Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
69bar	4,105,706	1.94	1.95	969,330	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
69bar	63.0%	1.92	68.8%	106,989	2,145

- 69bar .. `/ngc/projects2/gm/data/archive/2022/variants/snv/69barcraf-103807546809-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00441-211206_A01466_AH2CT3DSX3-EXT_LAB`
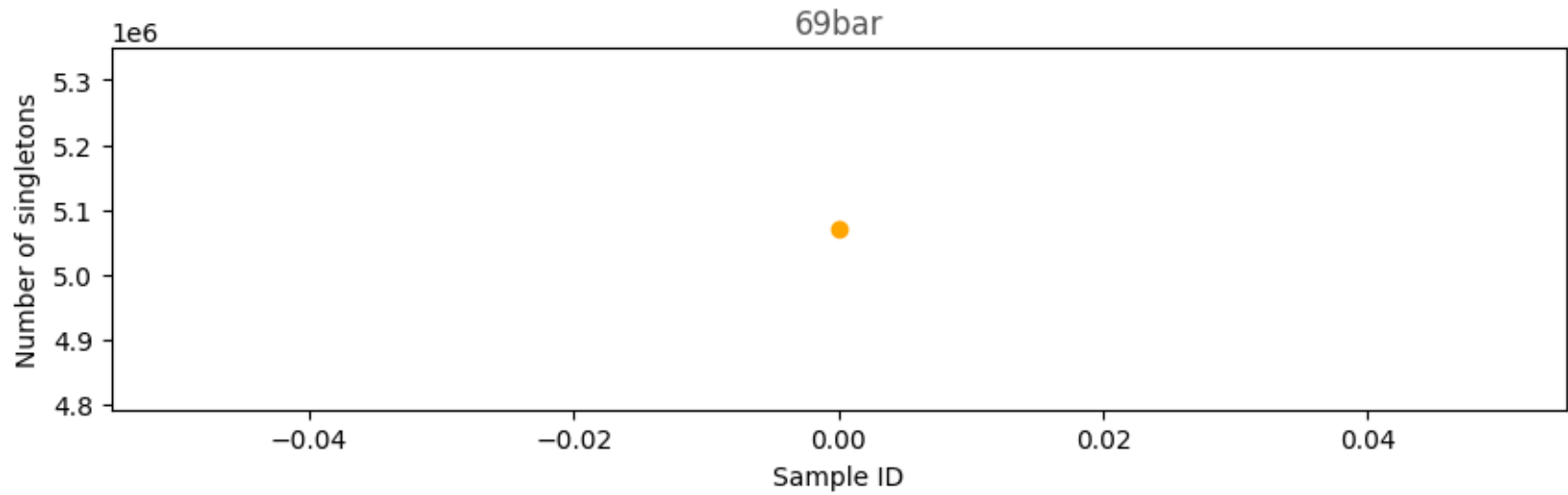 `KA_NGCWGS-NGCWGS03463_21RKG024029x01_snv_germline_raw.haplotype_caller.vcf.gz`
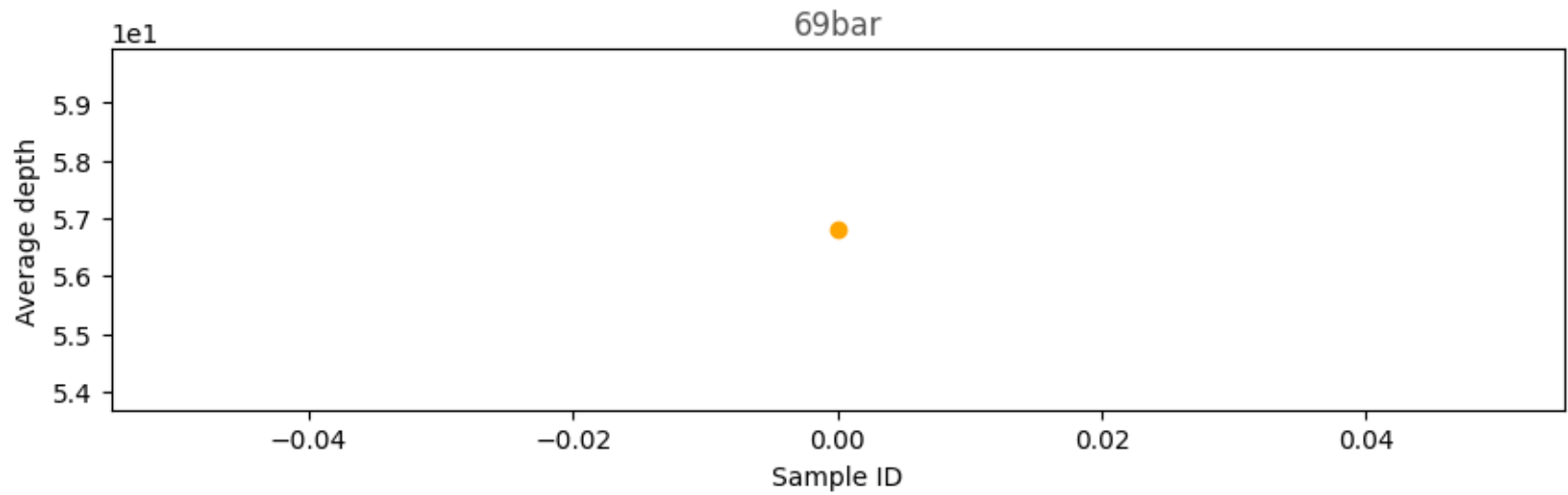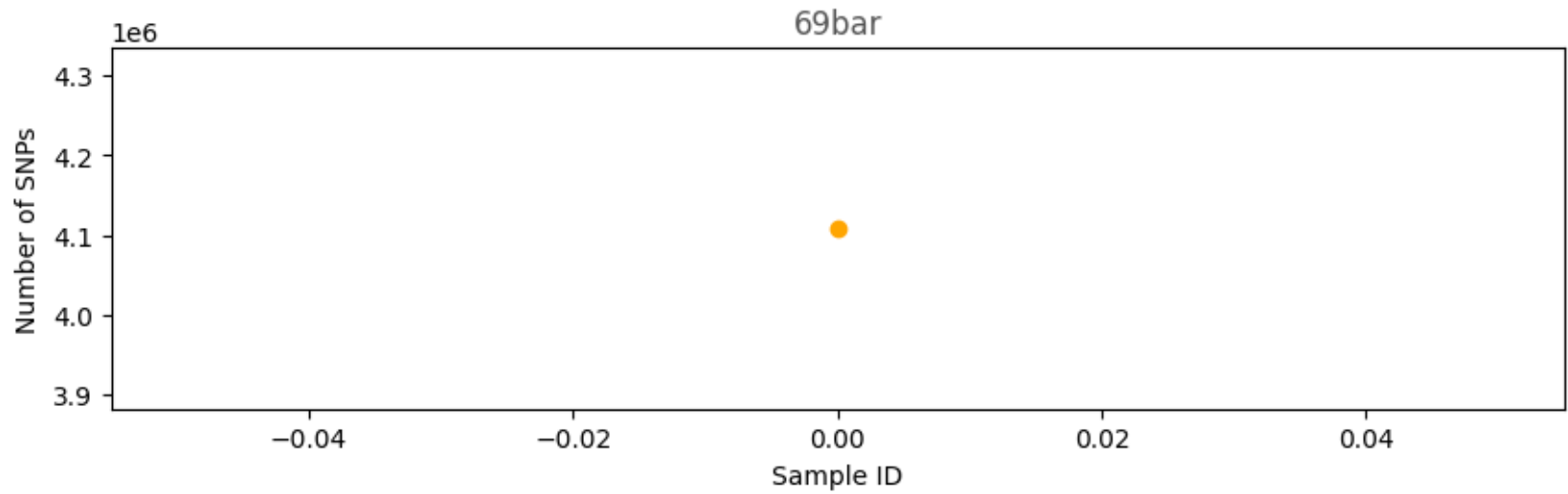
Ts/Tv by sample
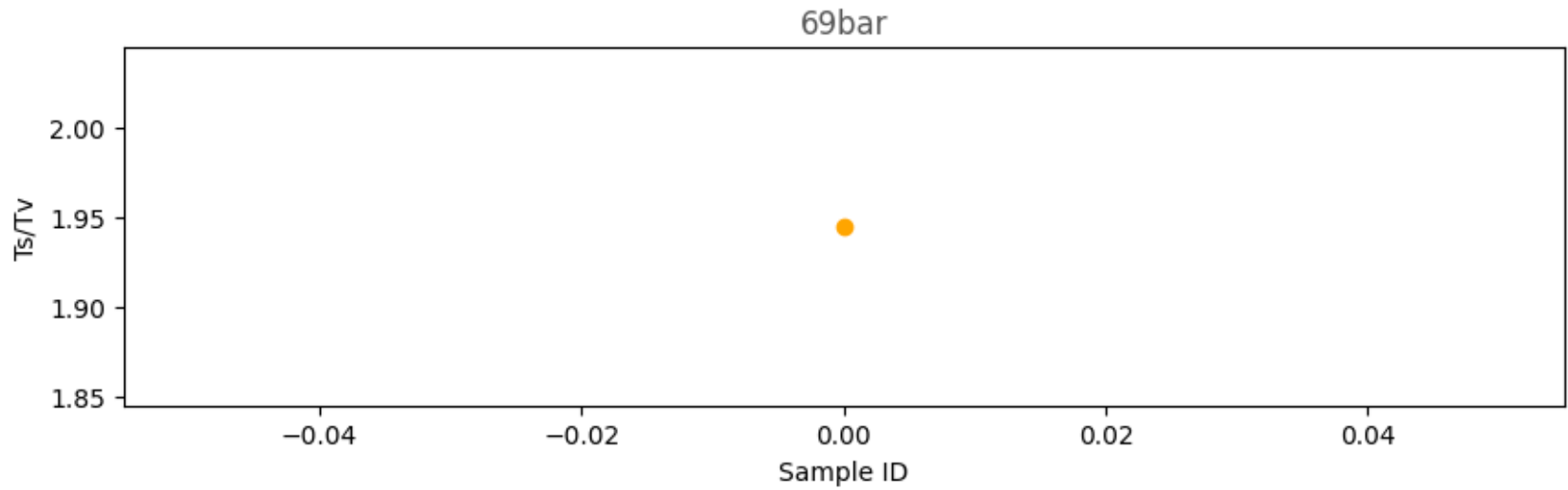
Hets vs non-ref Homs by sample
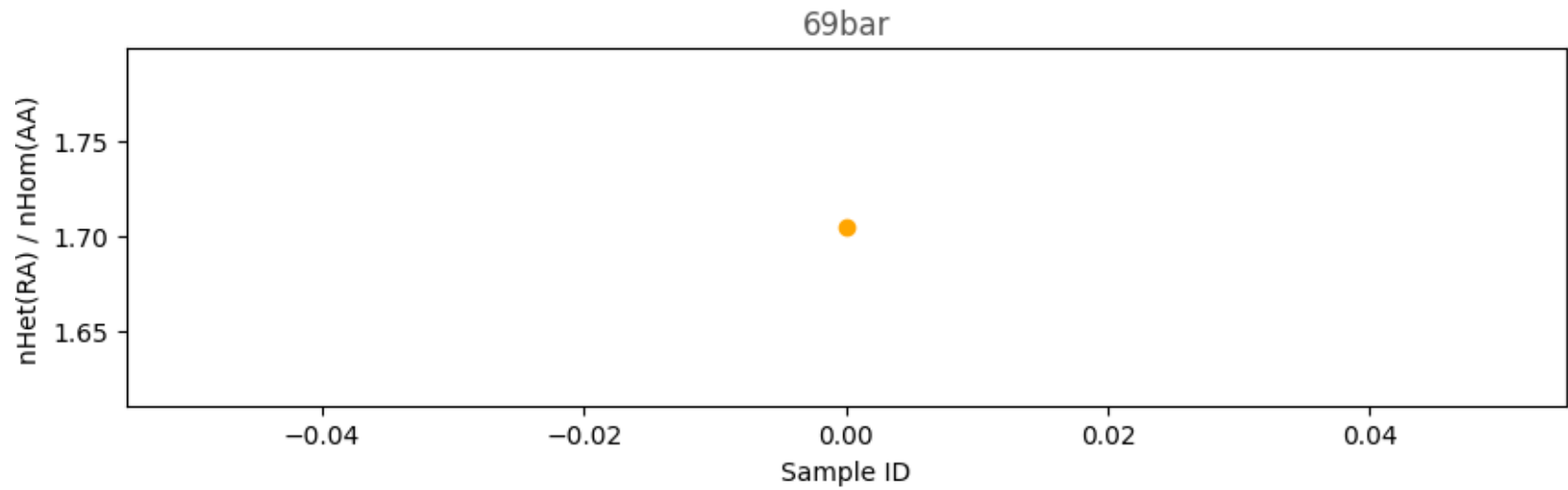
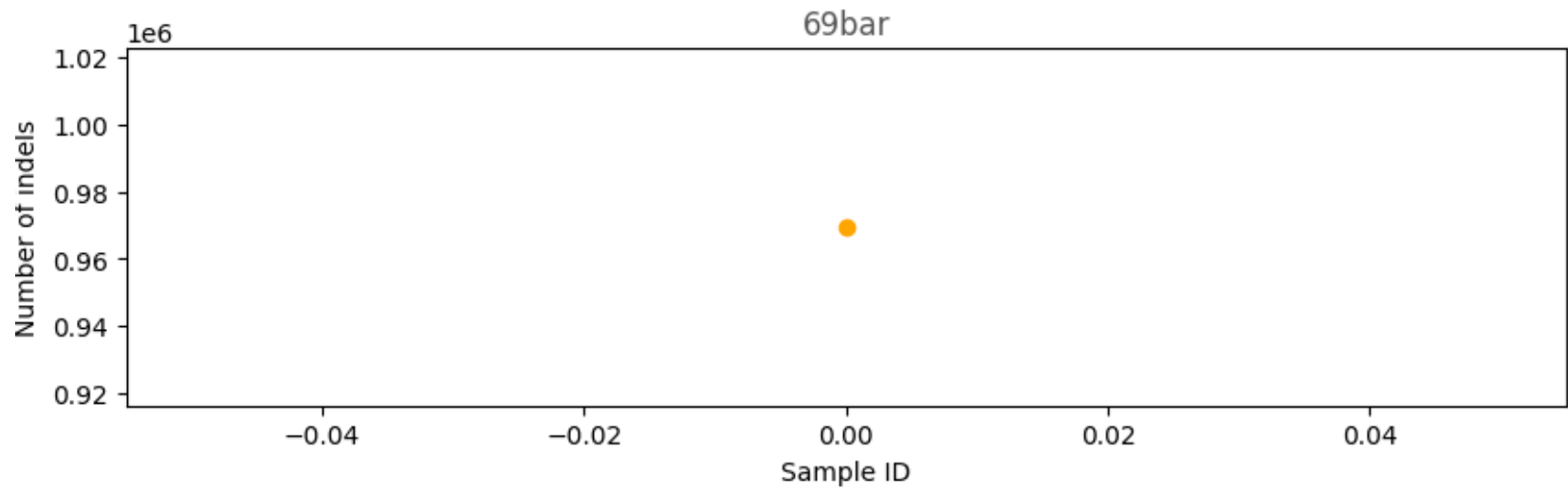# Singletons by sample (hets and homs)
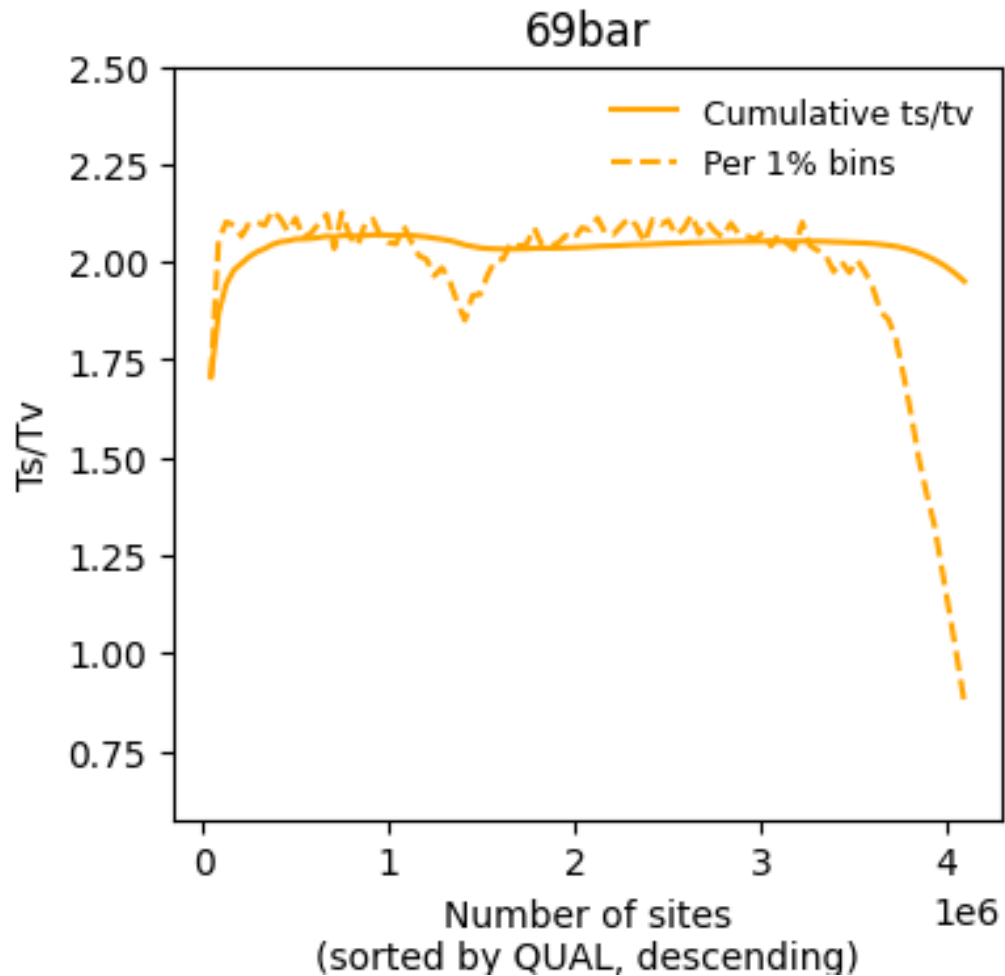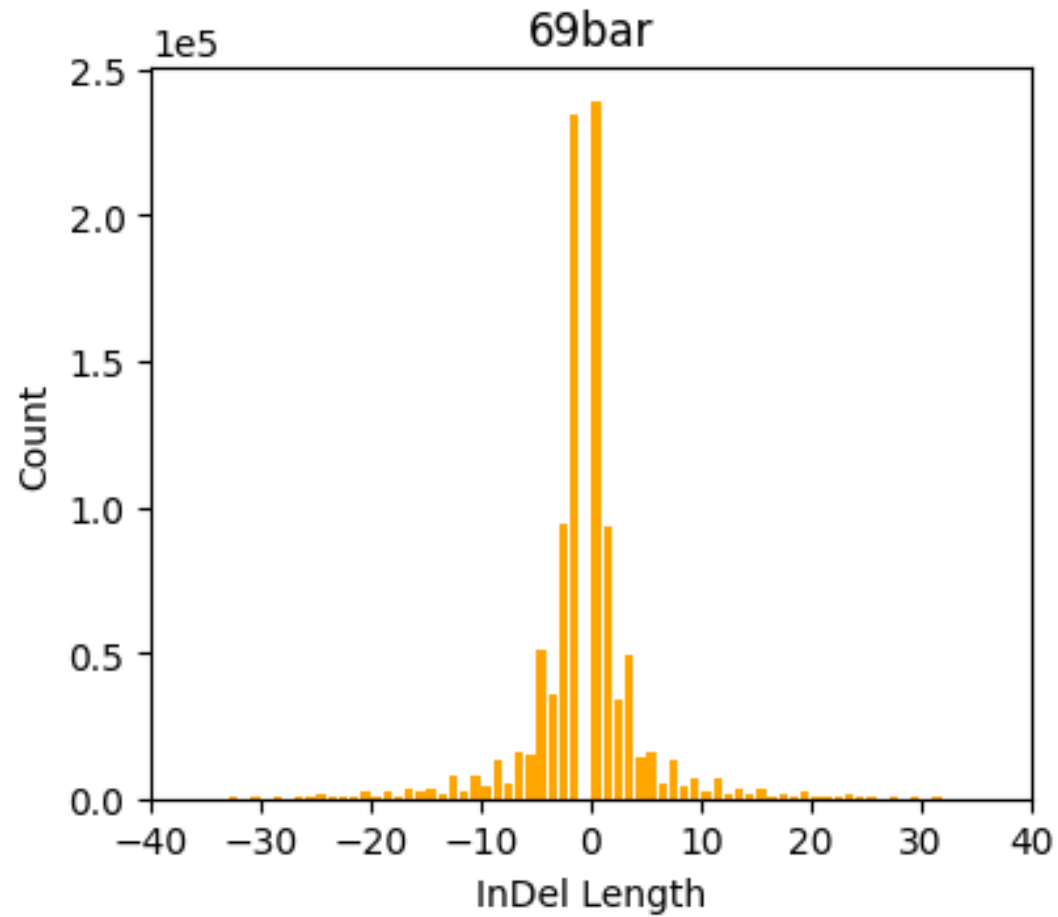
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

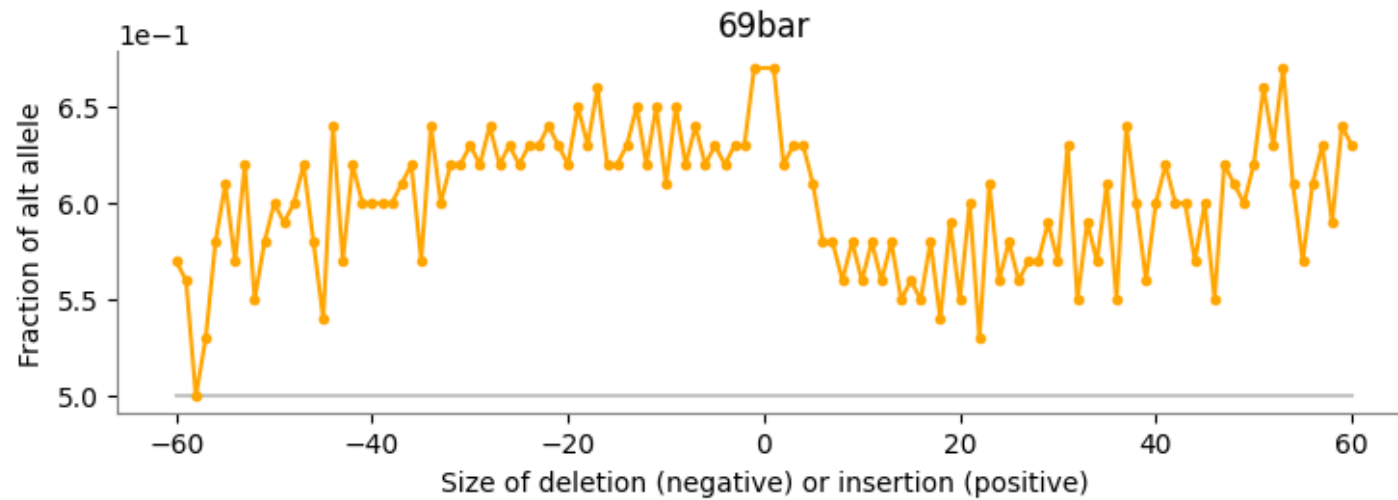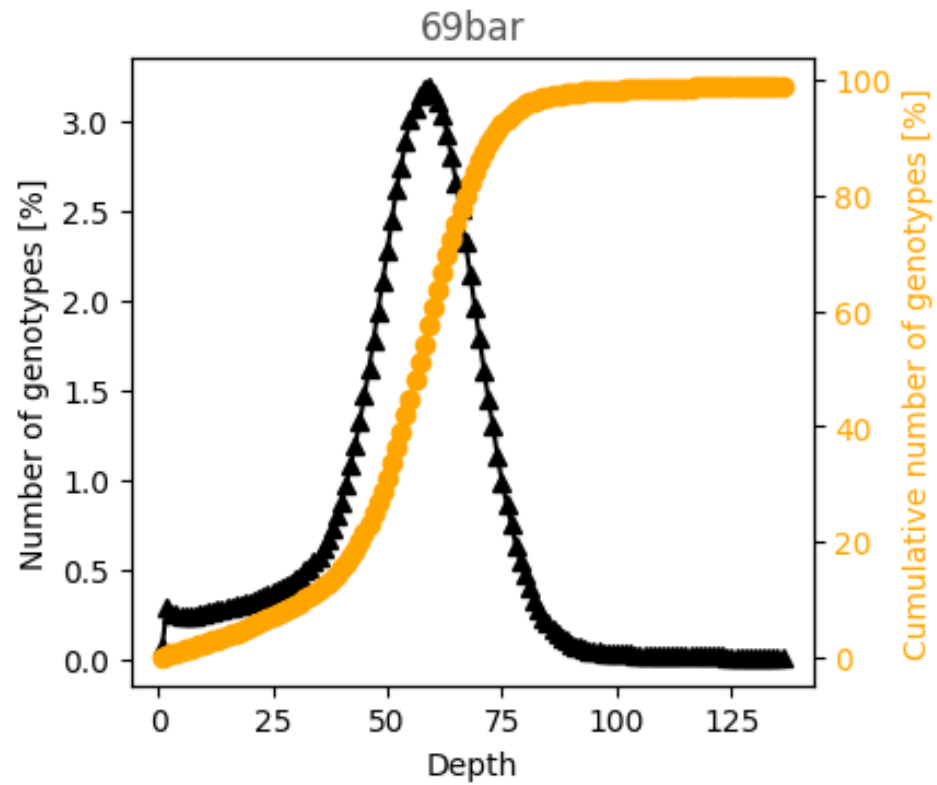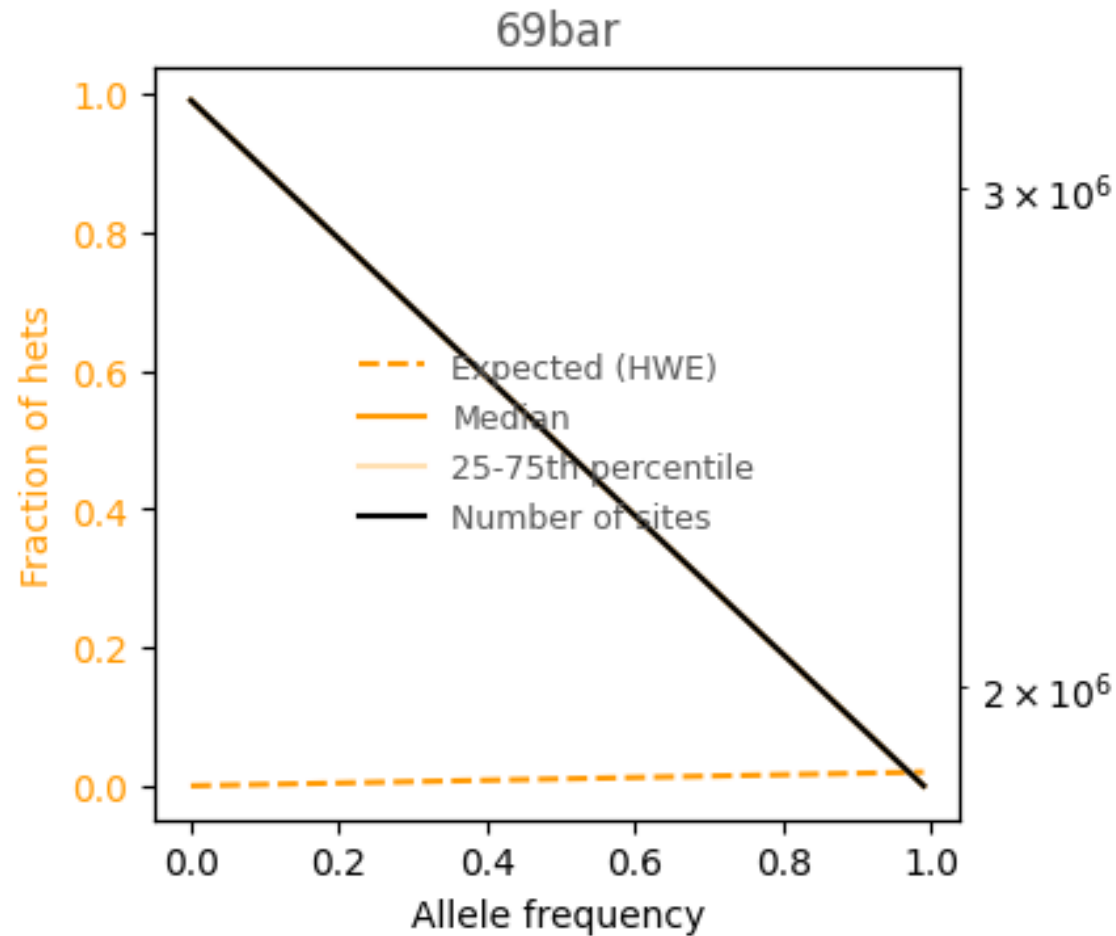# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types