Summary Numbers

| Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
16sor	4,106,176	1.92	1.93	961,194	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
16sor	63.4%	1.90	68.7%	104,462	2,007

- 16sor .. `/ngc/projects2/gm/data/archive/2022/variants/snv/16sorsesf-103760980439-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00246-210720_A01411_AHGWH5DSX2-EXT_LABKA_NGCWGS-NGCWGS00139_163rxbqv8_snv_germline_raw.haplotype_caller.vcf.gz`
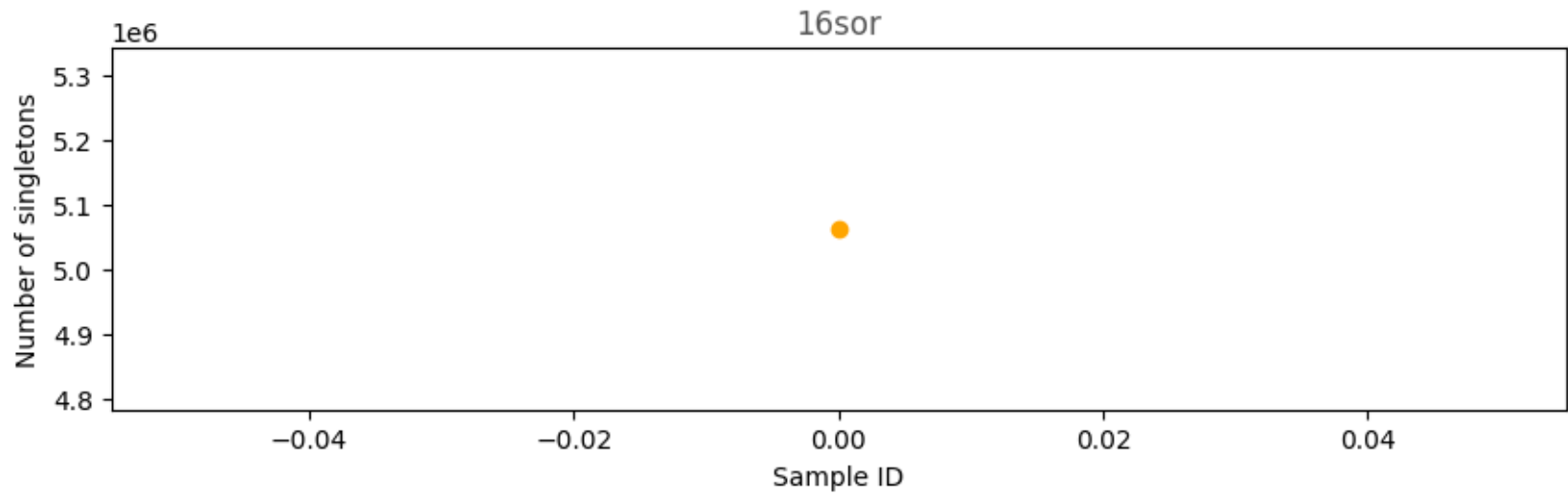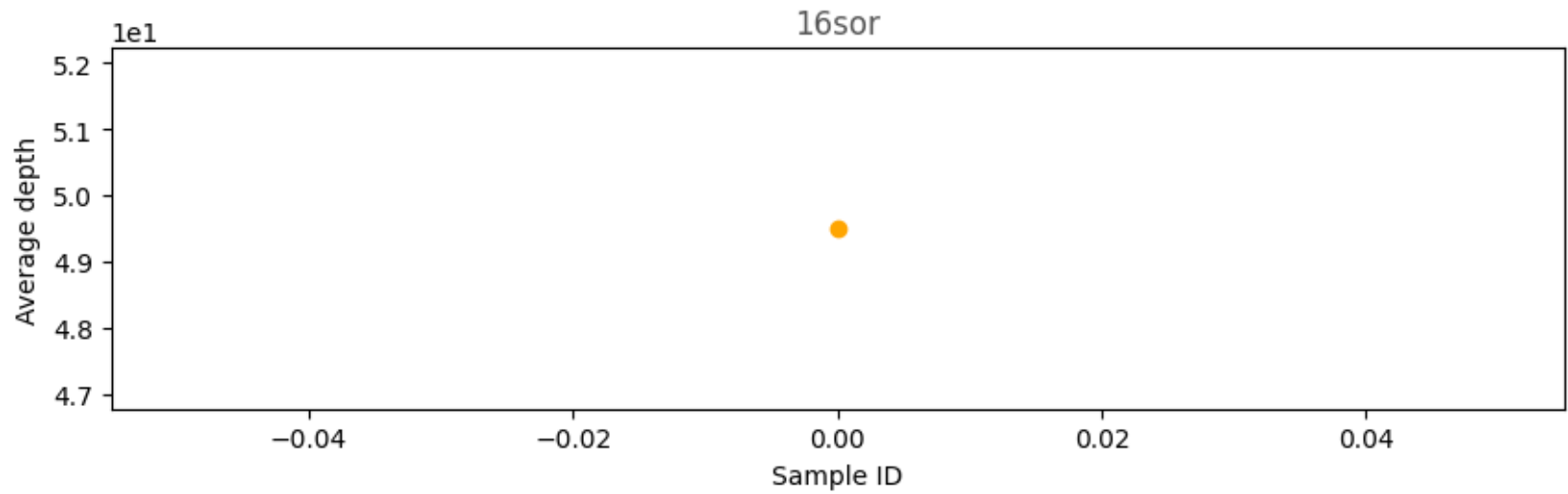
Ts/Tv by sample
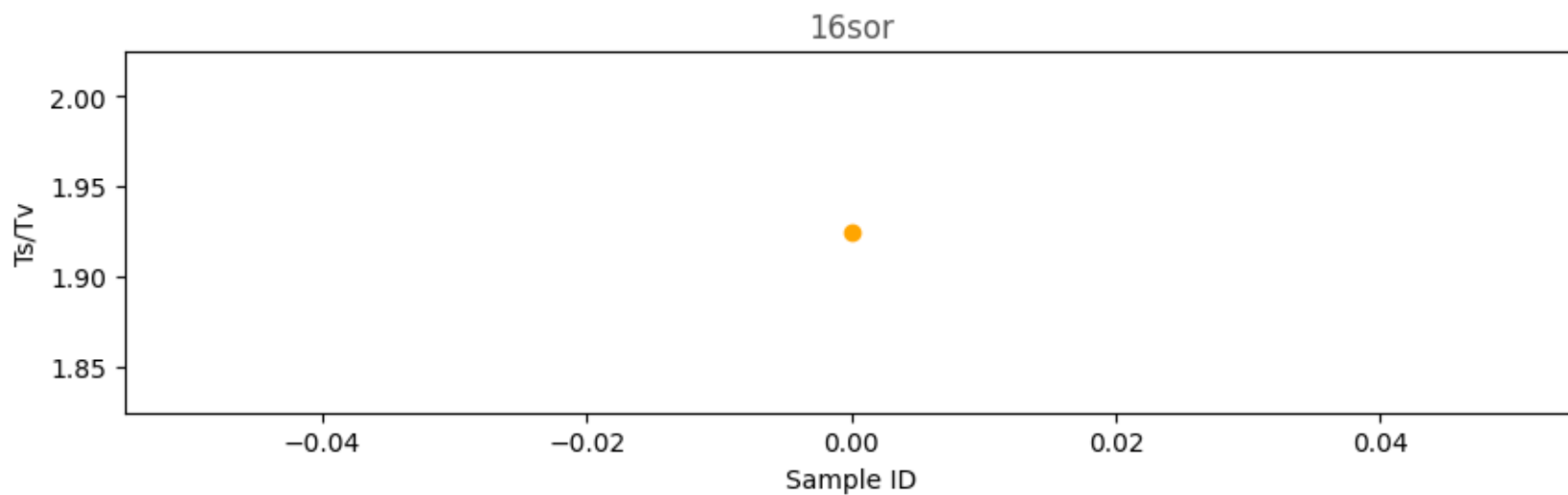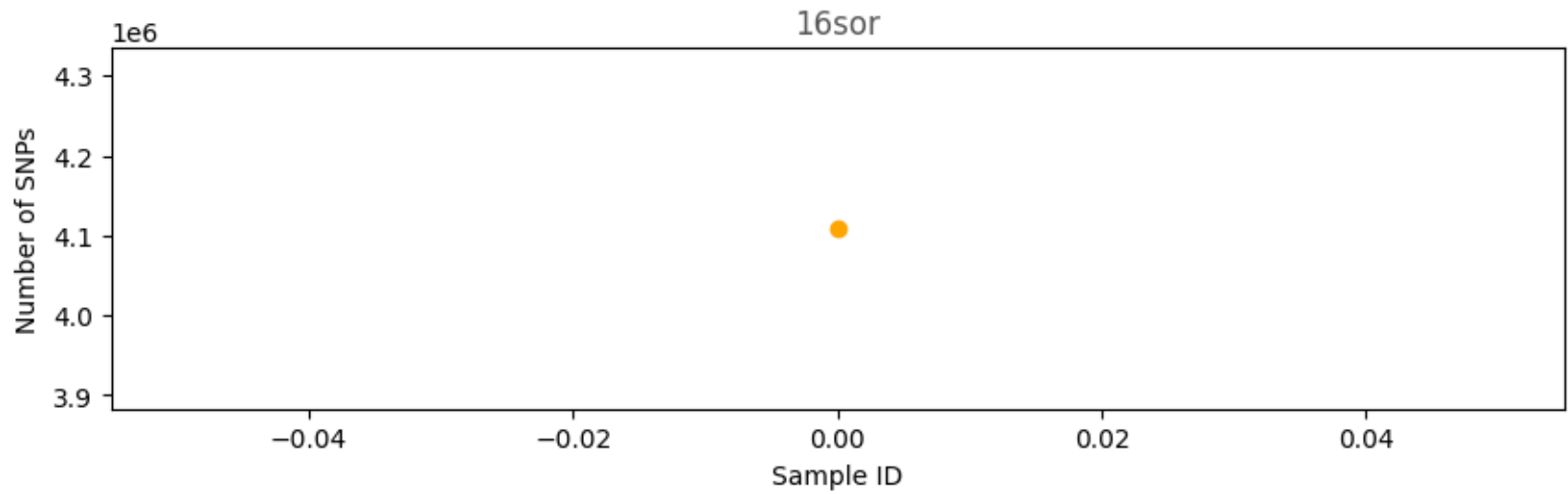
Hets vs non-ref Homs by sample
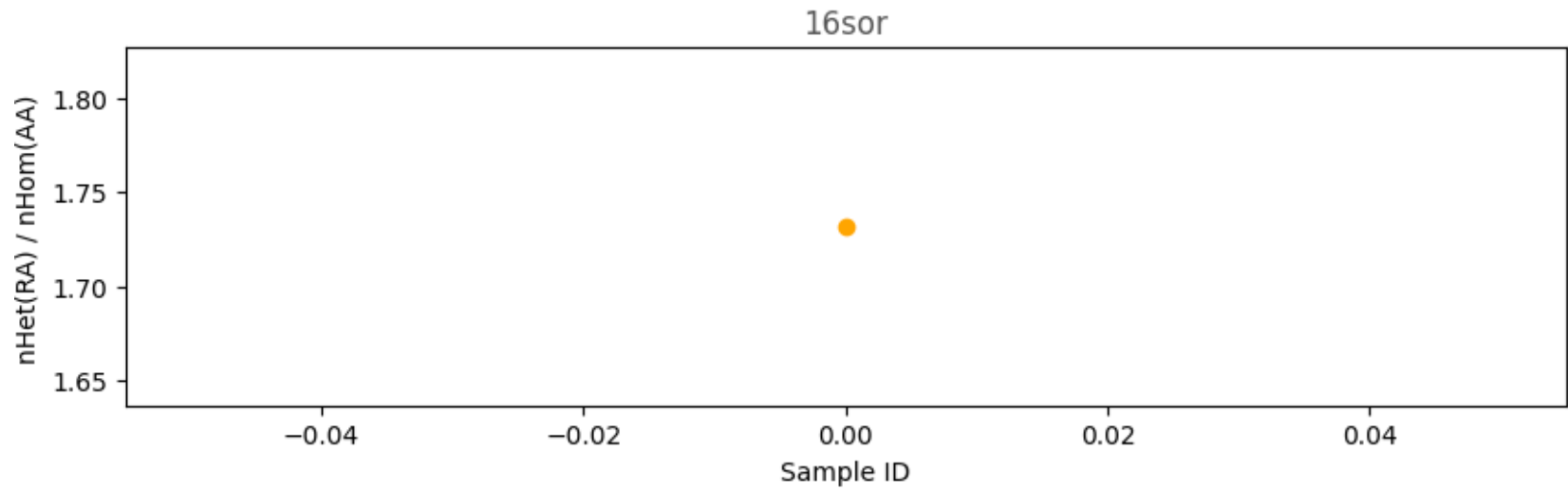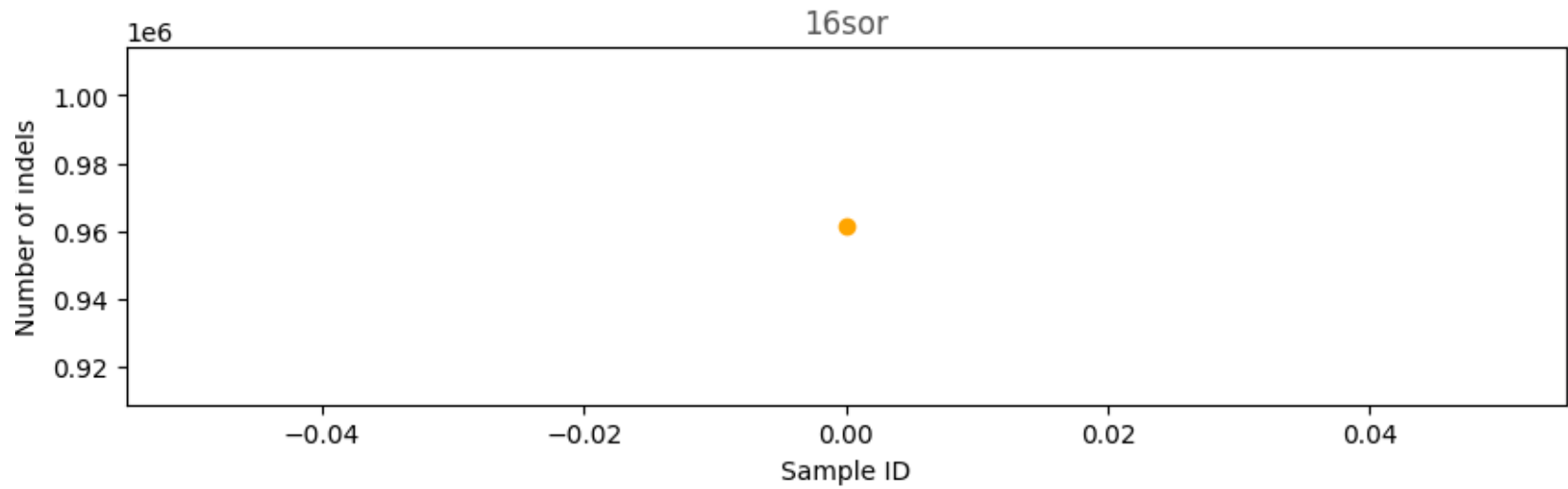
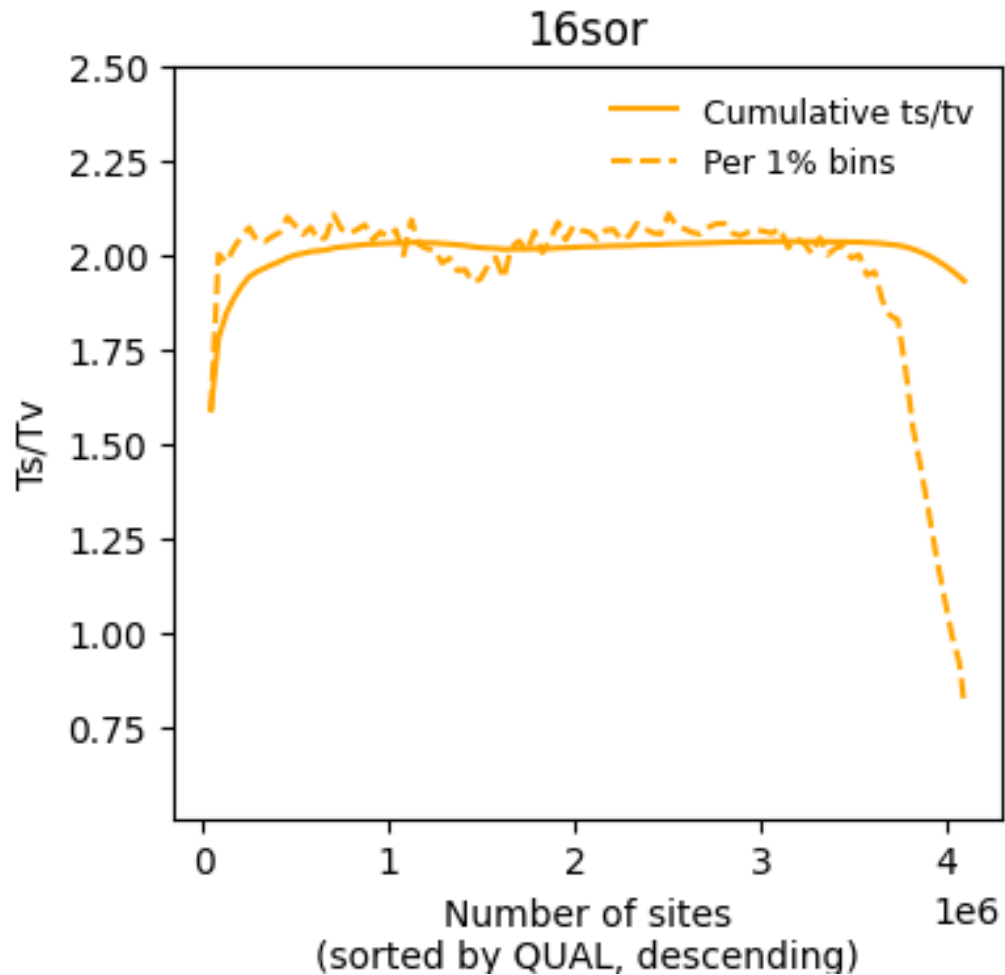# Singletons by sample (hets and homs)

Average depth by sample

Number of SNPs by sample
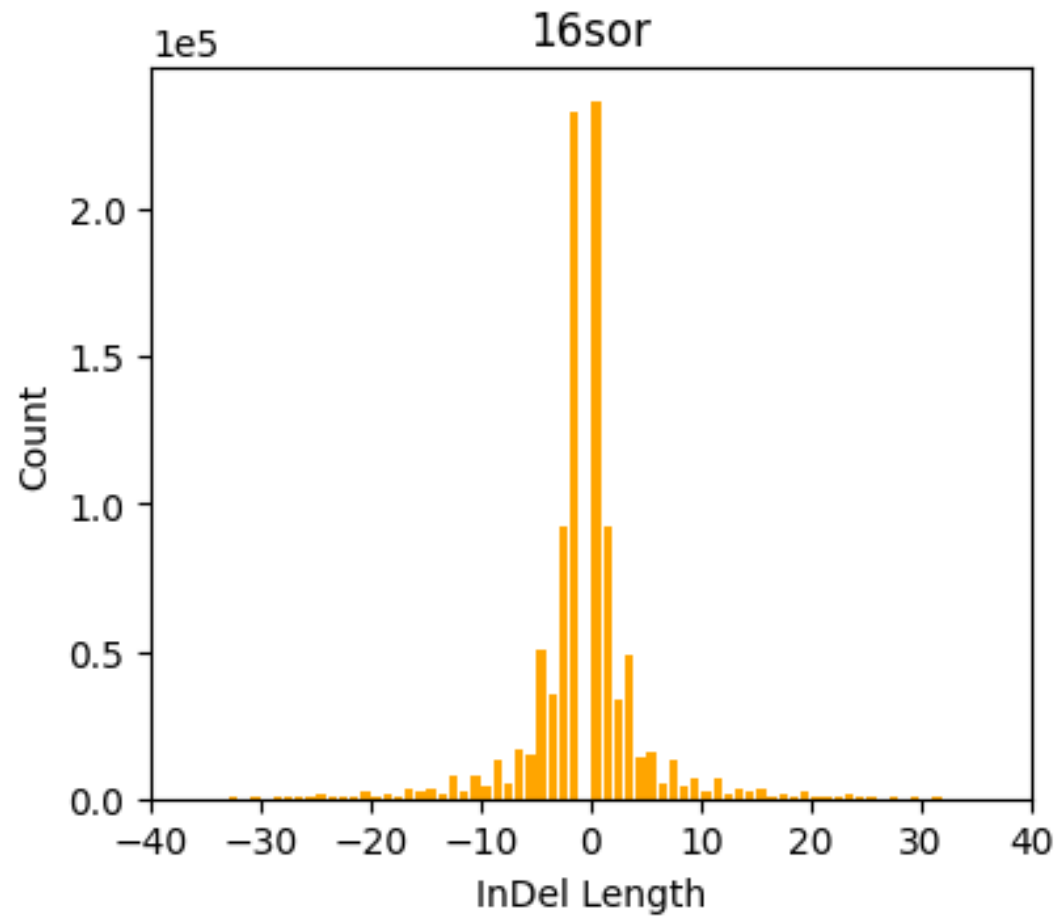
Number of indels by sample
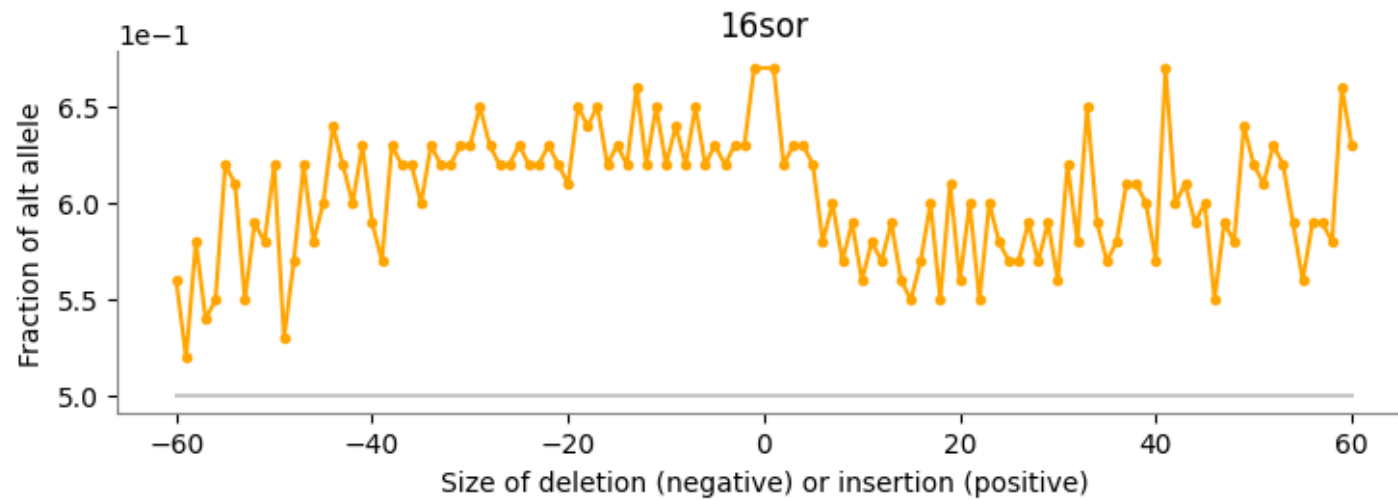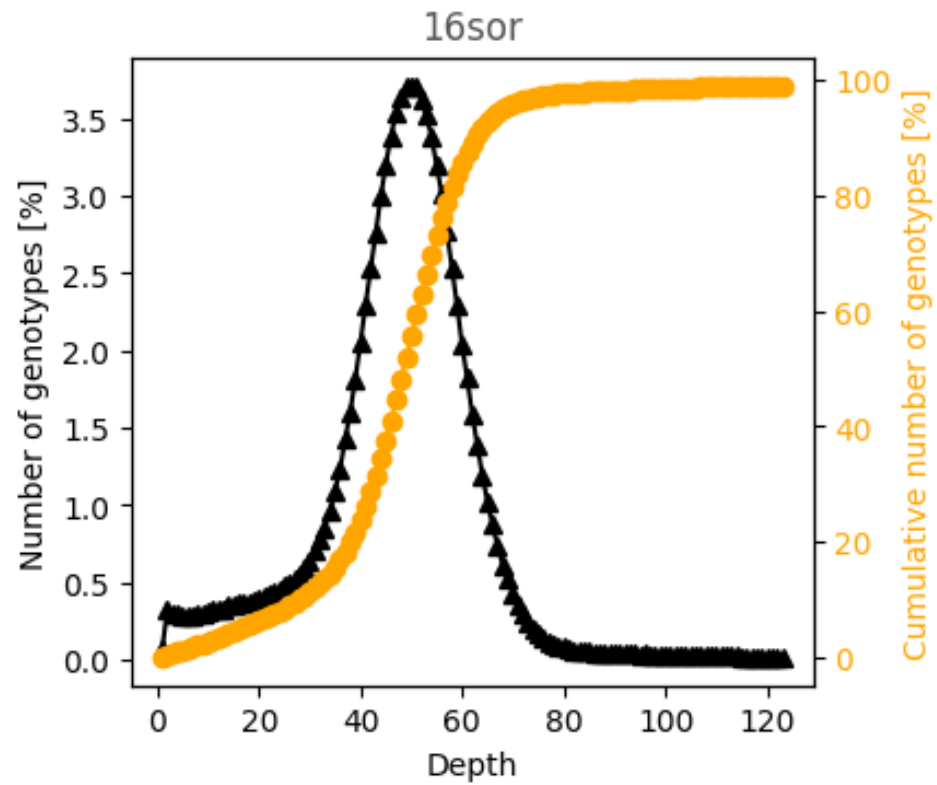
Ts/Tv stratified by QUAL
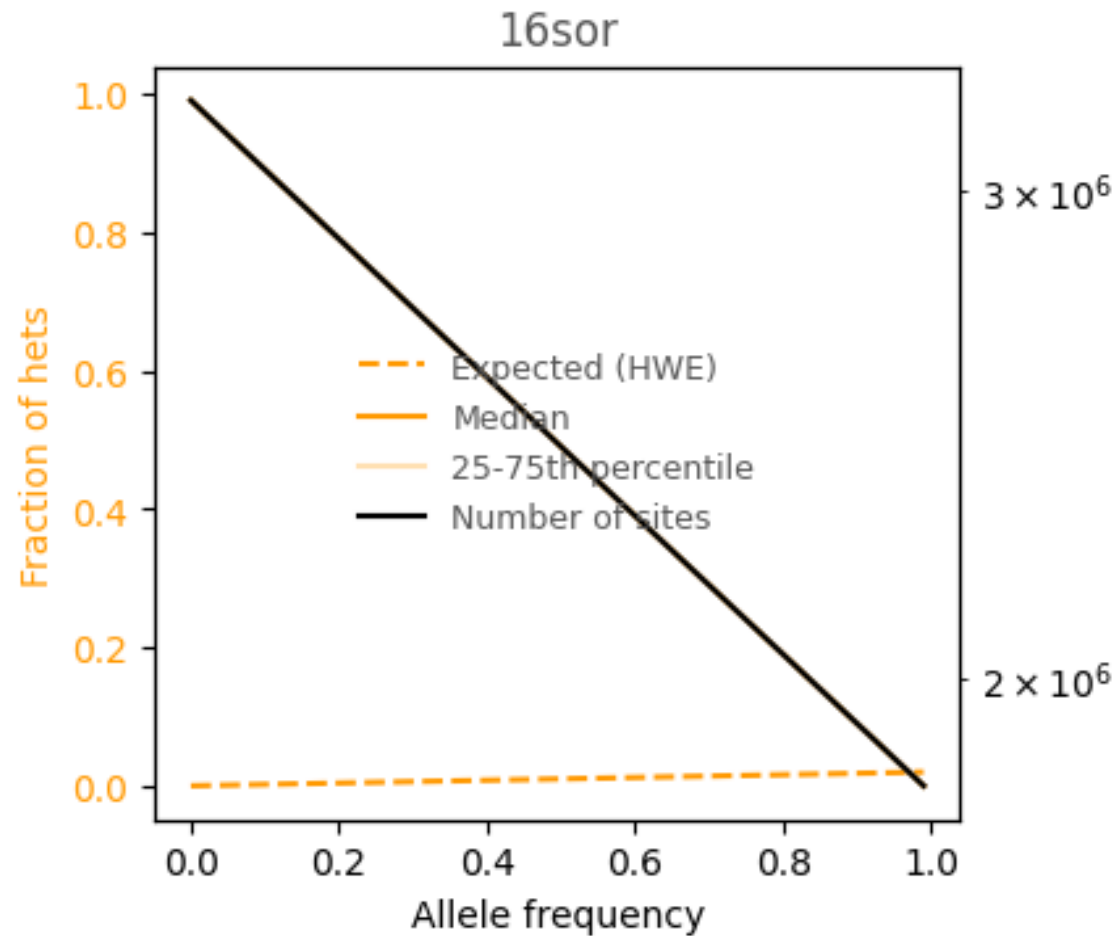

16sor

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types