Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
01tor	4,036,814	1.93	1.93	942,225	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
01tor	59.5%	1.88	64.1%	95,219	1,986

- 01tor .. `/ngc/projects2/gm/data/archive/2022/variants/snv/01torebef-103906972608-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01731-220923_A01176_BHTNN3DSX3-EXT_LAB KA_NGCWGS-NGCWGS05492_22RKG020979_snv_germline_raw.haplotype_caller.vcf.gz`
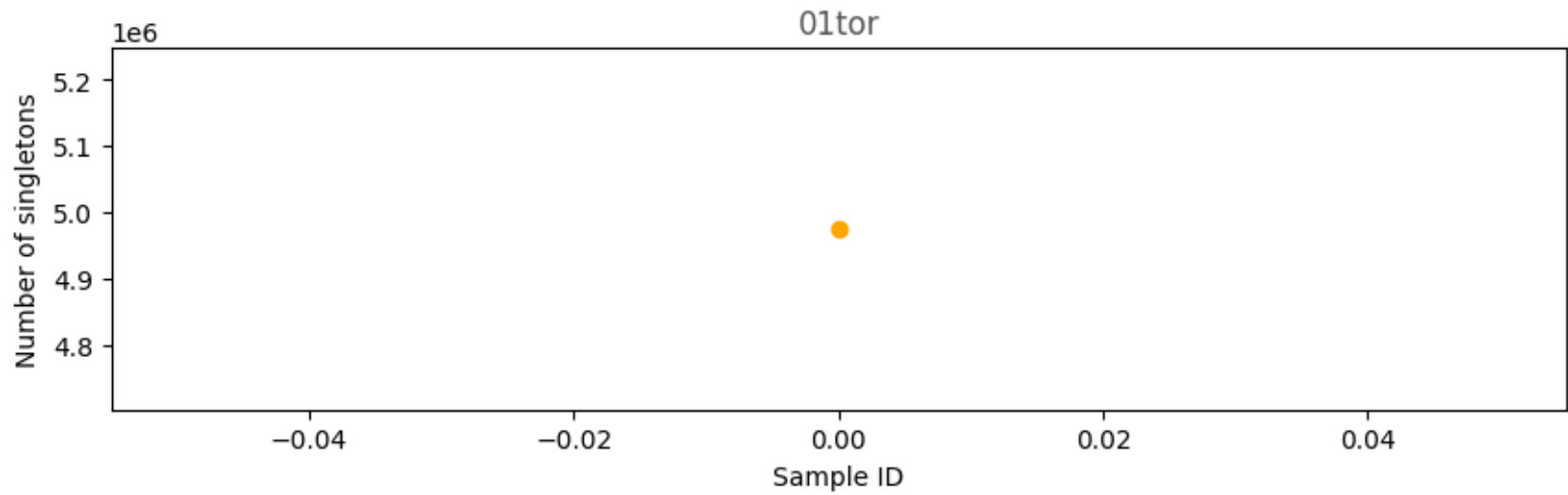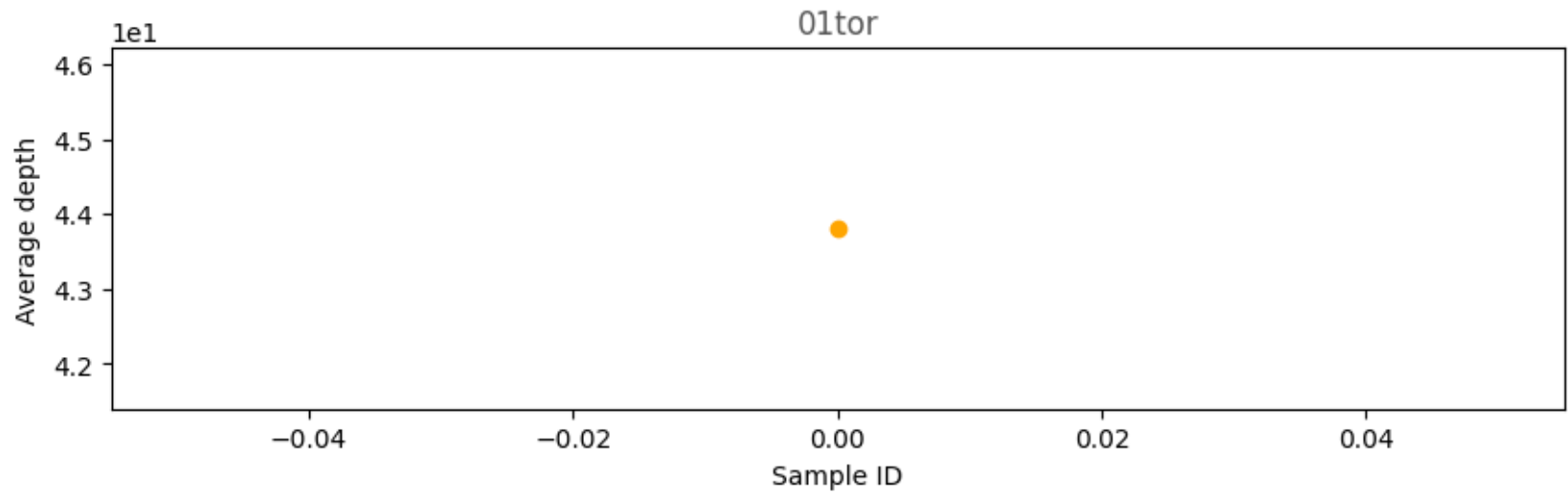
Ts/Tv by sample
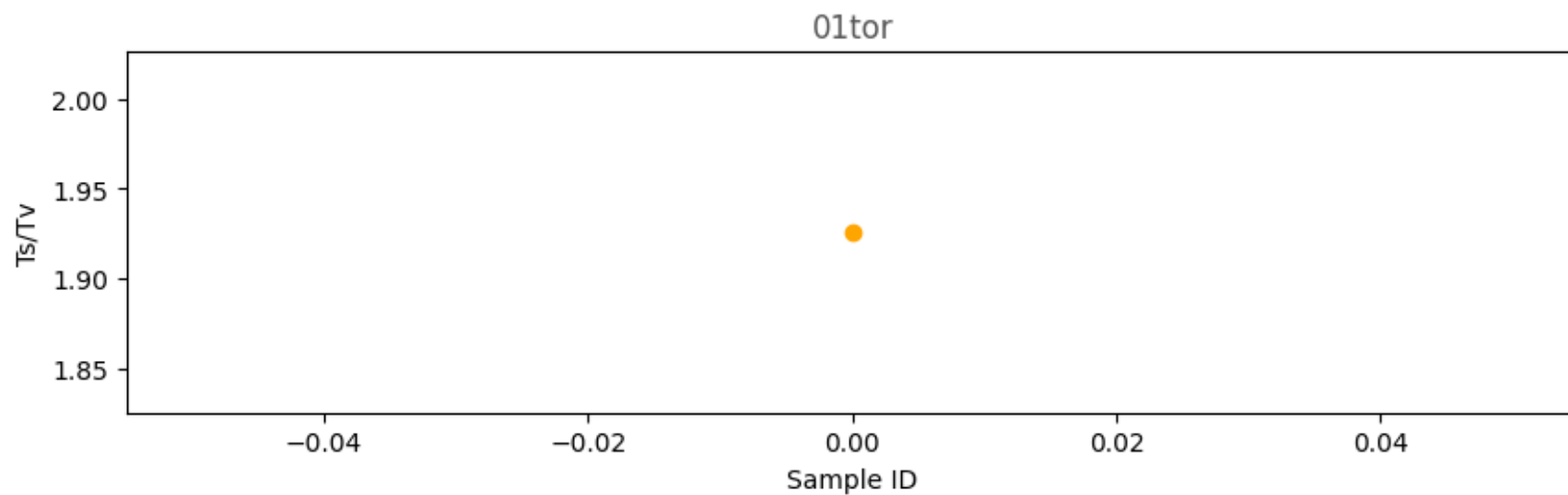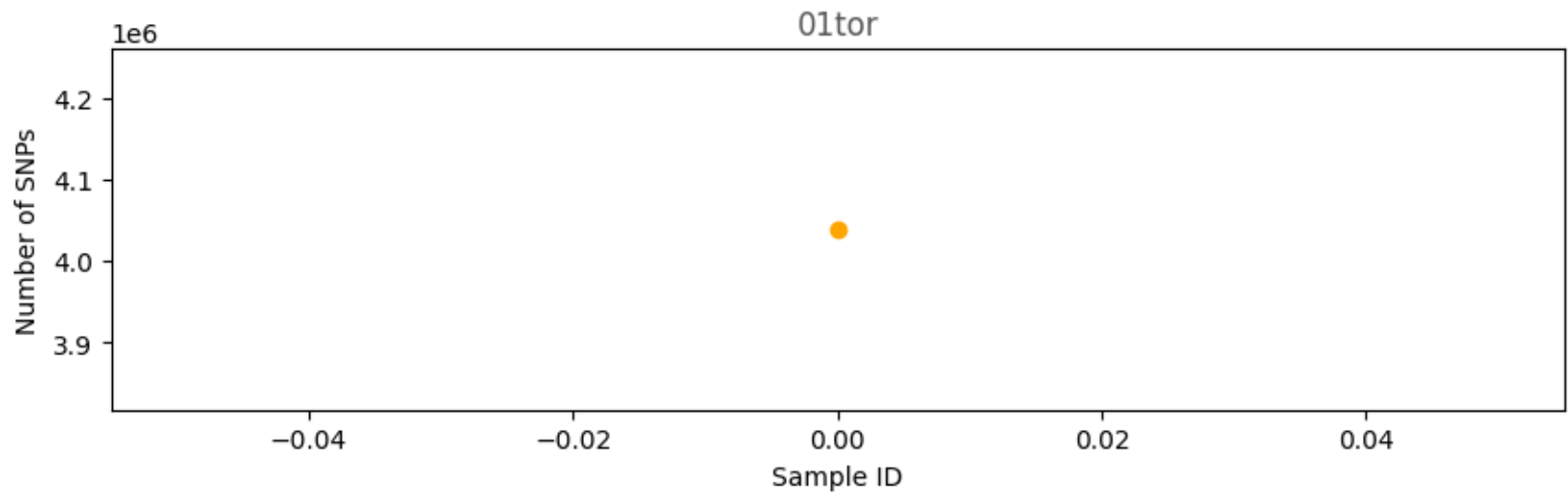
Hets vs non-ref Homs by sample
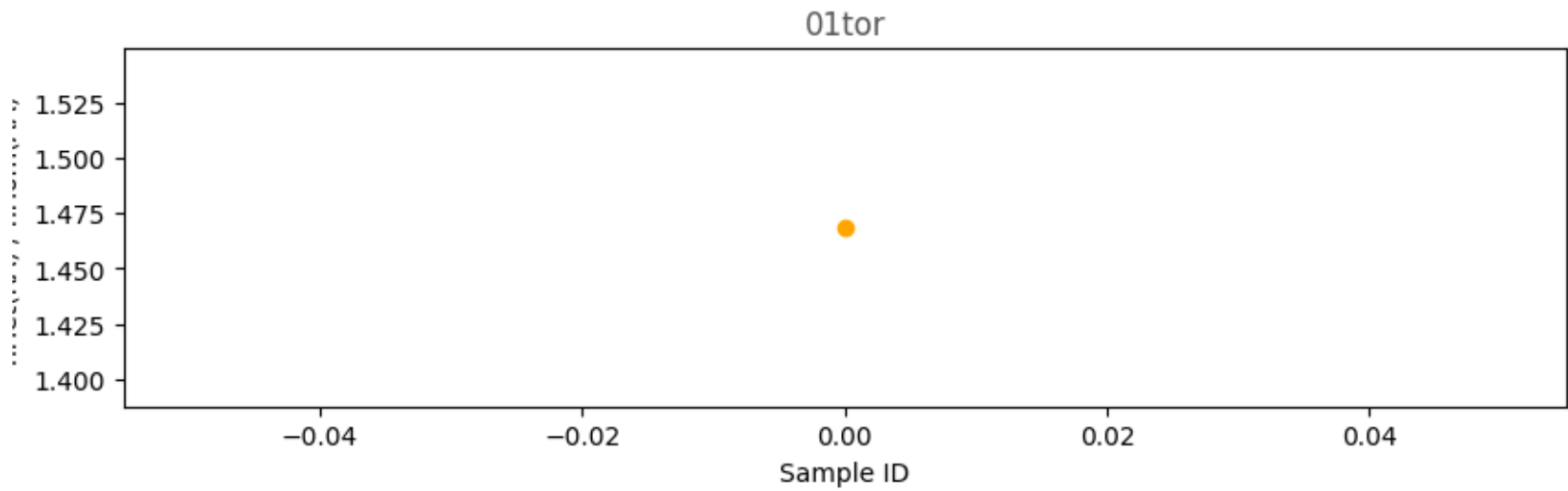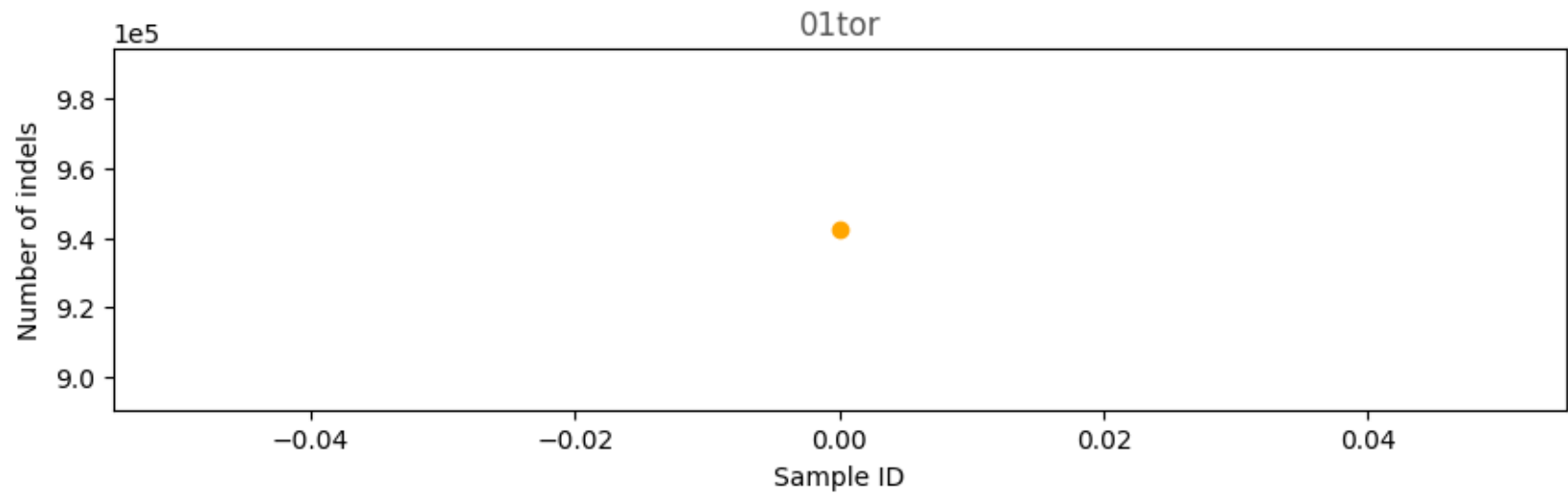
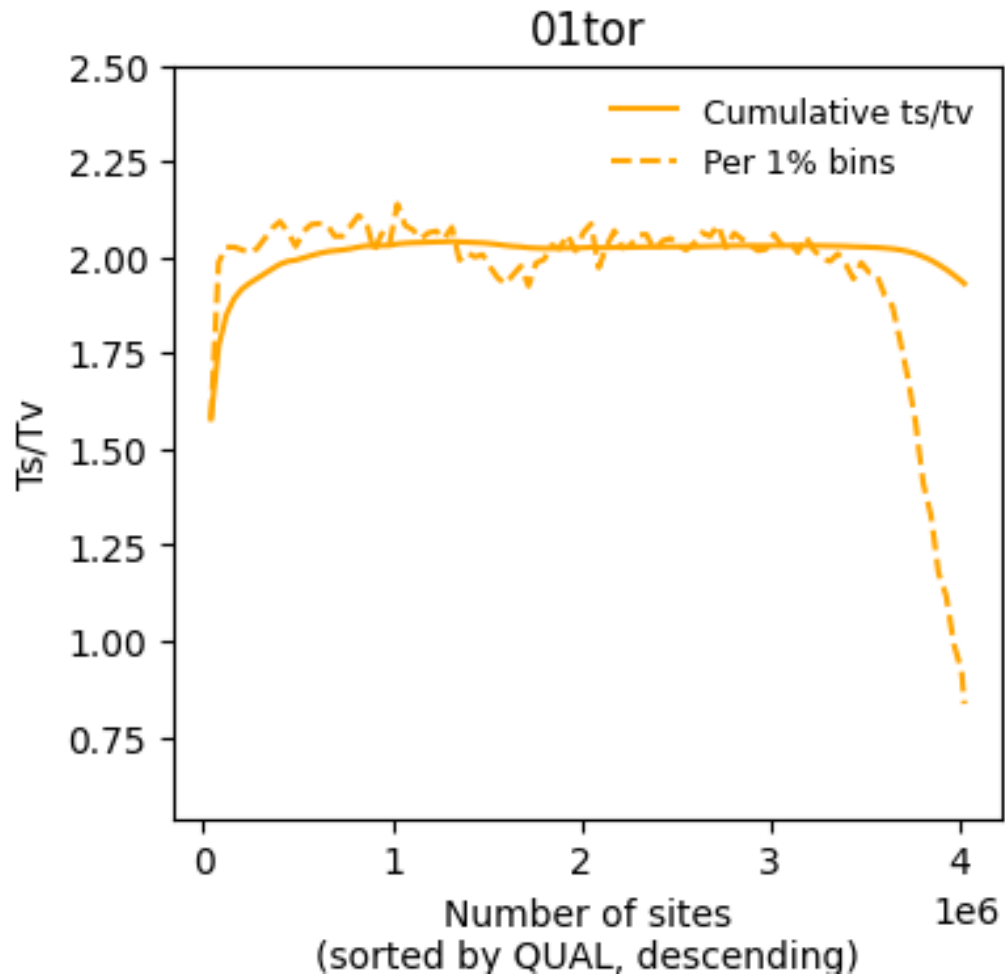# Singletons by sample (hets and homs)
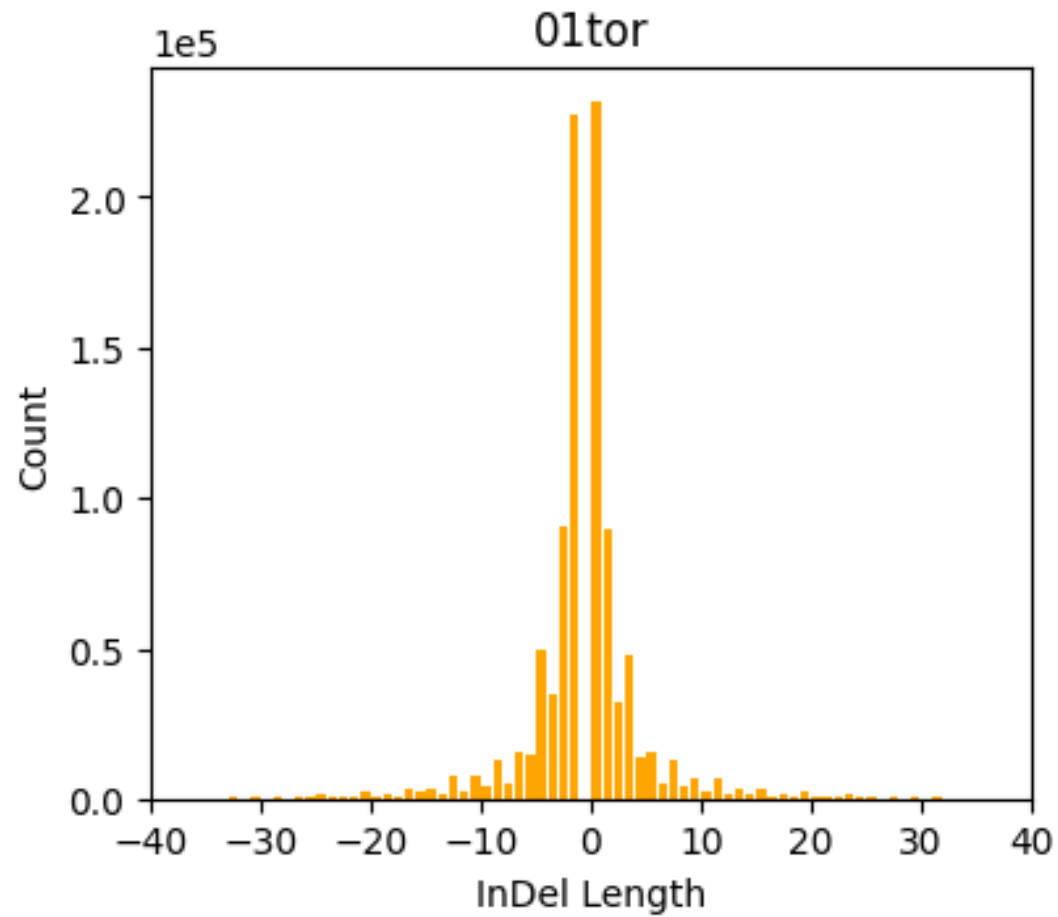
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

01tor

Indel distribution

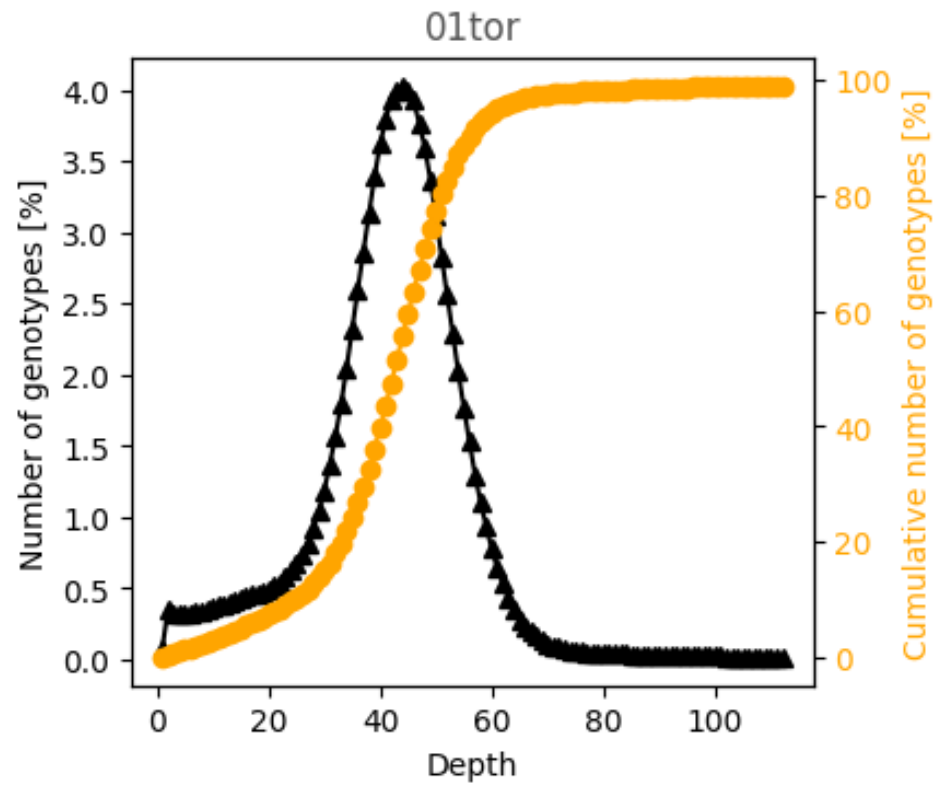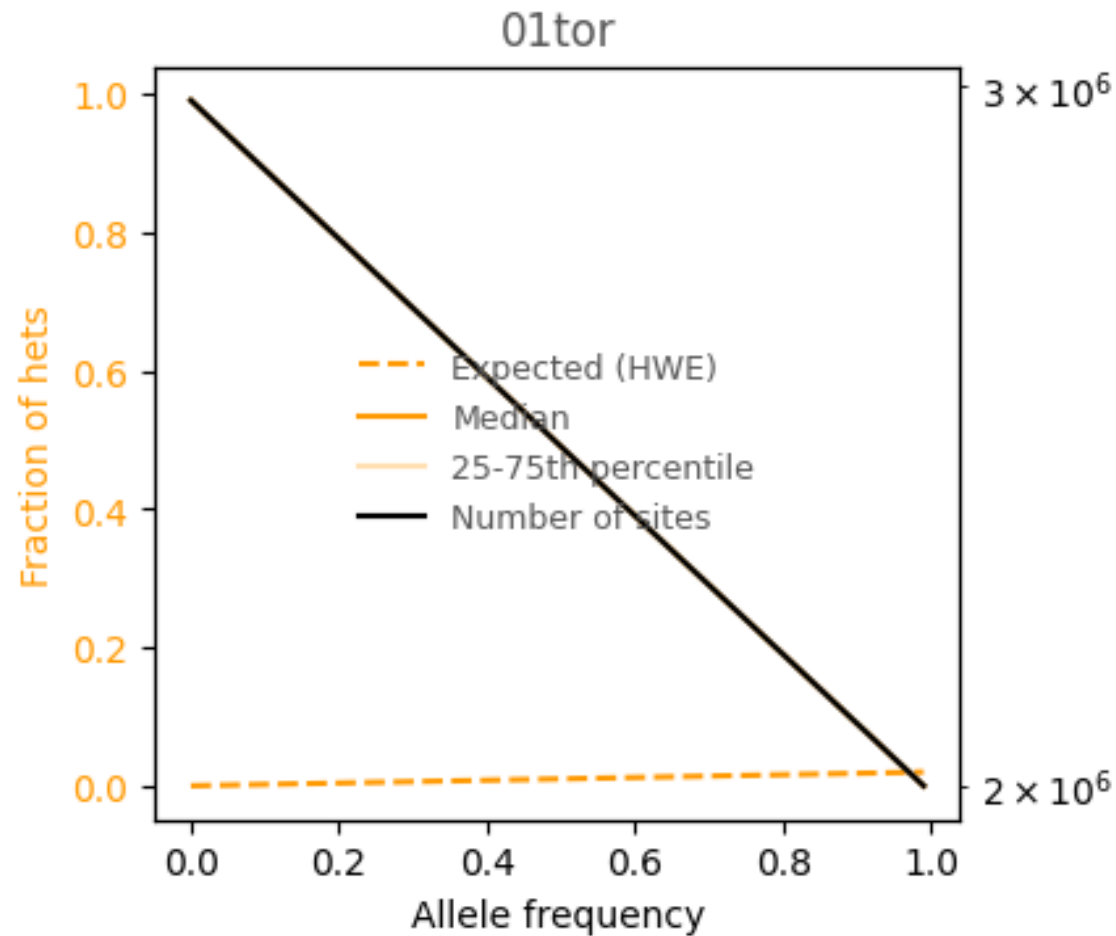# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types