Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
69sto	4,107,831	1.93	1.94	963,721	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
69sto	63.2%	1.90	68.8%	105,712	2,081

- 69sto .. `/ngc/projects2/gm/data/archive/2022/variants/snv/69stovedf-103827616969-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220119_A01176_BH2CT5DSX3-RHGM_LABKA_WGS-`
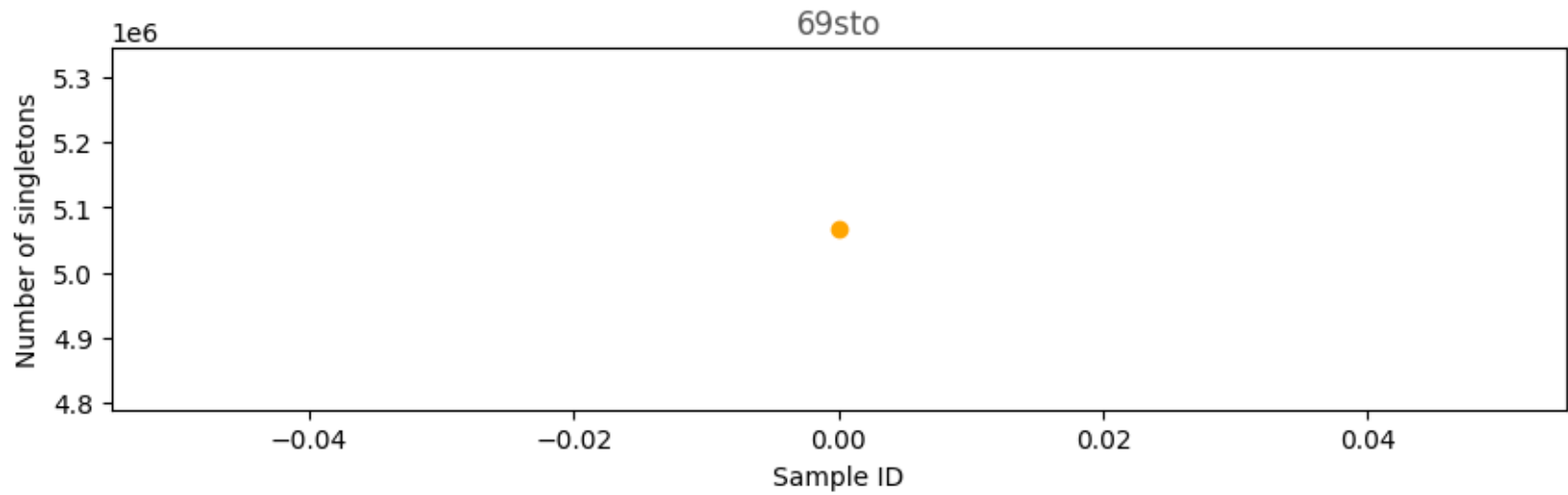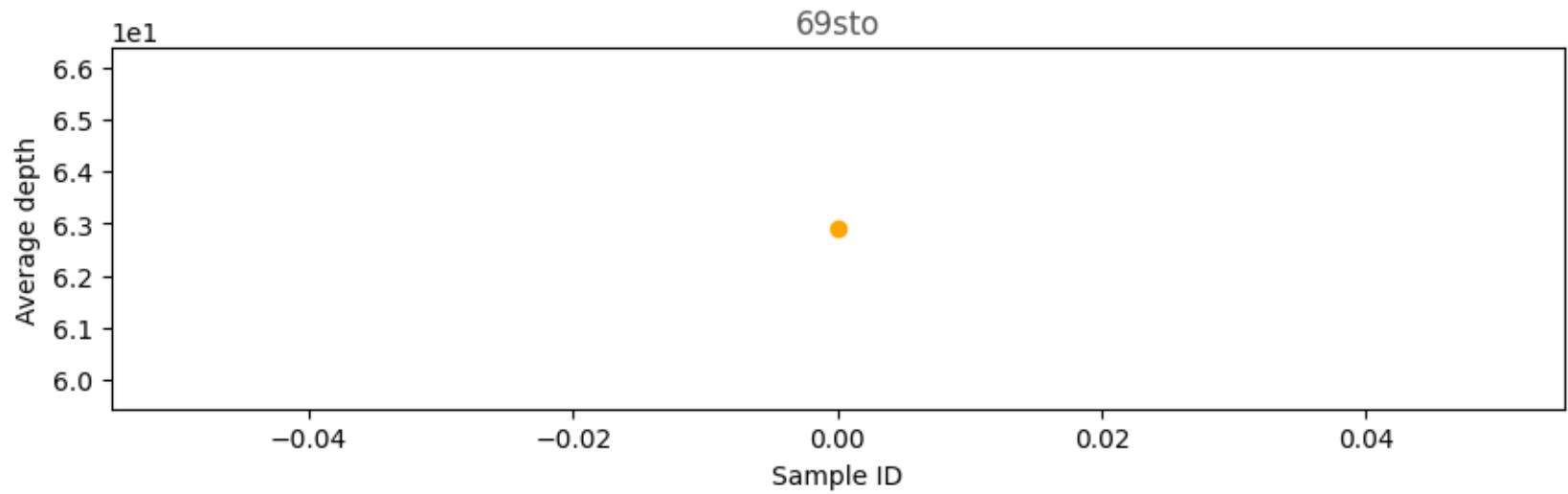 `WGS03696_snv_germline_raw.haplotype_caller.vcf.gz`
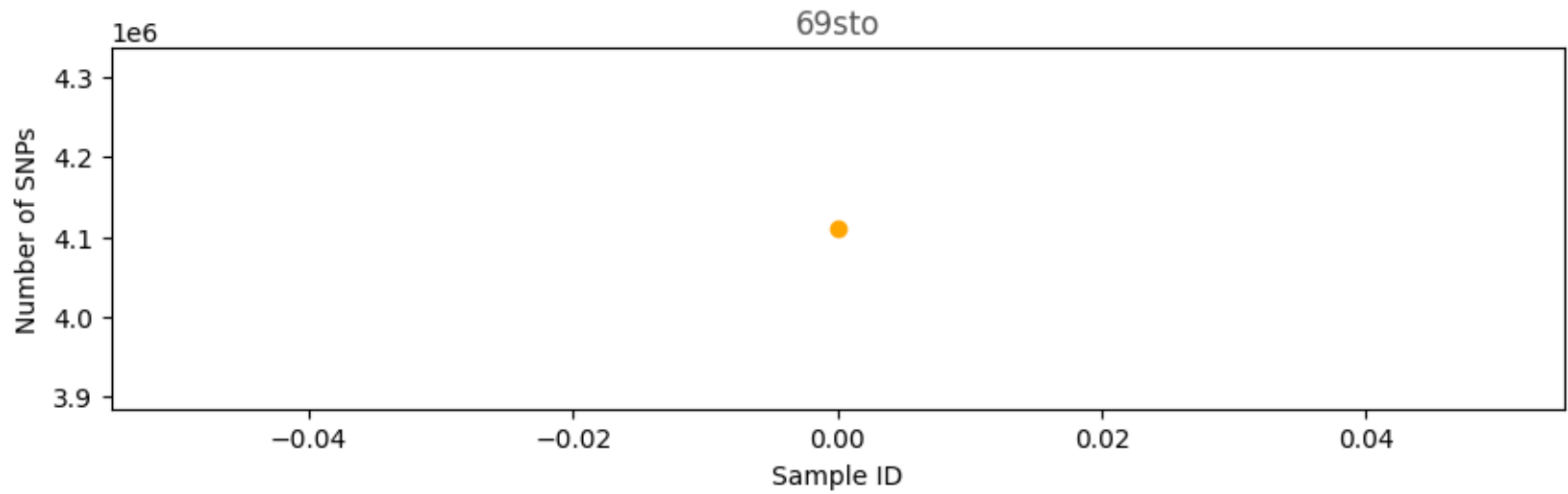
Ts/Tv by sample
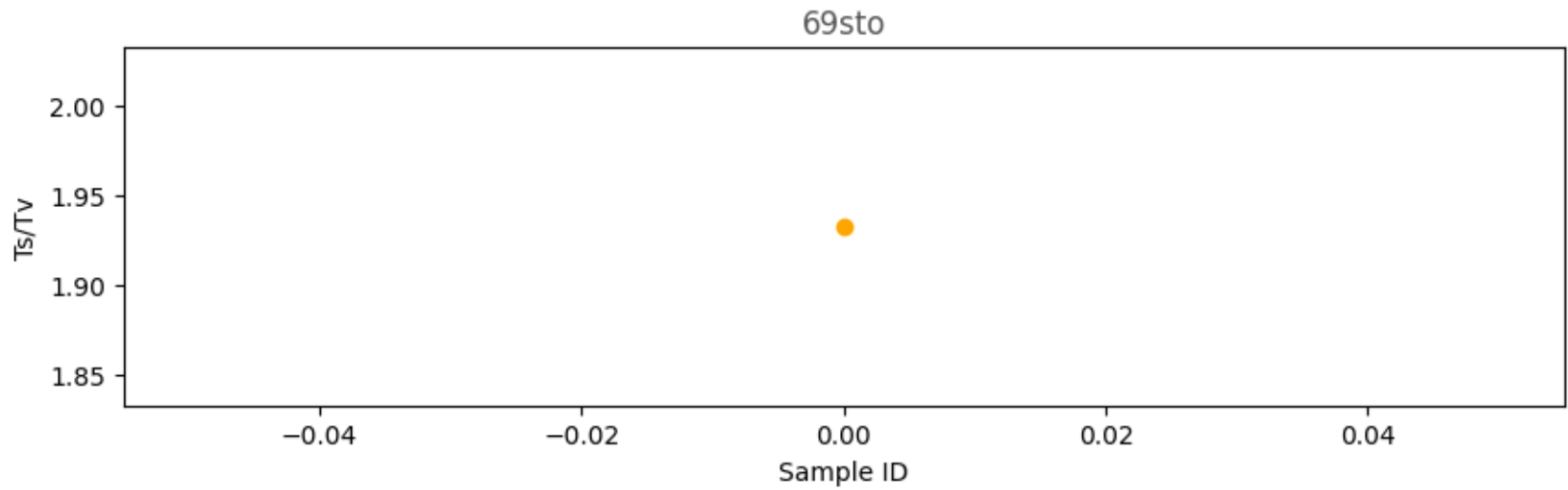
Hets vs non-ref Homs by sample
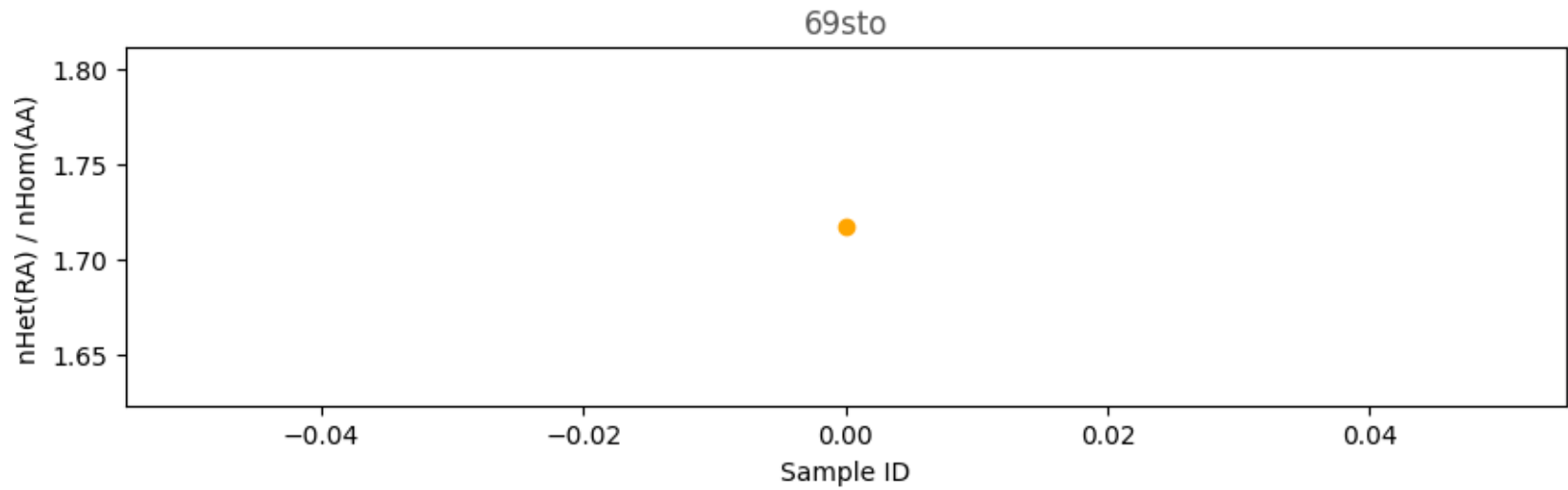
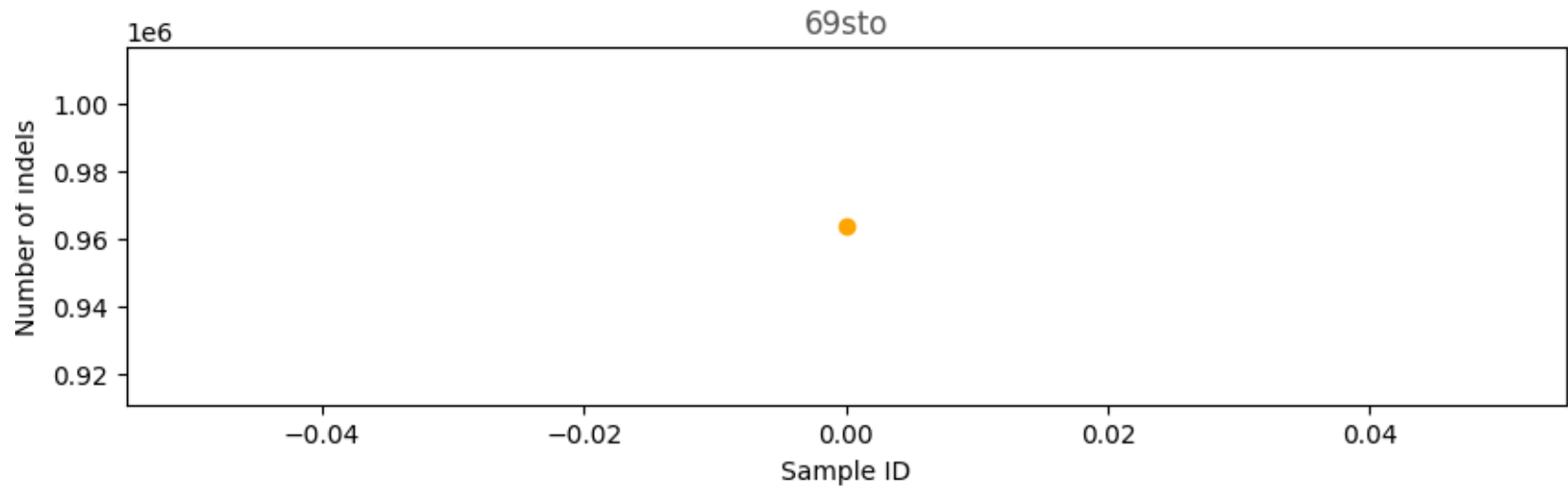# Singletons by sample (hets and homs)
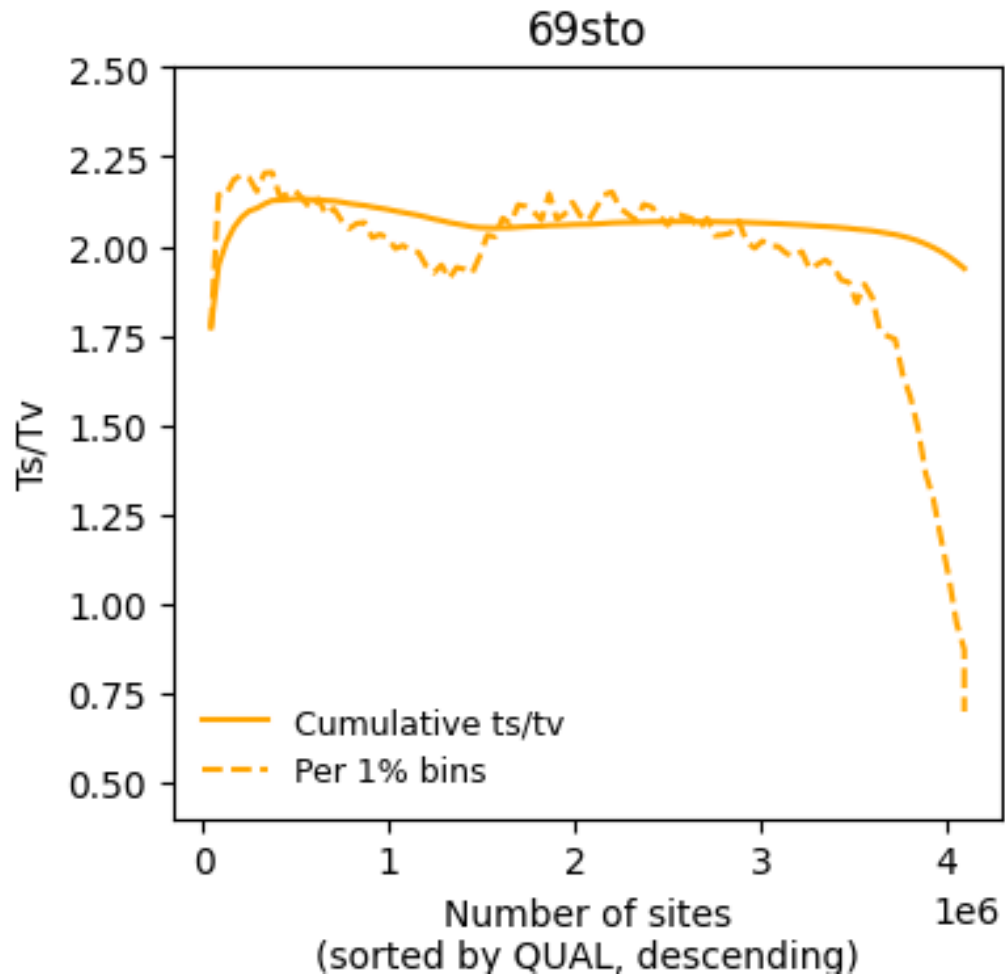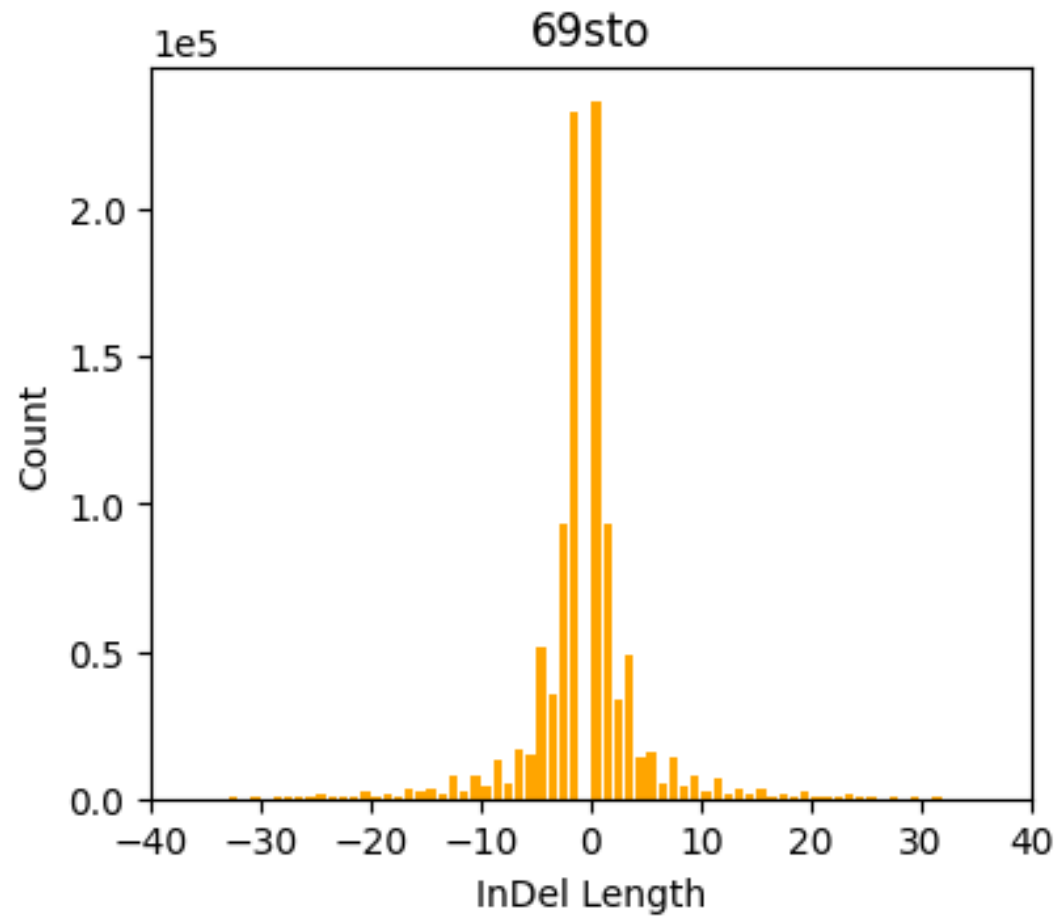
Average depth by sample

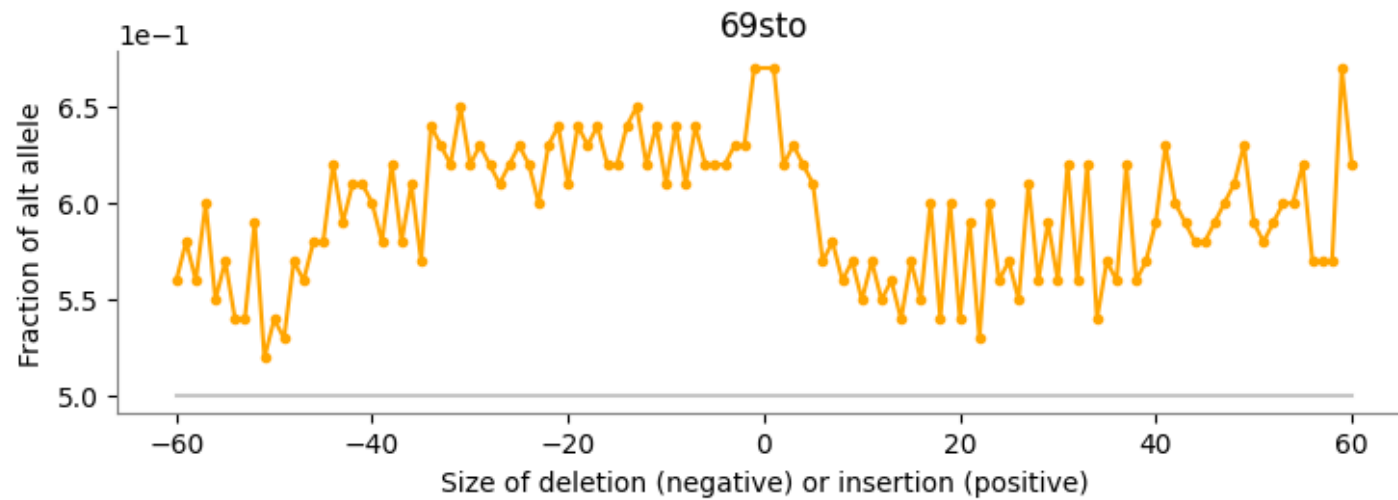# Number of SNPs by sample

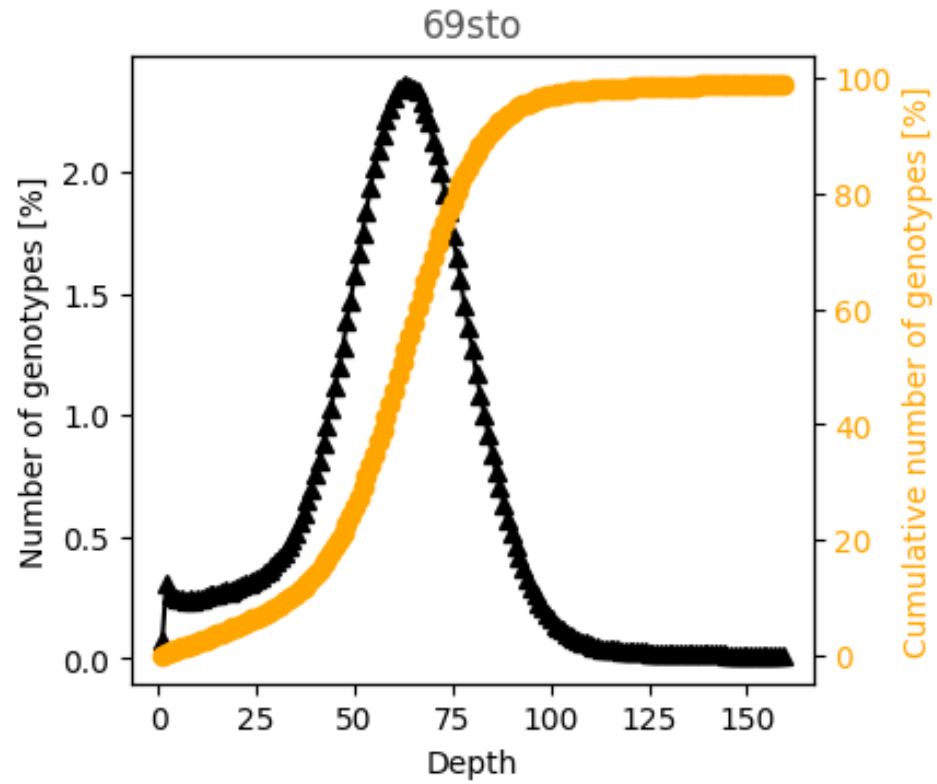# Number of indels by sample

Ts/Tv stratified by QUAL

69sto

Indel distribution

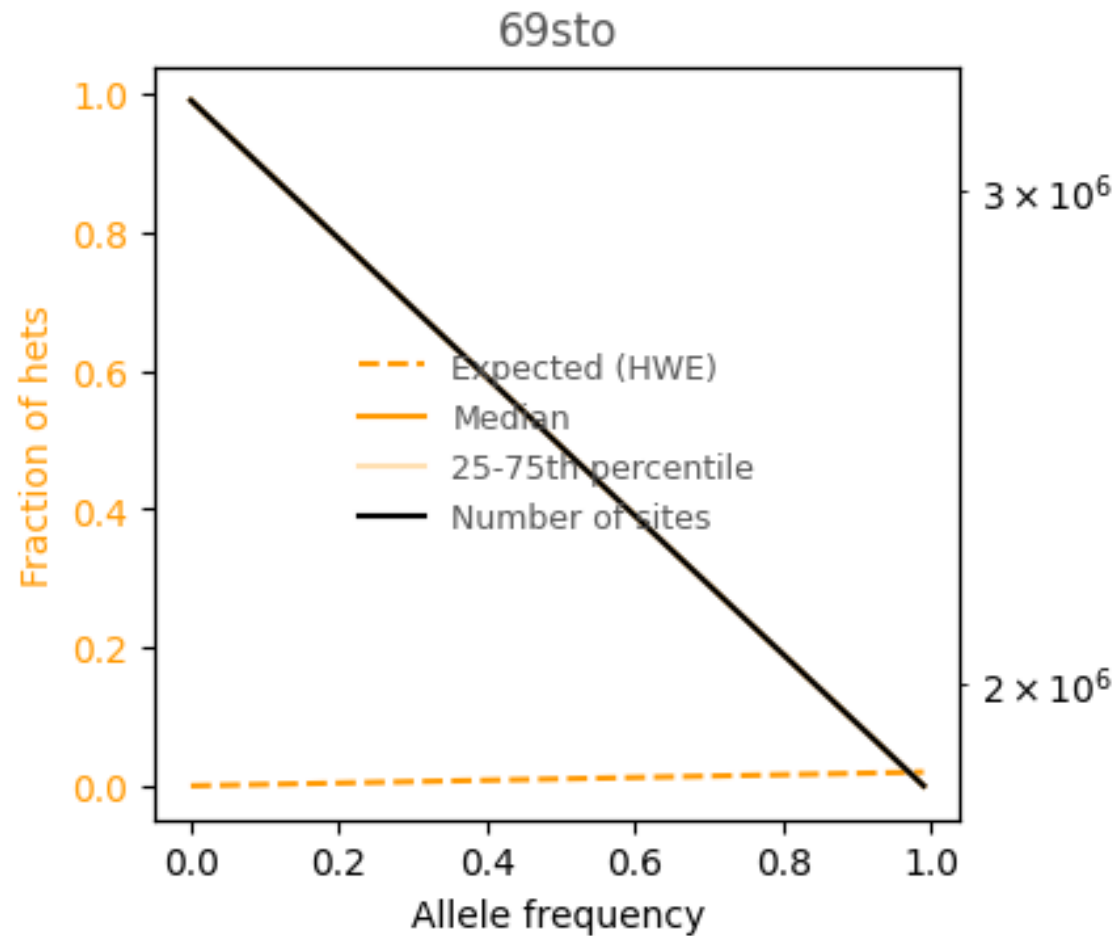# Fraction of alternate indel allele

Depth distribution

69sto

Number of HETs by AF

Substitution types