Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78ote	4,128,263	1.92	1.93	970,052	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78ote	63.0%	1.89	68.8%	109,333	2,327

- 78ote .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78otedvef-110297573040-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02015-221021_A01411_AH3MJYDSX5-EXT_LAB`
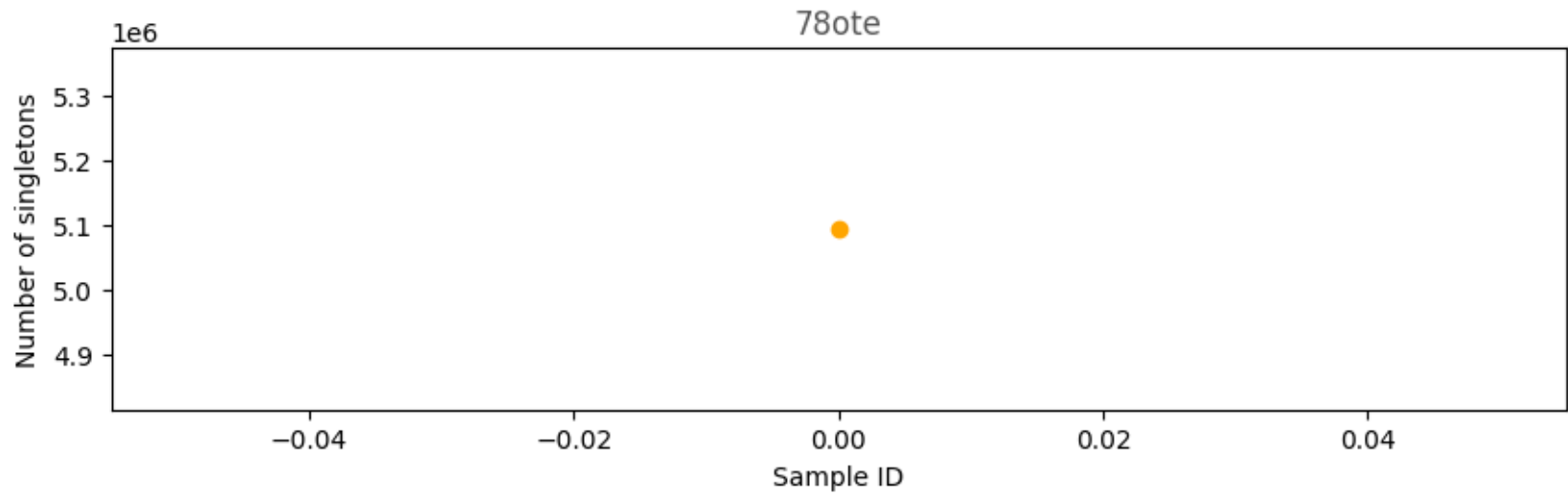 `KA_NGCWGS-NGCWGS05888_22RKG023952_snv_germline_raw.haplotype_caller.vcf.gz`
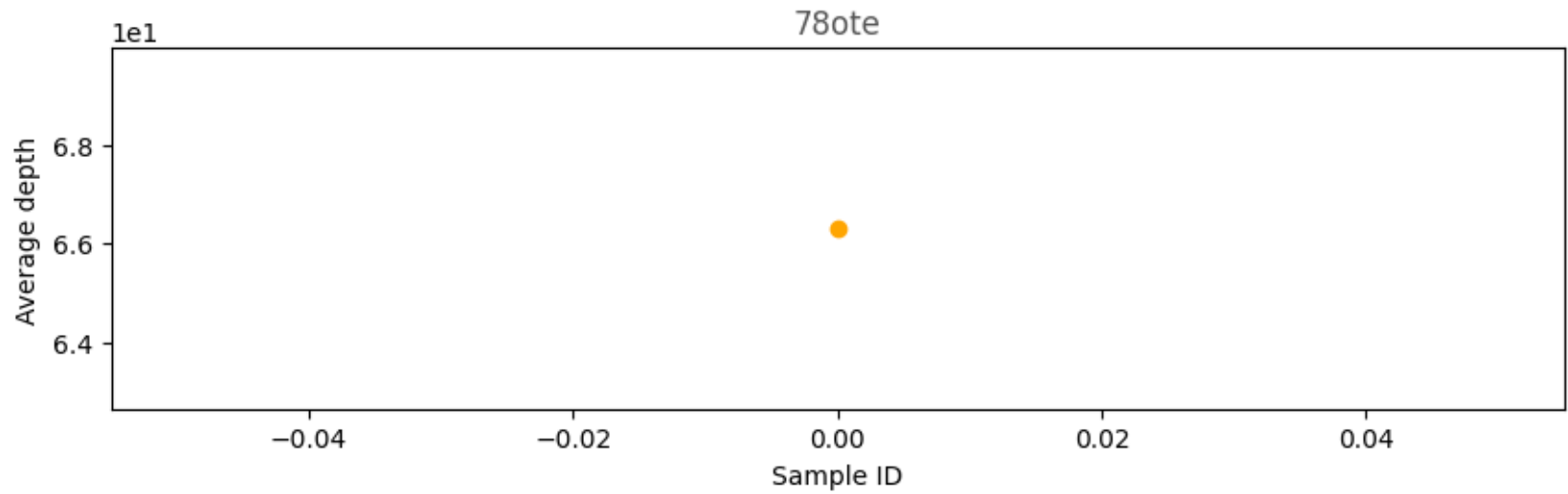
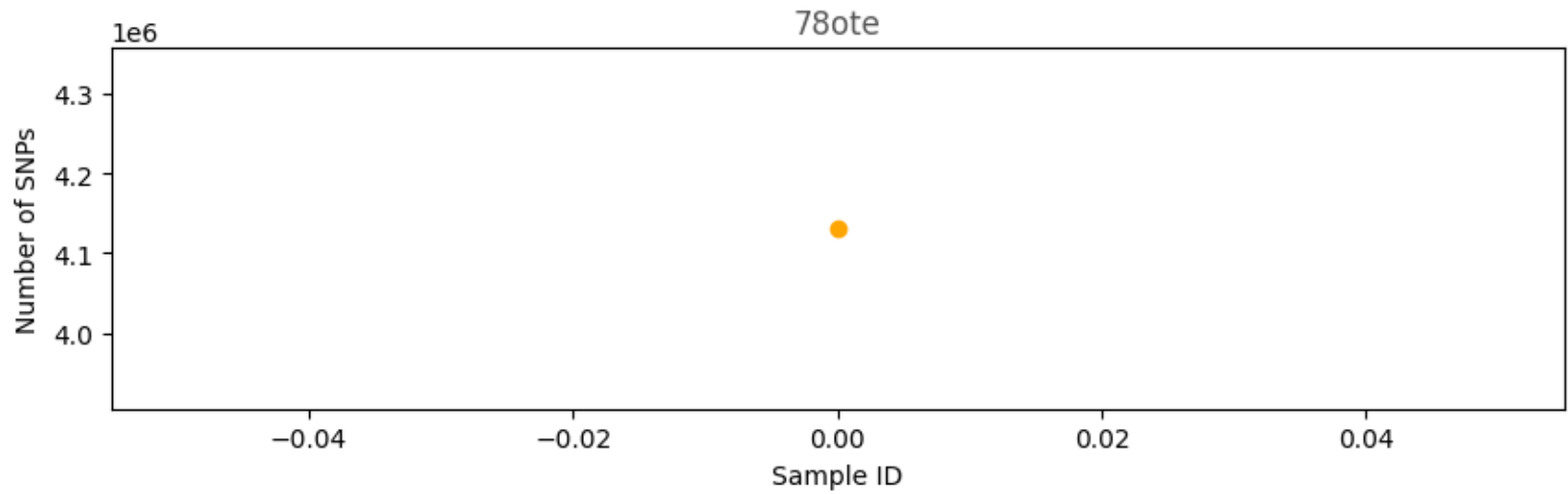# Ts/Tv by sample
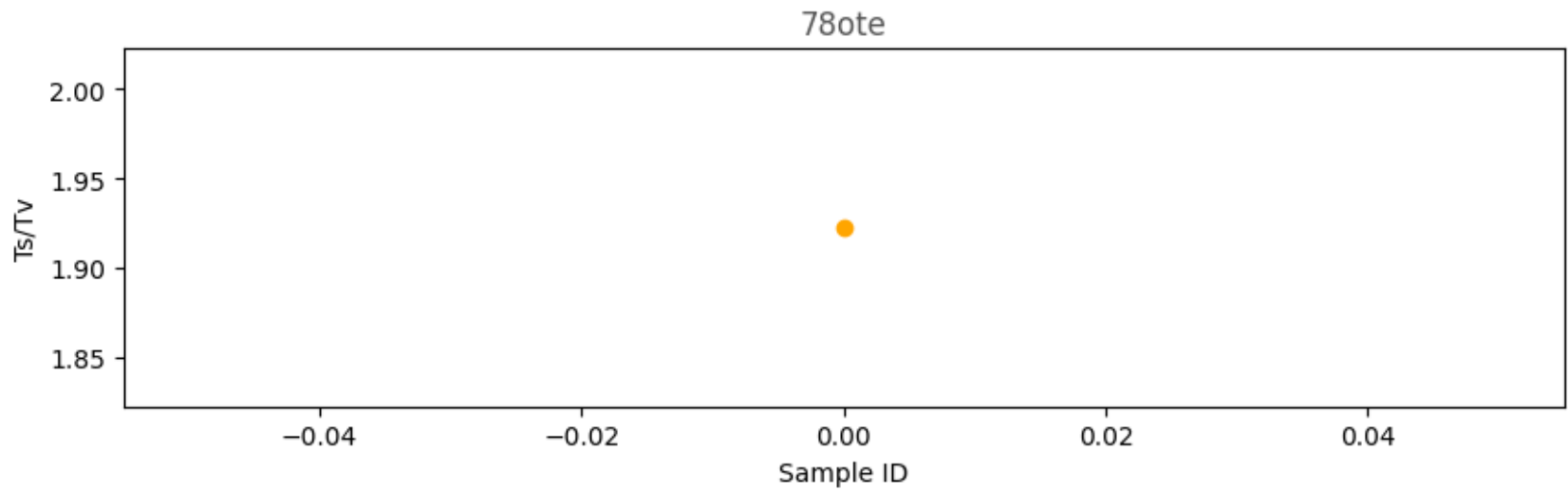
Hets vs non-ref Homs by sample
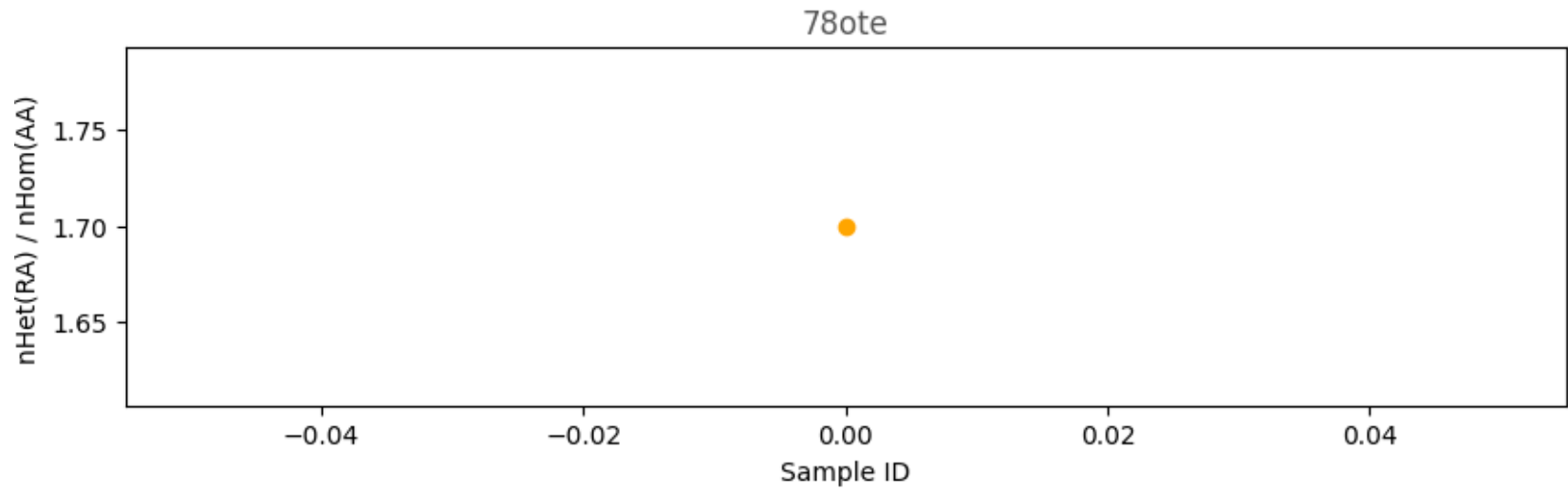
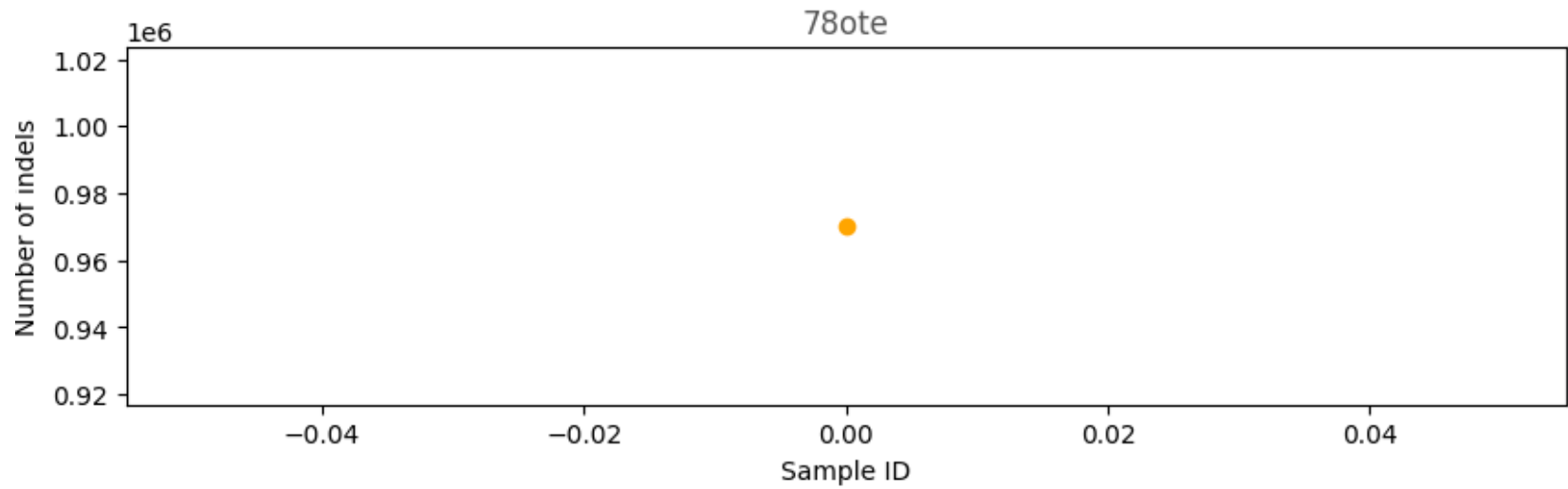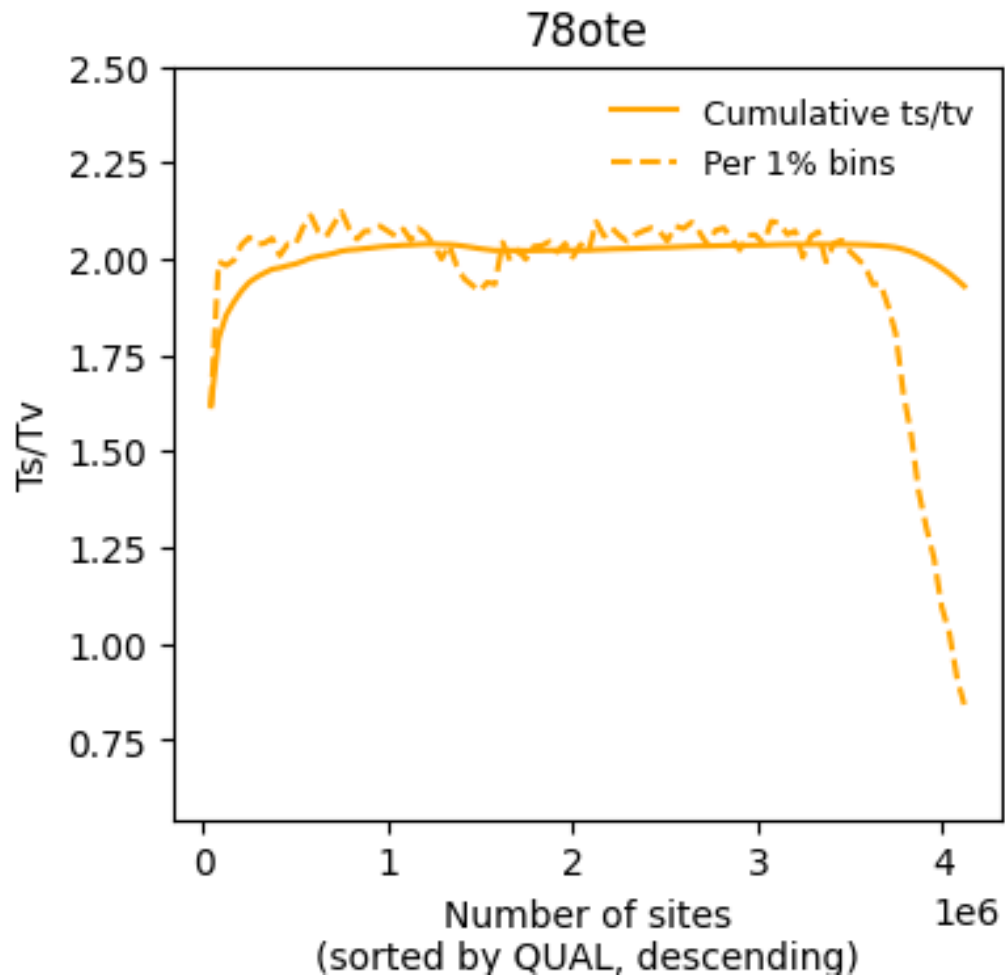# Singletons by sample (hets and homs)
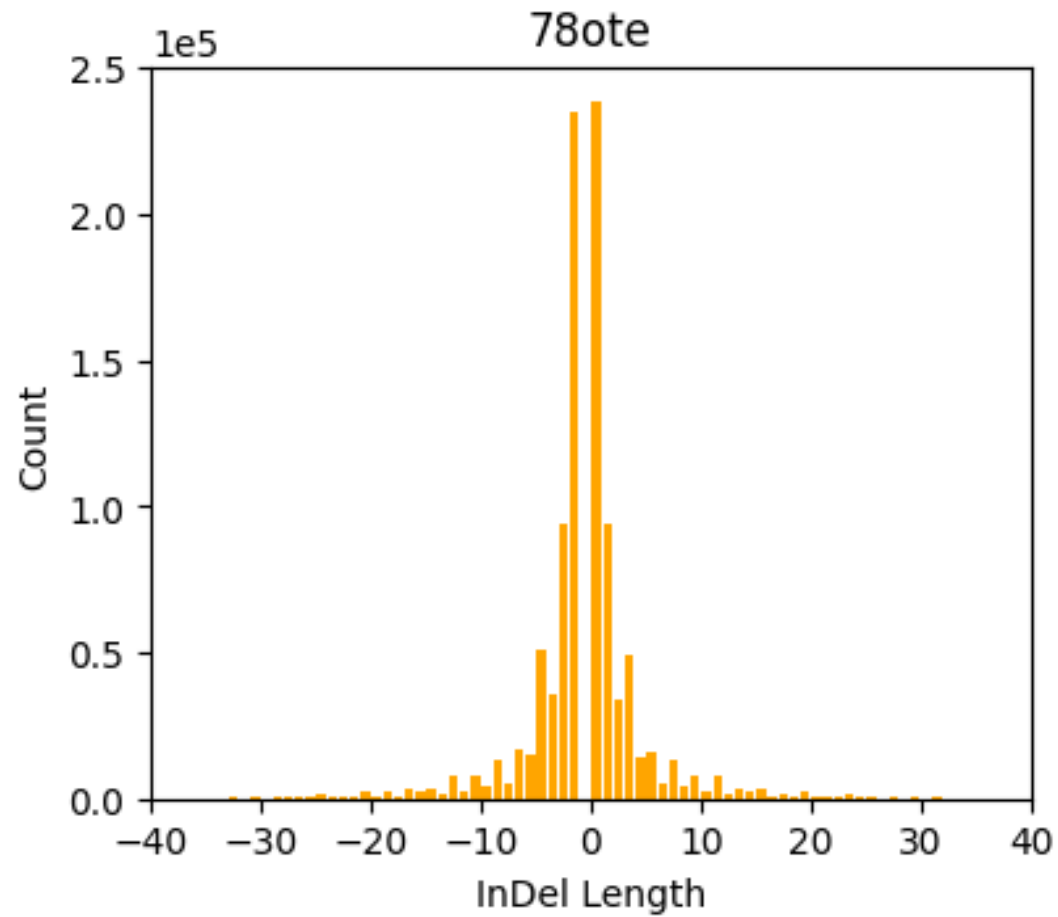
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

8

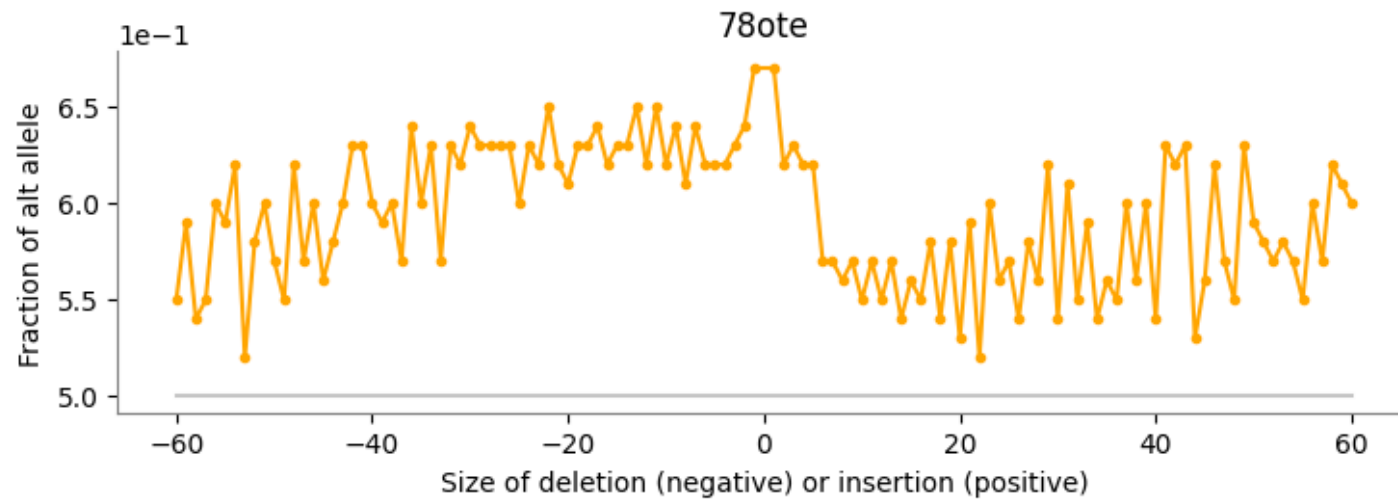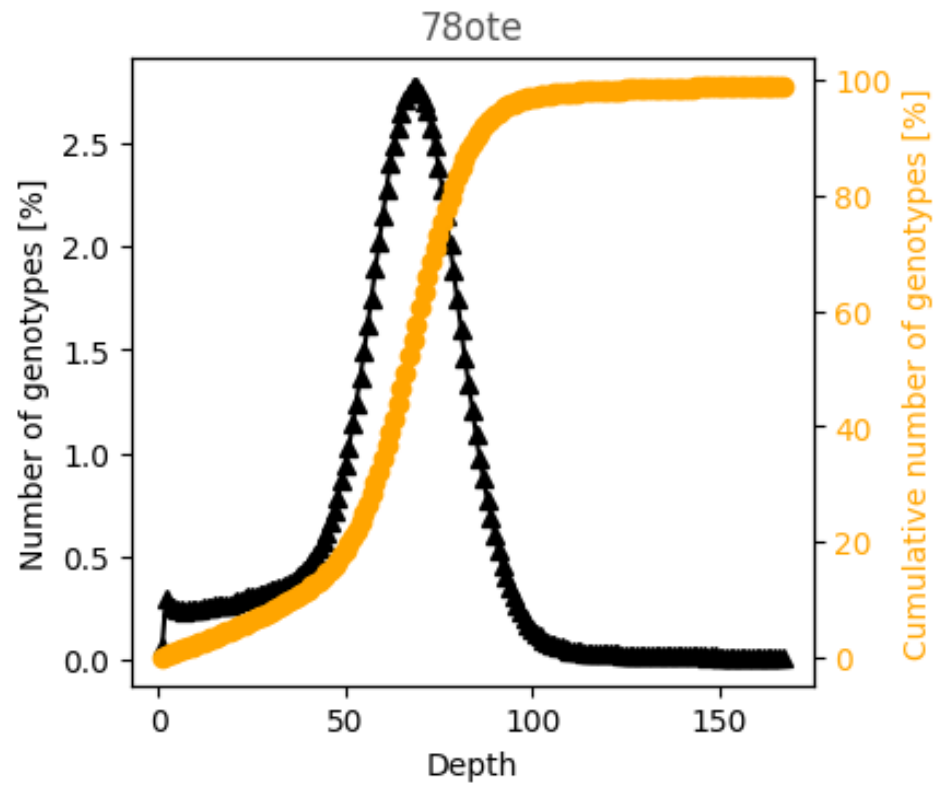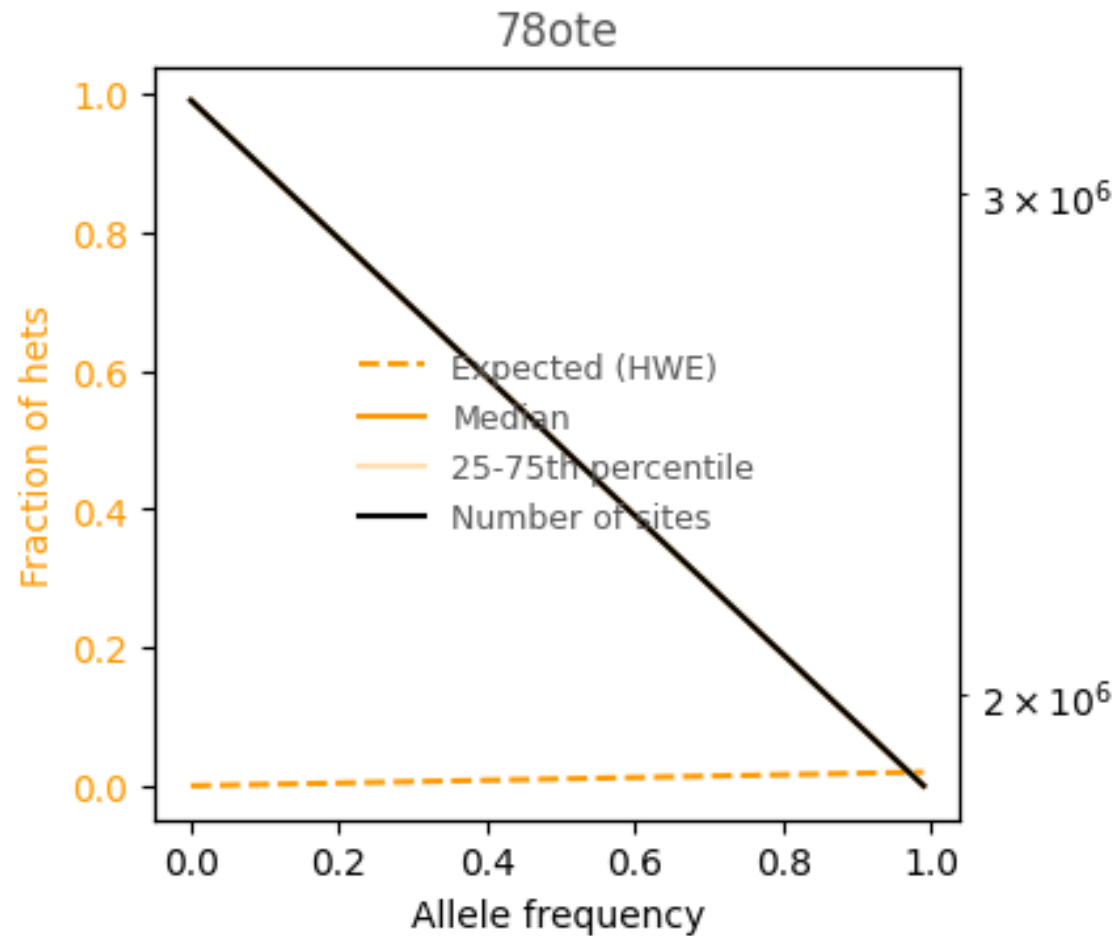# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types