Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
04iva	4,097,733	1.93	1.94	960,406	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
04iva	63.1%	1.90	68.5%	103,082	2,022

- 04iva .. `/ngc/projects2/gm/data/archive/2022/variants/snv/04ivatbef-103911275316-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01549-220916_A01411_AHTJ7JDSX3-EXT_LAB`
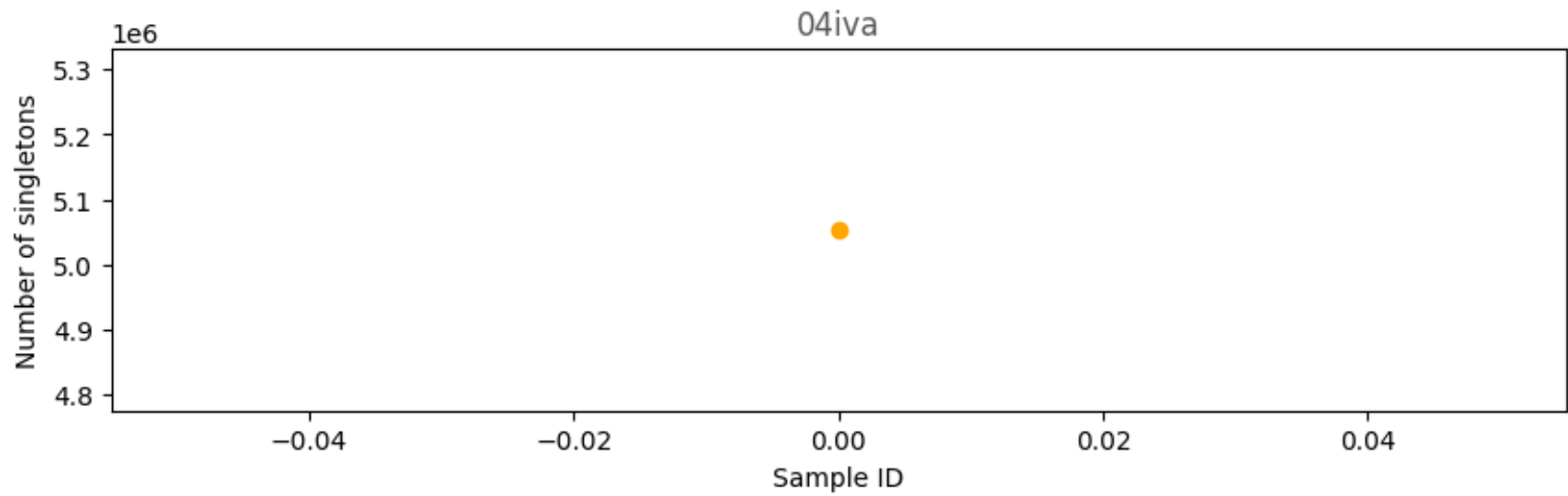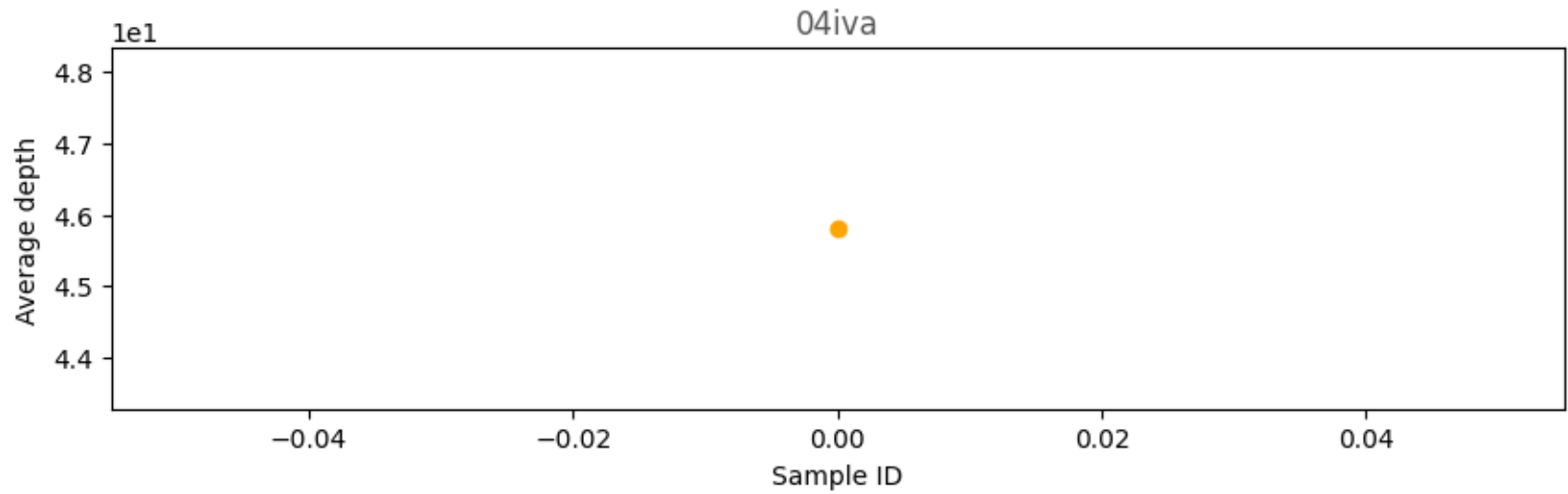 `KA_NGCWGS-NGCWGS05392_snv_germline_raw.haplotype_caller.vcf.gz`

Ts/Tv by sample
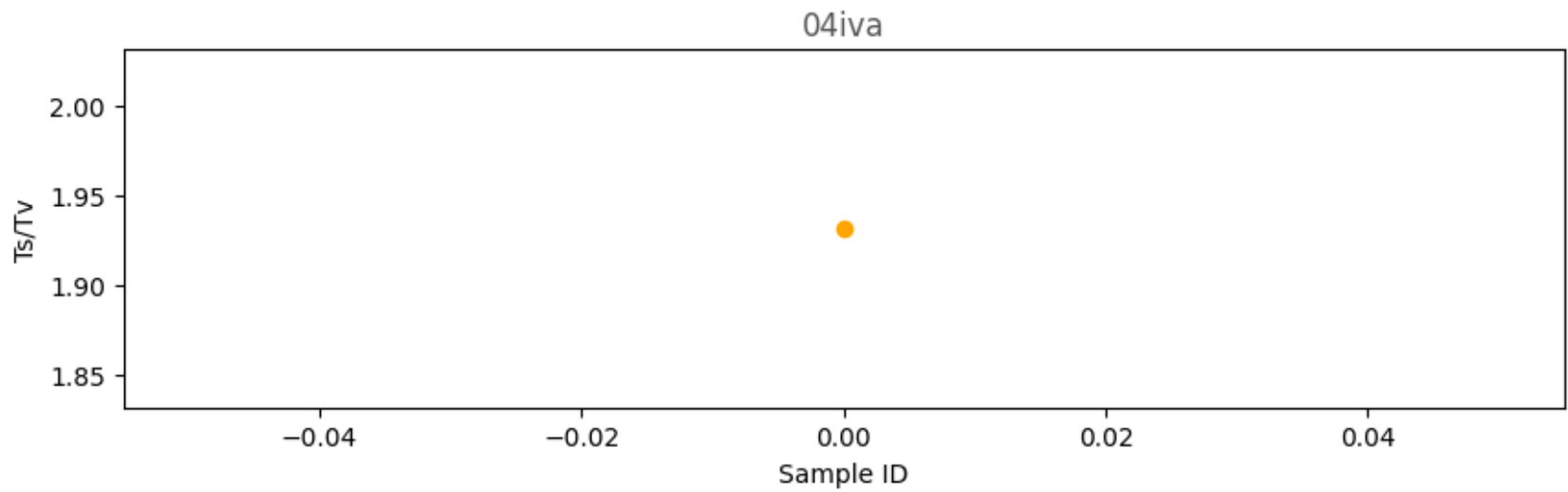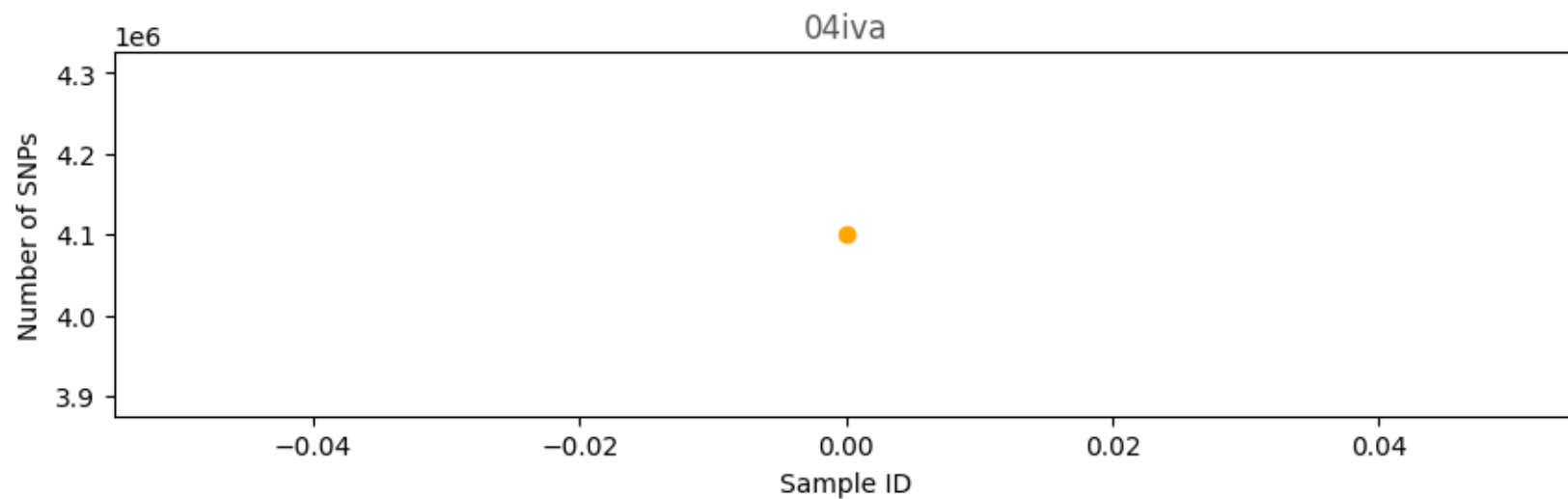
Hets vs non-ref Homs by sample
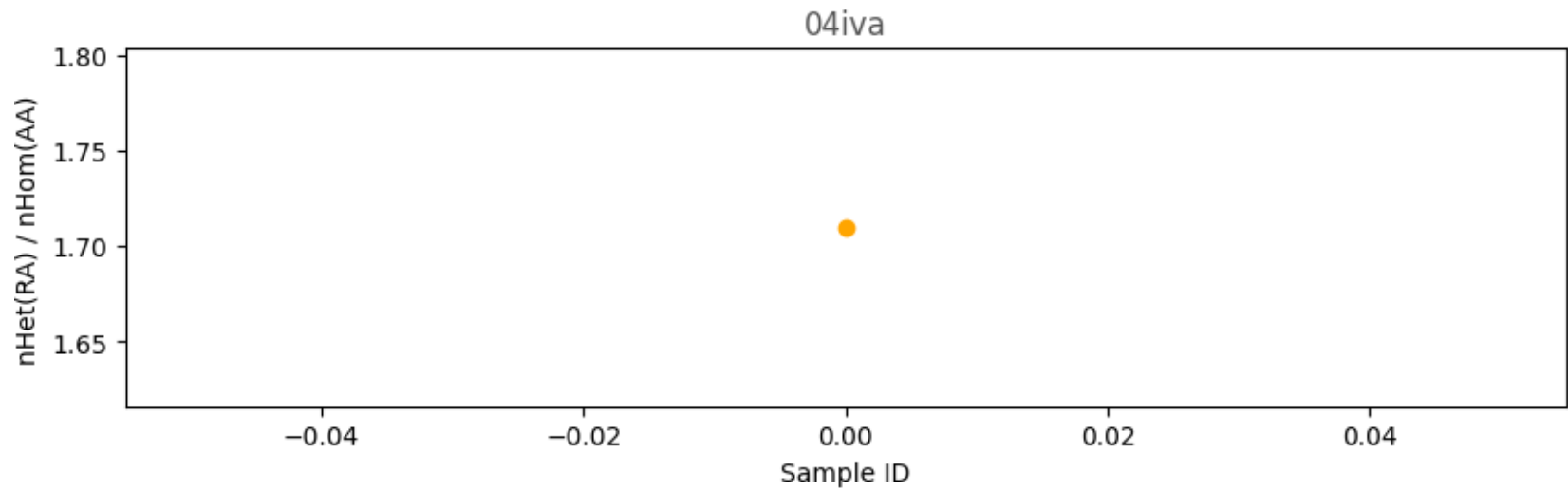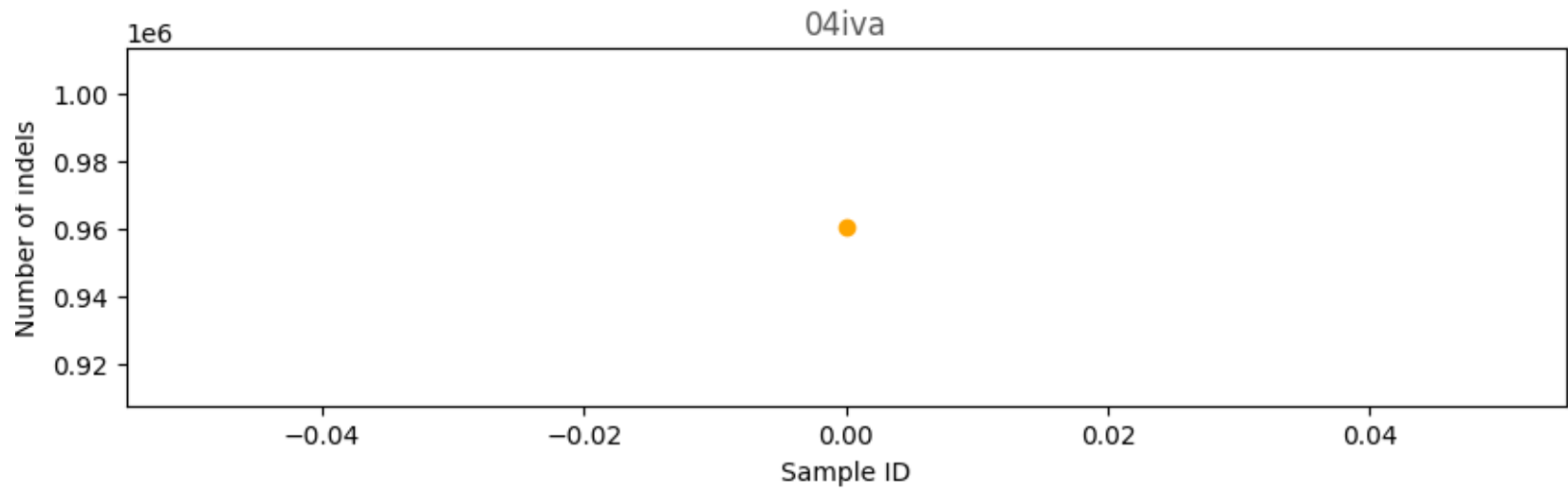
Singletons by sample (hets and homs)
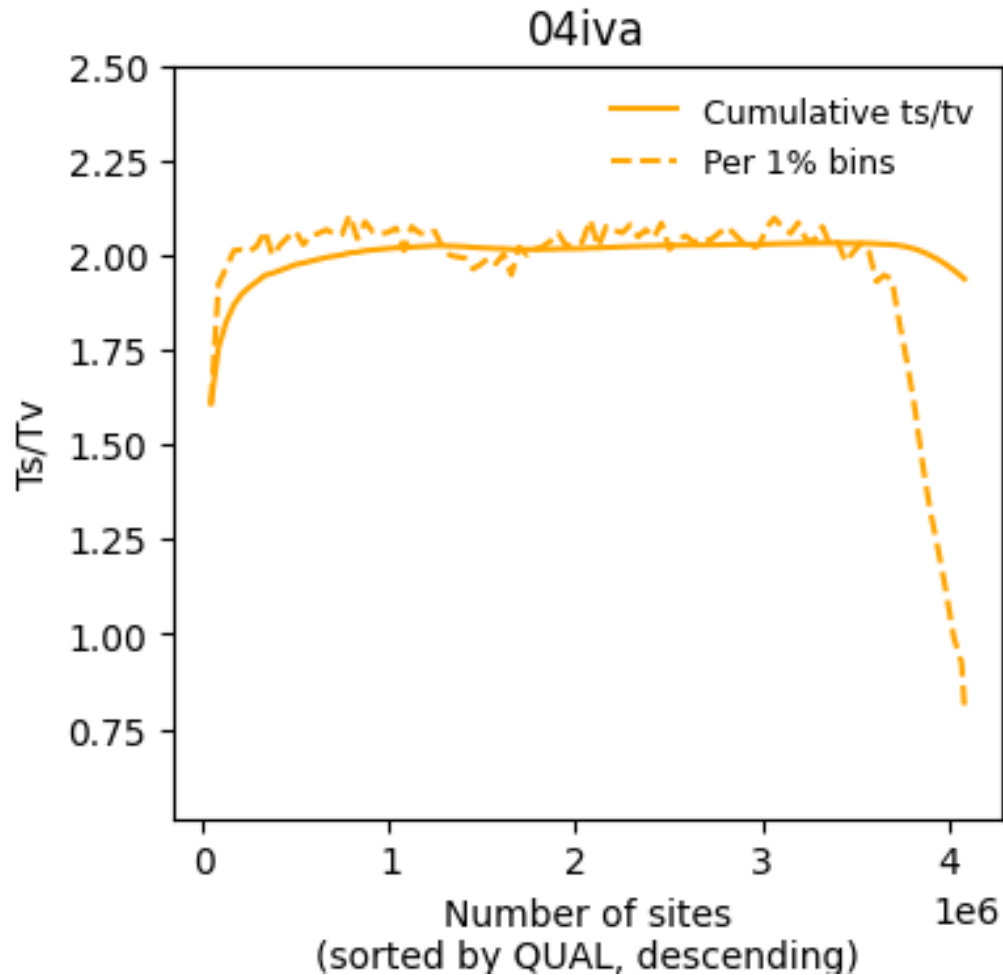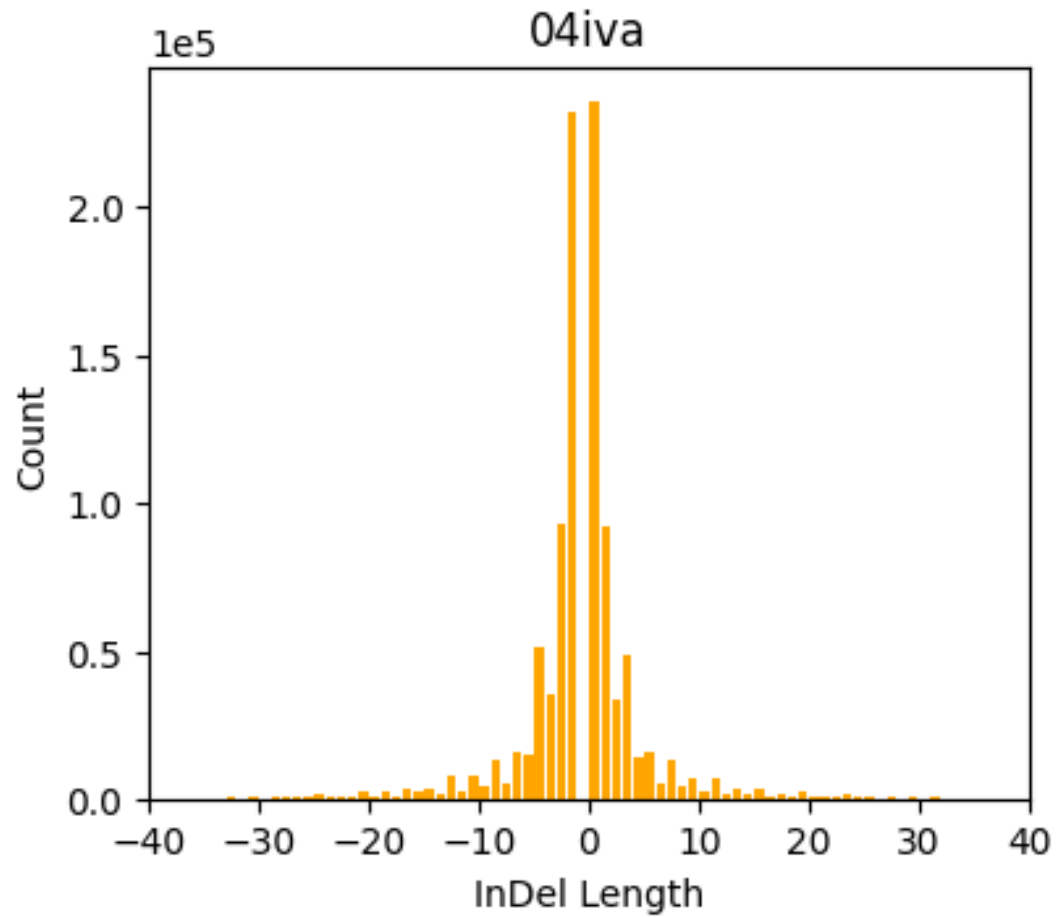
Average depth by sample
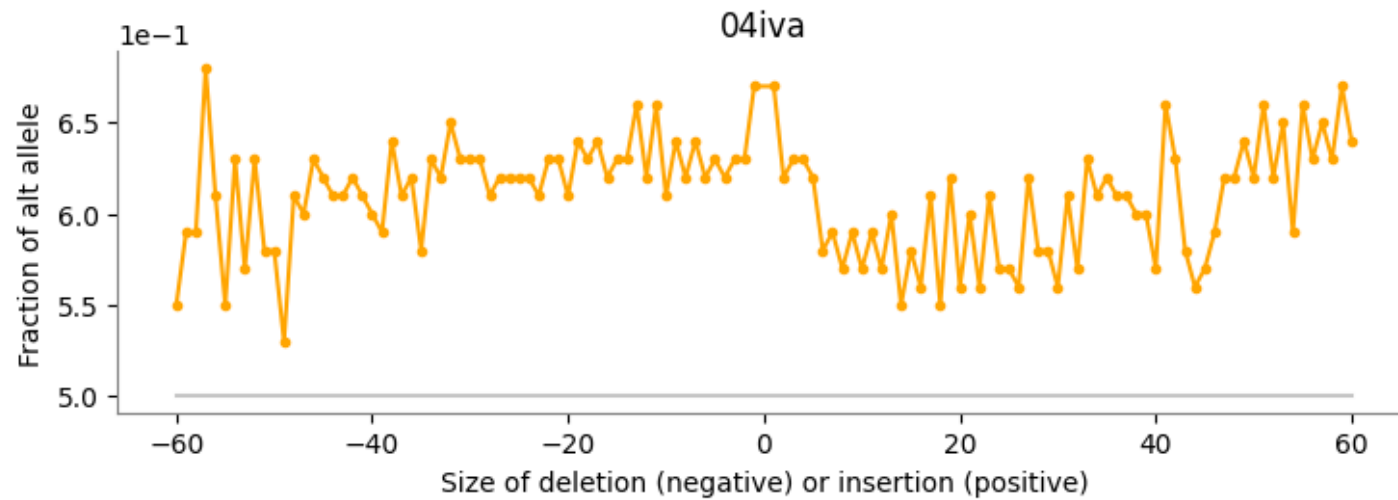
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

04iva

Indel distribution

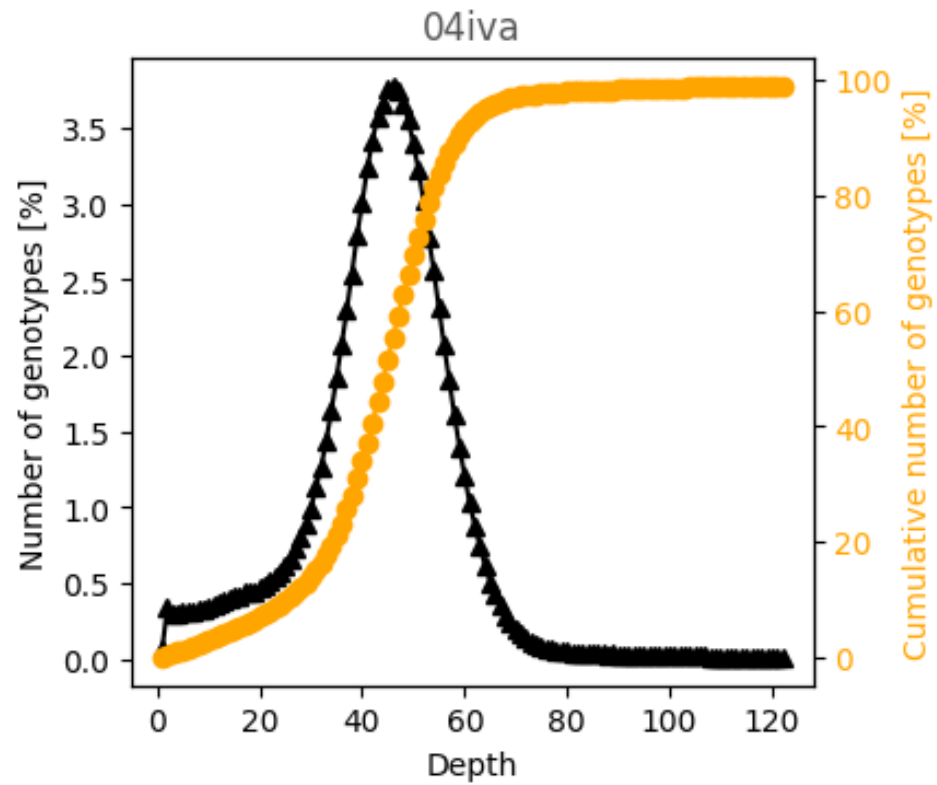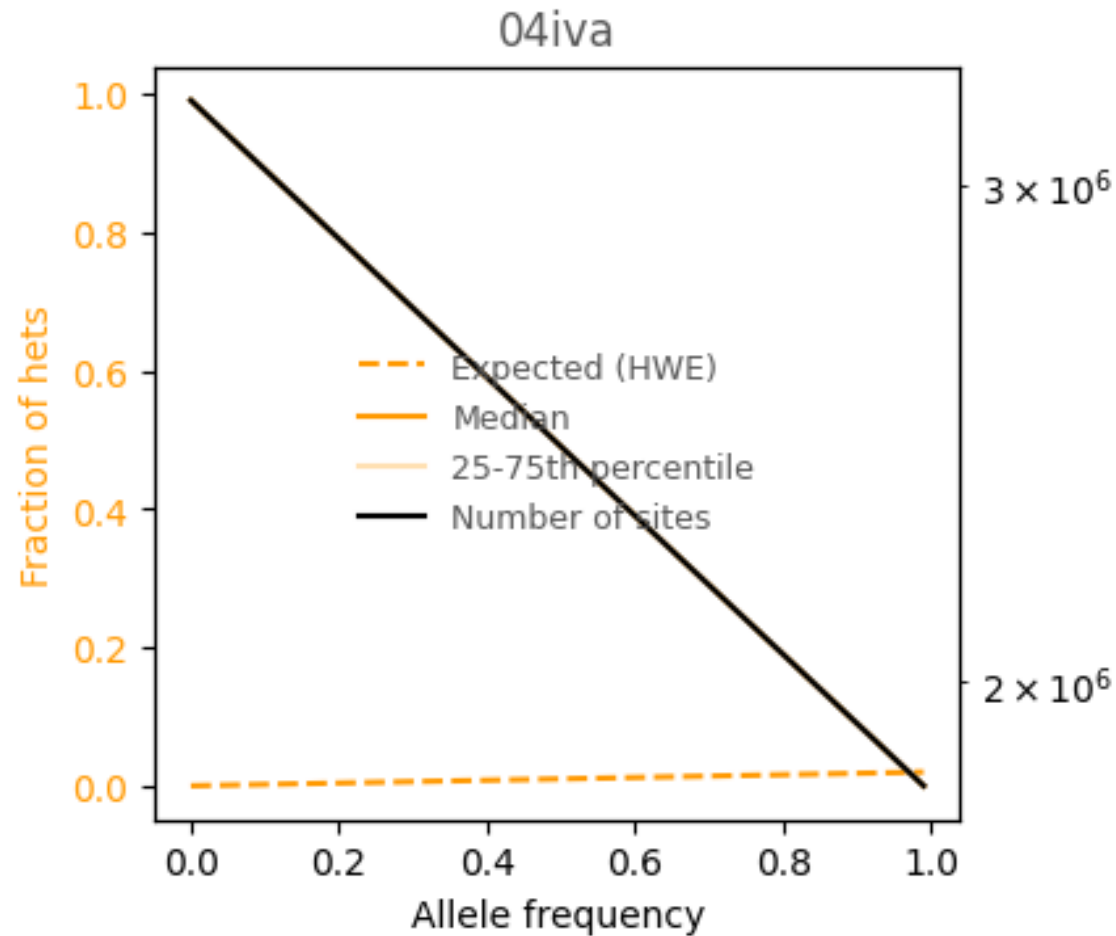# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types