Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
66ato	4,079,123	1.94	1.94	944,264	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
66ato	63.1%	1.91	67.7%	97,698	2,245

- 66ato .. `/ngc/projects2/gm/data/archive/2022/variants/snv/66atoniof-110294830190-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01978-221012_A00559_BH2CMWDSX5-EXT_LAB`
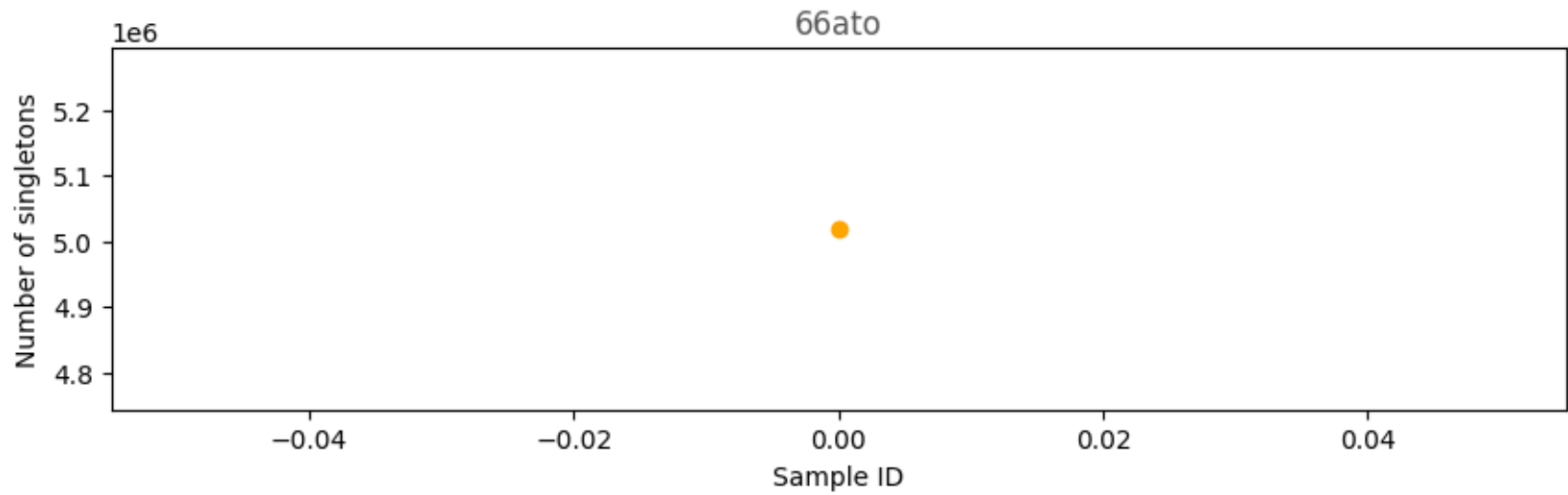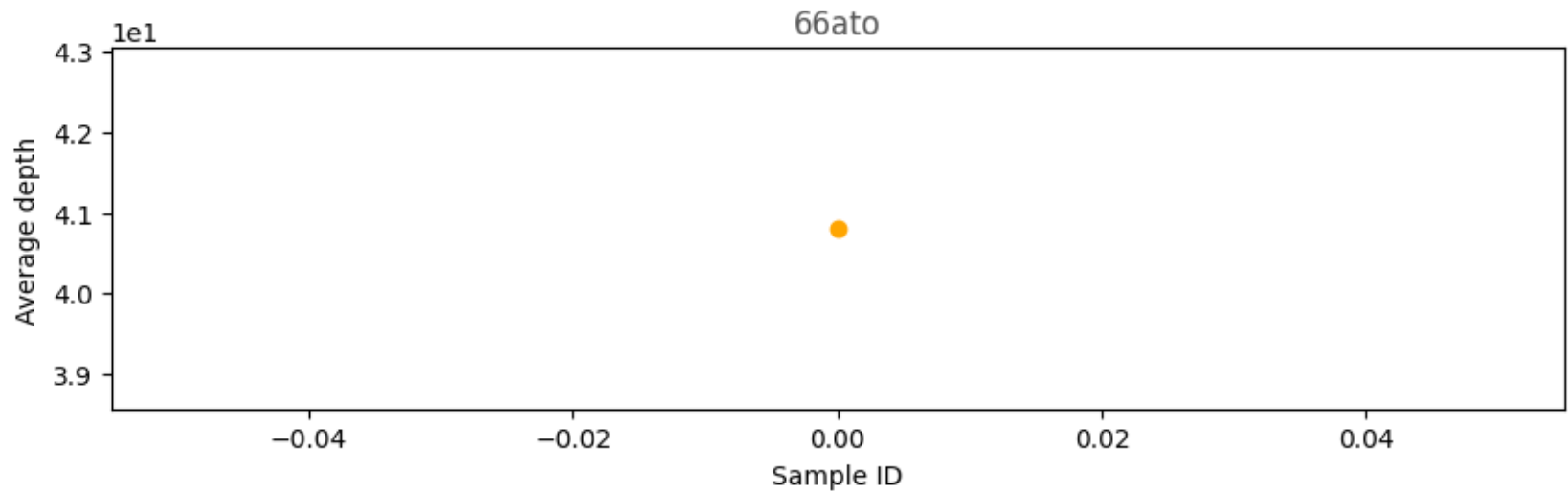 `KA_NGCWGS-NGCWGS05796_snv_germline_raw.haplotype_caller.vcf.gz`
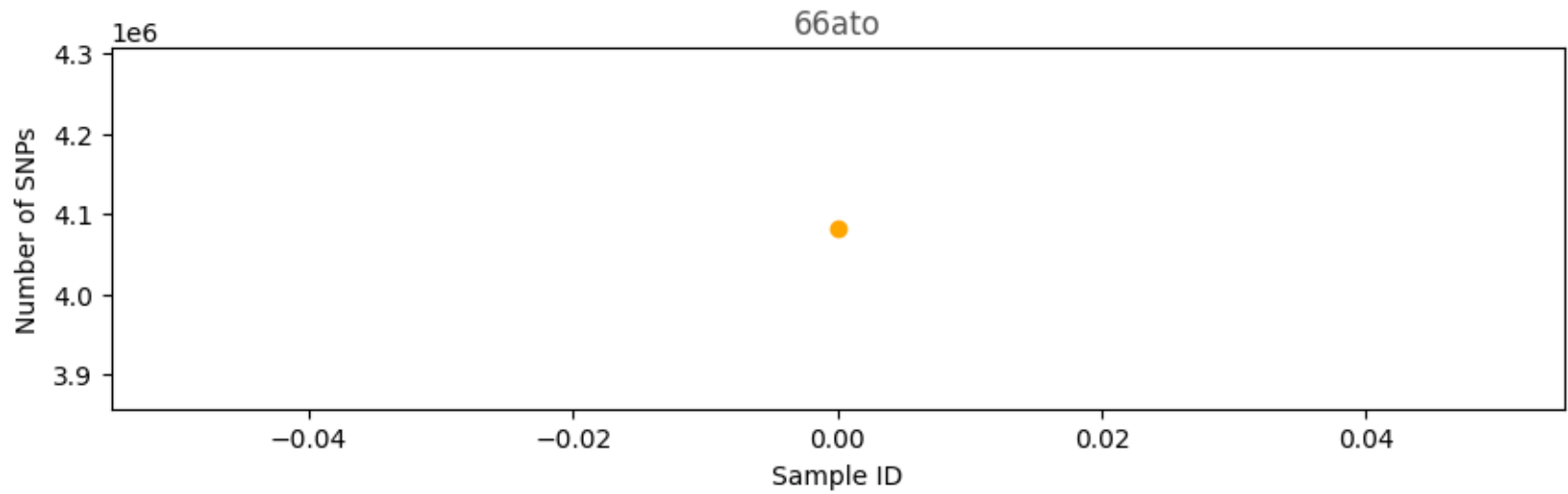
Ts/Tv by sample
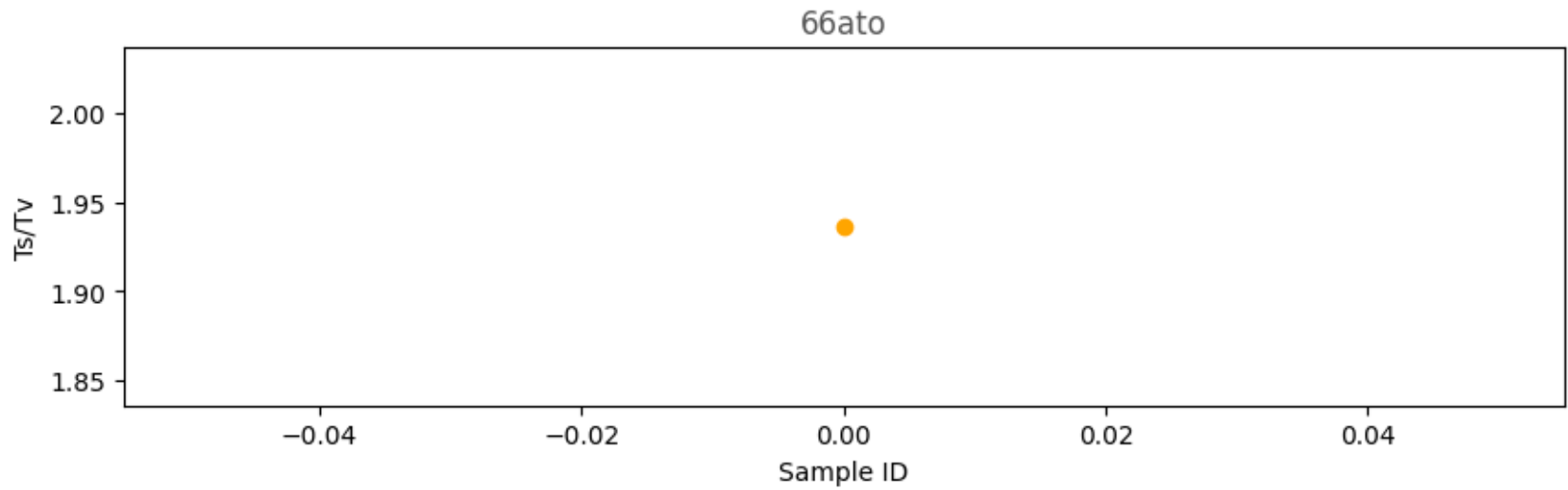
Hets vs non-ref Homs by sample
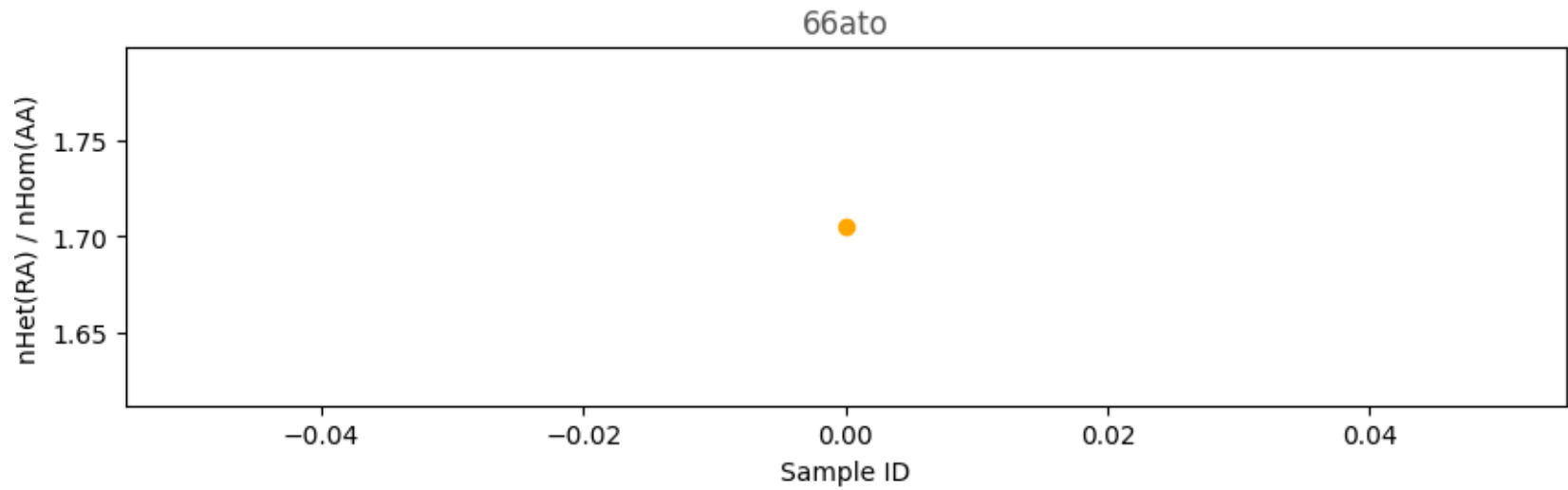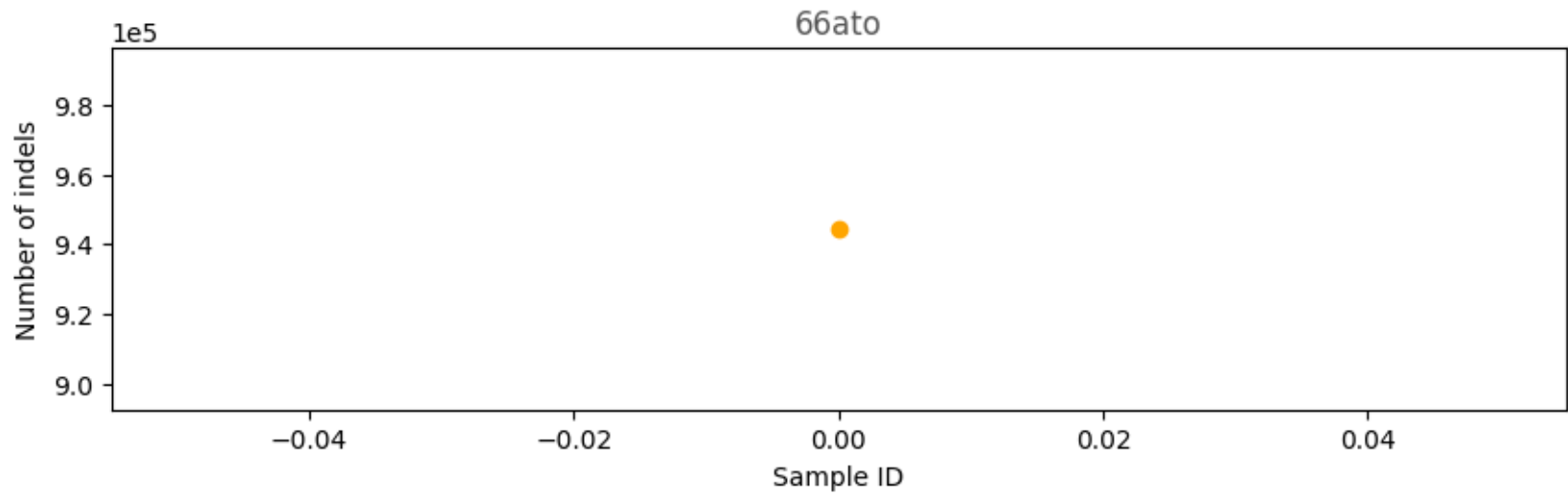
Singletons by sample (hets and homs)
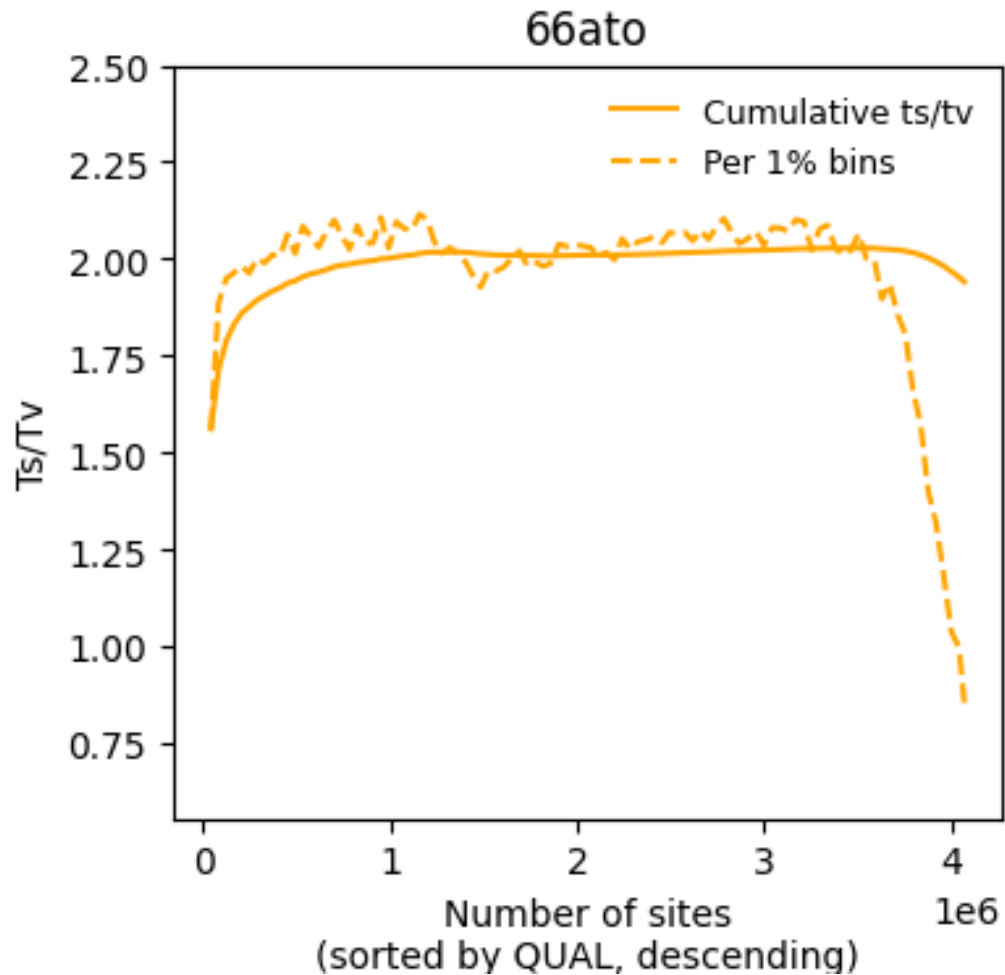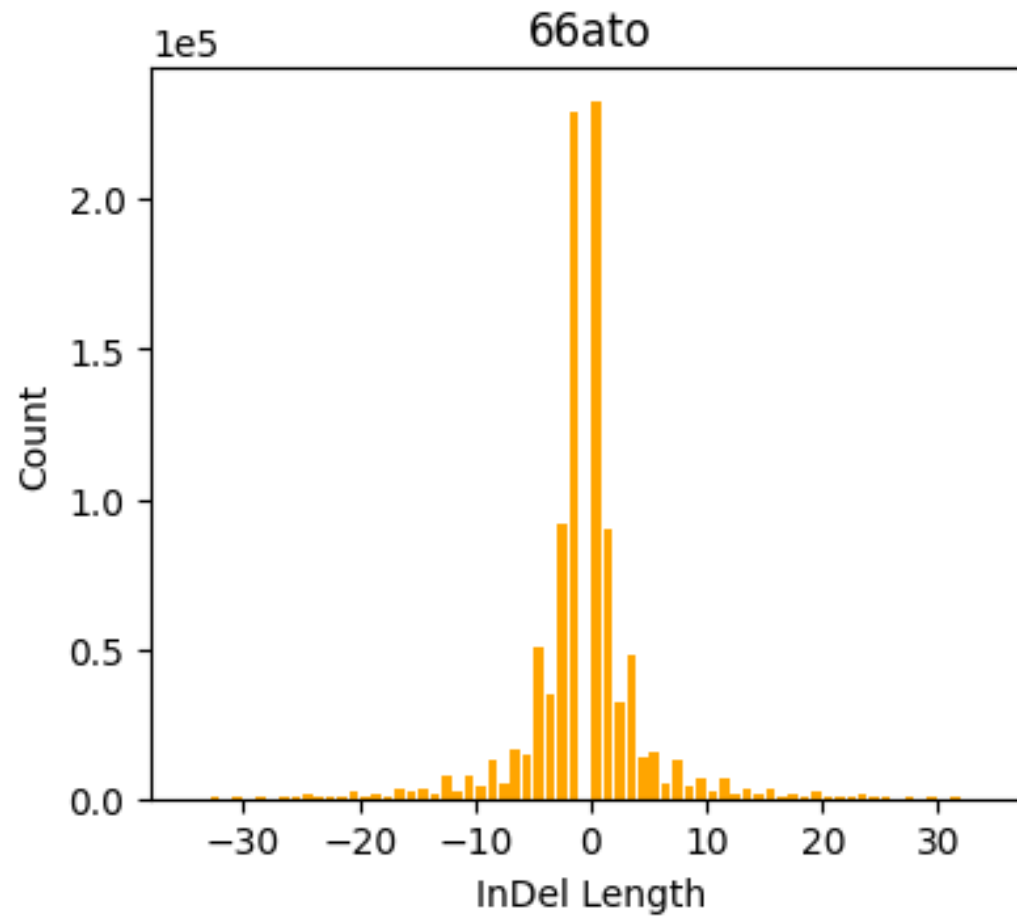
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

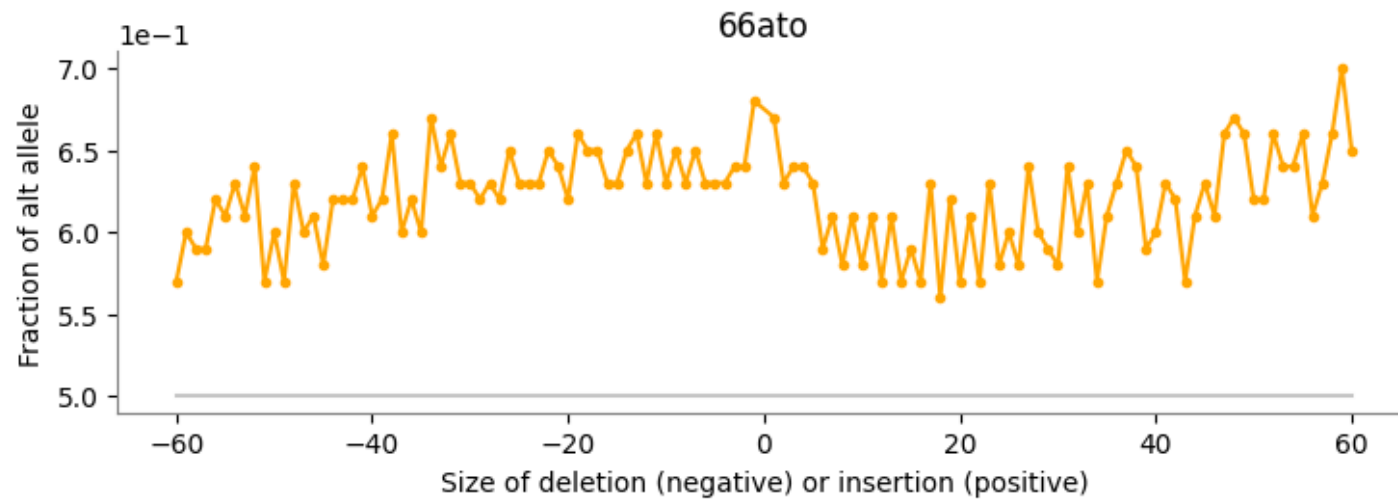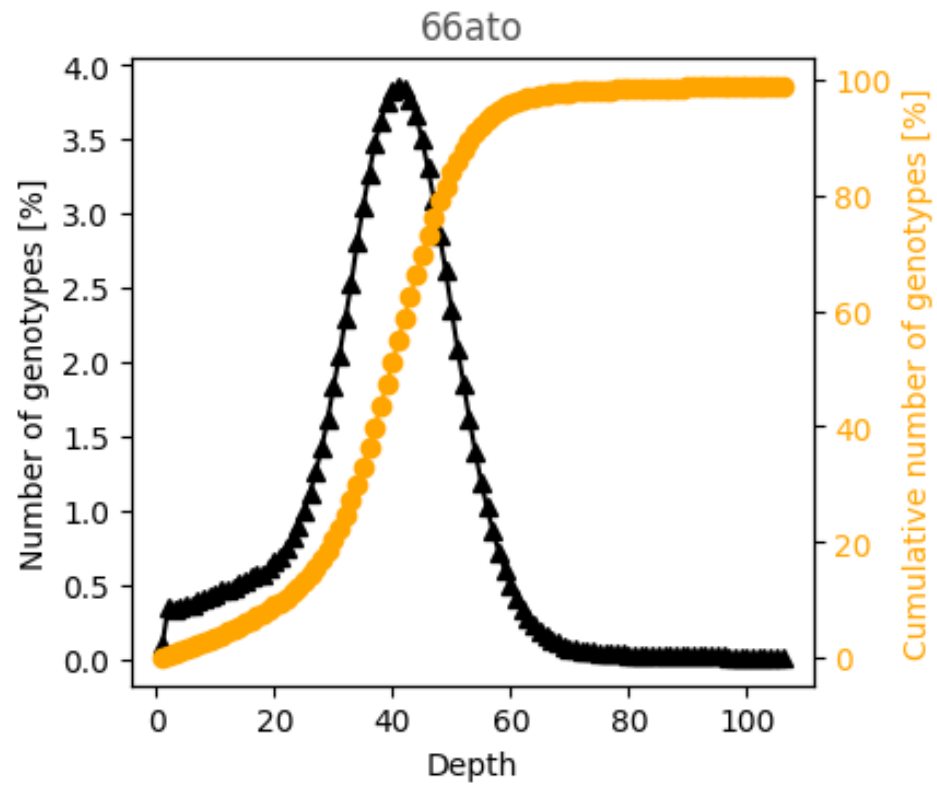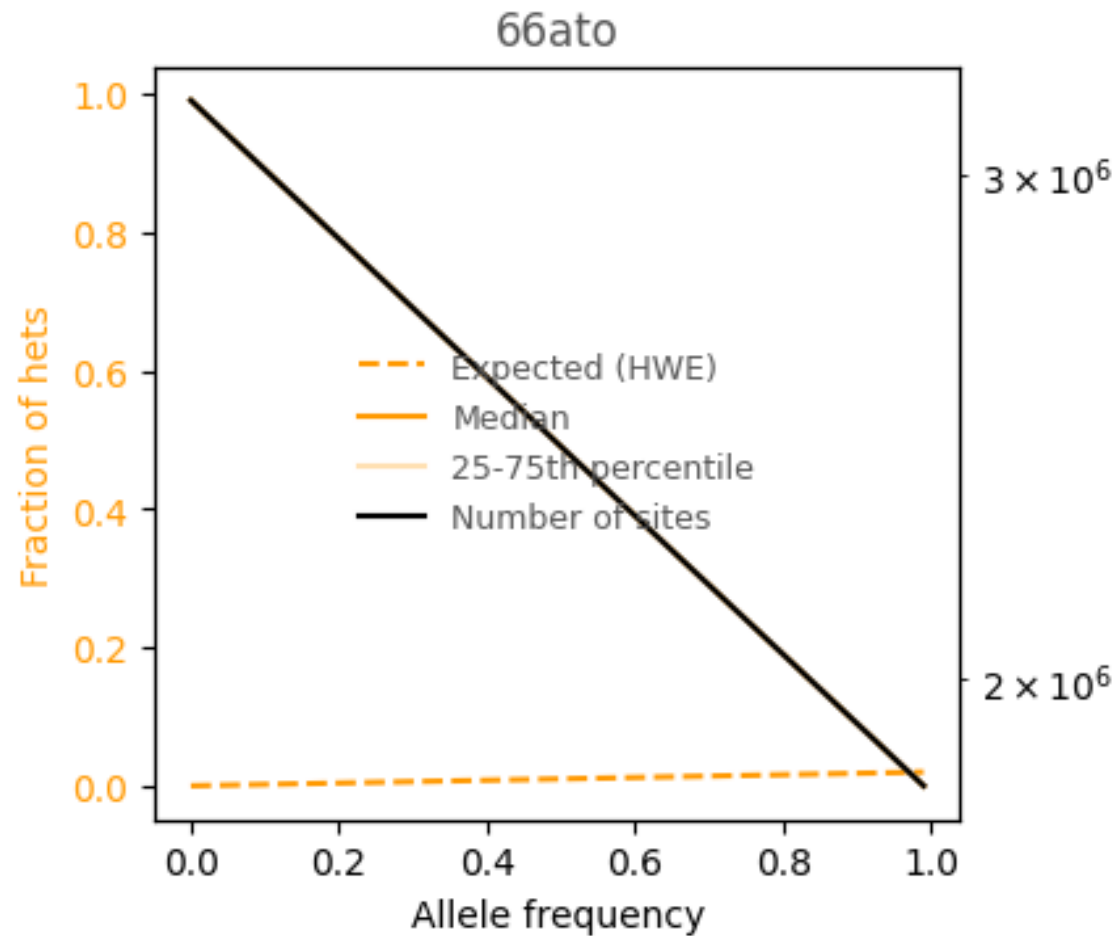# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types