Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
72mou	3,999,940	1.96	1.96	928,972	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
72mou	61.5%	1.93	65.6%	90,282	1,731

- 72mou .. `/ngc/projects2/gm/data/archive/2022/variants/snv/72mouprem-103904421728-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01186-220831_A01961_BHTNYFDSX3-EXT_LAB`
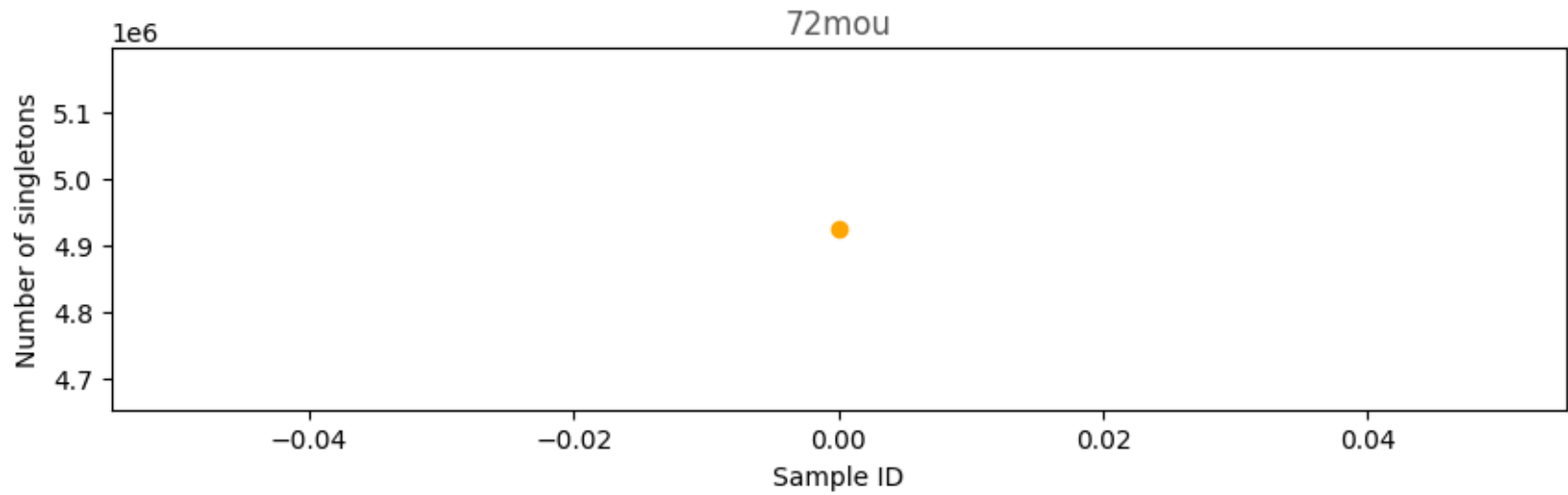 `KA_NGCWGS-NGCWGS04942_22RKG012402x01_snv_germline_raw.haplotype_caller.vcf.gz`
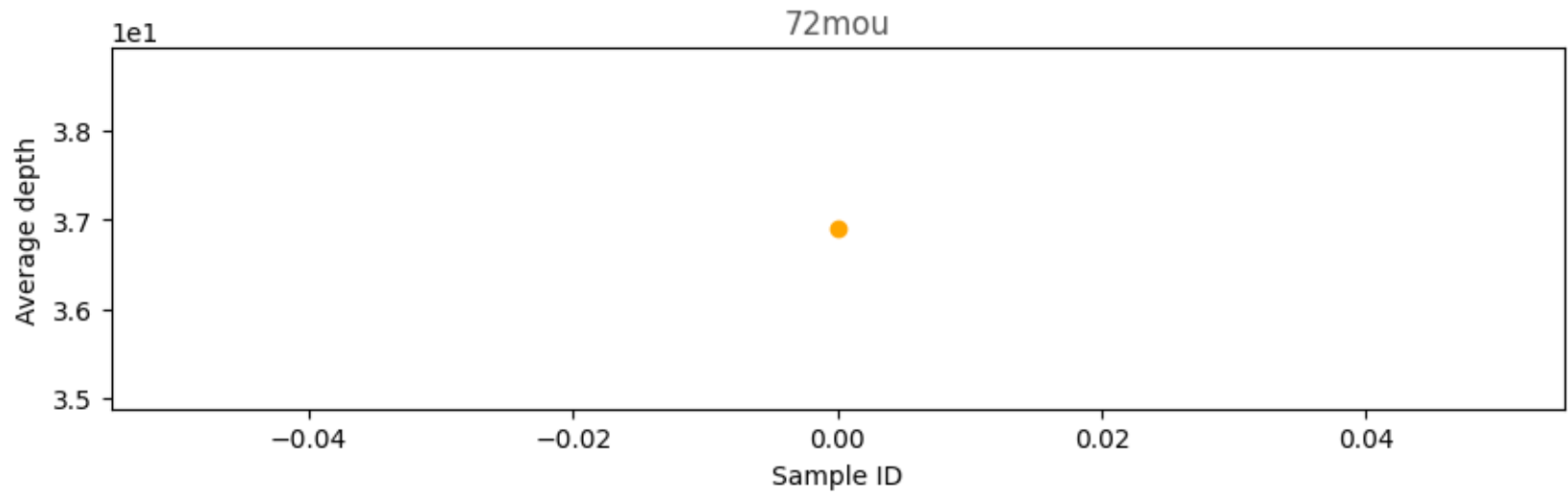
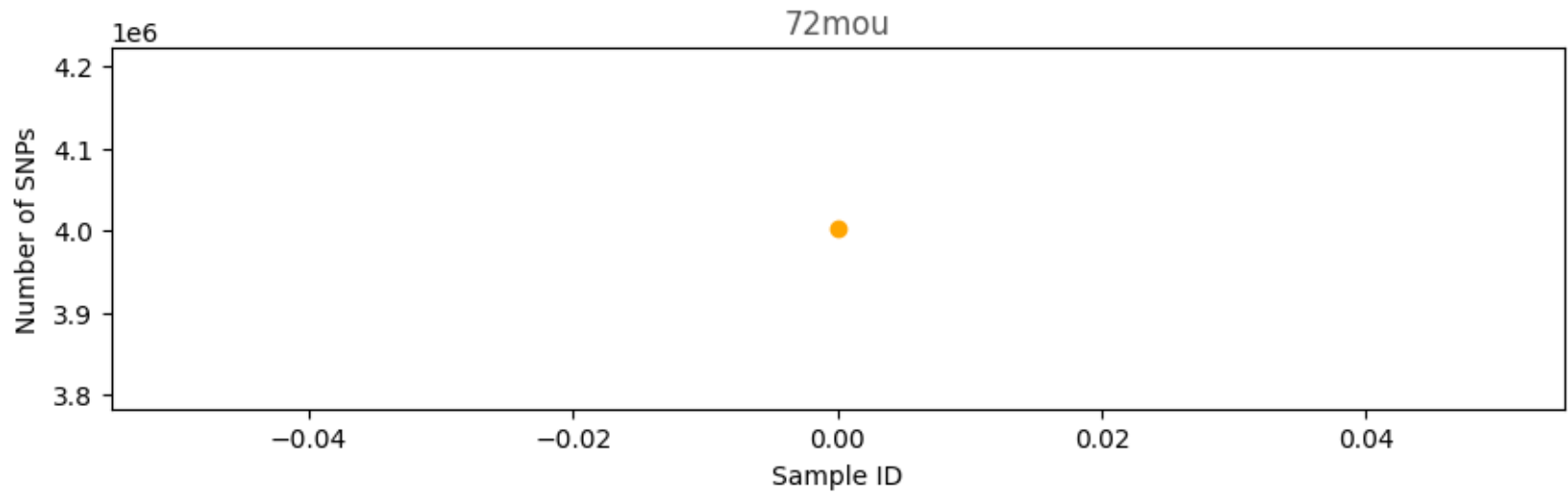# Ts/Tv by sample
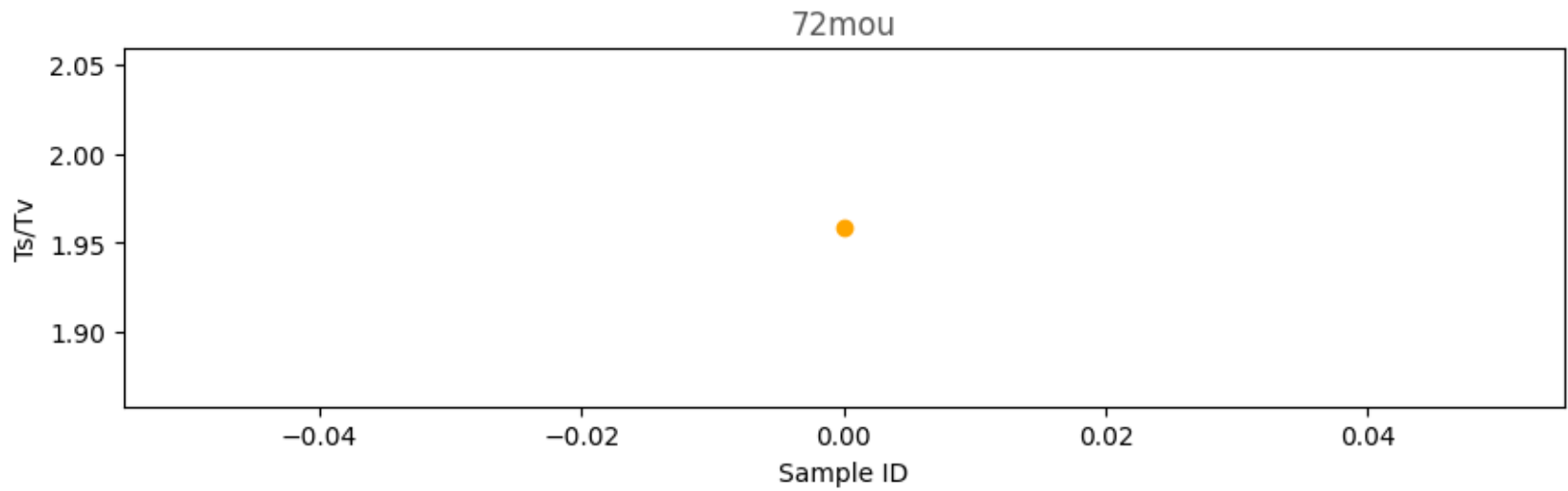
Hets vs non-ref Homs by sample
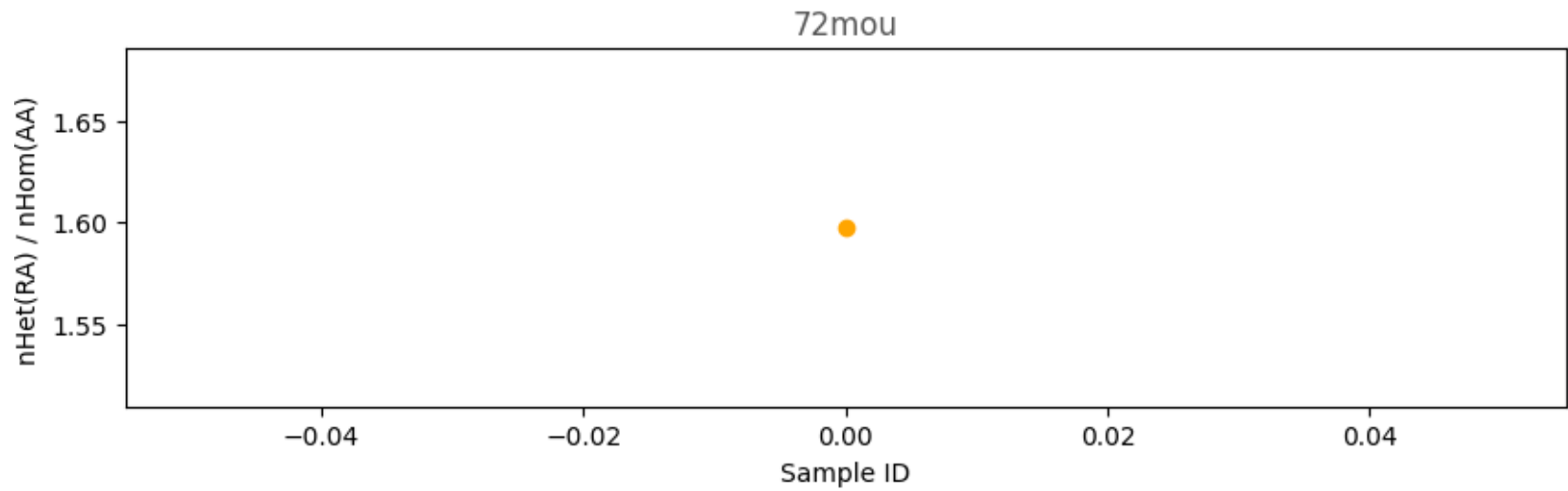
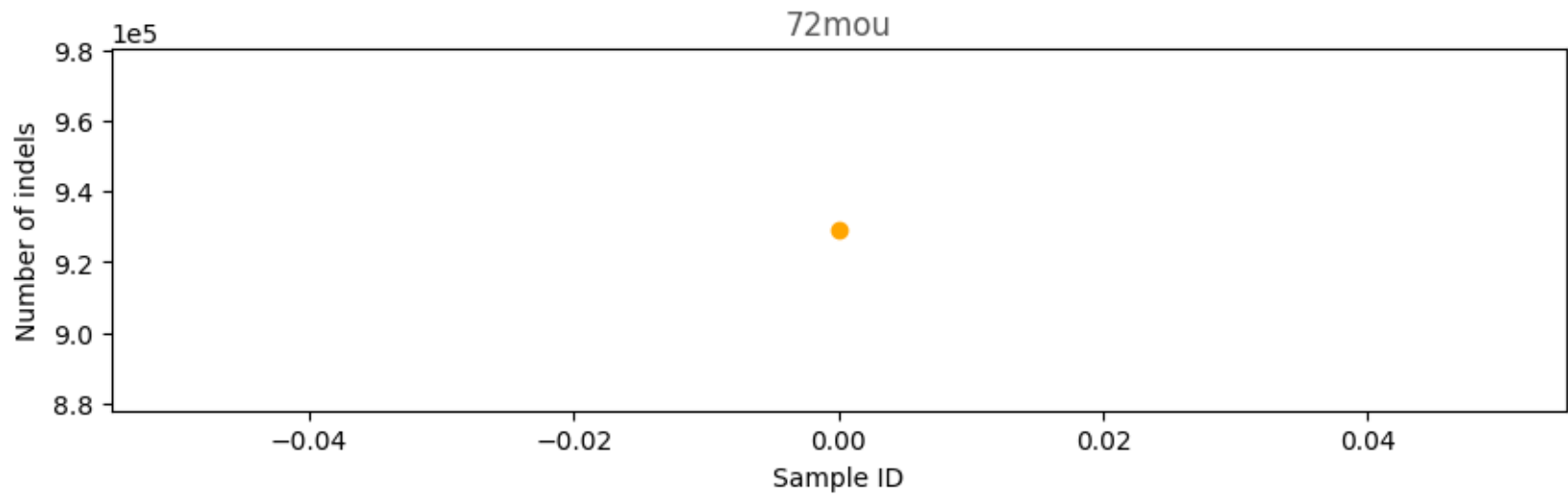# Singletons by sample (hets and homs)
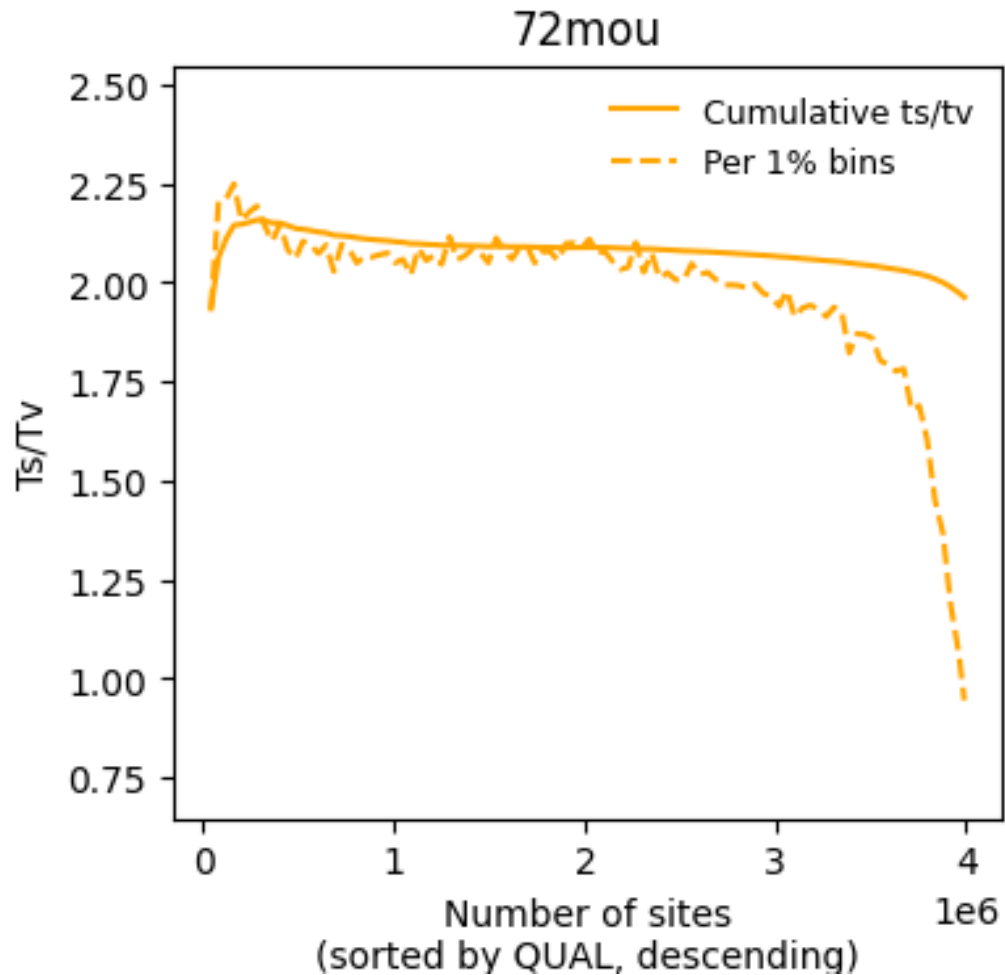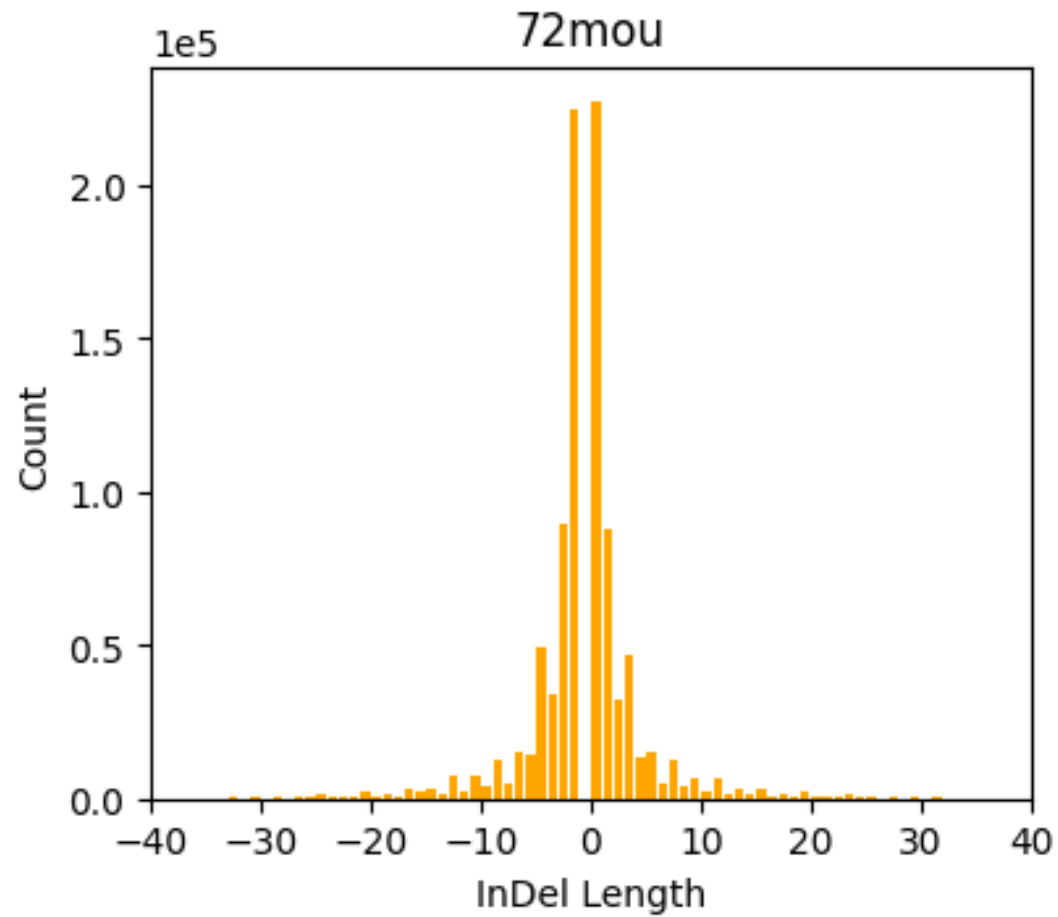
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

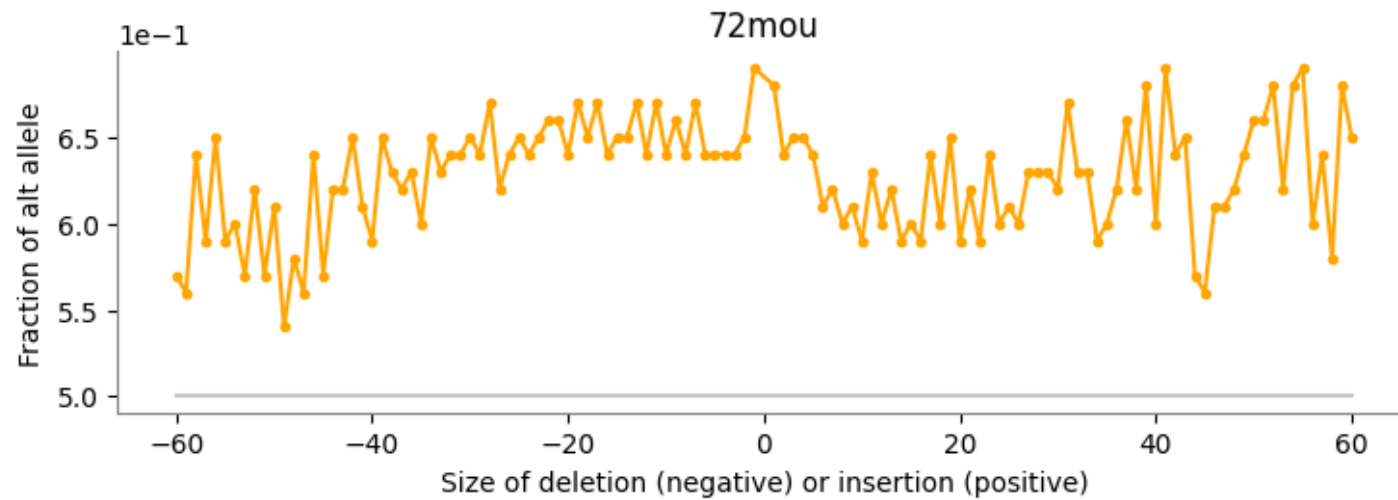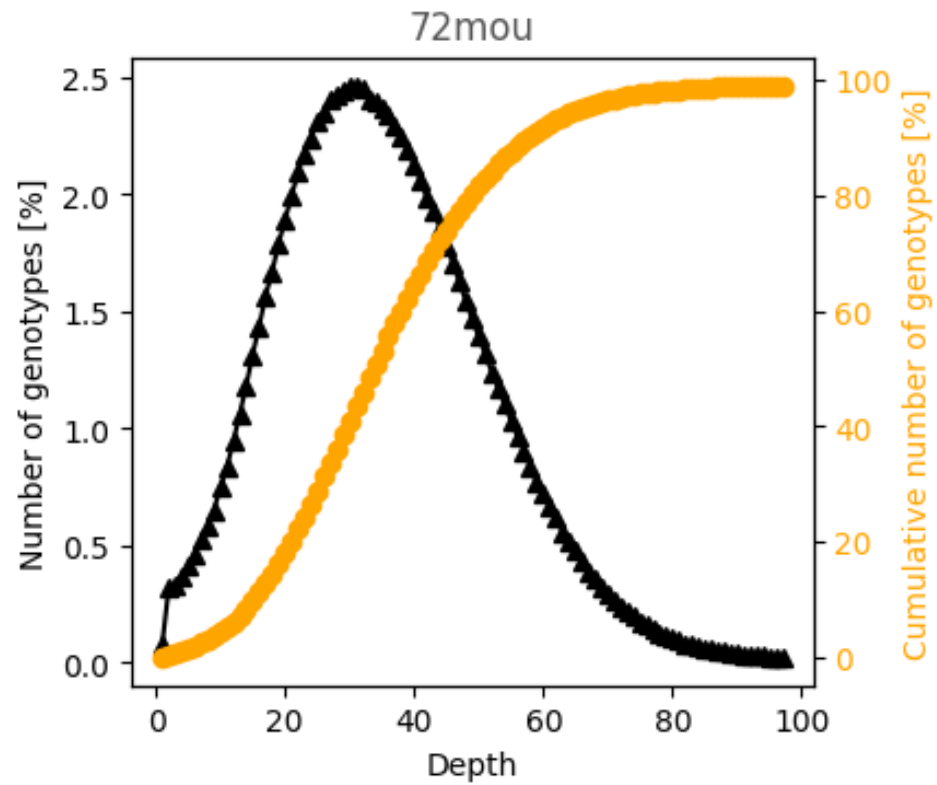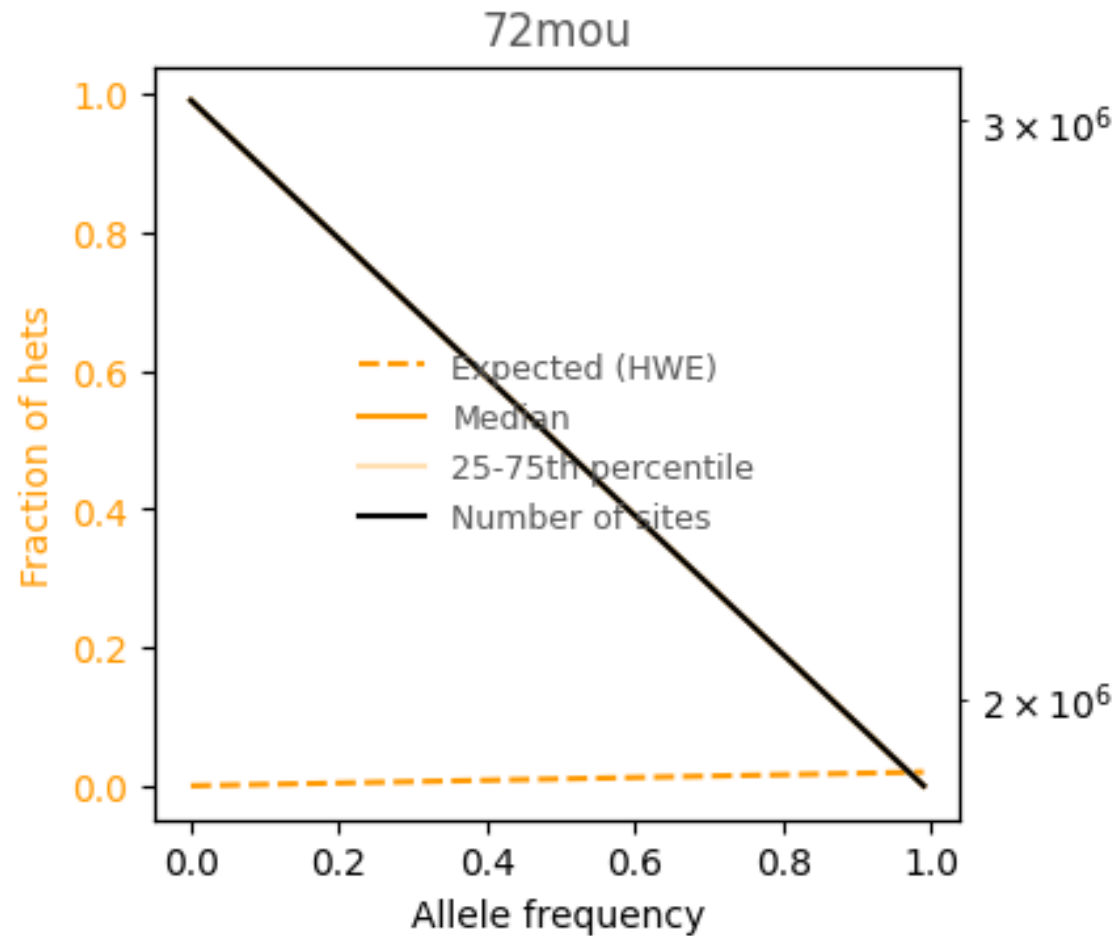# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types