Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
00kis	4,134,284	1.92	1.93	970,522	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
00kis	63.1%	1.90	68.8%	108,613	2,336

- 00kis .. `/ngc/projects2/gm/data/archive/2022/variants/snv/00kistfrf-103914475588-Normal_B`
    `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01762-220928_A01961_AH35JJDSX5-EXT_LAB`
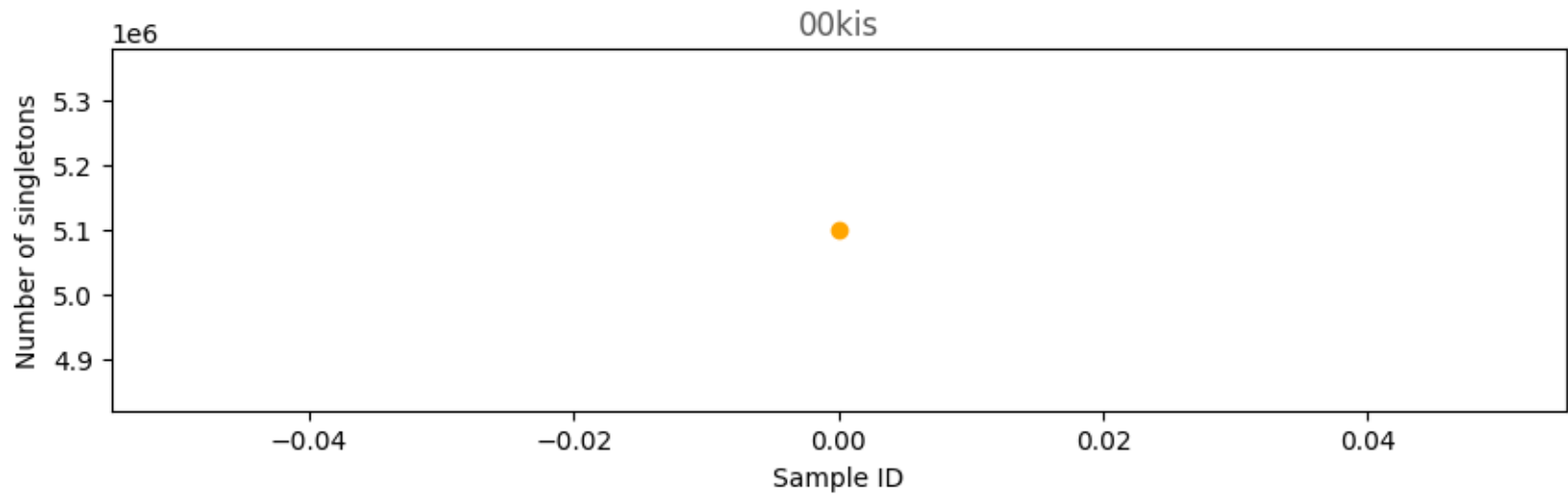   `KA_NGCWGS-NGCWGS05626_22RKG013369_snv_germline_raw.haplotype_caller.vcf.gz`
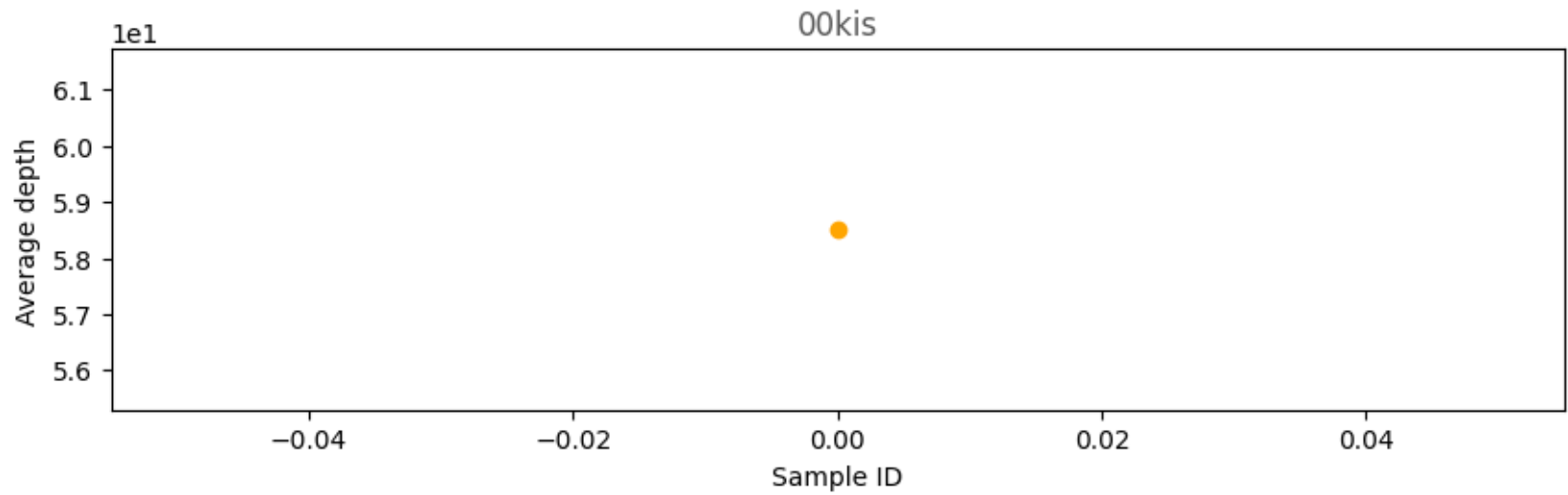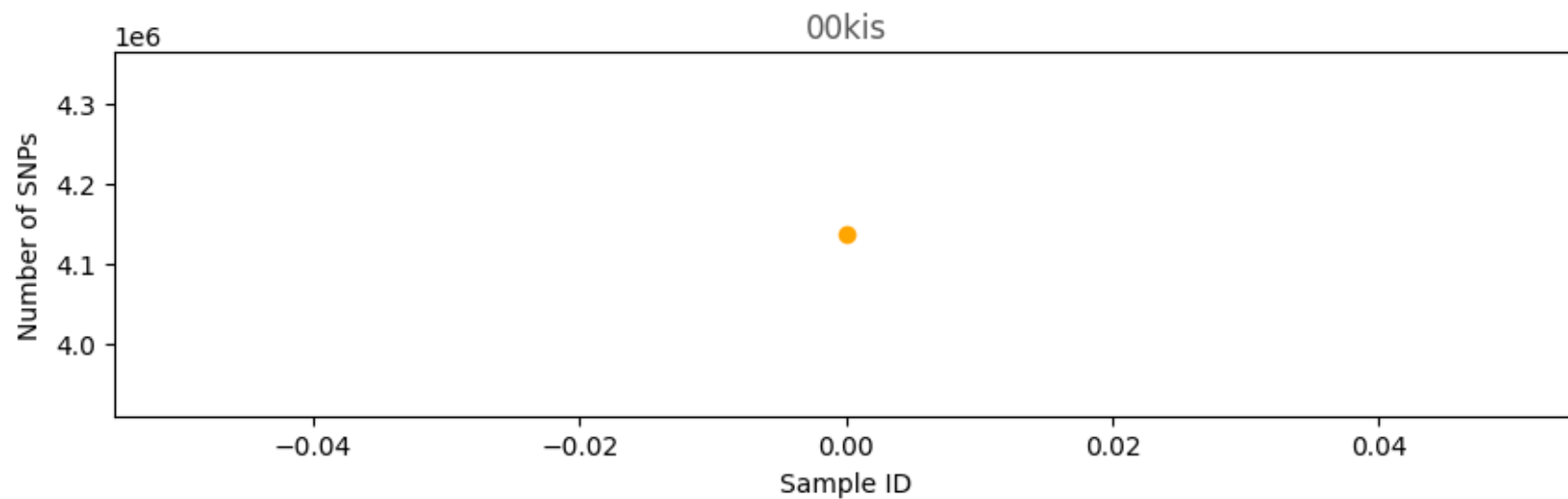
Ts/Tv by sample
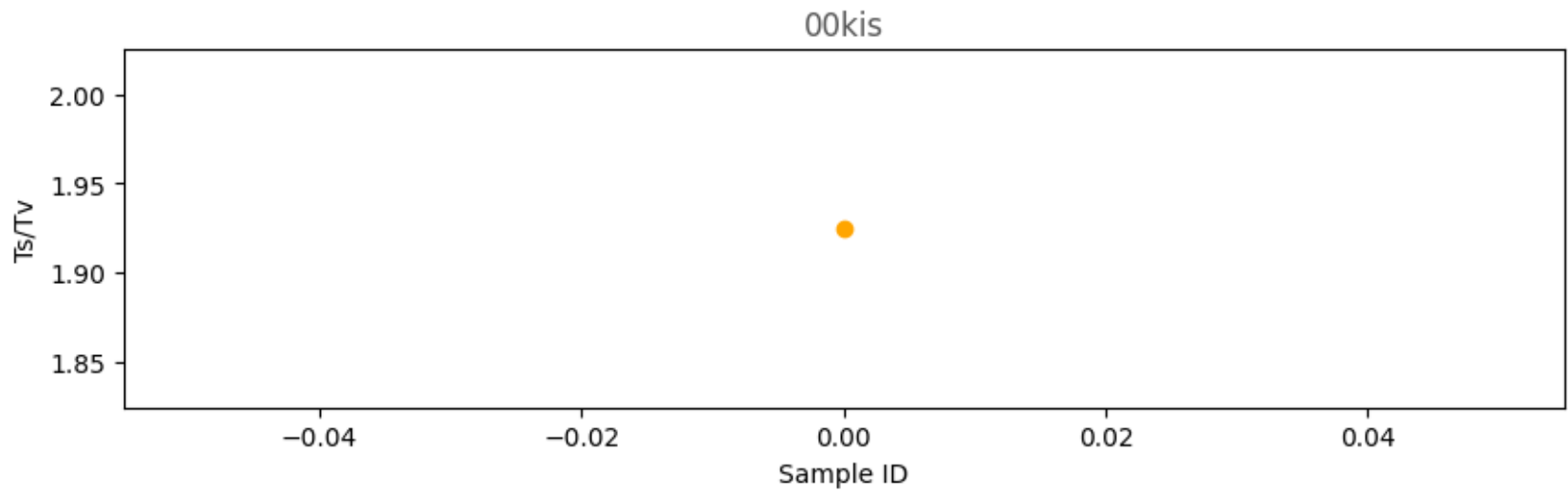
Hets vs non-ref Homs by sample
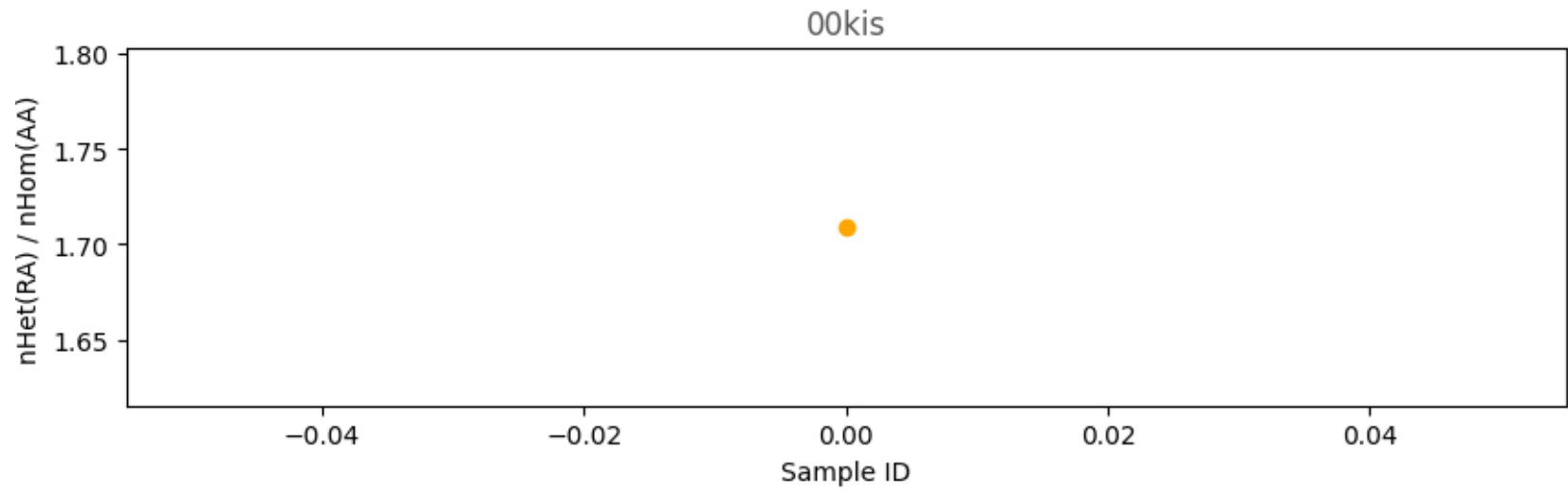
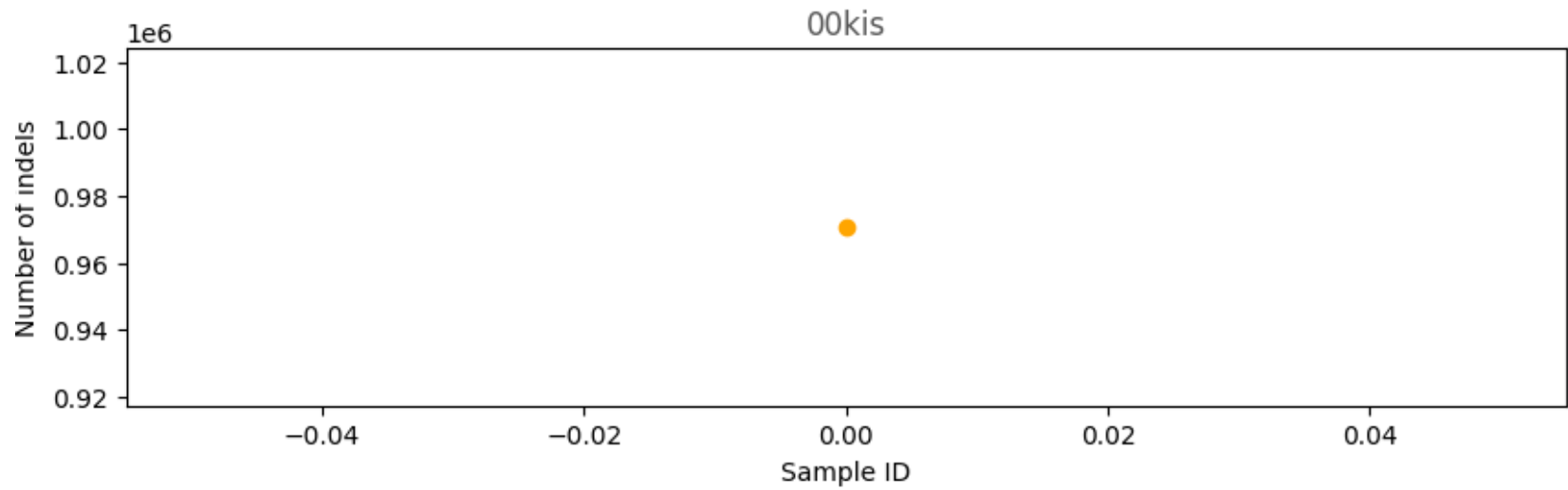# Singletons by sample (hets and homs)
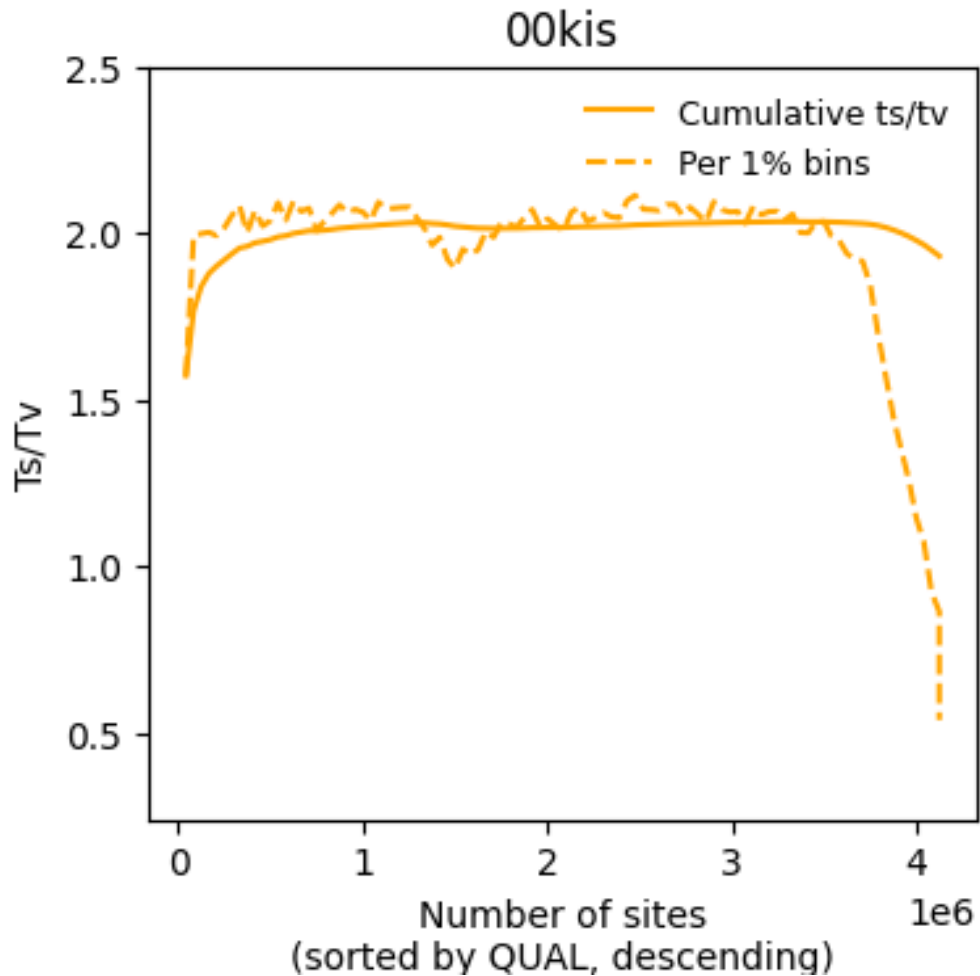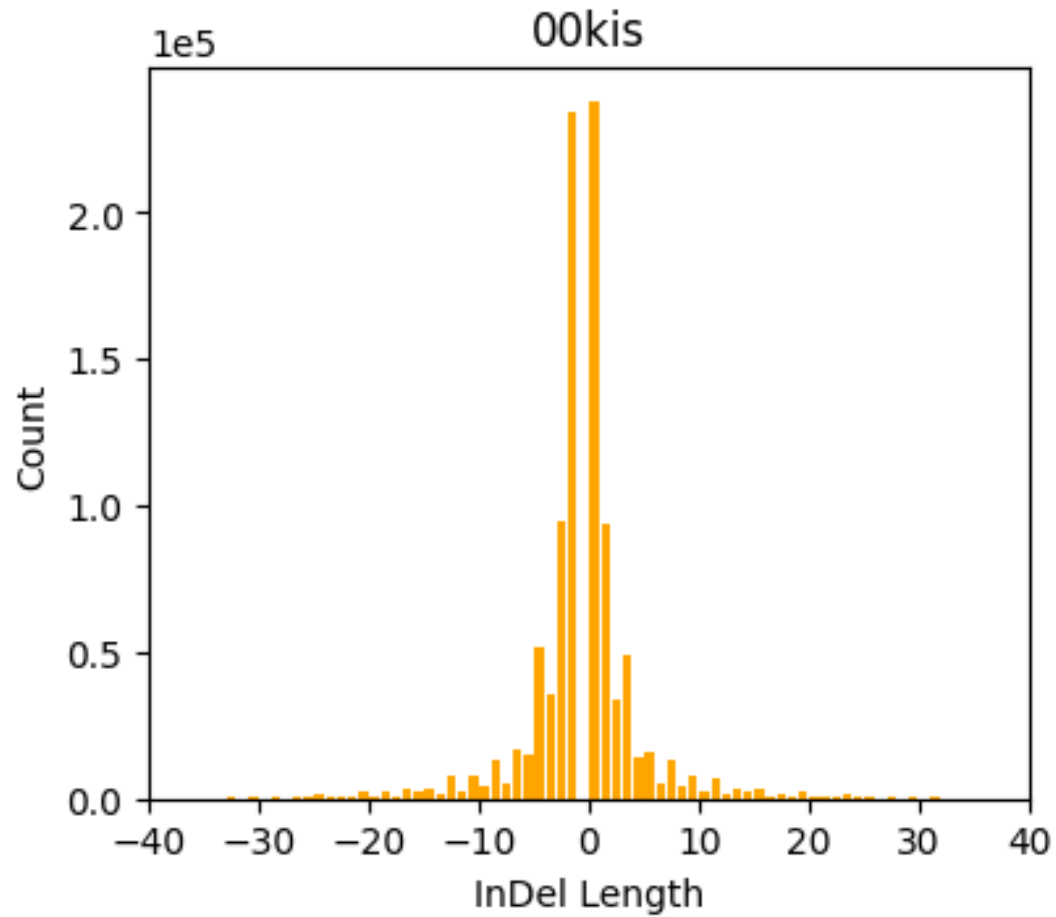
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

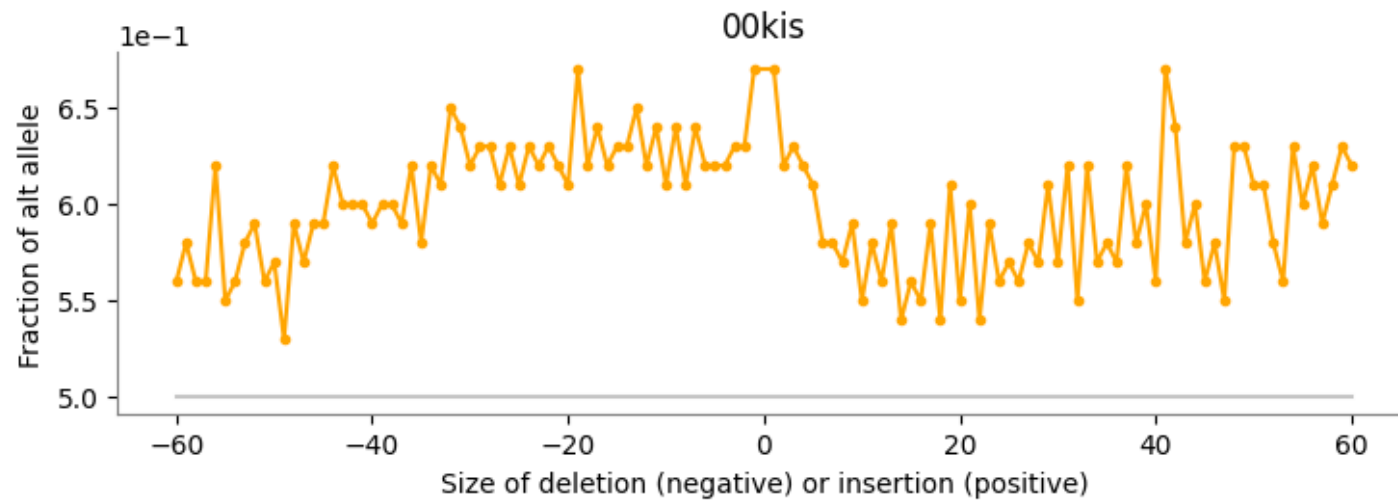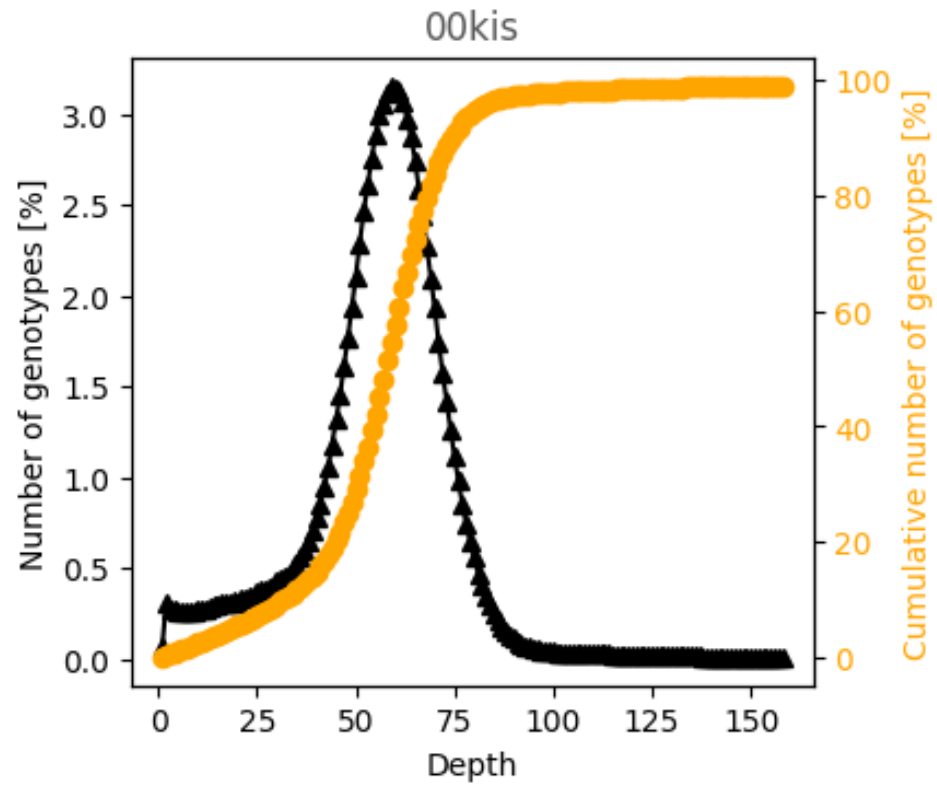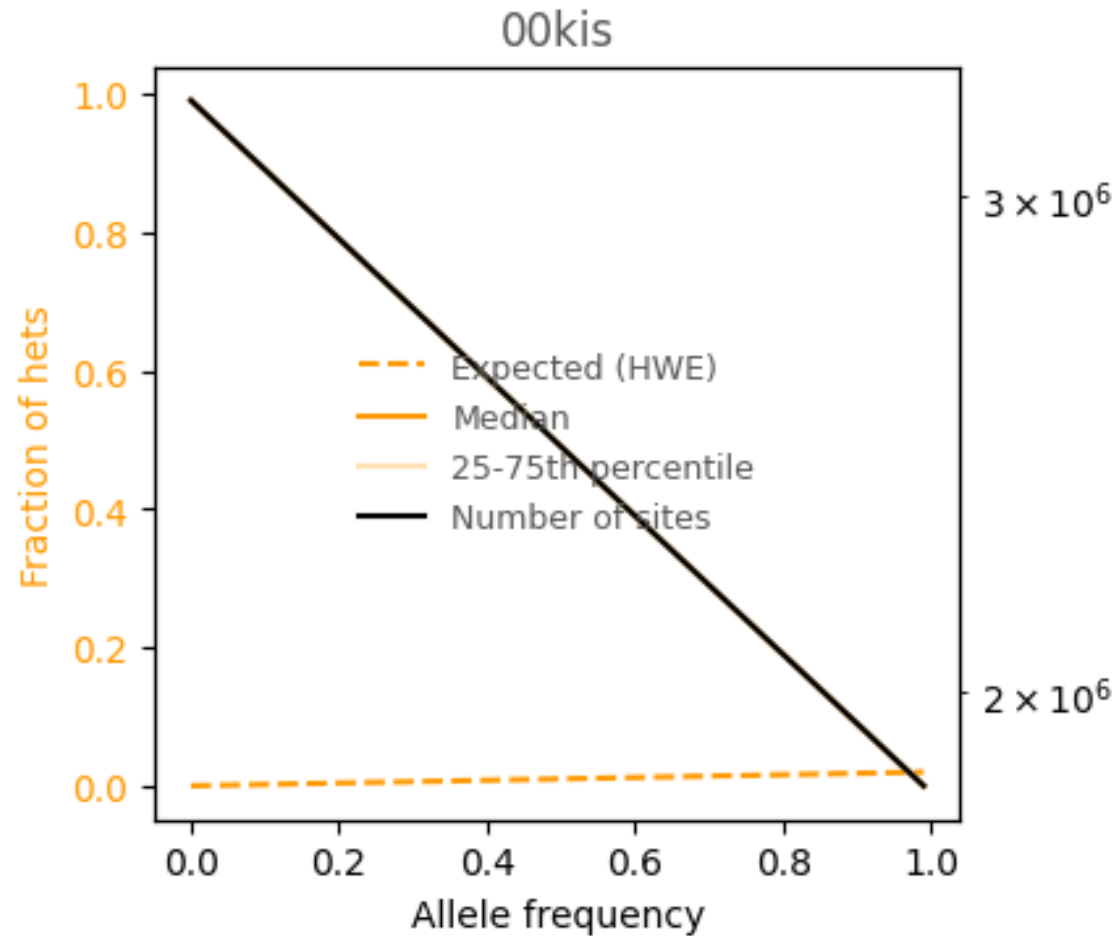# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types

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