Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
00lef	4,060,443	1.95	1.96	949,980	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
00lef	60.3%	1.92	64.7%	89,839	1,693

- 00lef .. `/ngc/projects2/gm/data/archive/2022/variants/snv/00lefrtom-103799506339-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211103_A01466_BHT575DSX2-RHGM_LABKA_WGS-`
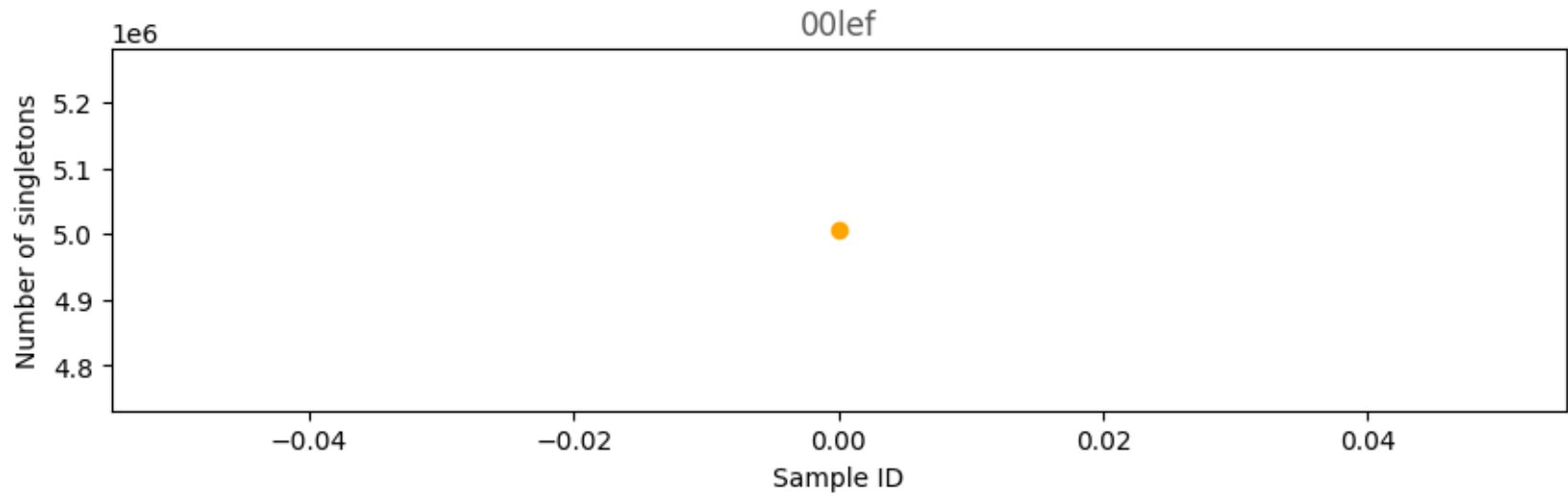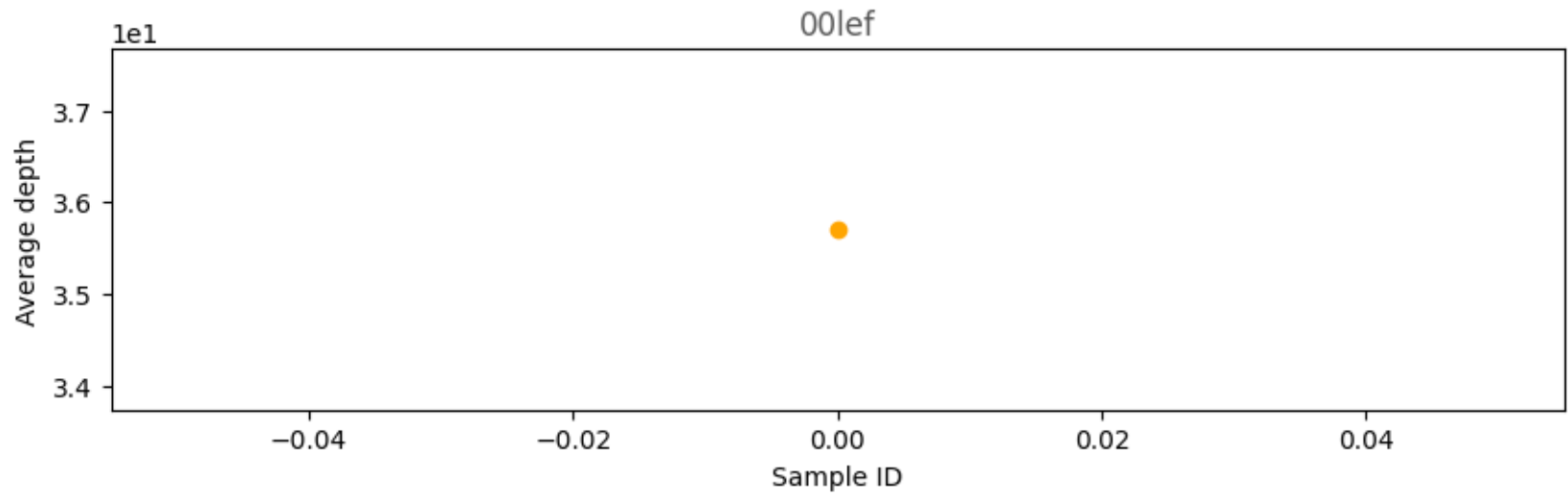 `WGS03296_snv_germline_raw.haplotype_caller.vcf.gz`
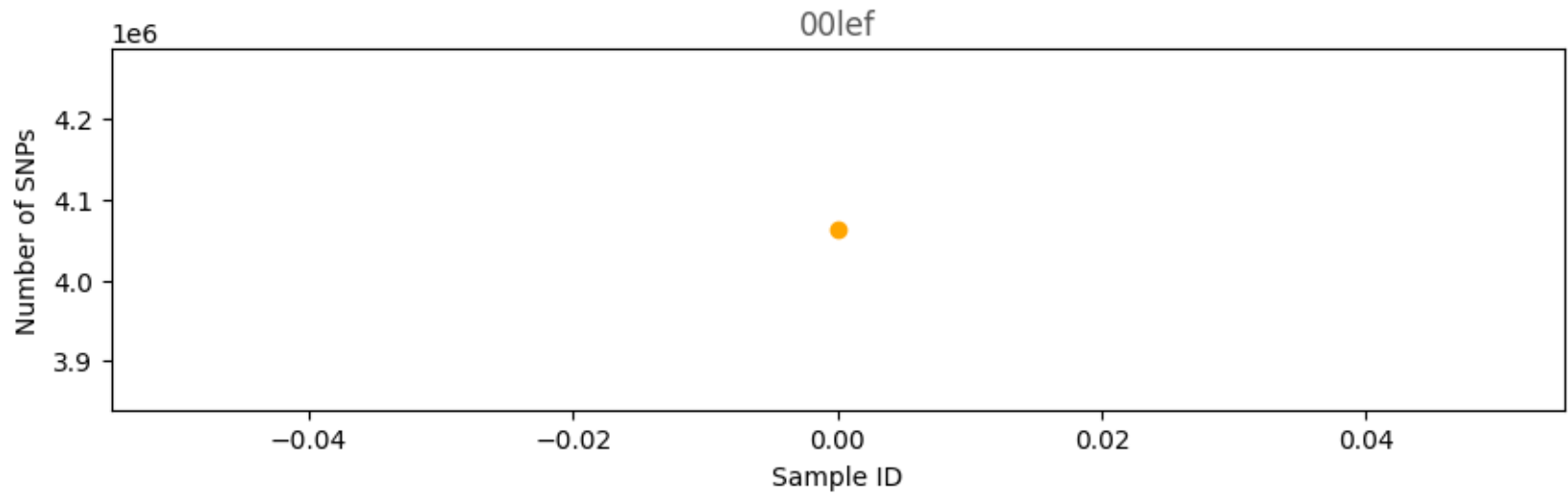
Ts/Tv by sample
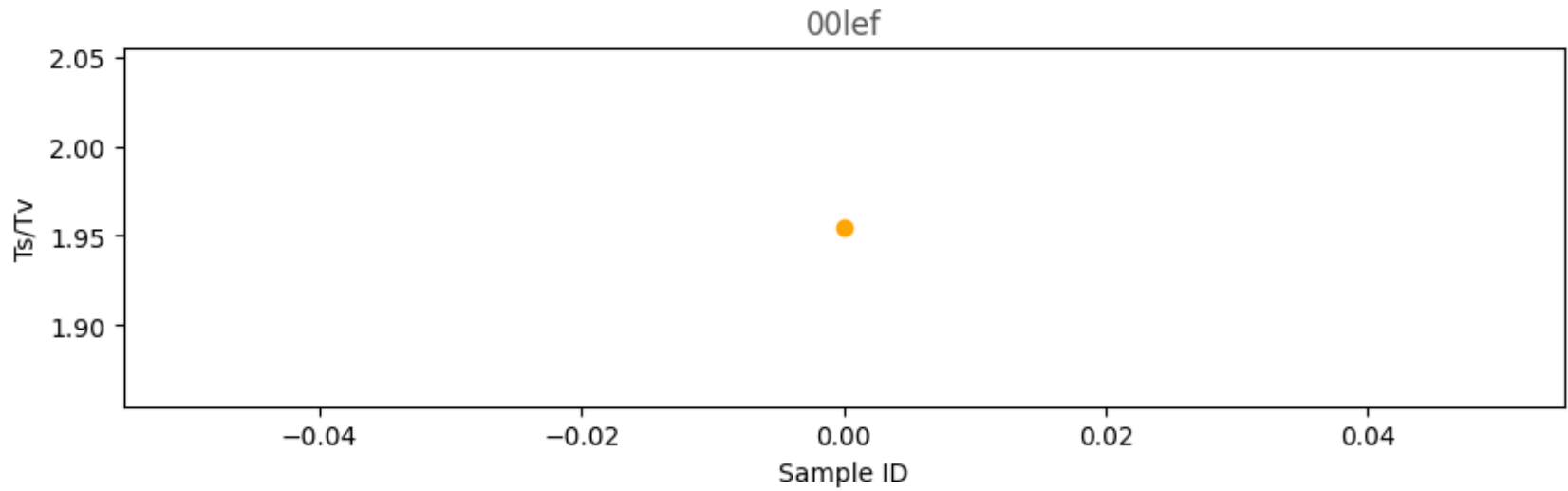
Hets vs non-ref Homs by sample
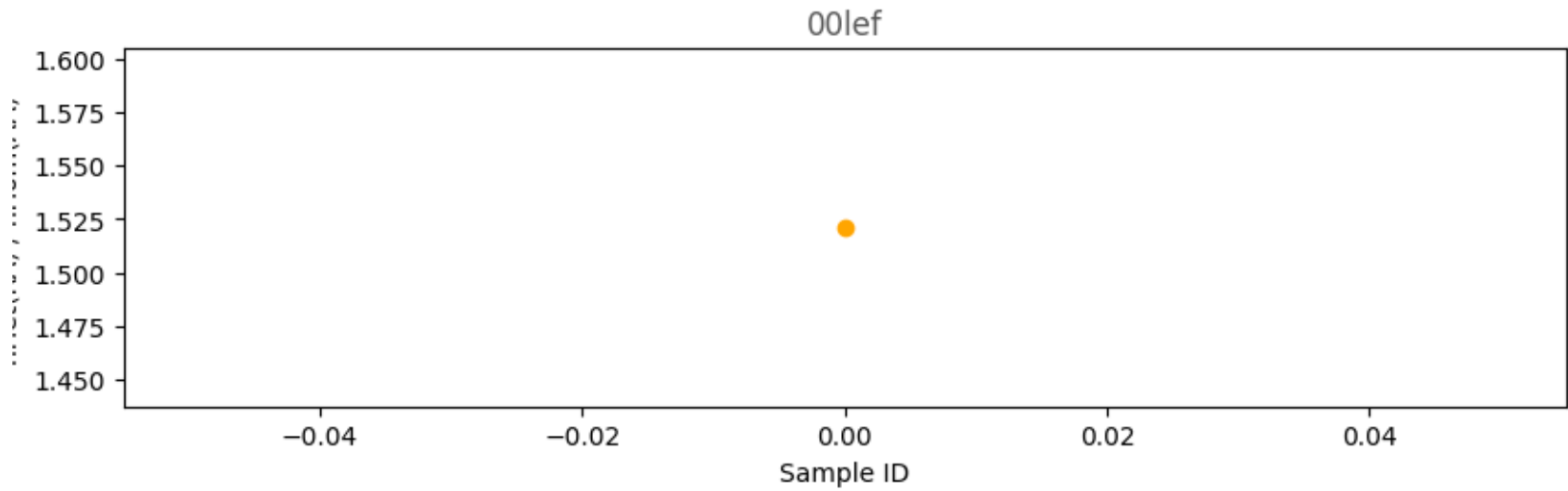
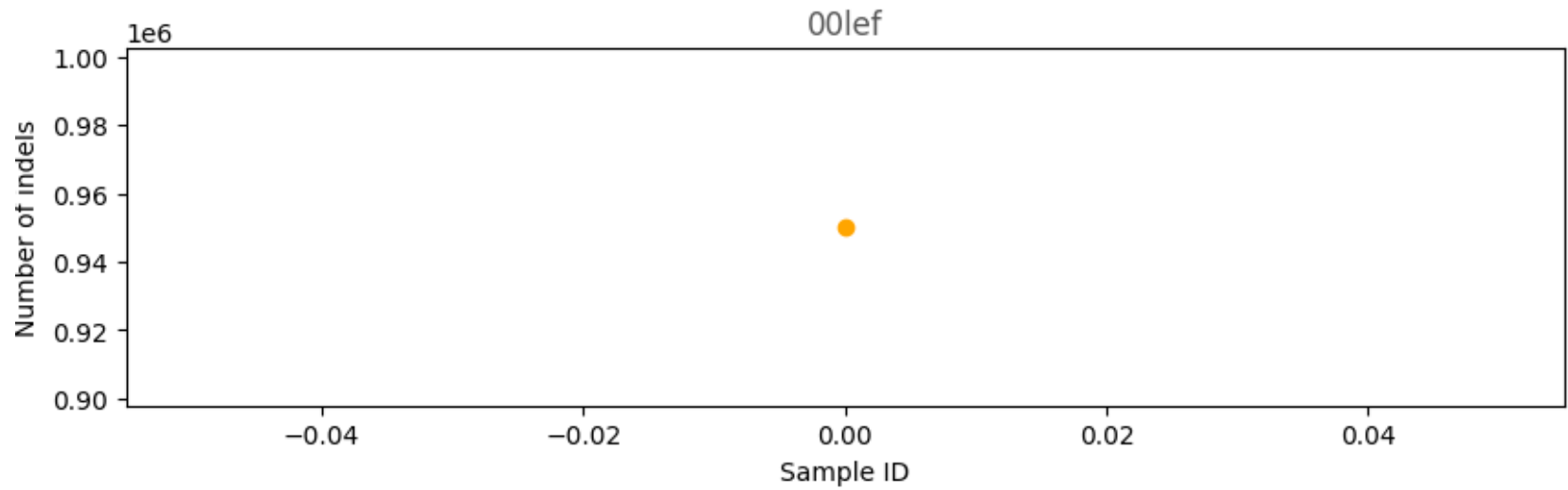# Singletons by sample (hets and homs)
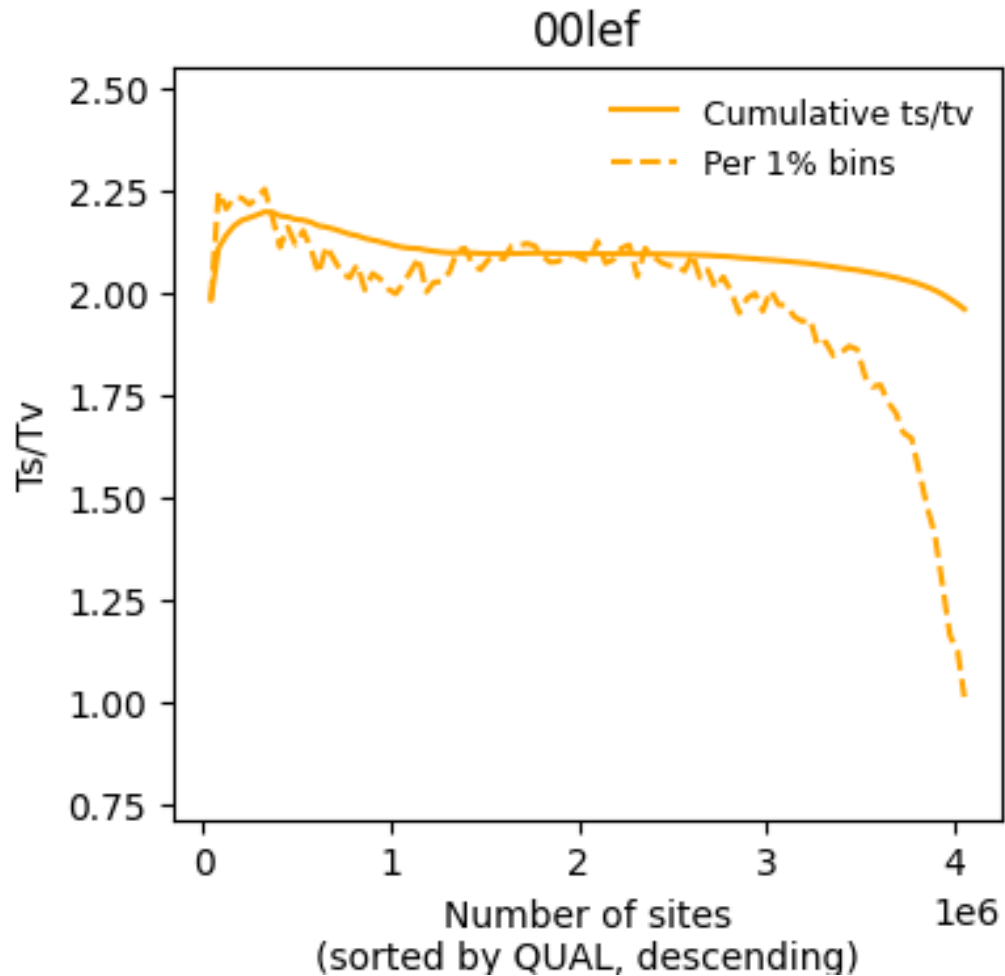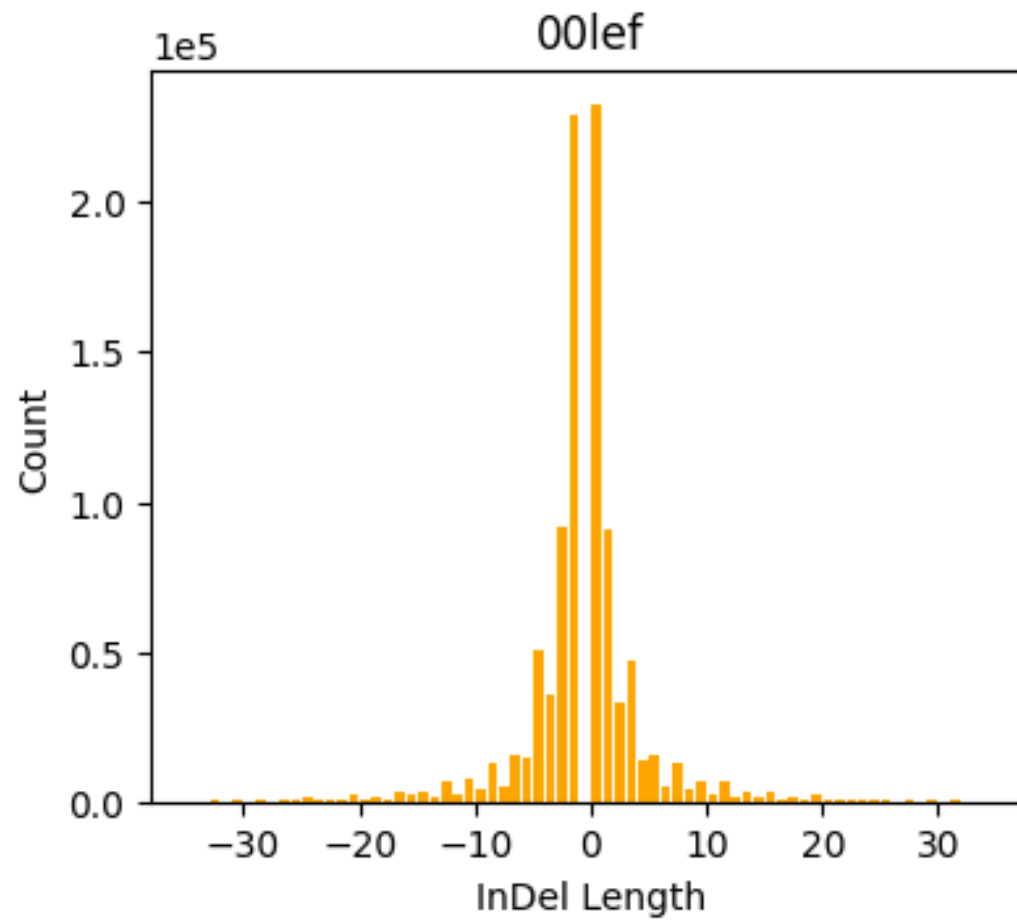
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

00lef

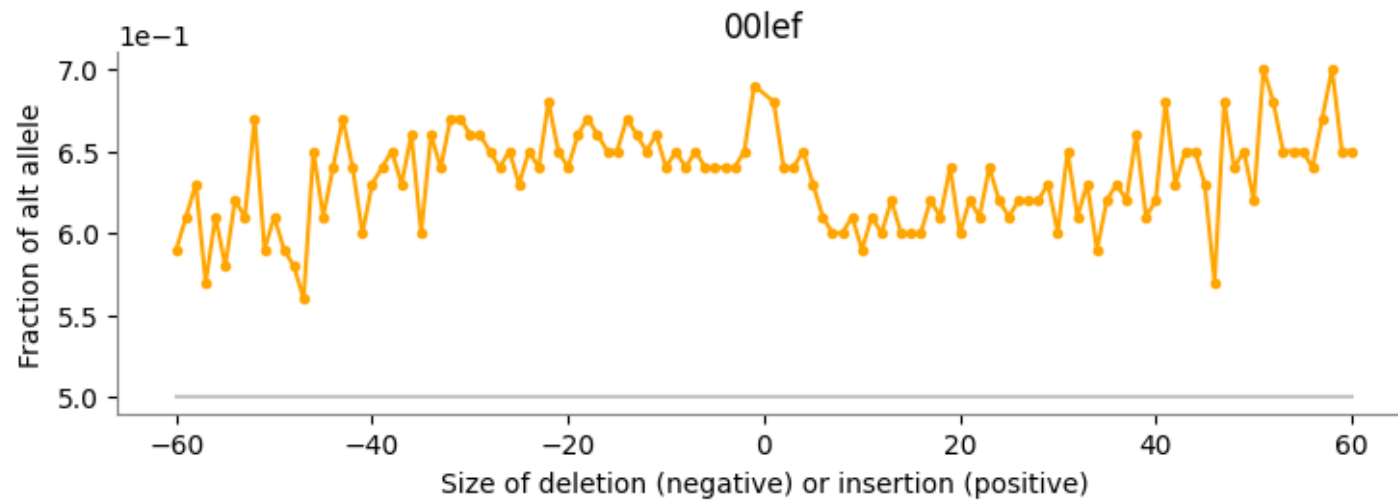# Indel distribution

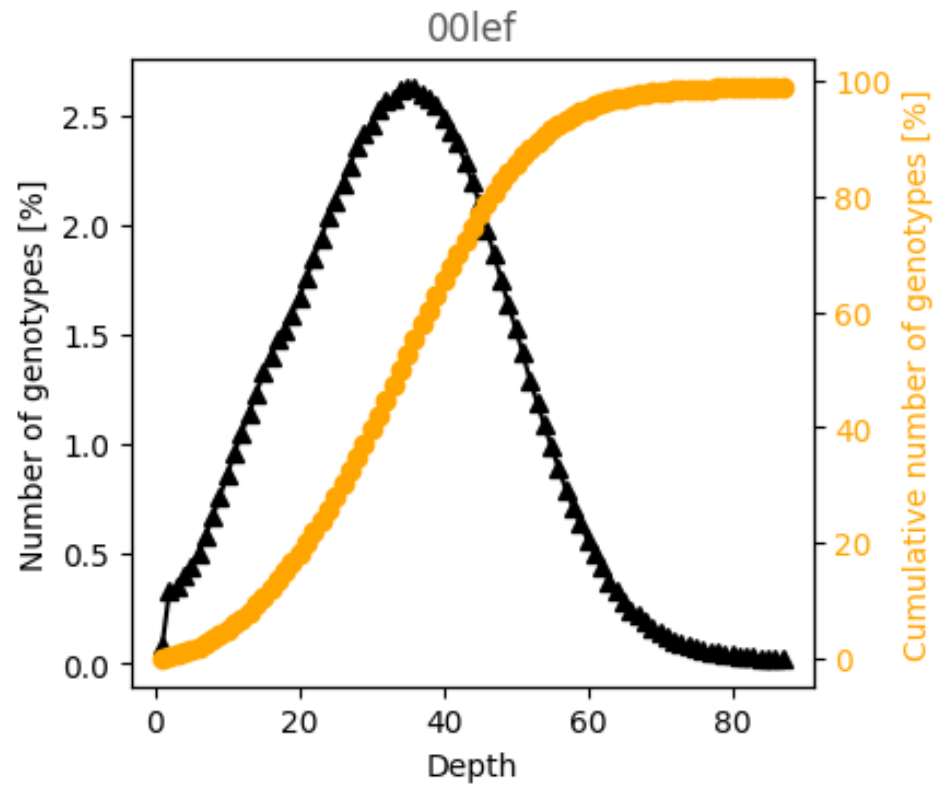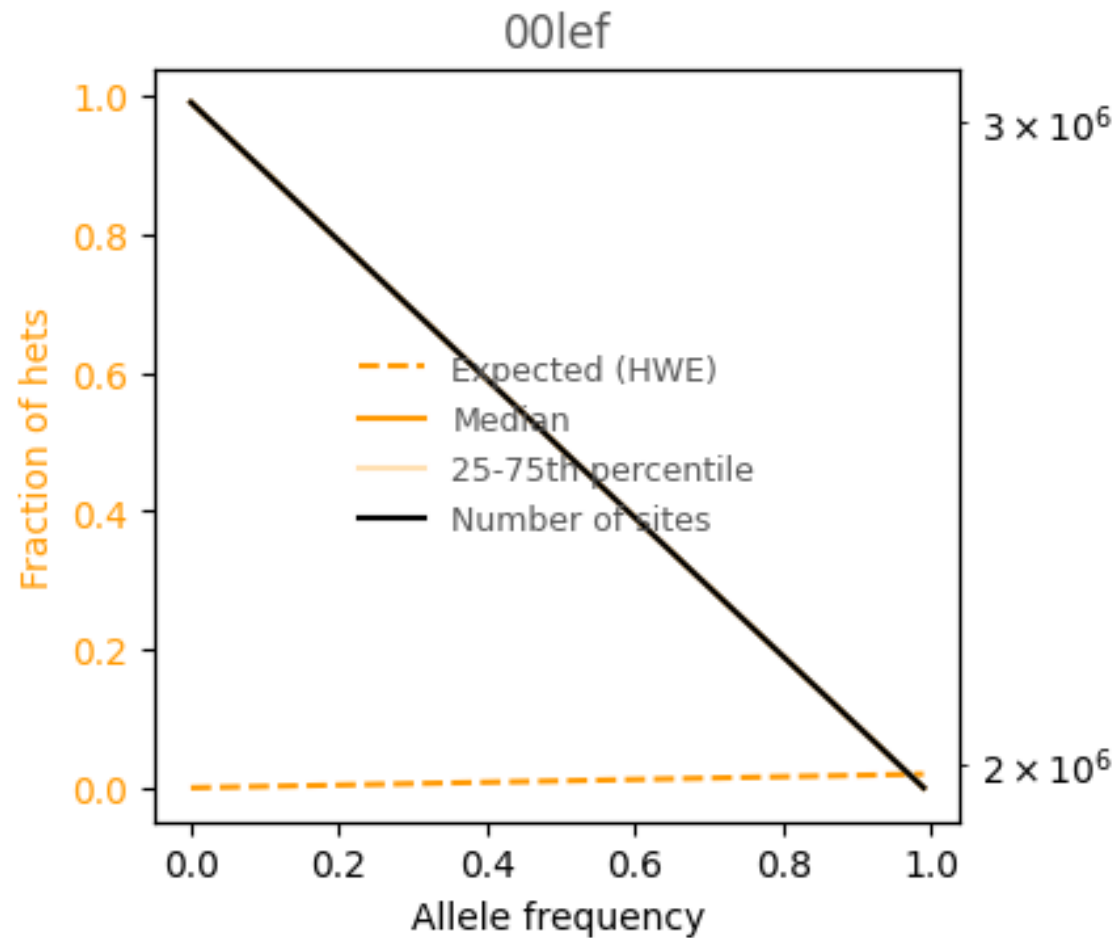# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types