Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
74mul	4,084,731	1.93	1.93	951,146	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
74mul	62.8%	1.90	67.9%	101,064	2,170

- 74mul .. `/ngc/projects2/gm/data/archive/2022/variants/snv/74mullulf-110297773007-Normal_B`
`lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02099-221026_A00559_AH3MCCDSX5-EXT_LAB`
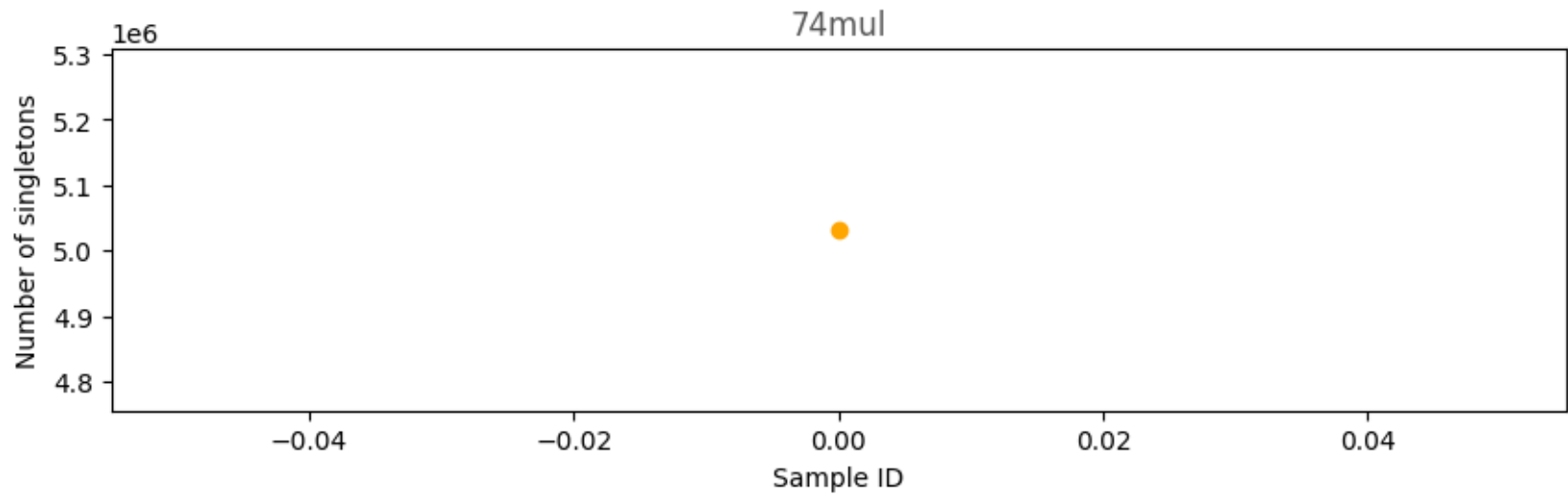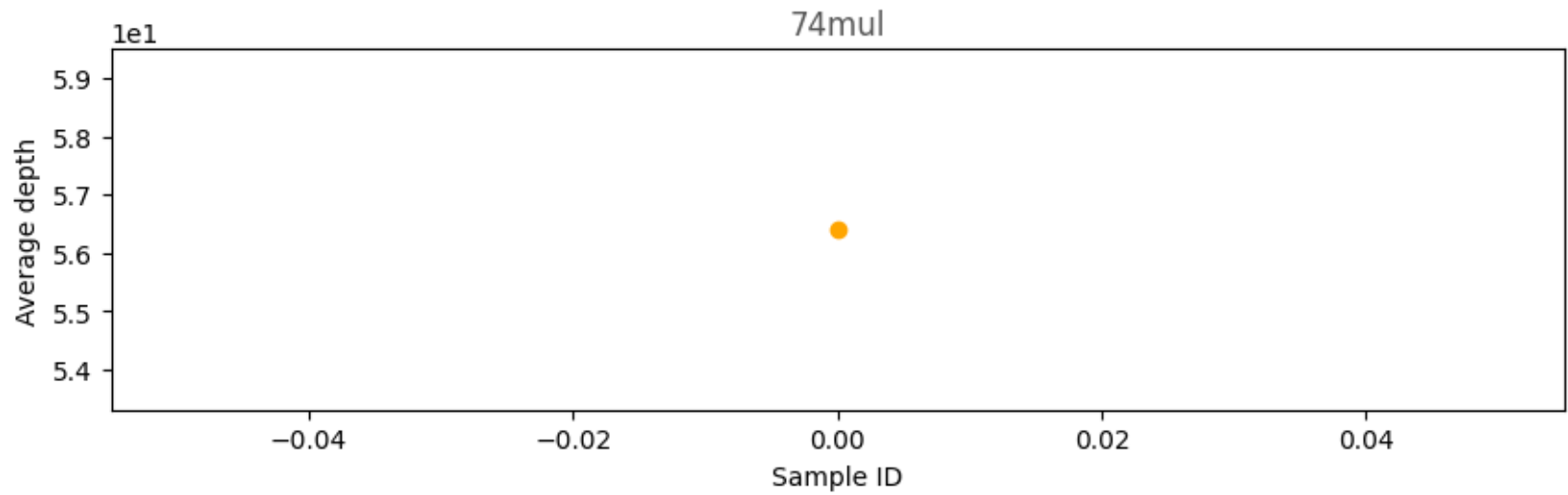`KA_NGCWGS-NGCWGS05960_snv_germline_raw.haplotype_caller.vcf.gz`
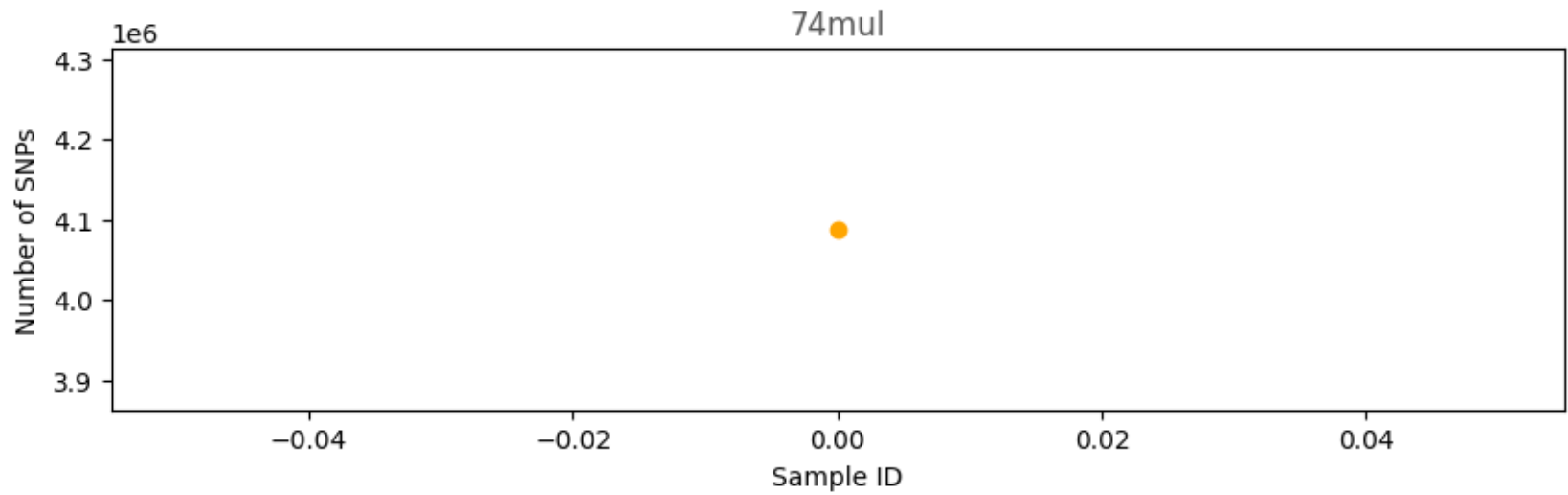
Ts/Tv by sample
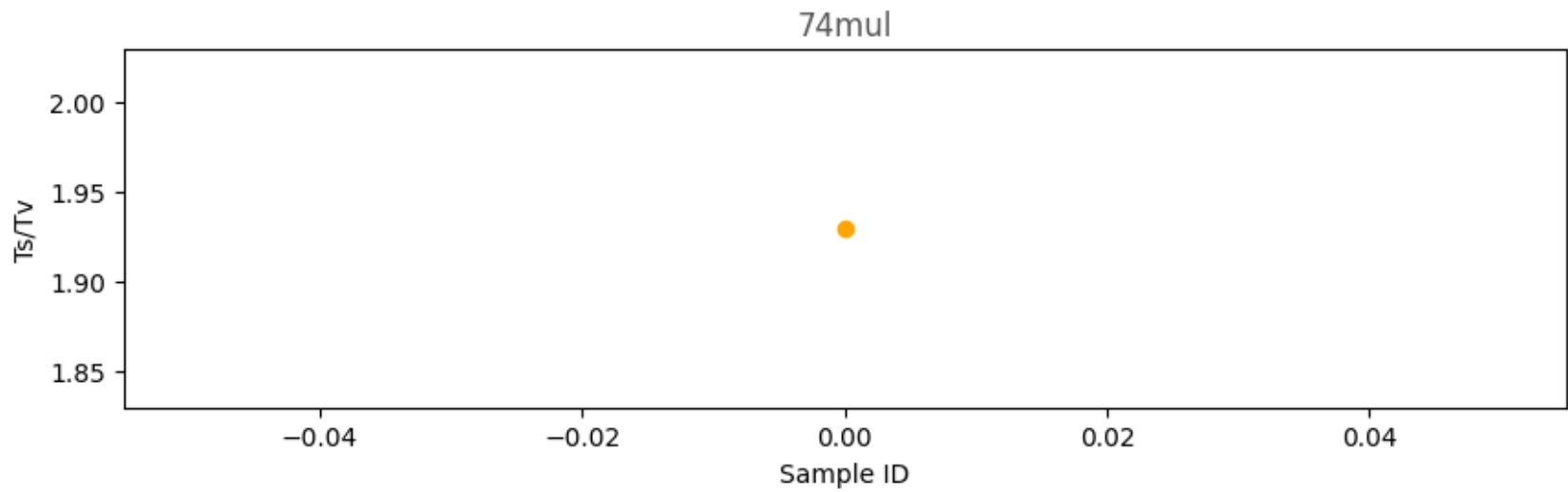
Hets vs non-ref Homs by sample
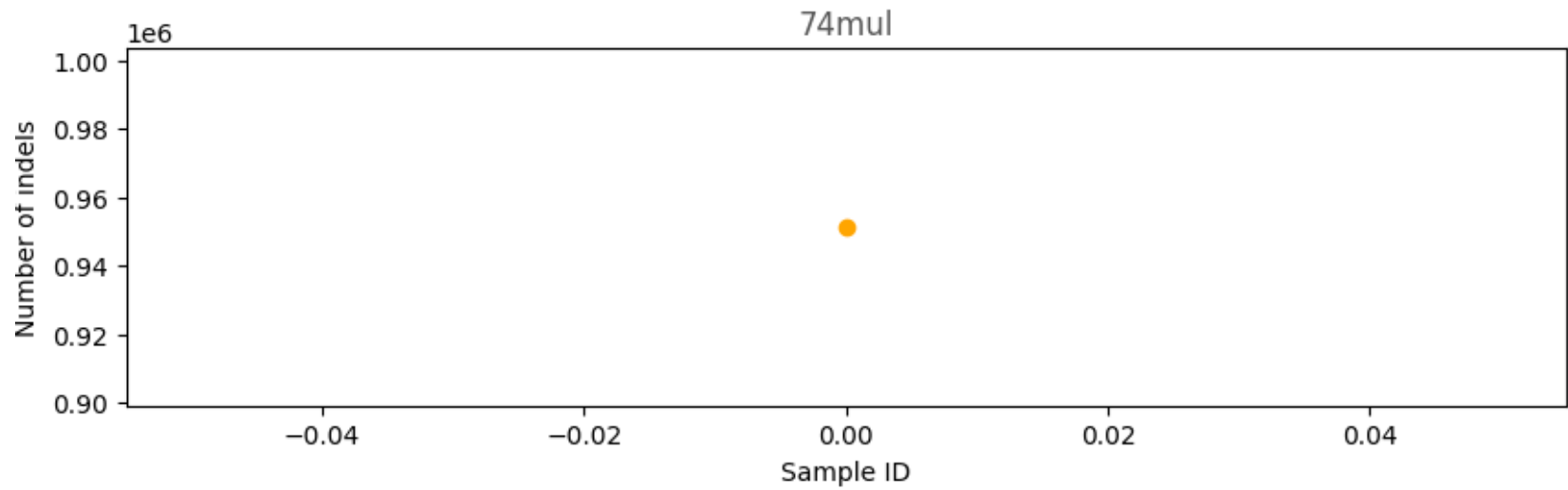

3
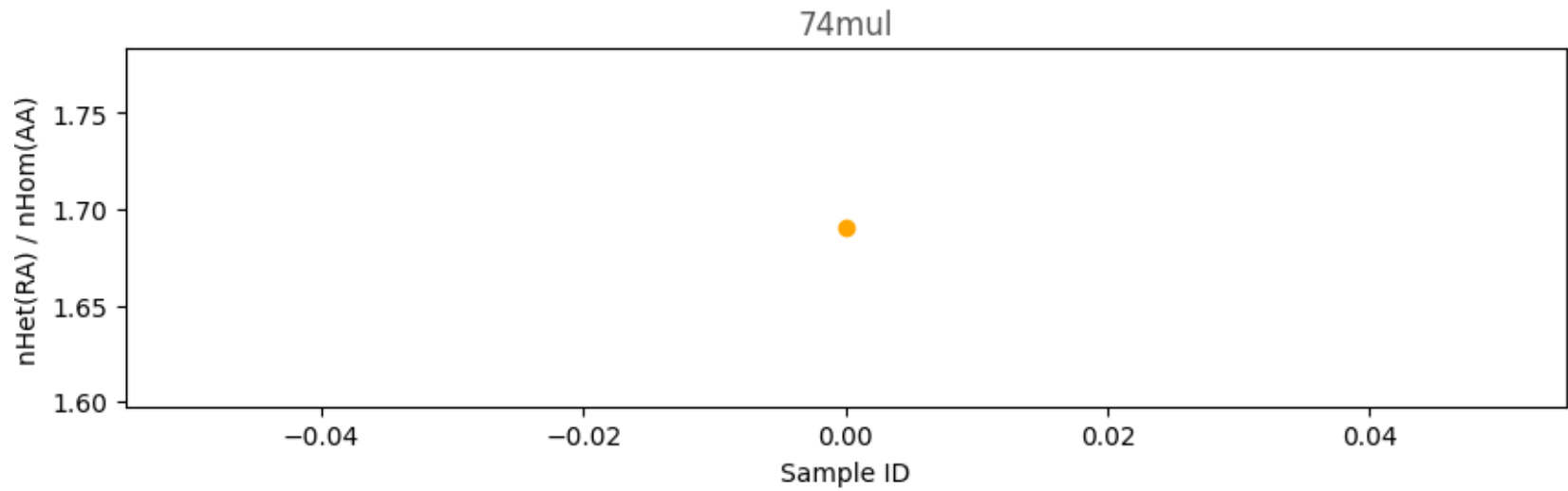
Singletons by sample (hets and homs)
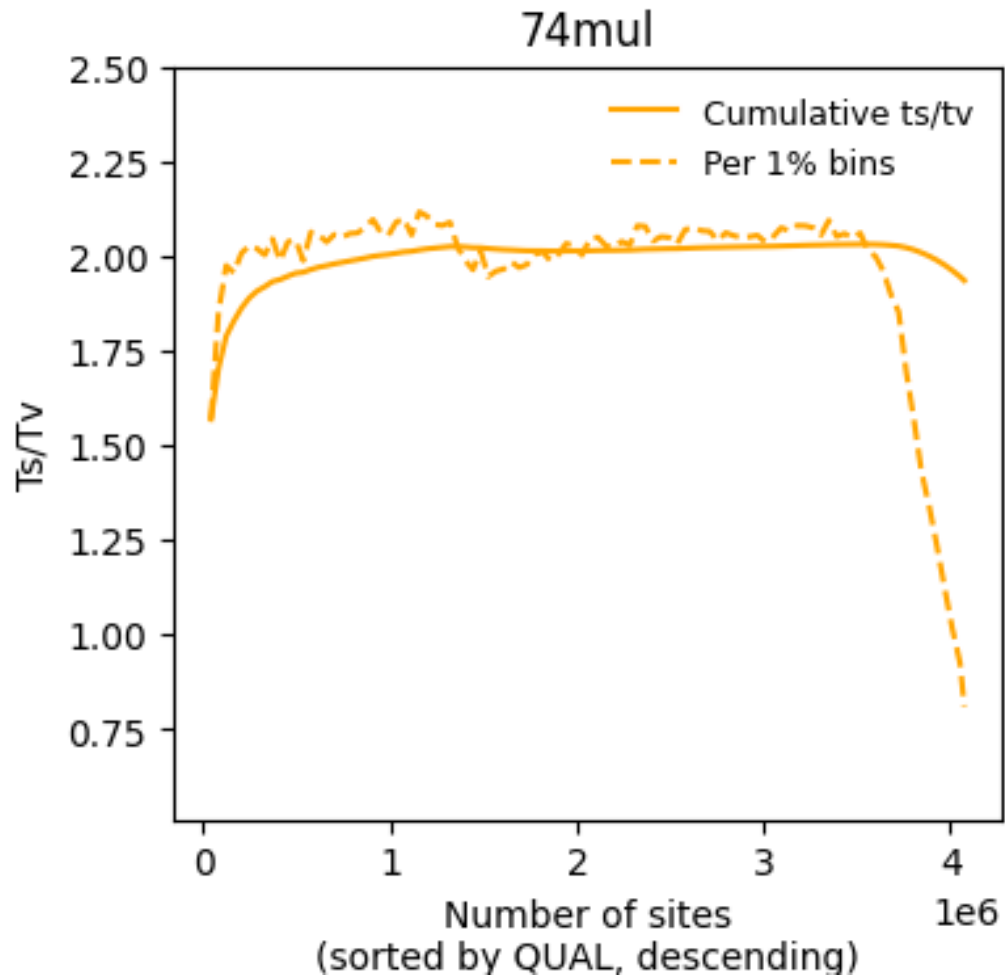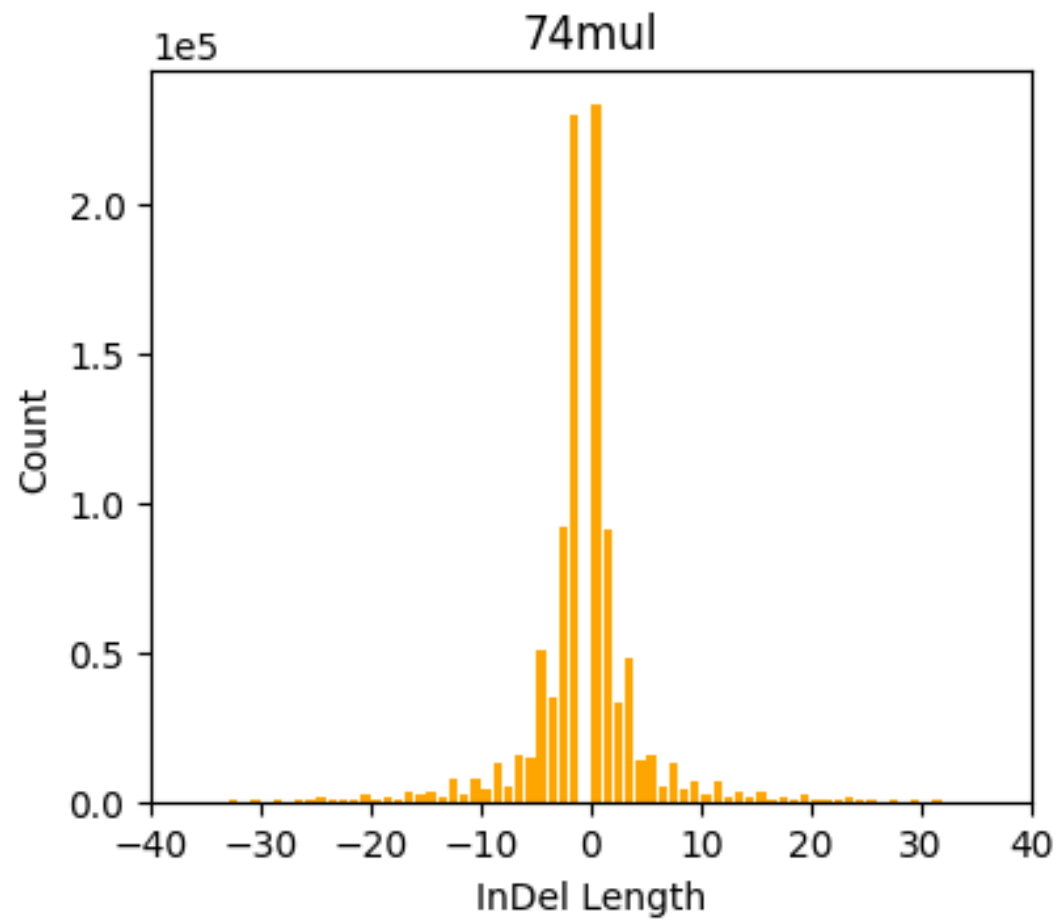
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

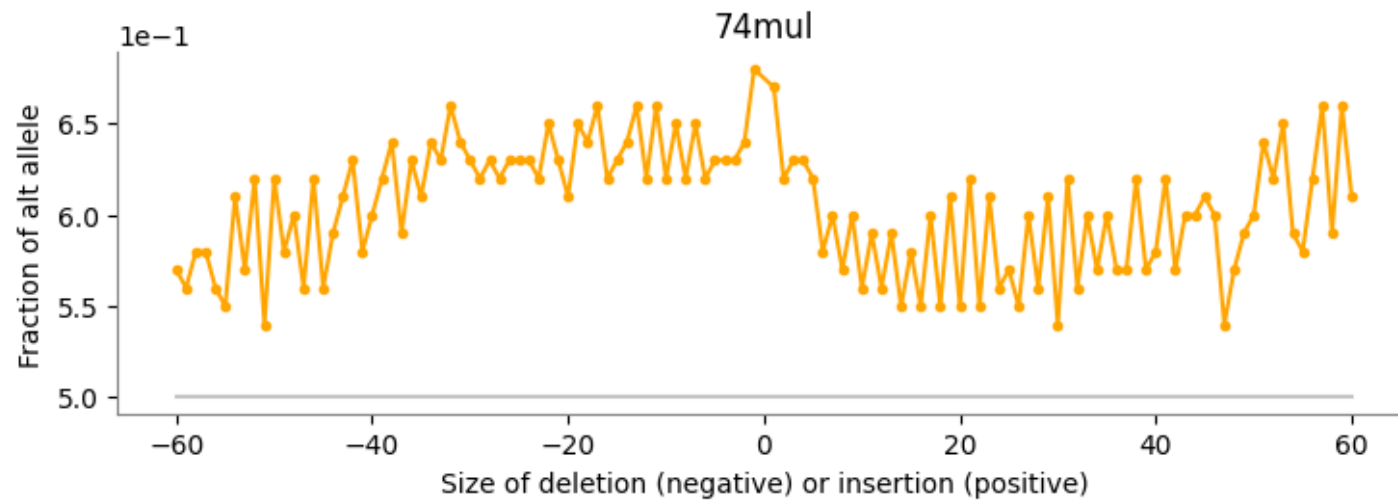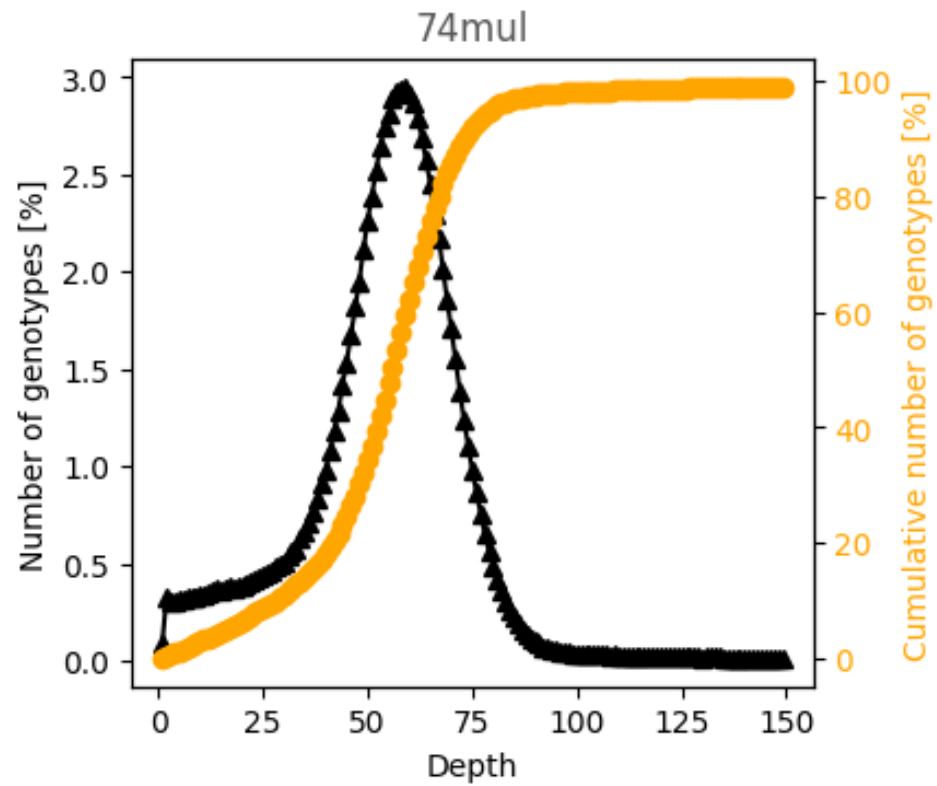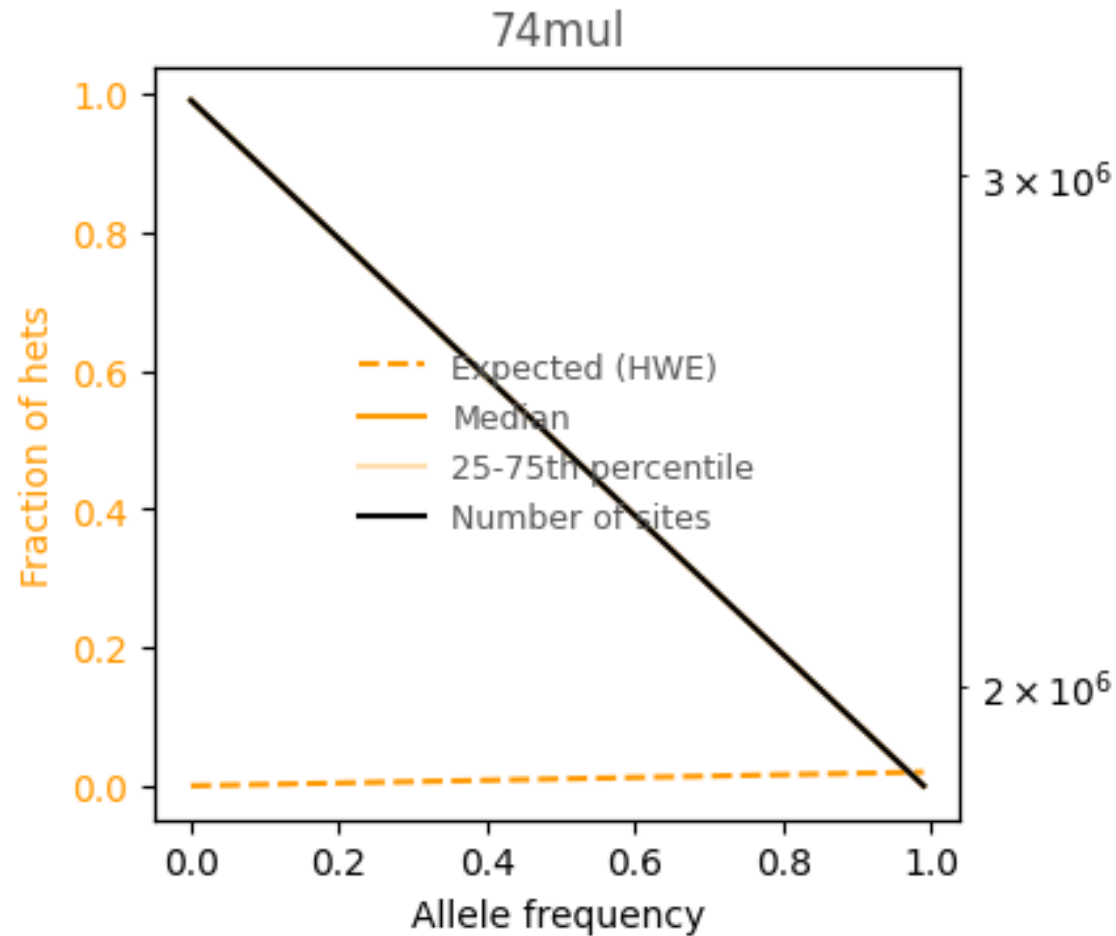# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types