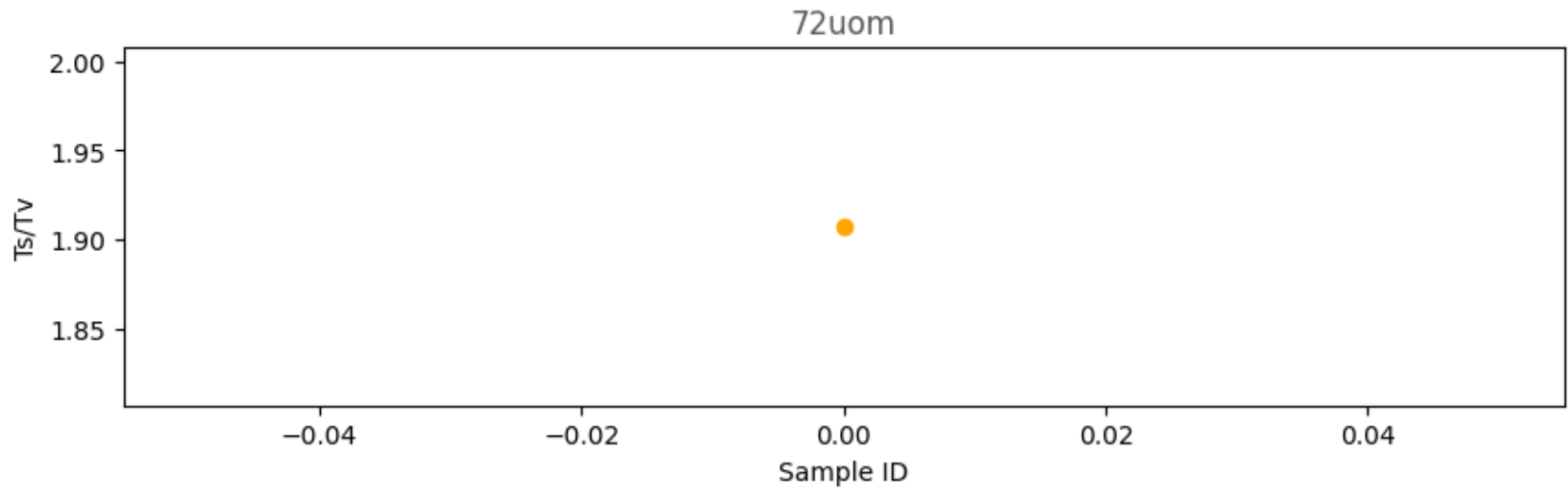
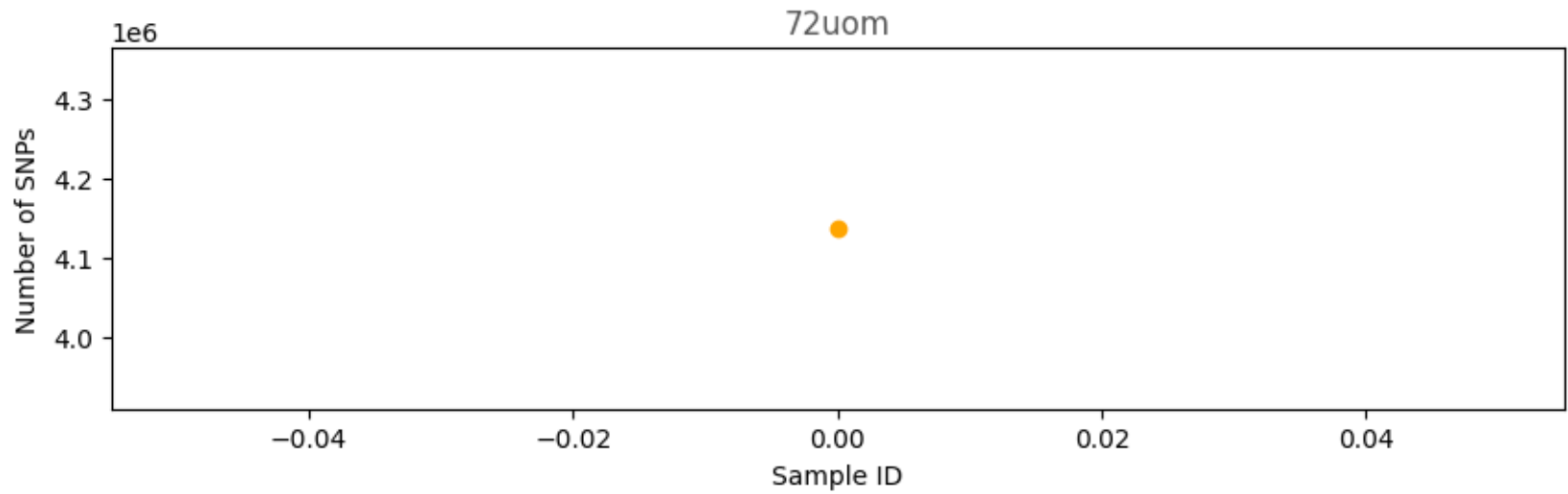# Hets vs non-ref Homs by sample
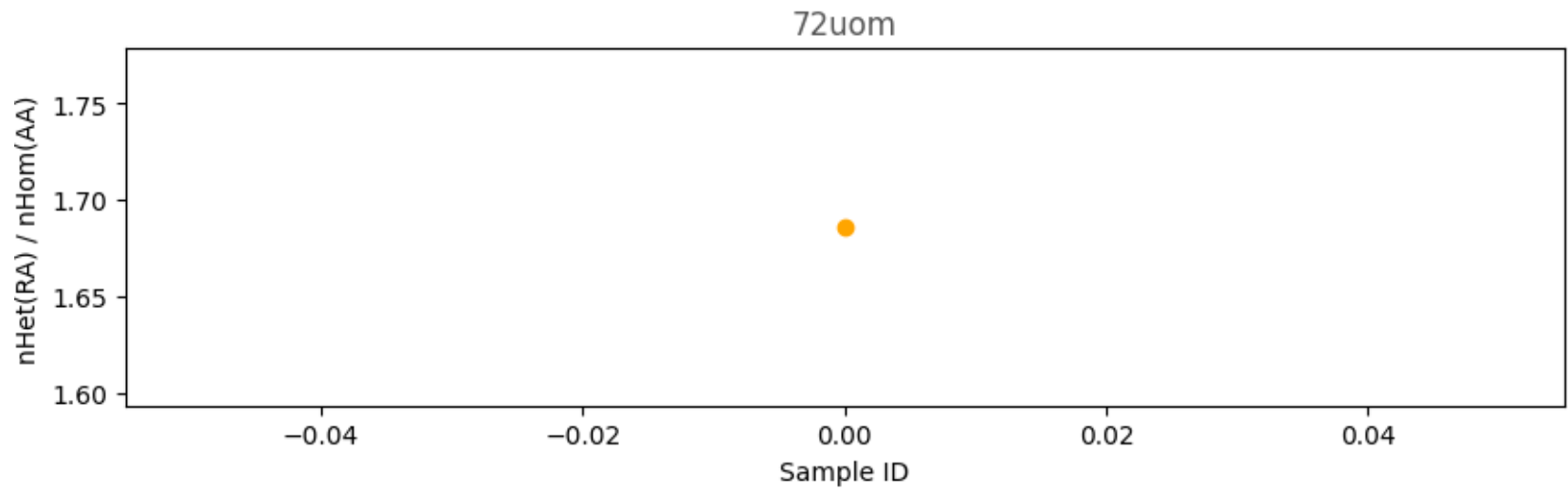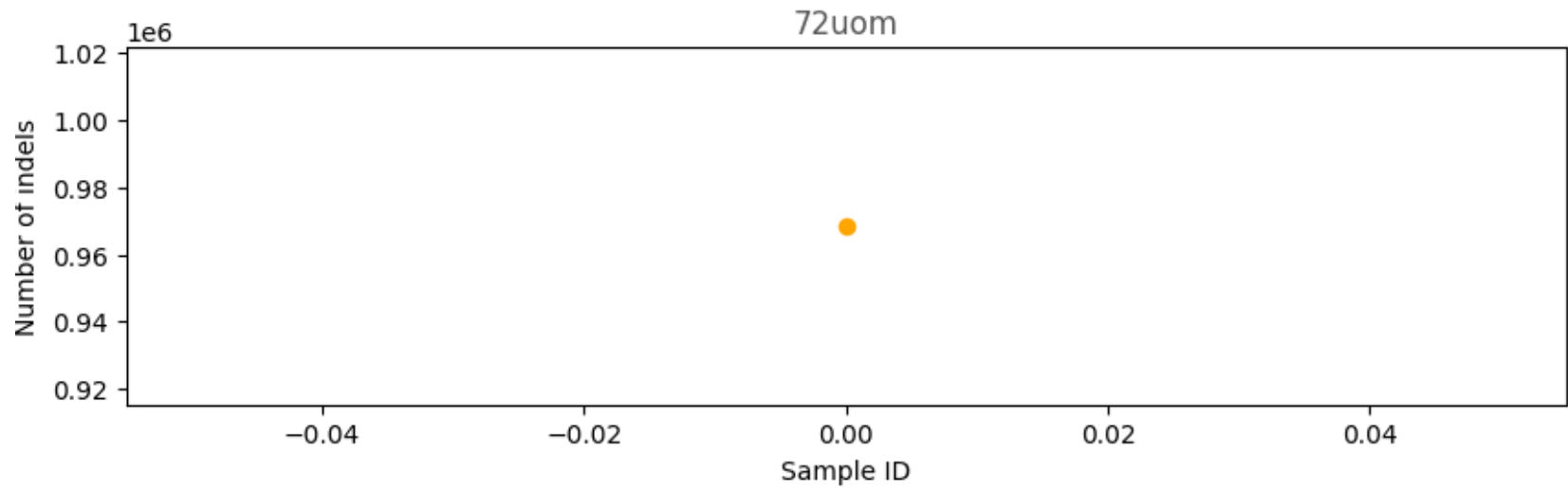
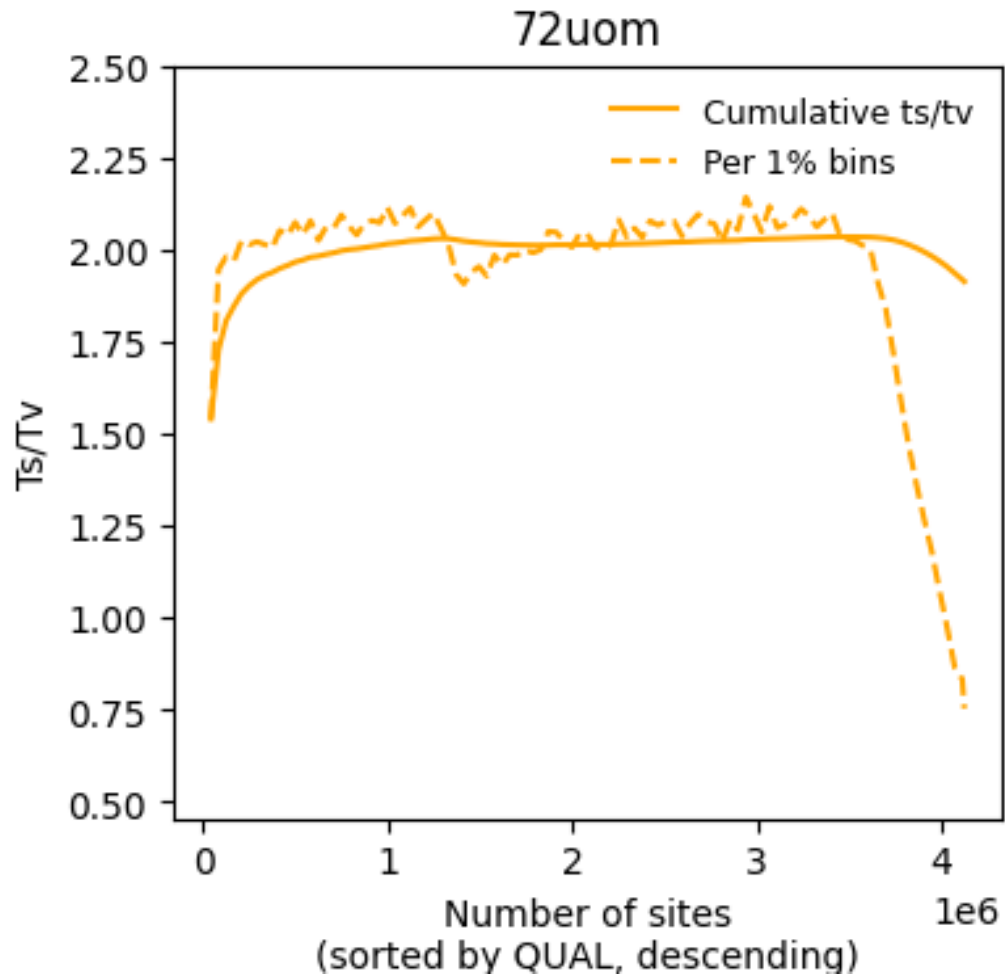# Singletons by sample (hets and homs)

Average depth by sample

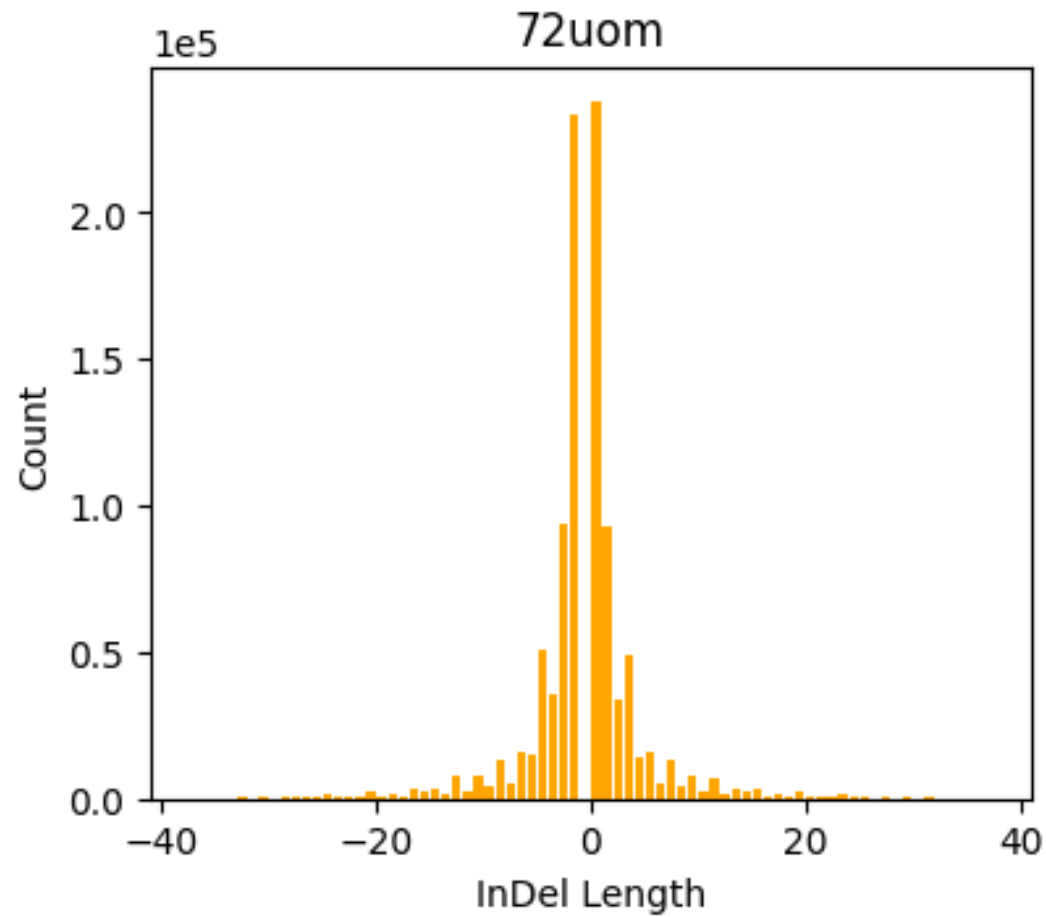# Number of SNPs by sample

Number of indels by sample
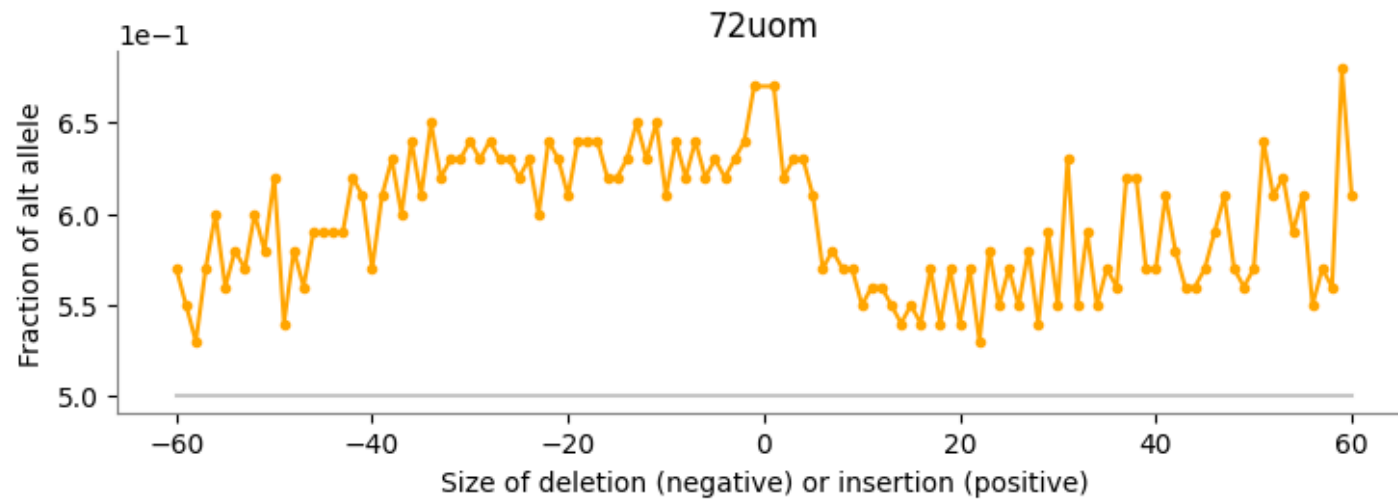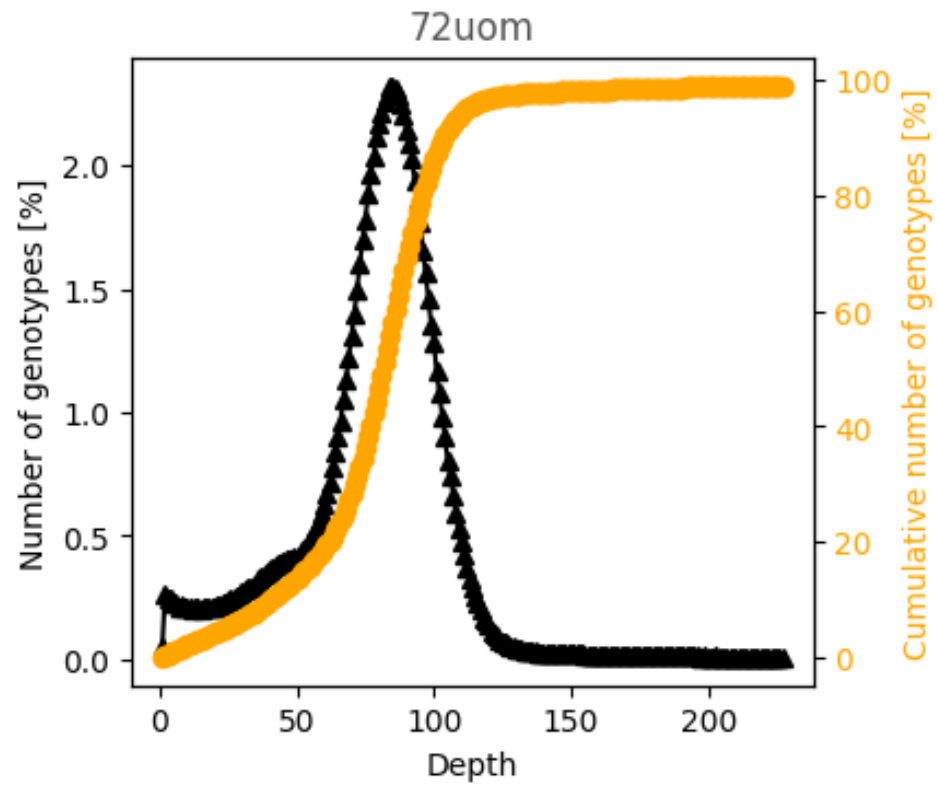
Ts/Tv stratified by QUAL

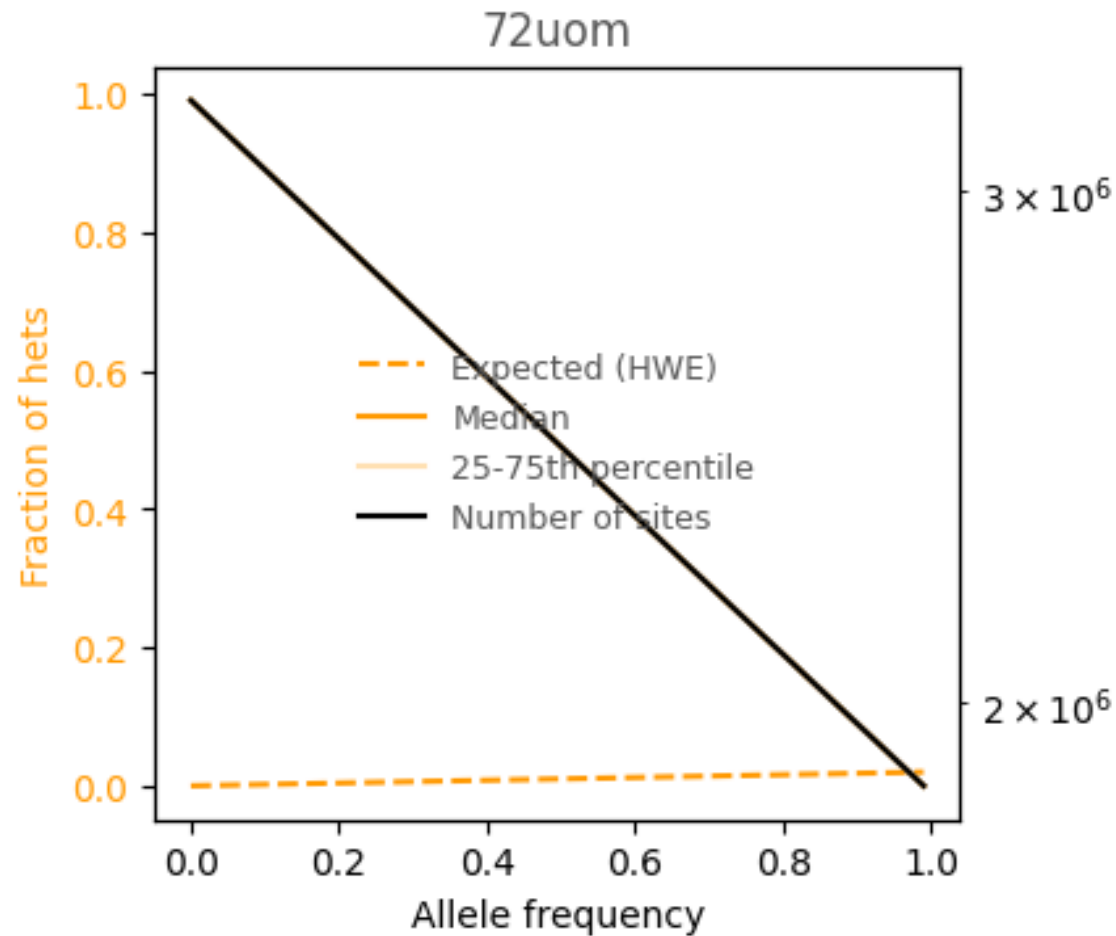

72uom

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types