Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
16ode	4,223,031	1.94	1.94	985,760	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
16ode	64.5%	1.91	69.7%	109,968	2,204

- 16ode .. `/ngc/projects2/gm/data/archive/2022/variants/snv/16odereim-103794068341-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00369-211019_A01411_BHHHY3DSX2-EXT_LAB`
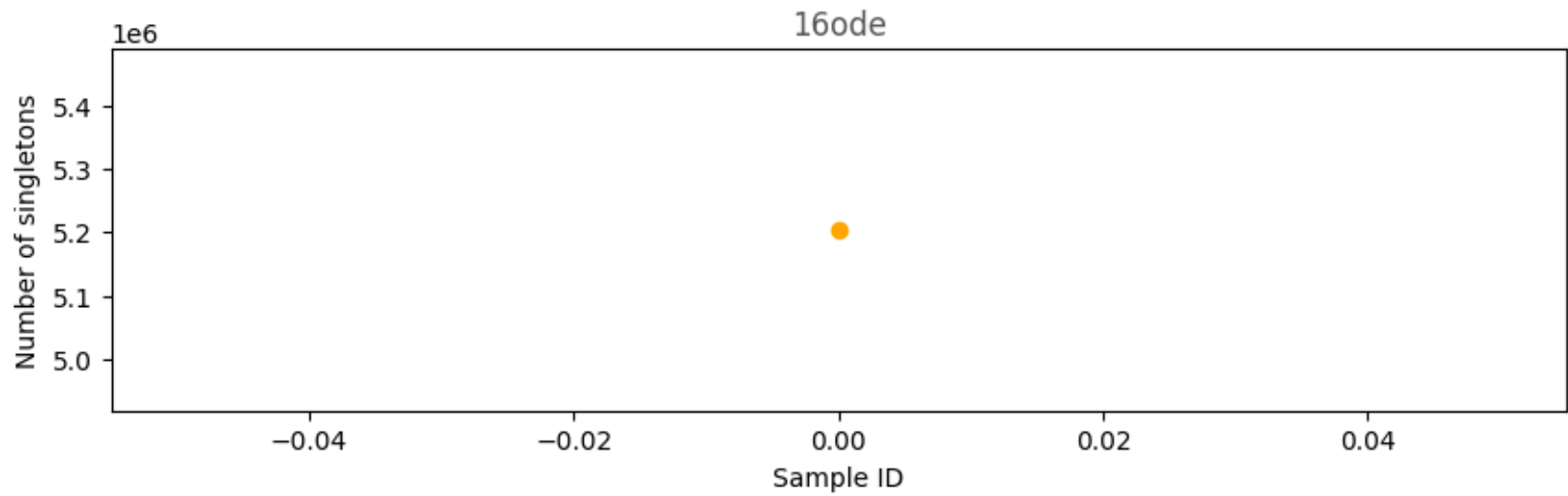 `KA_NGCWGS-NGCWGS00268_21RKG020241x01_snv_germline_raw.haplotype_caller.vcf.gz`
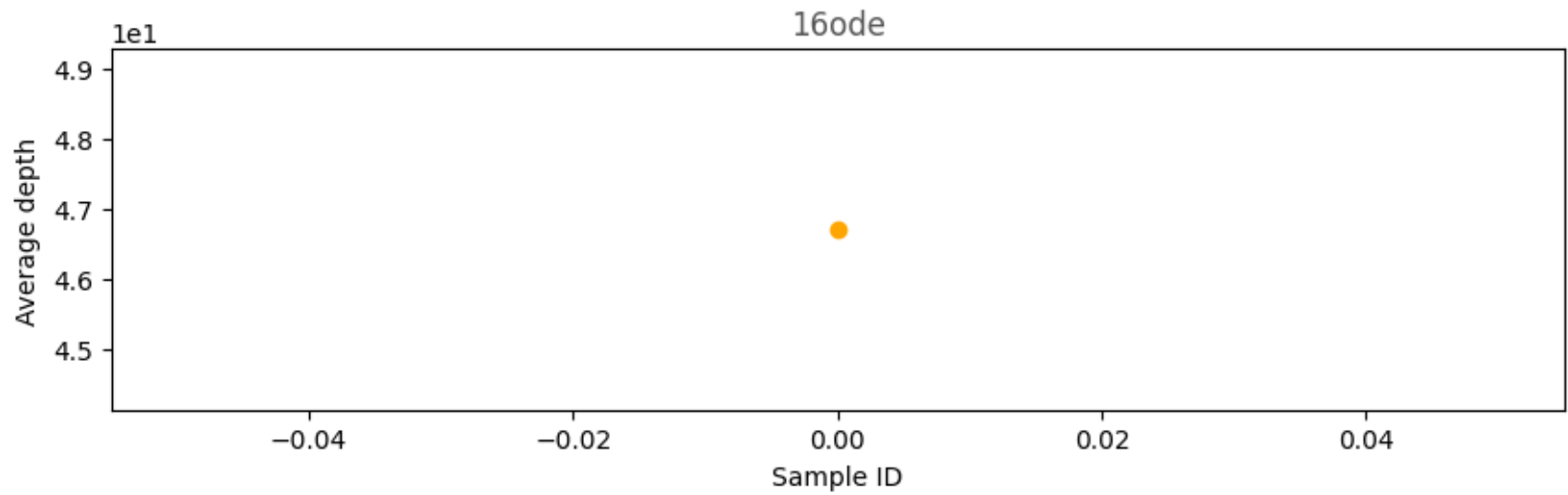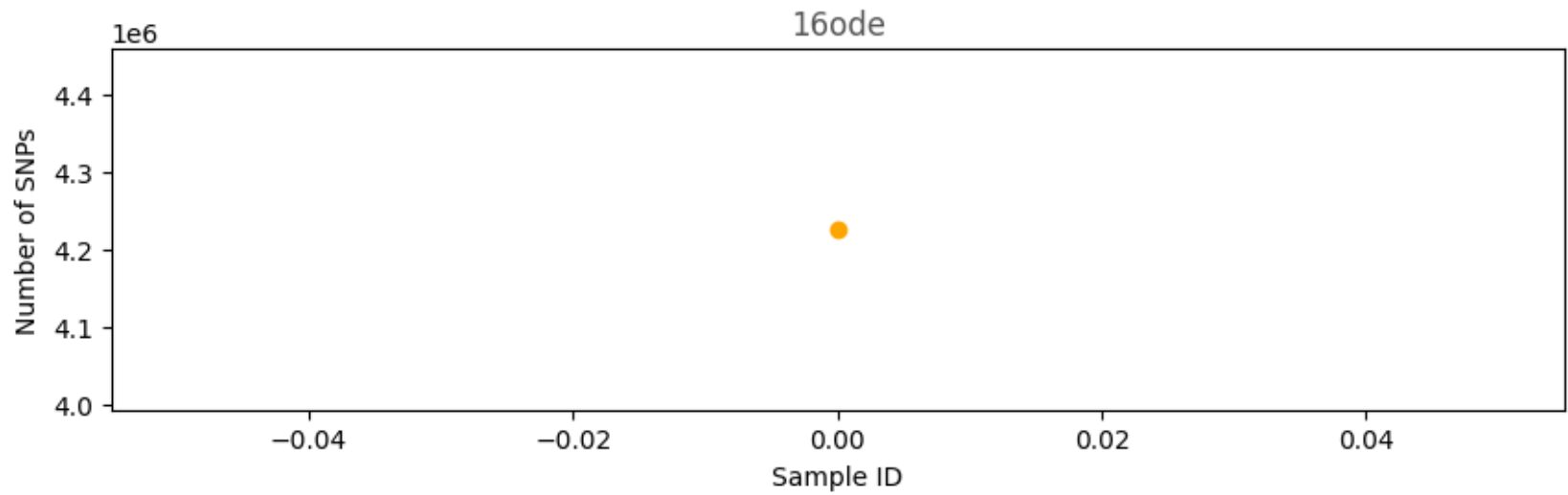
Ts/Tv by sample
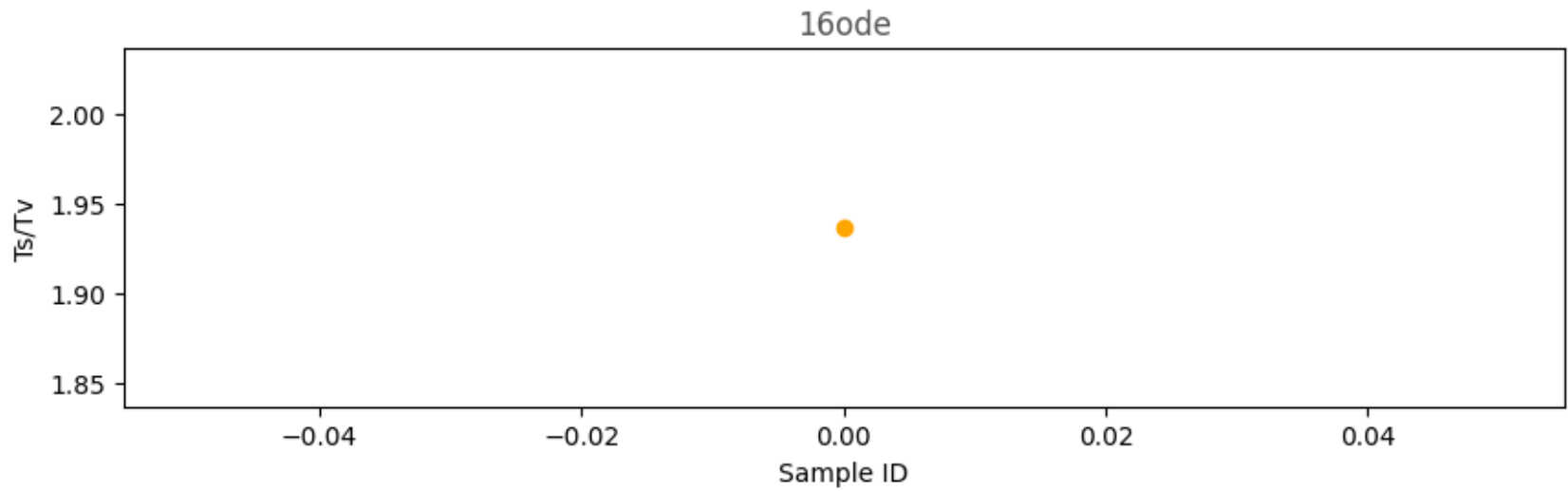
Hets vs non-ref Homs by sample
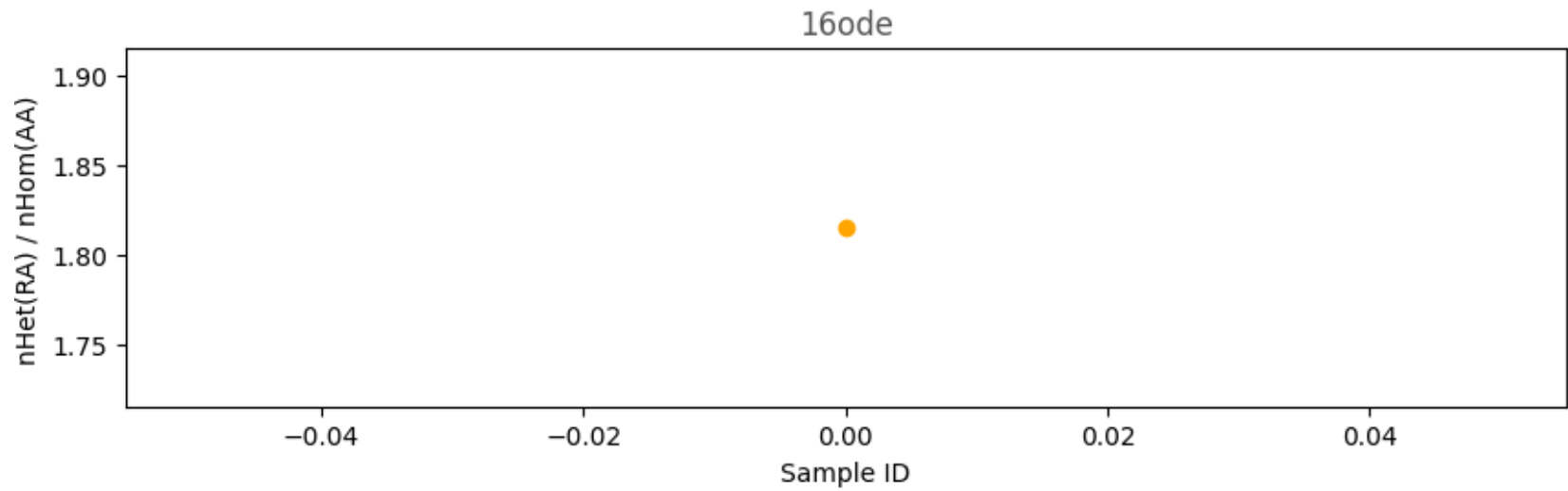
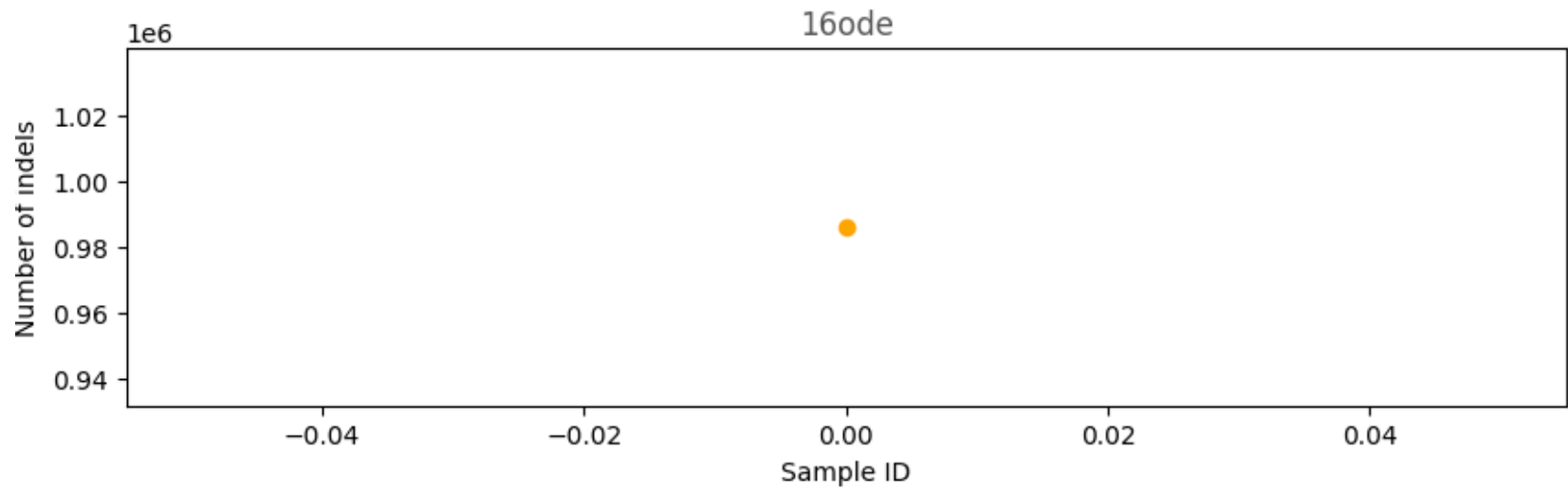# Singletons by sample (hets and homs)
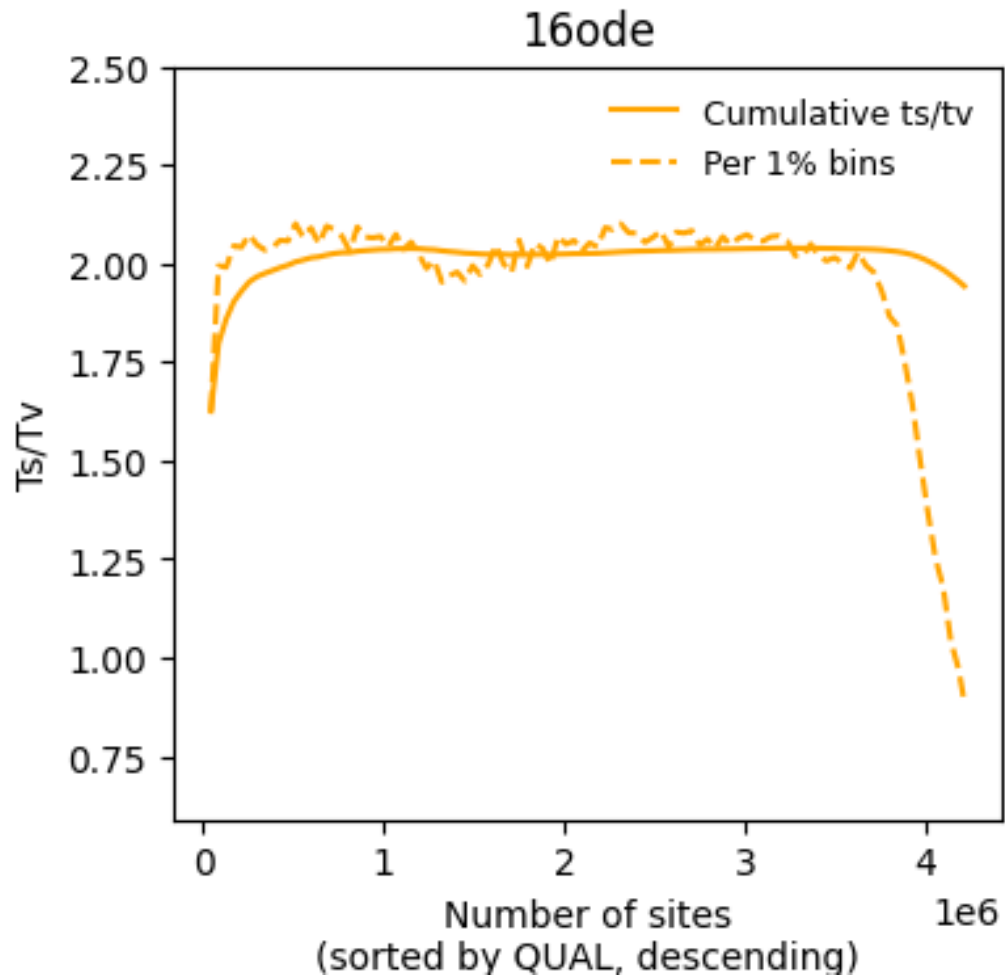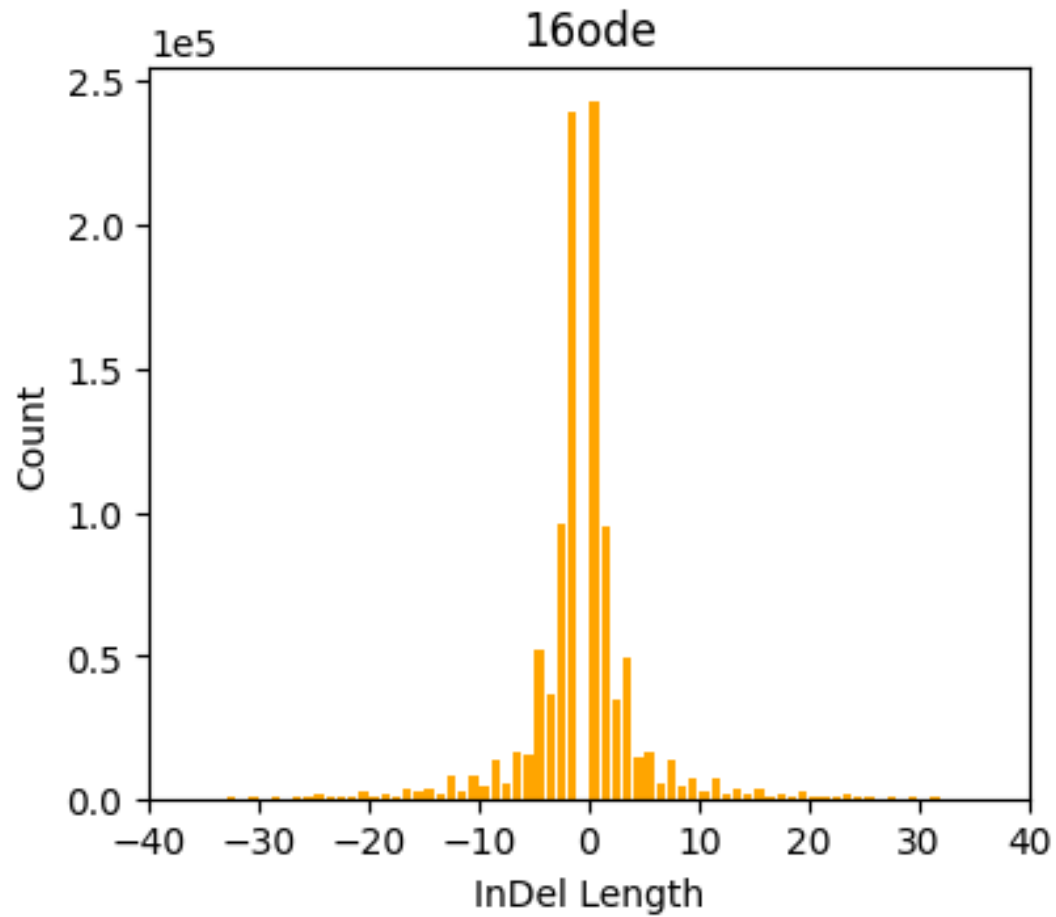
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

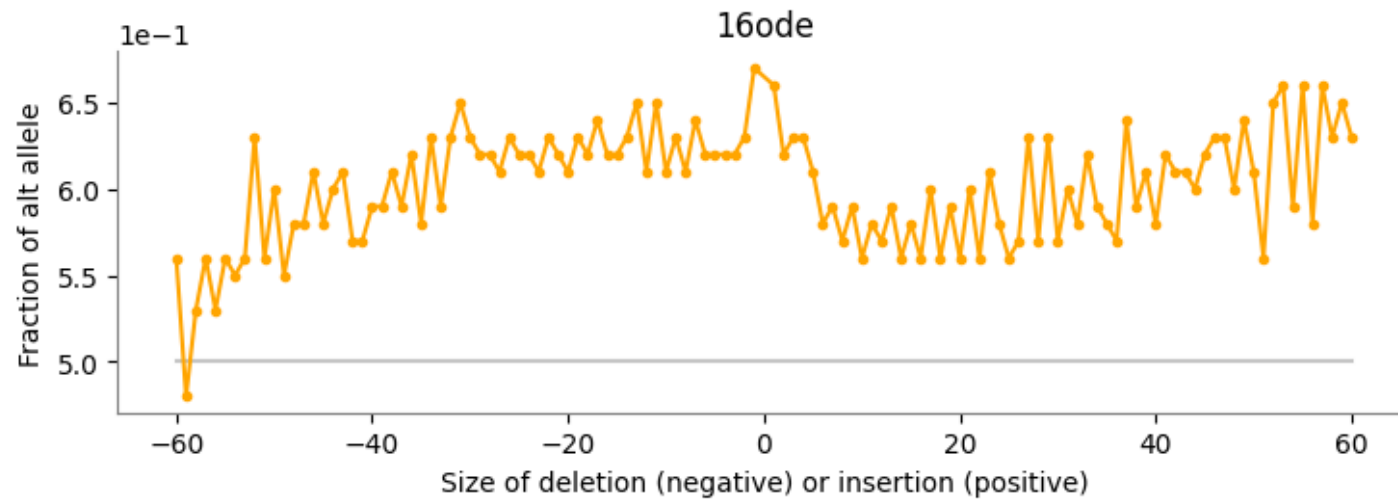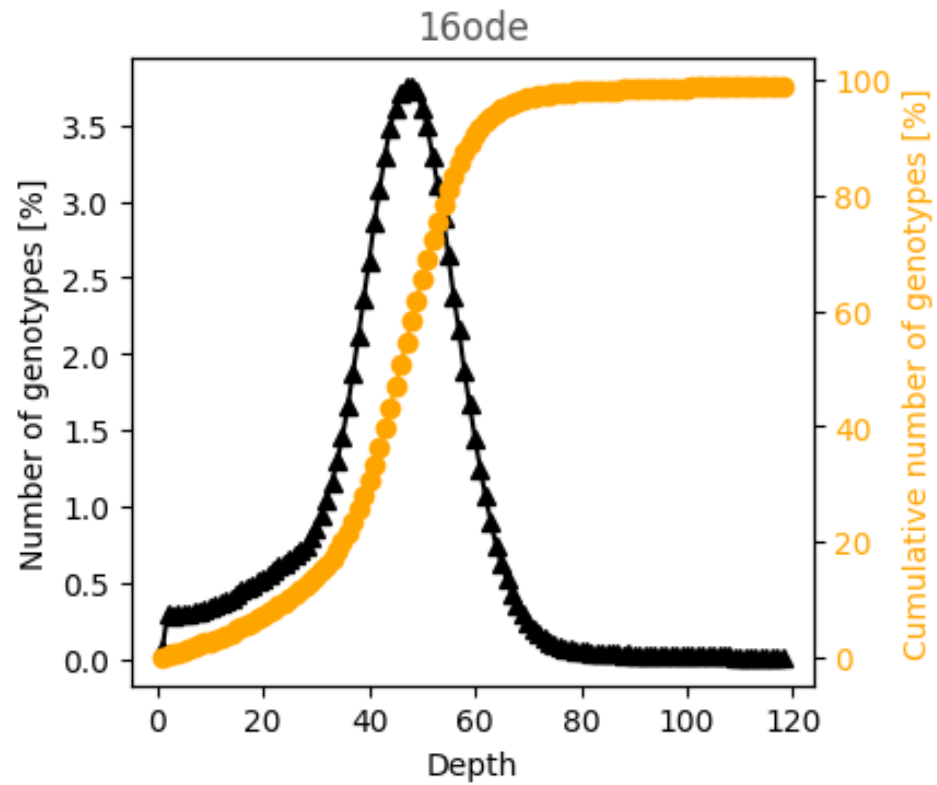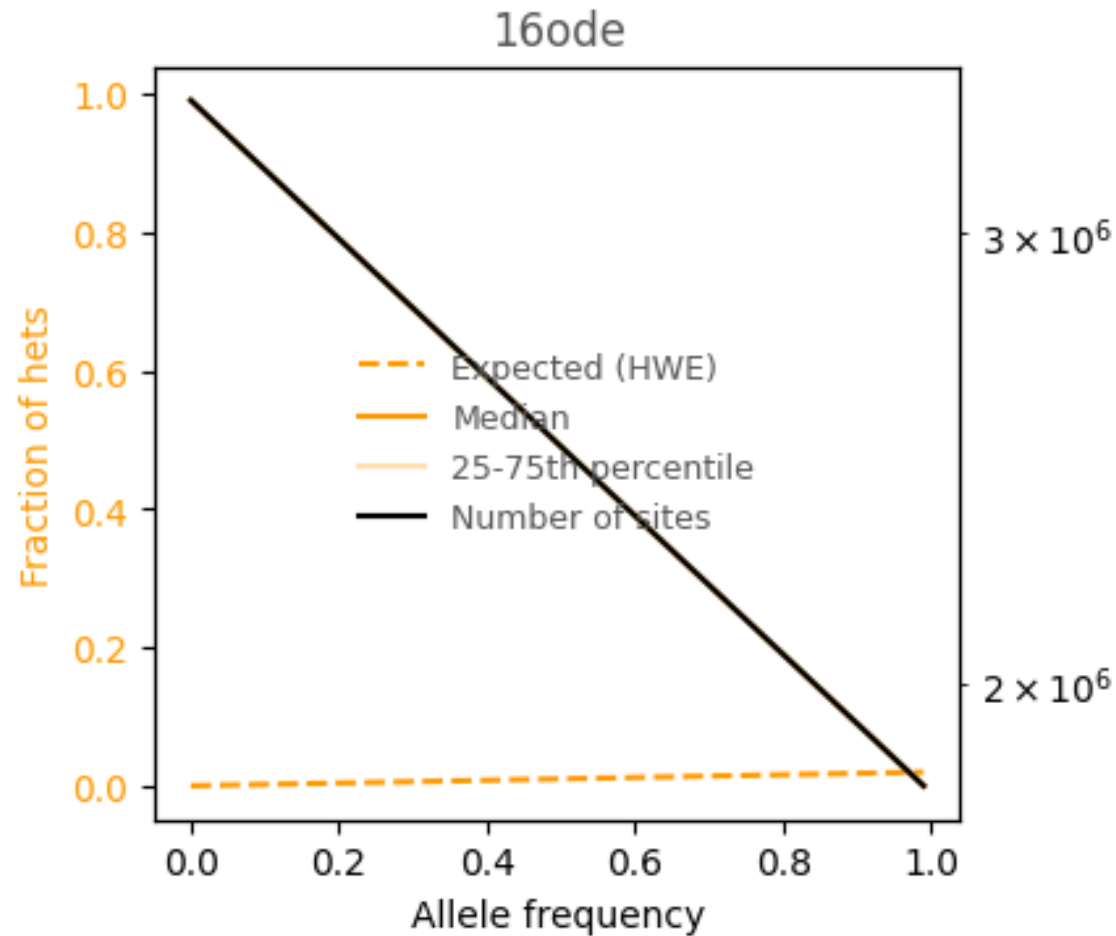# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types