Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
86oti	4,034,937	1.95	1.95	947,960	–	0	0

* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
86oti	62.1%	1.92	66.9%	99,496	2,000

- 86oti .. `/ngc/projects2/gm/data/archive/2022/variants/snv/86otienim-103885495826-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220706_A01176_AHKCJ2DSX3-RHGM_LABKA_WGS-` `WGS04587_snv_germline_raw.haplotype_caller.vcf.gz`
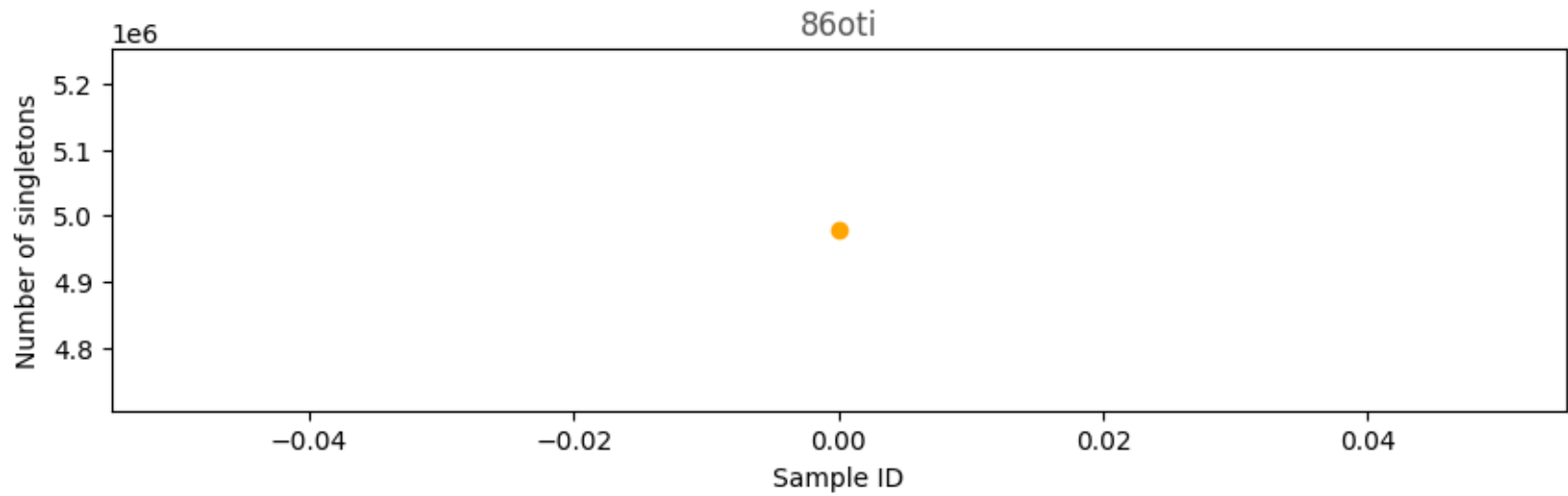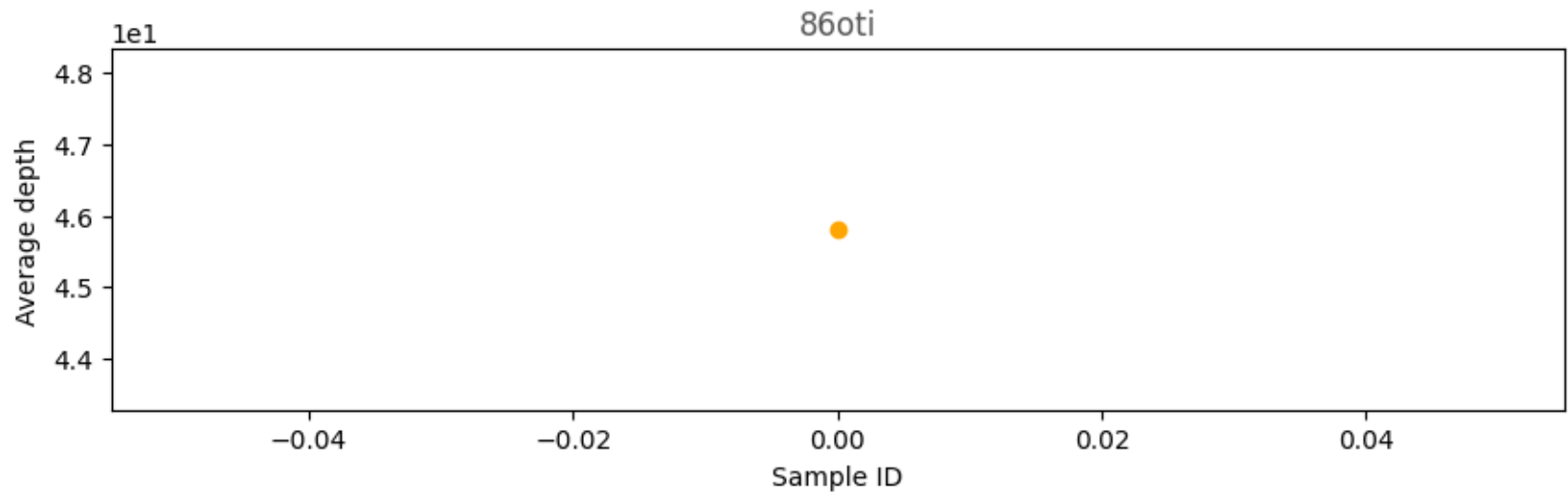
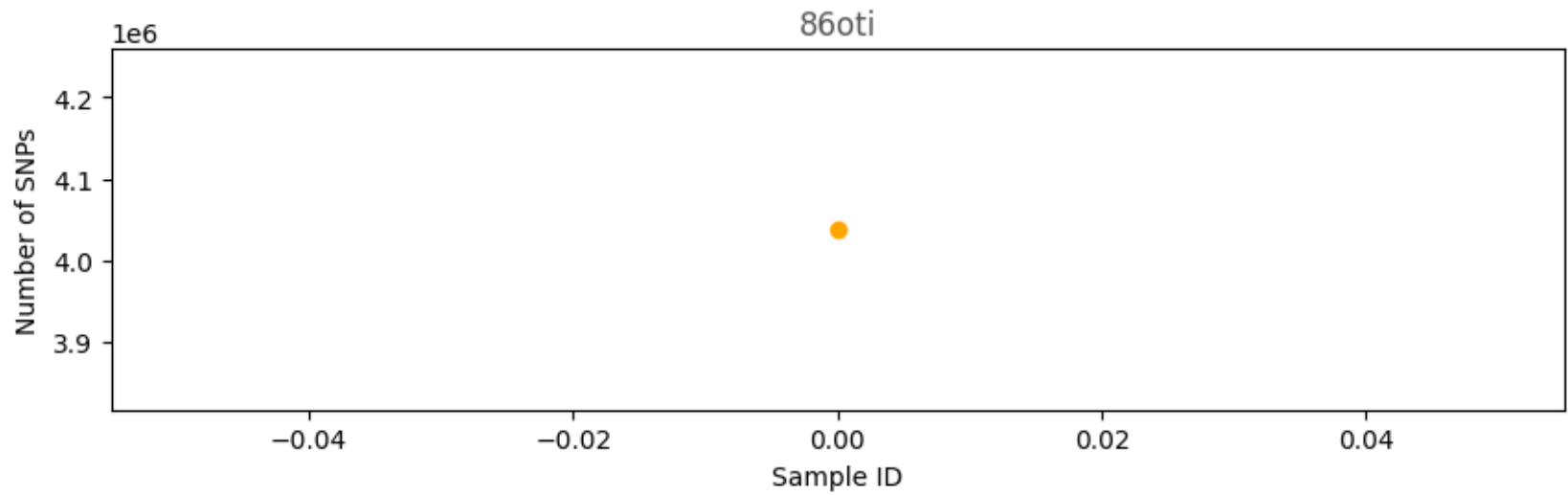# Ts/Tv by sample

86oti
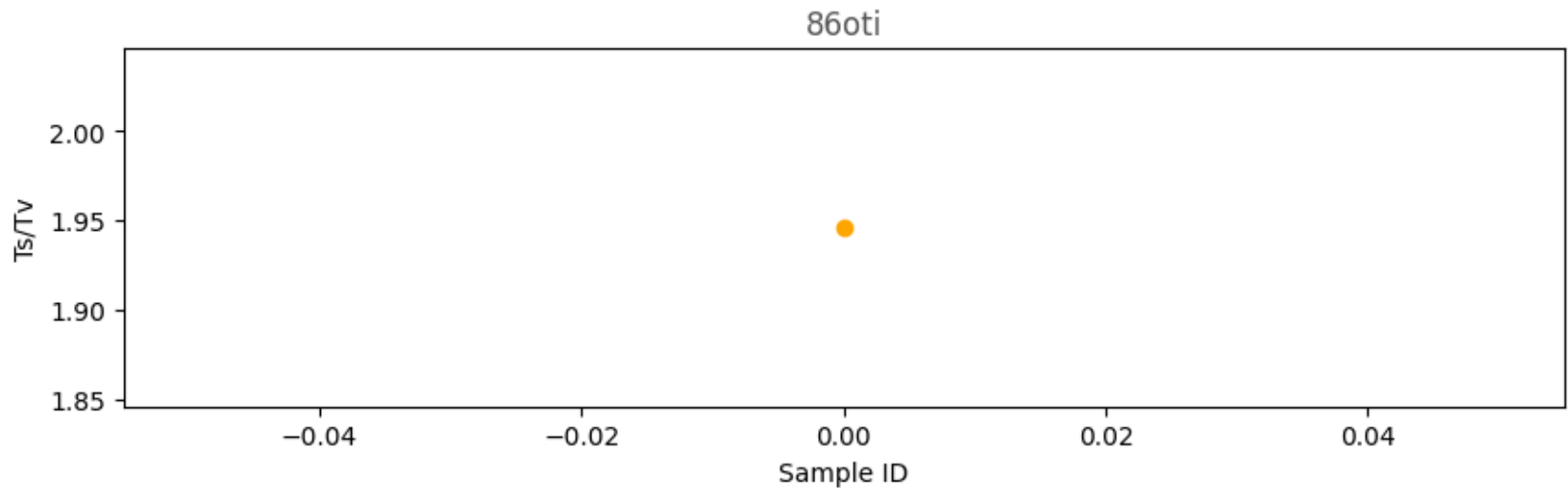
Hets vs non-ref Homs by sample
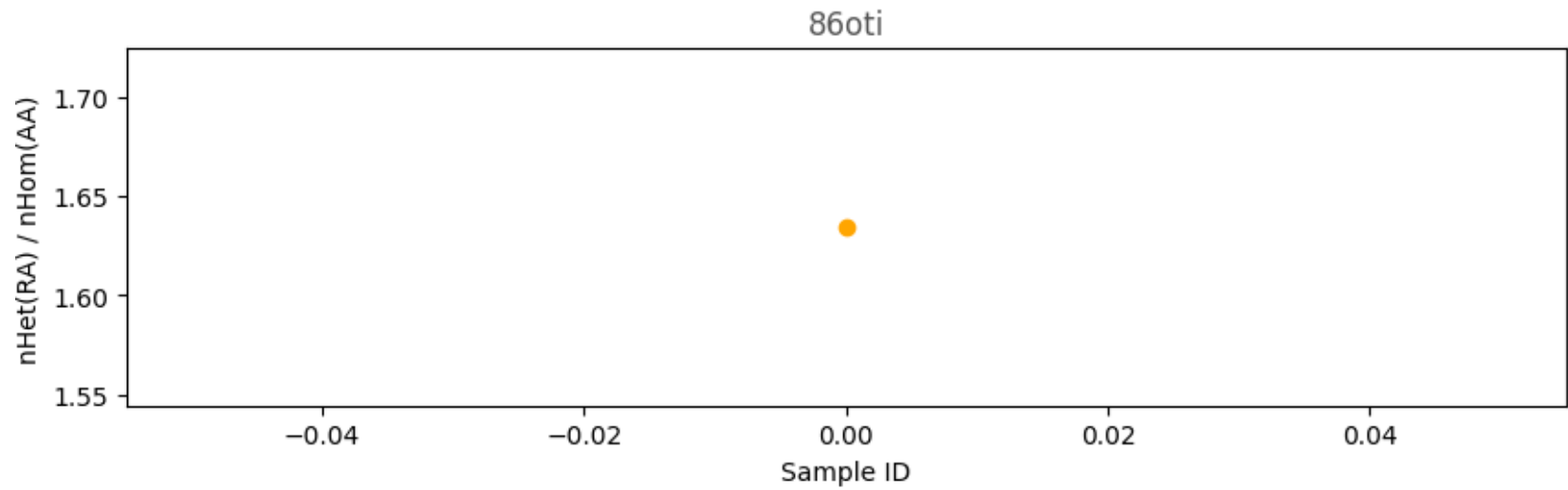
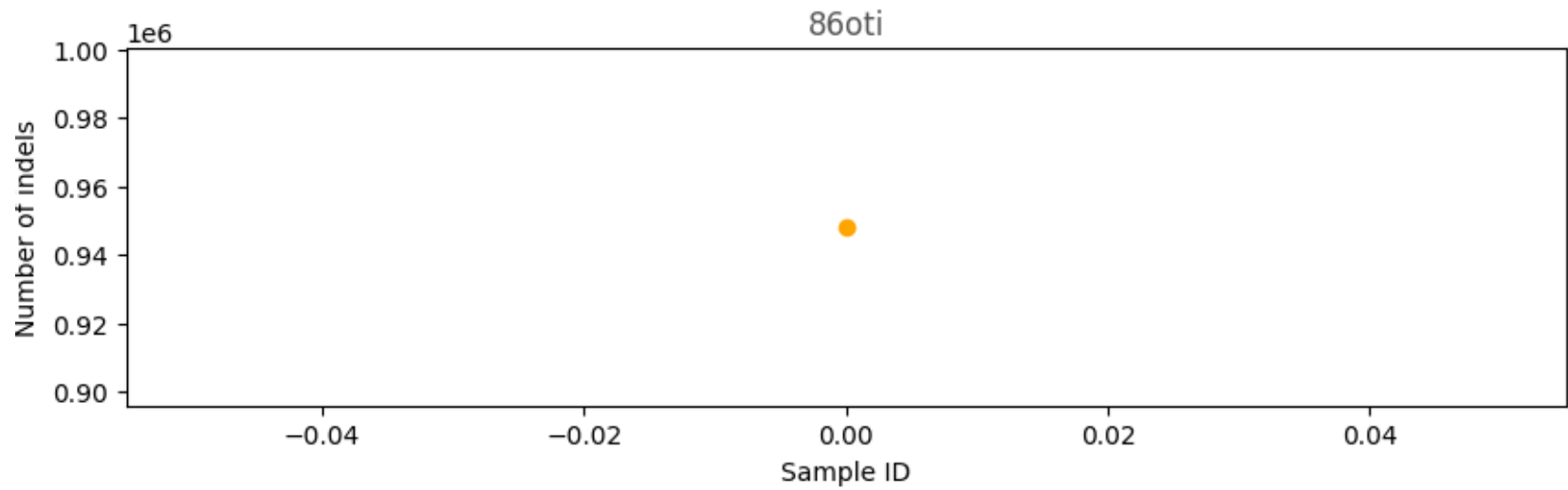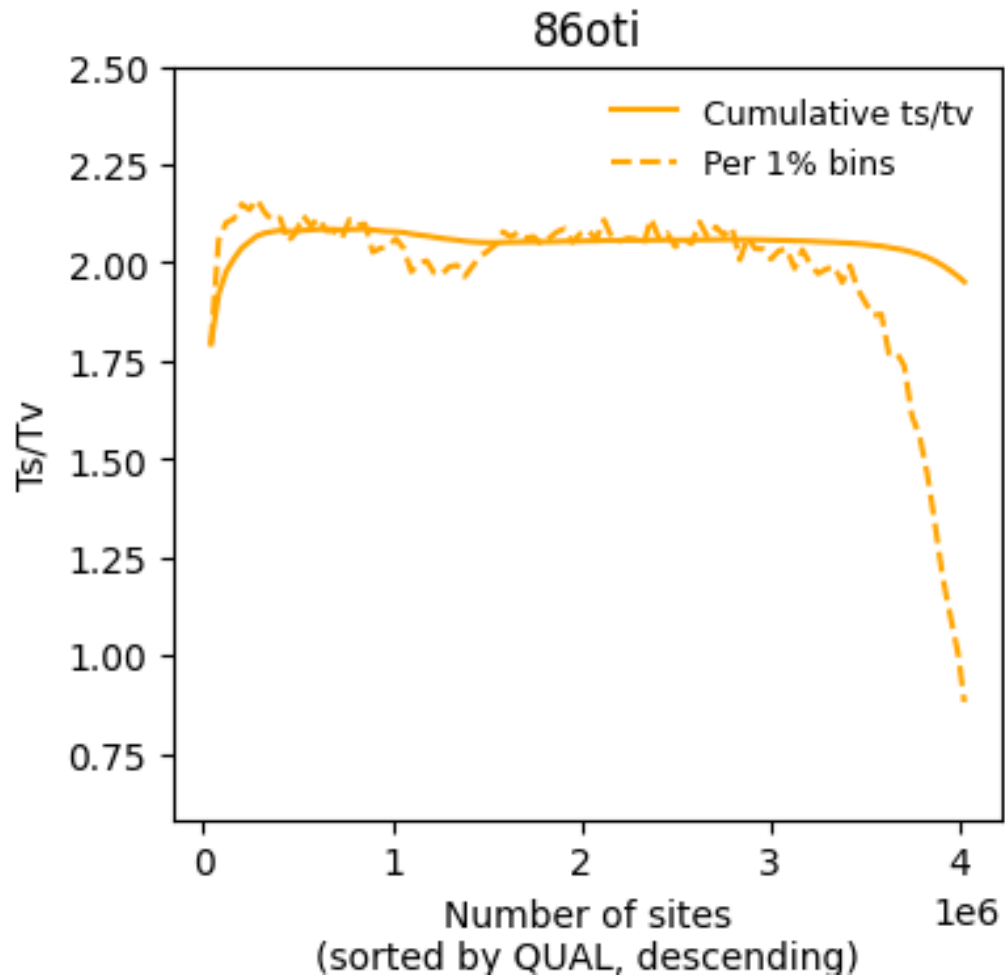# Singletons by sample (hets and homs)
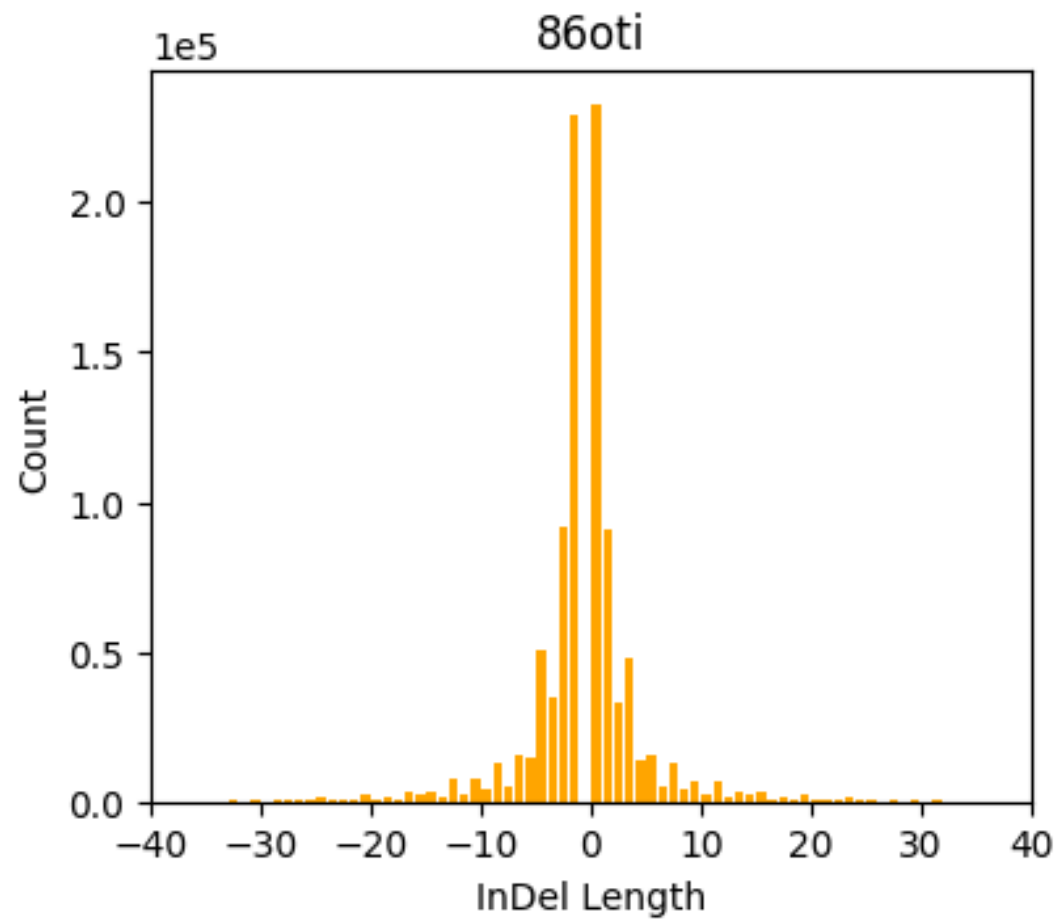
Average depth by sample
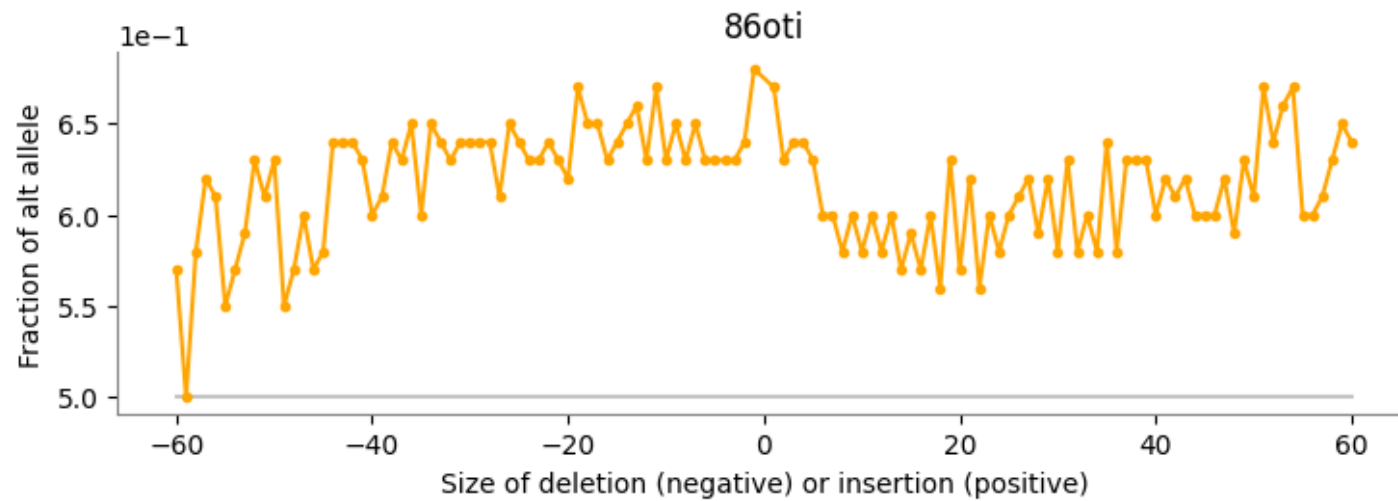
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

86oti

Indel distribution

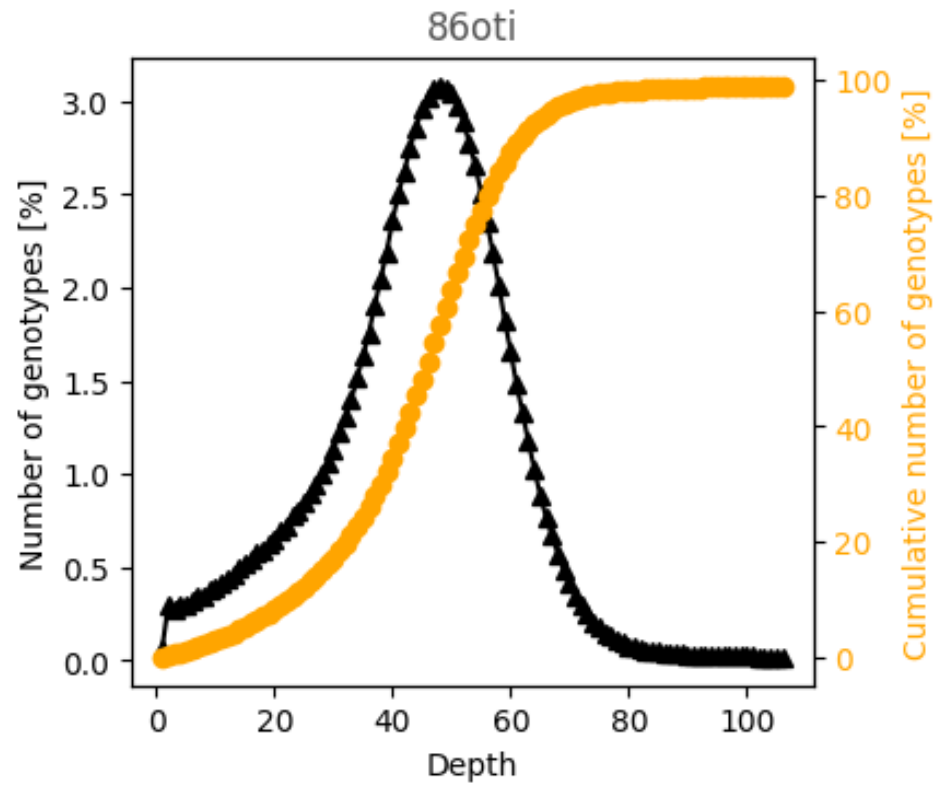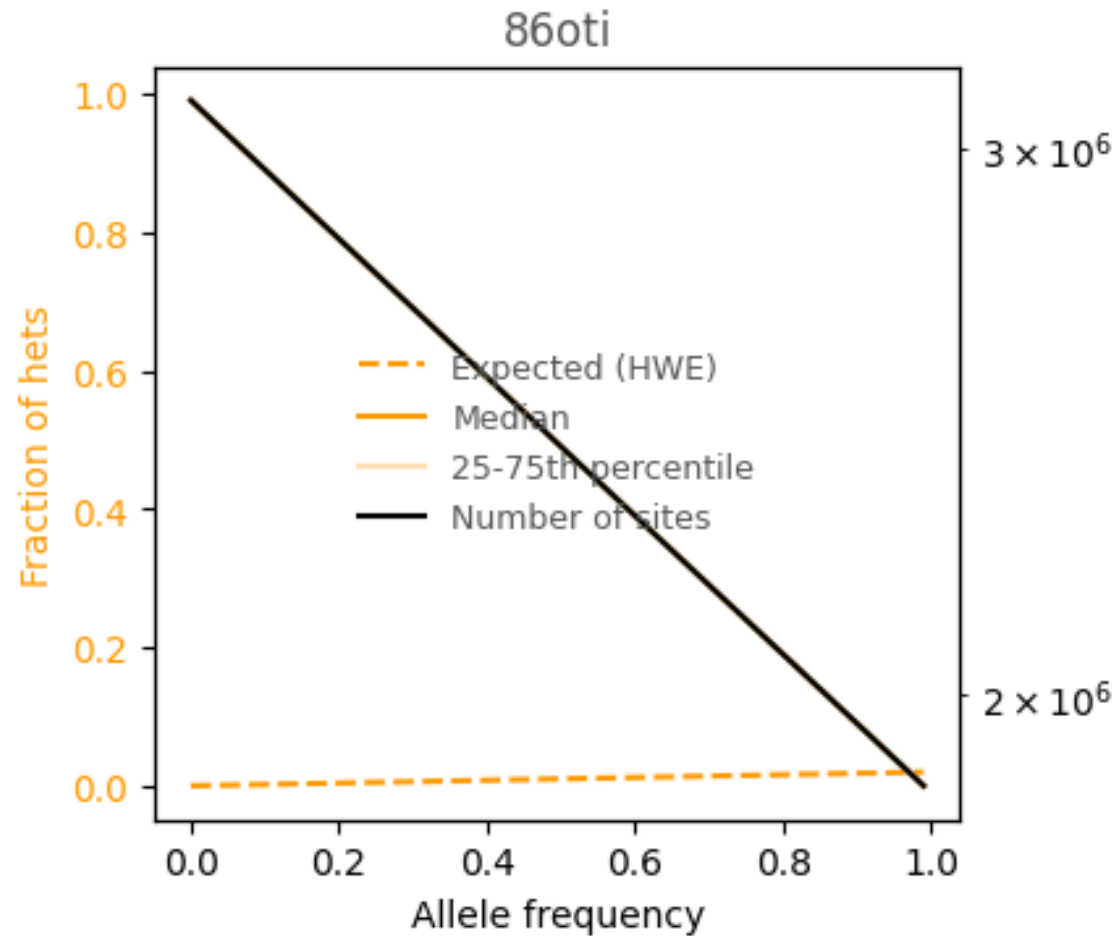# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types