Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
69sad	4,050,091	1.93	1.94	947,118	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
69sad	62.1%	1.90	66.7%	99,137	1,832

- 69sad .. `/ngc/projects2/gm/data/archive/2022/variants/snv/69sadchem-103907604173-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01241-220907_A01176_AHYN73DSX3-EXT_LAB`
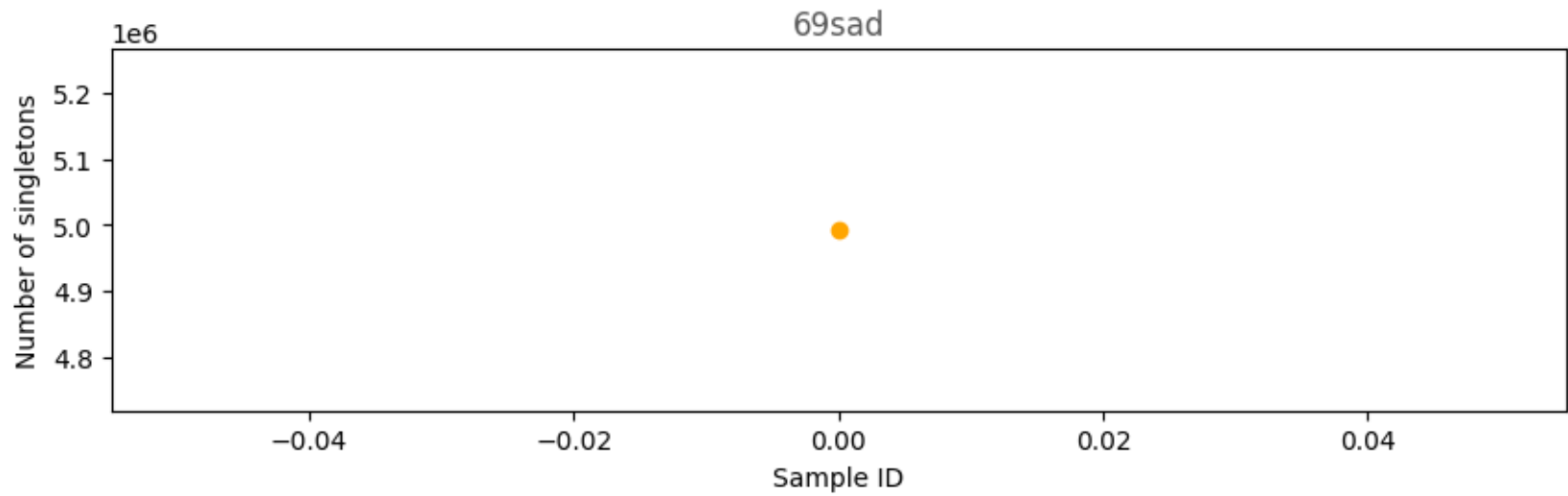 `KA_NGCWGS-NGCWGS05013_22RKG006602x01_snv_germline_raw.haplotype_caller.vcf.gz`
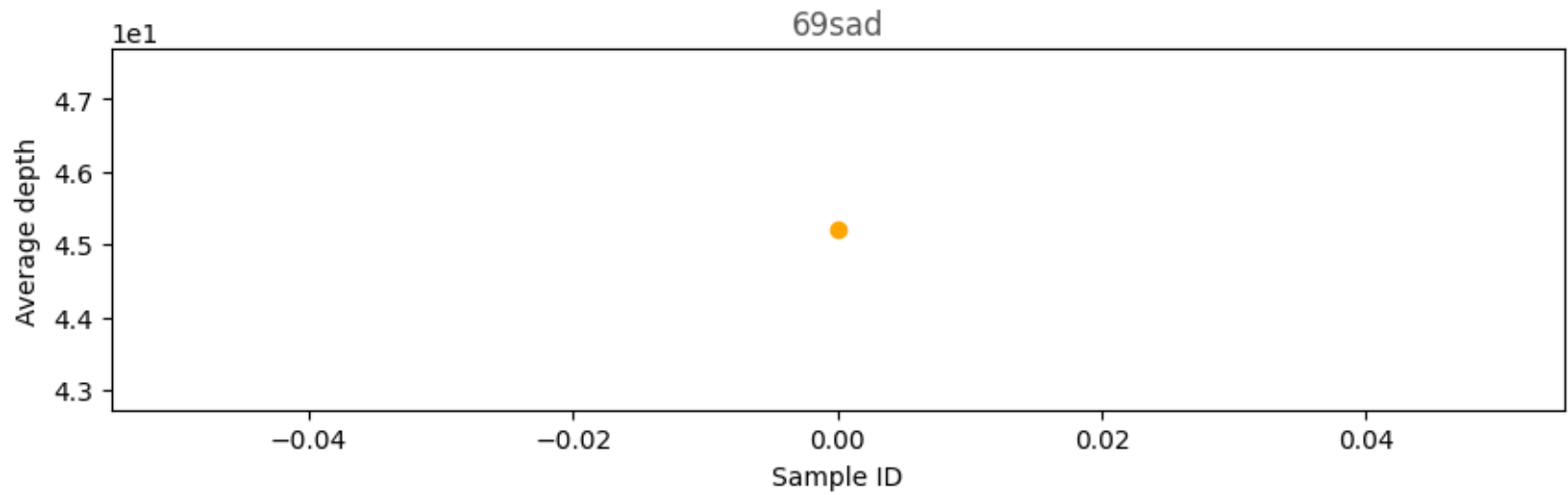
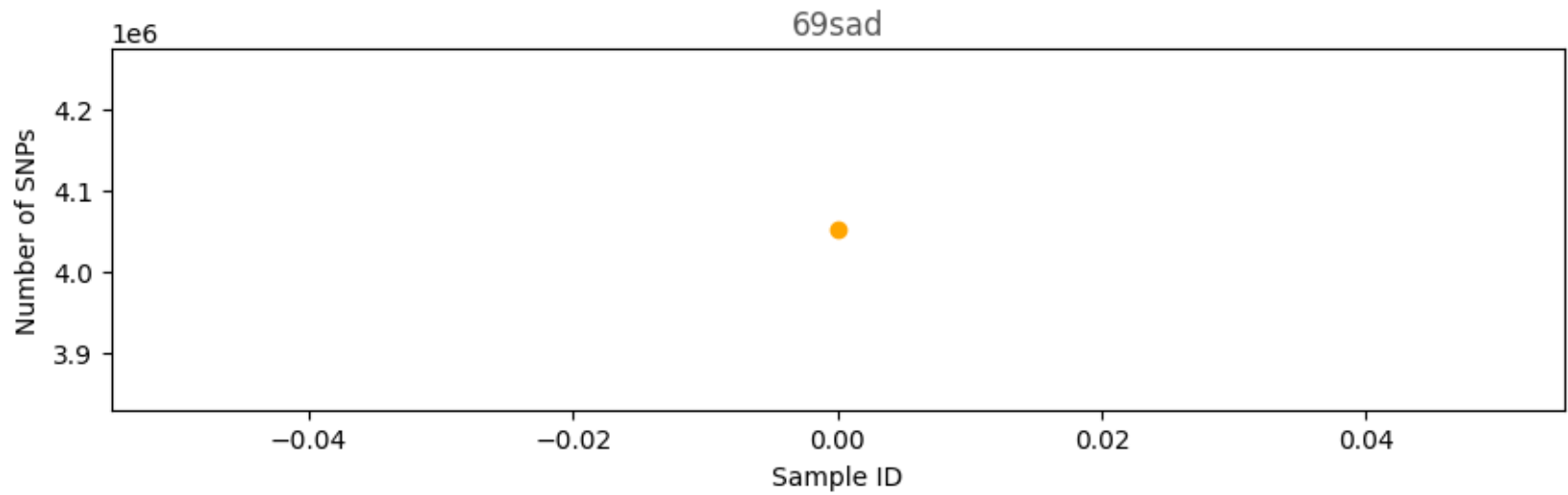# Ts/Tv by sample
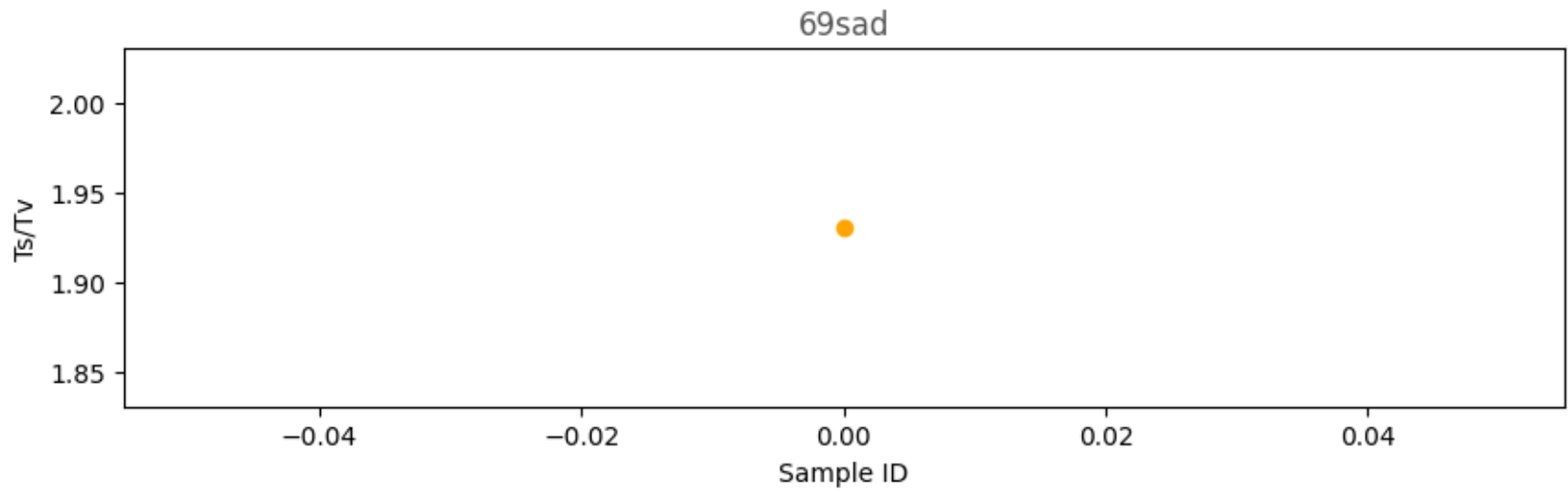
Hets vs non-ref Homs by sample
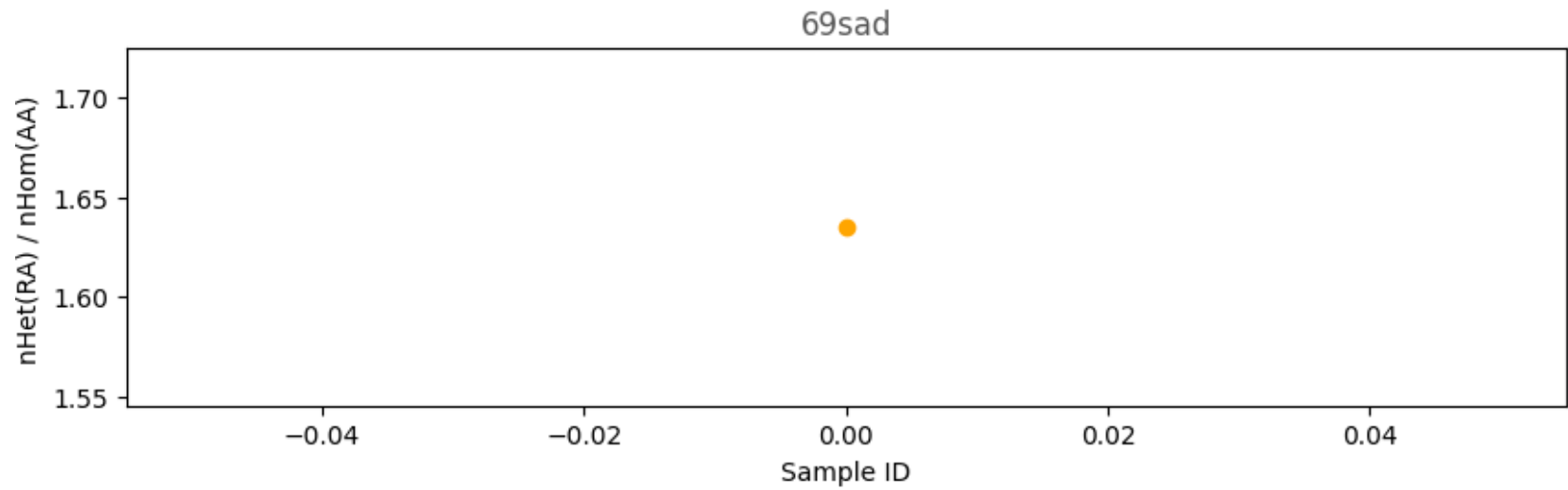
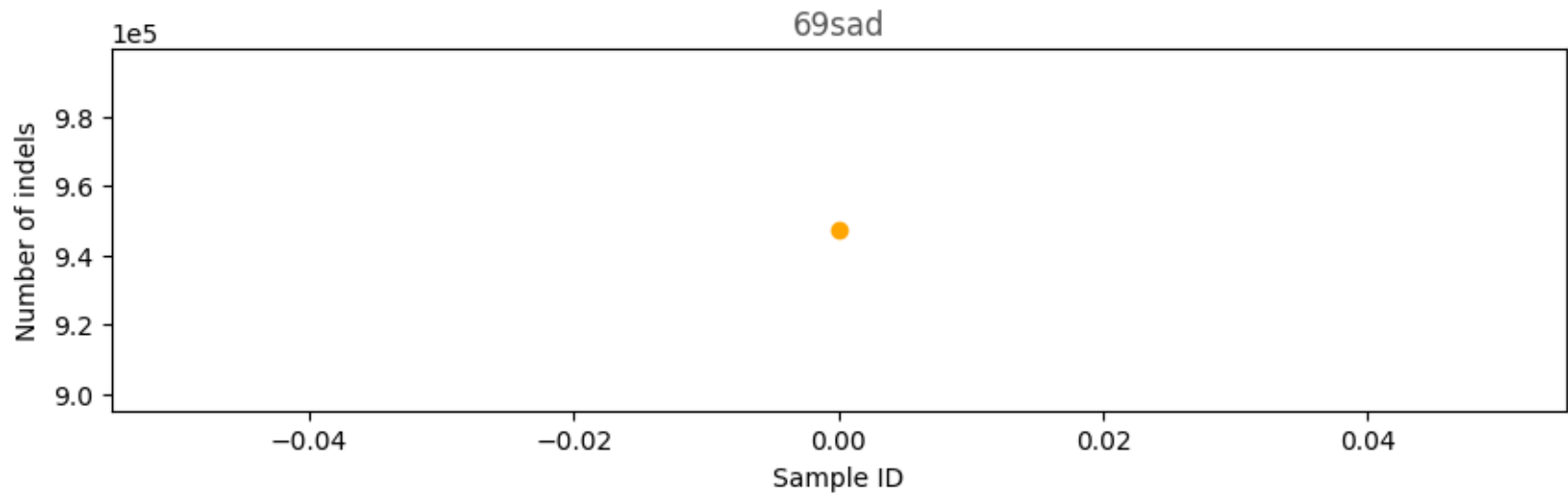# Singletons by sample (hets and homs)
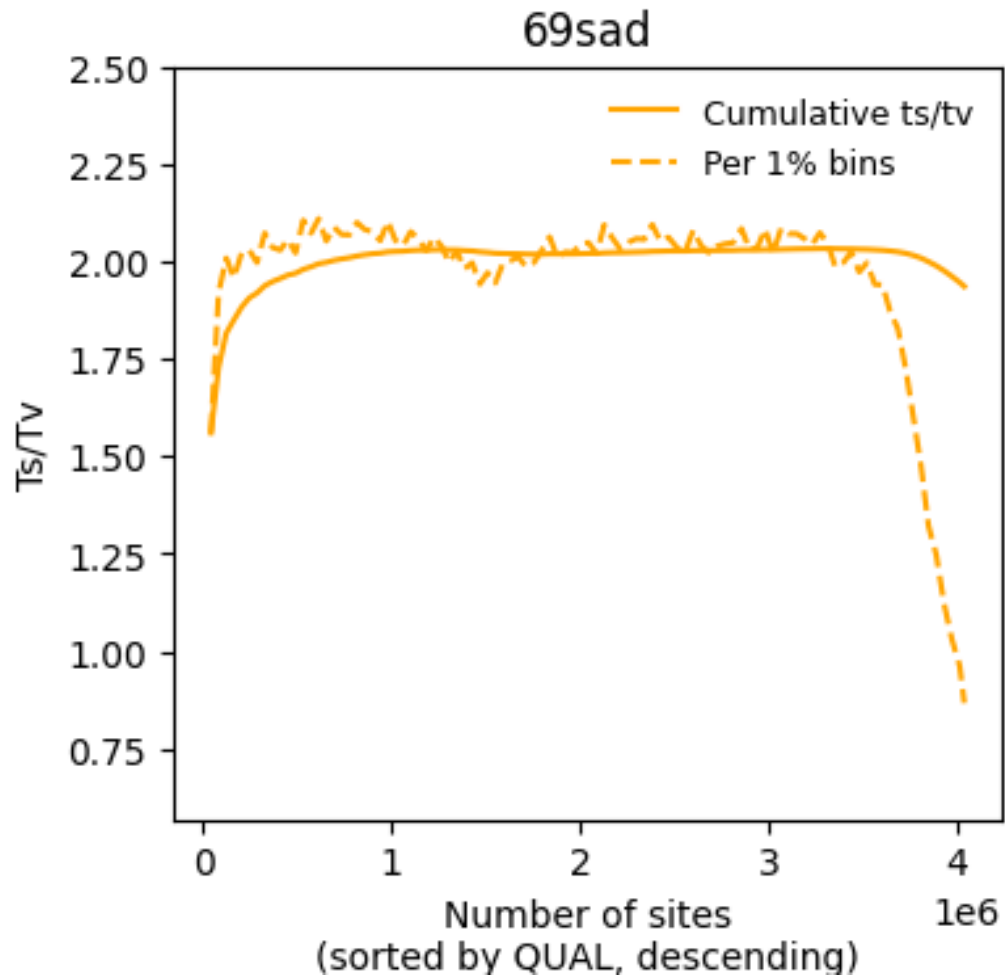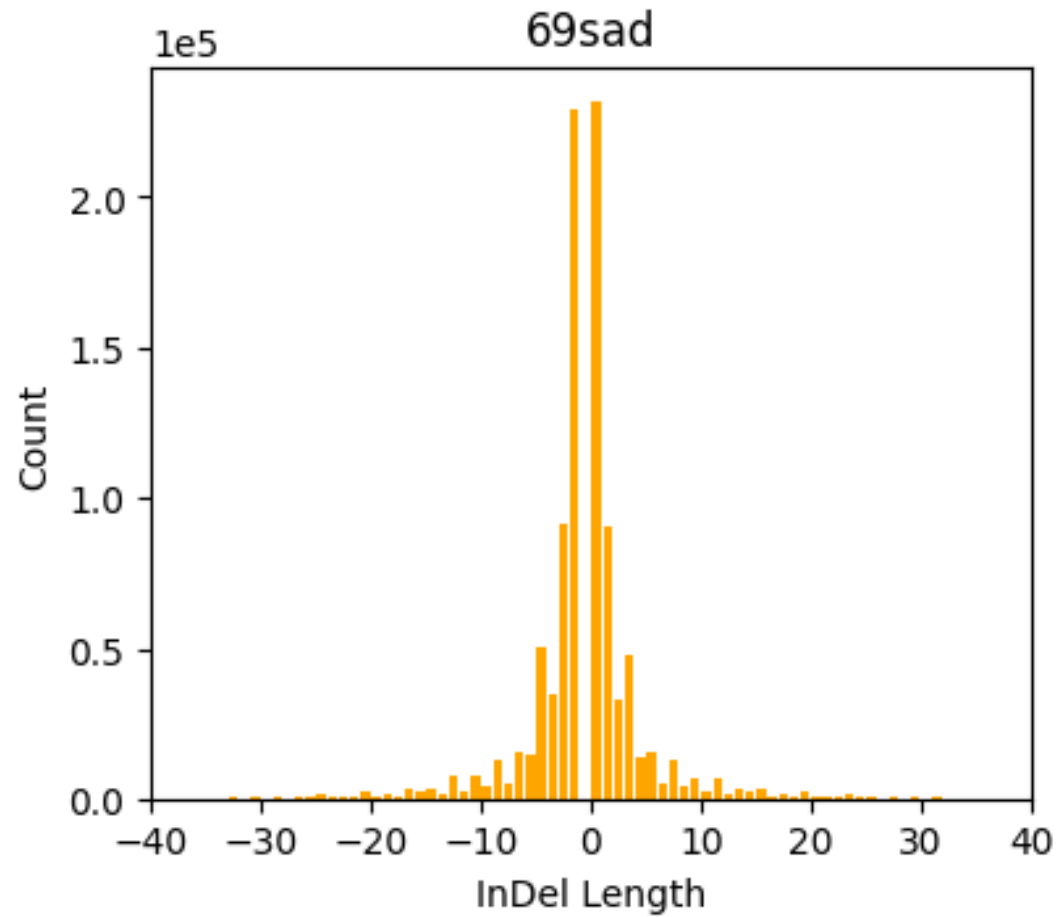
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

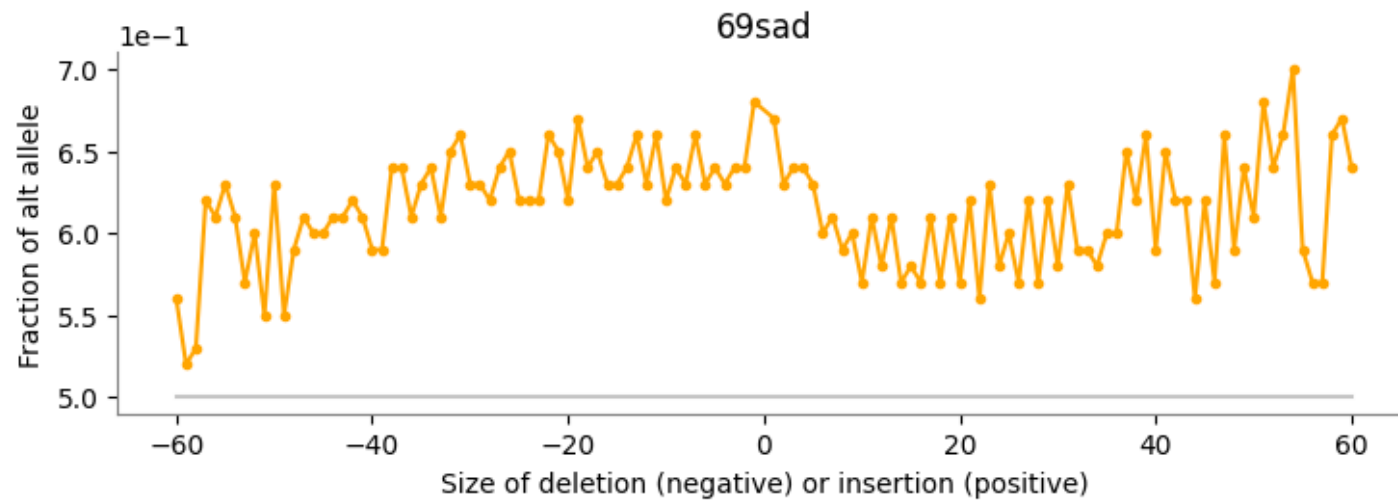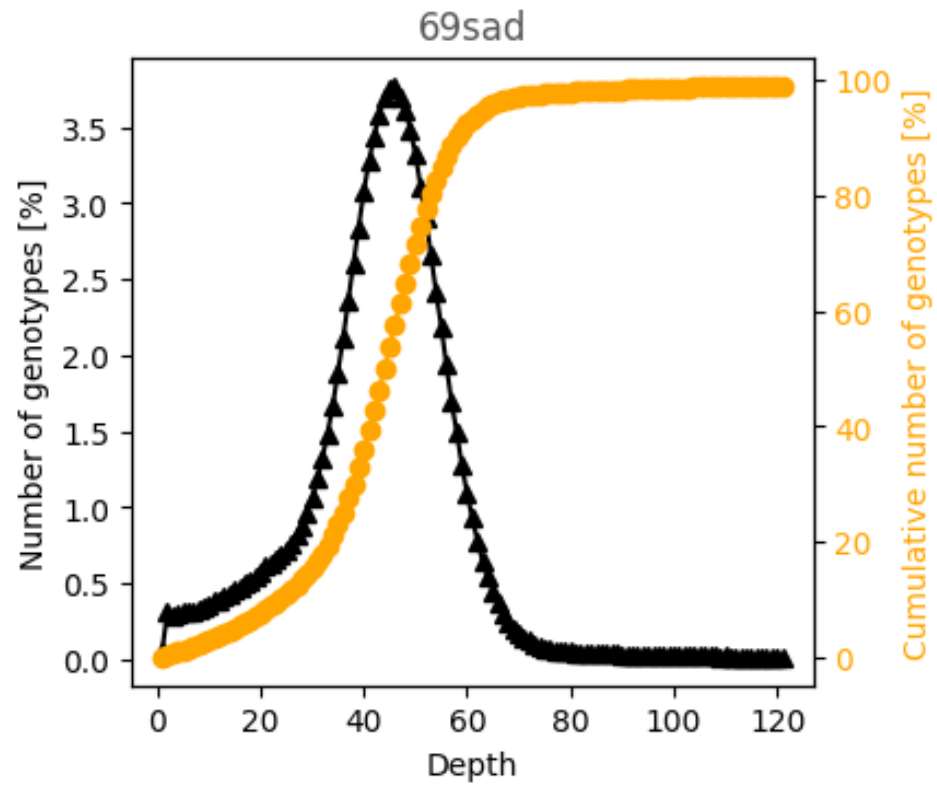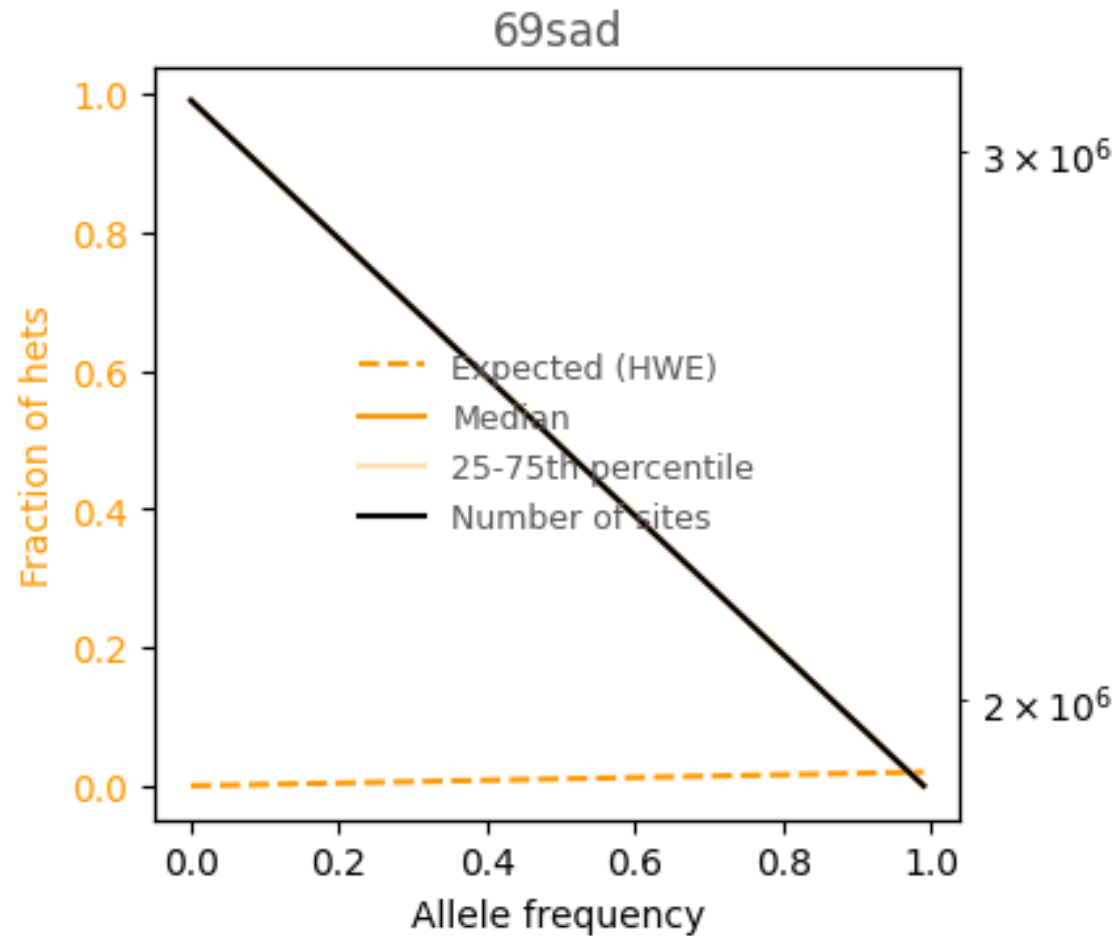# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types