Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78fof	4,131,898	1.92	1.93	959,318	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78fof	62.6%	1.88	67.1%	100,496	2,188

- 78fof .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78foftugm-103836010571-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00574-220209_A01176_BHCWJ5DSX3-EXT_LAB KA_NGCWGS-NGCWGS03802_22RKG001983x01_snv_germline_raw.haplotype_caller.vcf.gz`
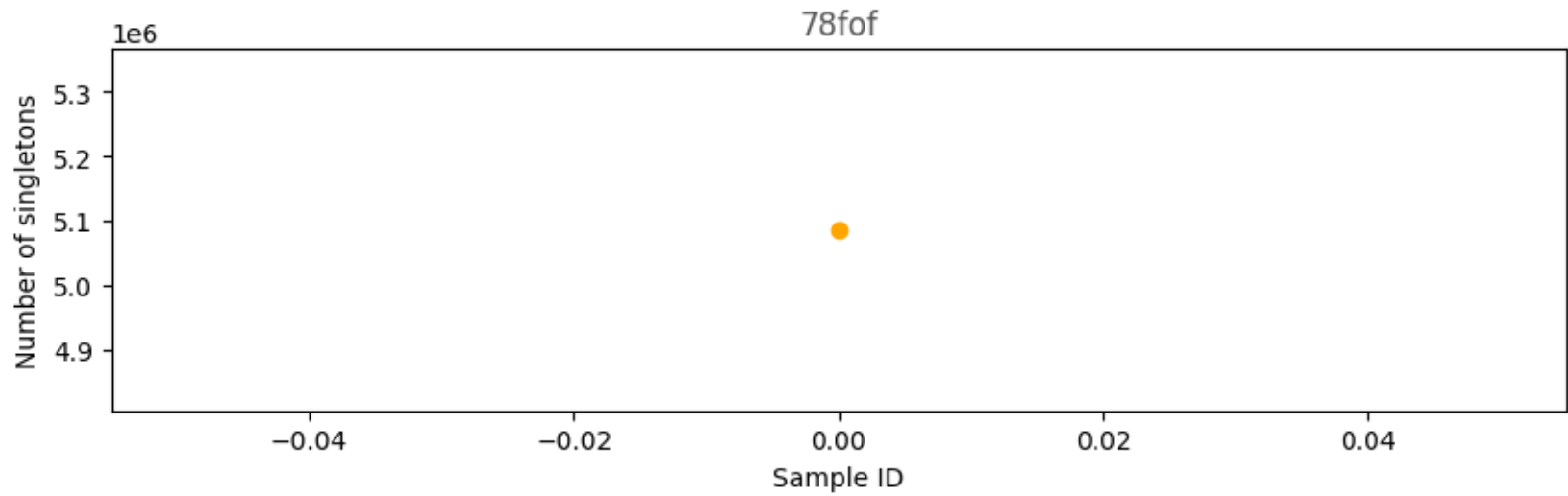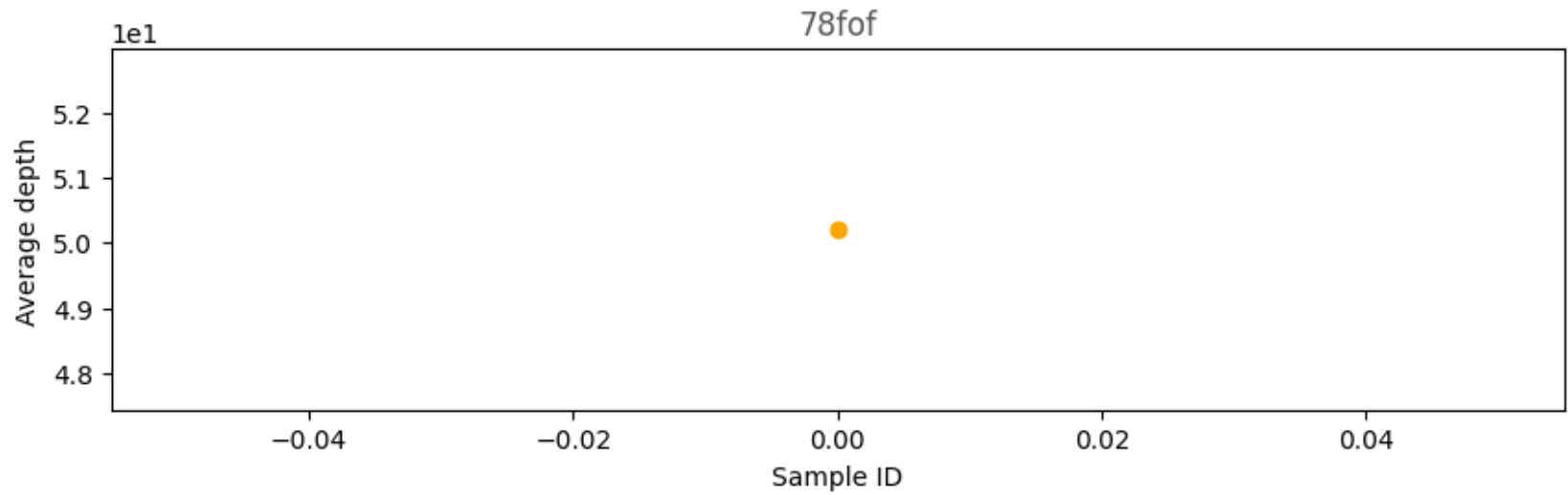
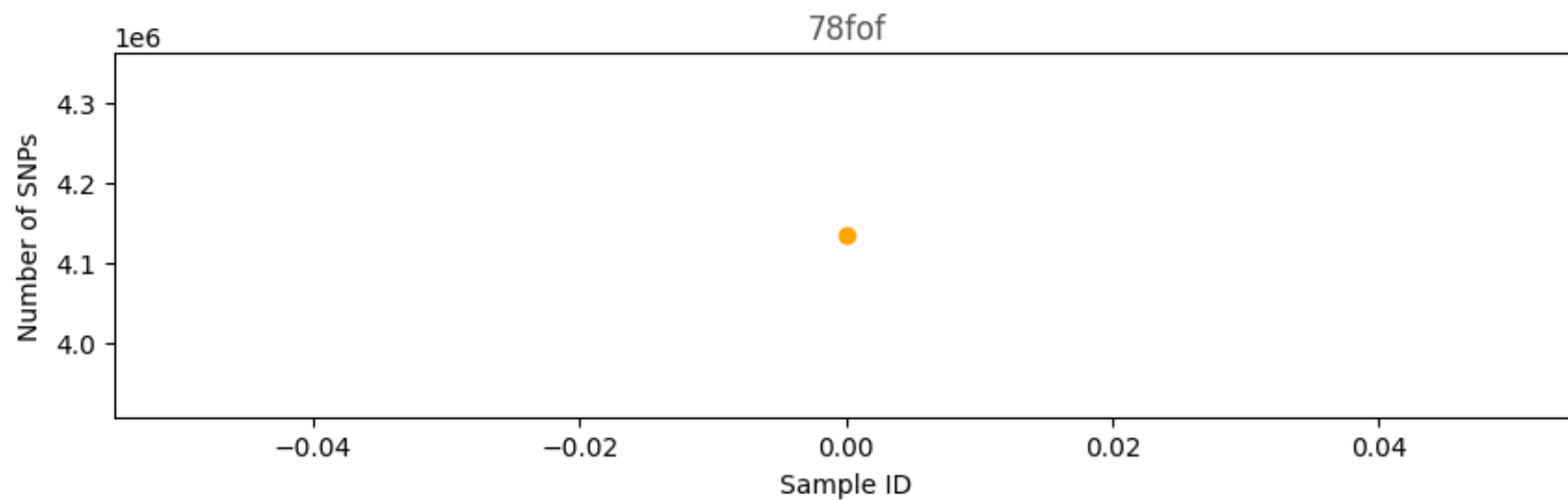# Ts/Tv by sample
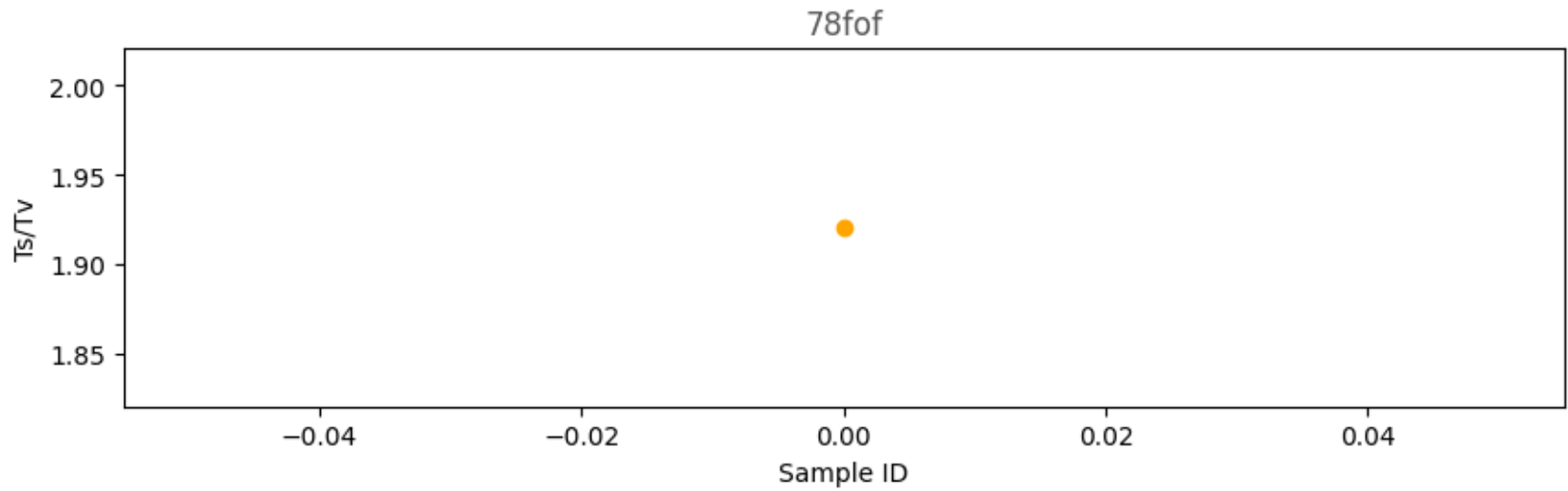
Hets vs non-ref Homs by sample
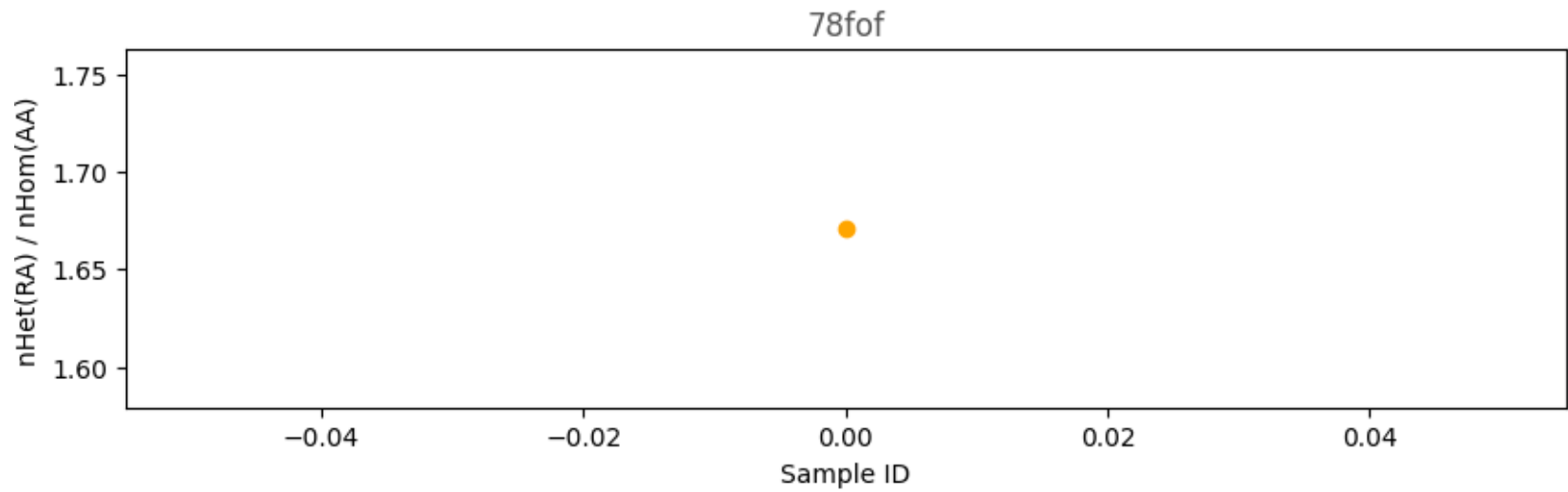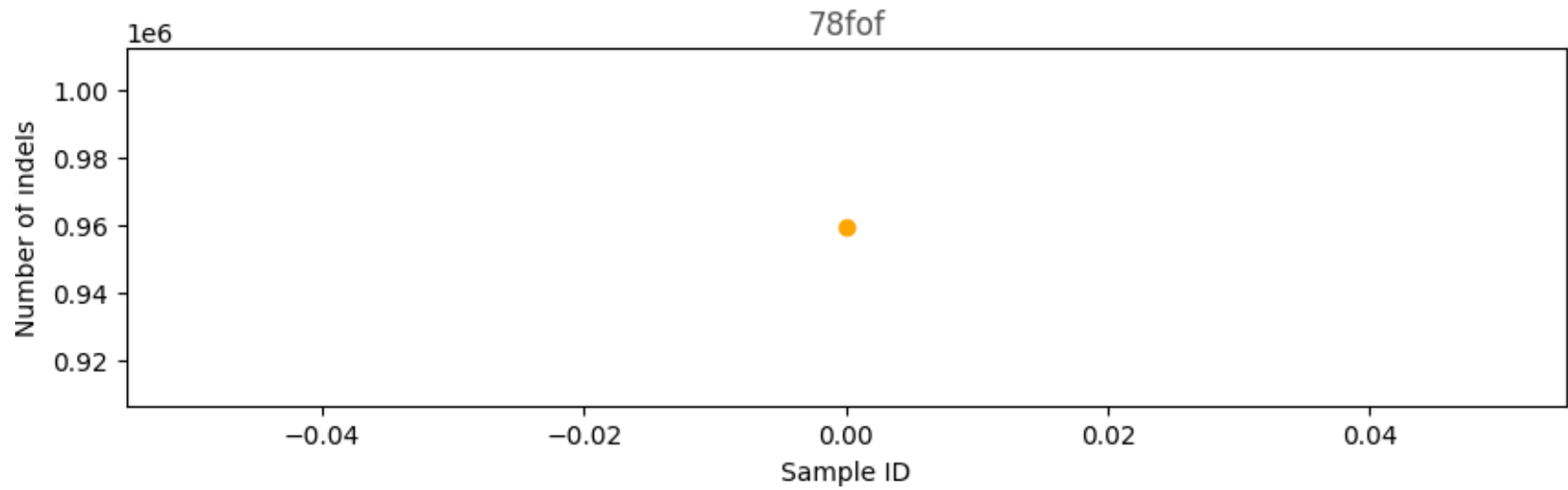
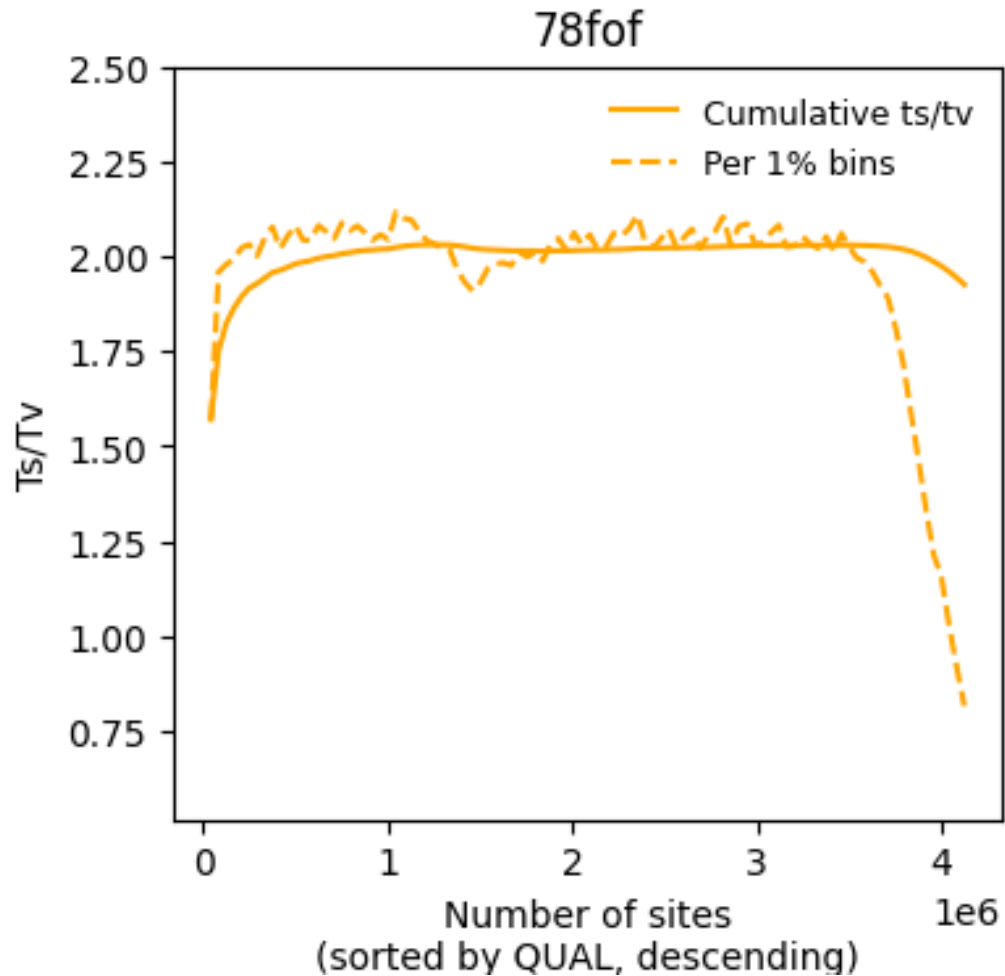# Singletons by sample (hets and homs)
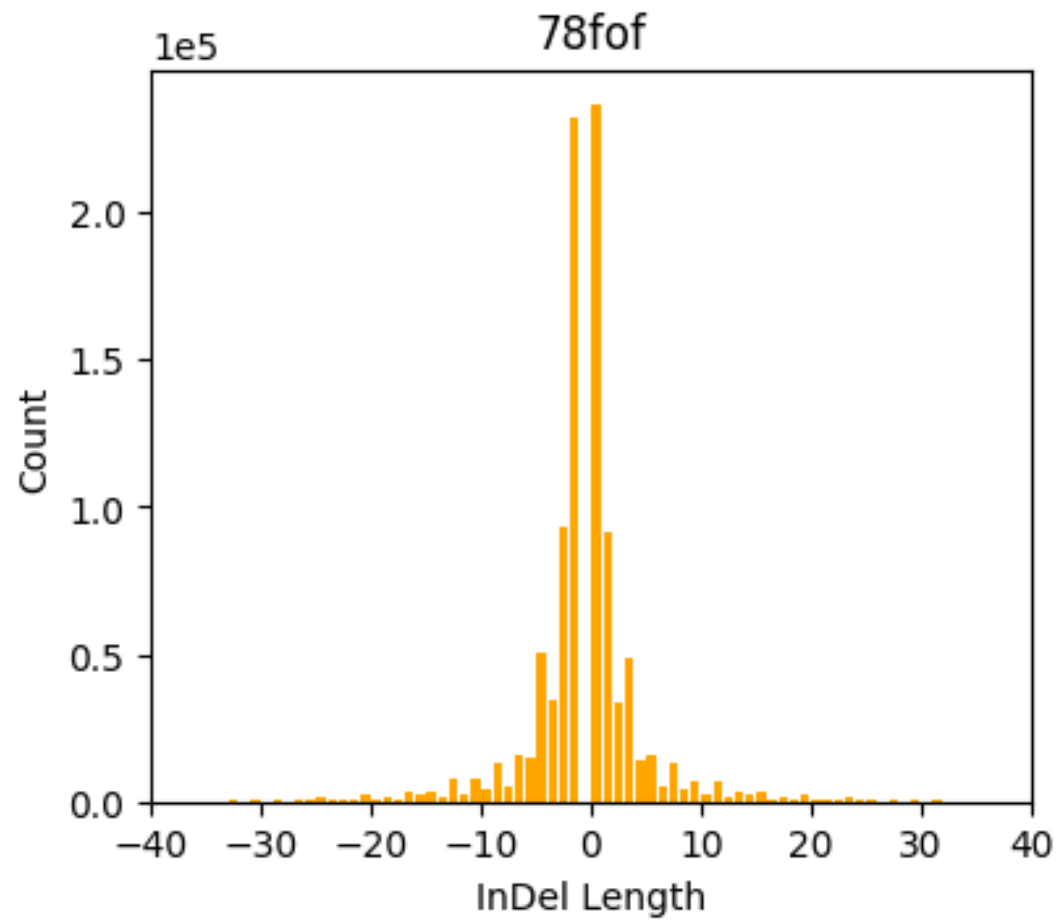
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

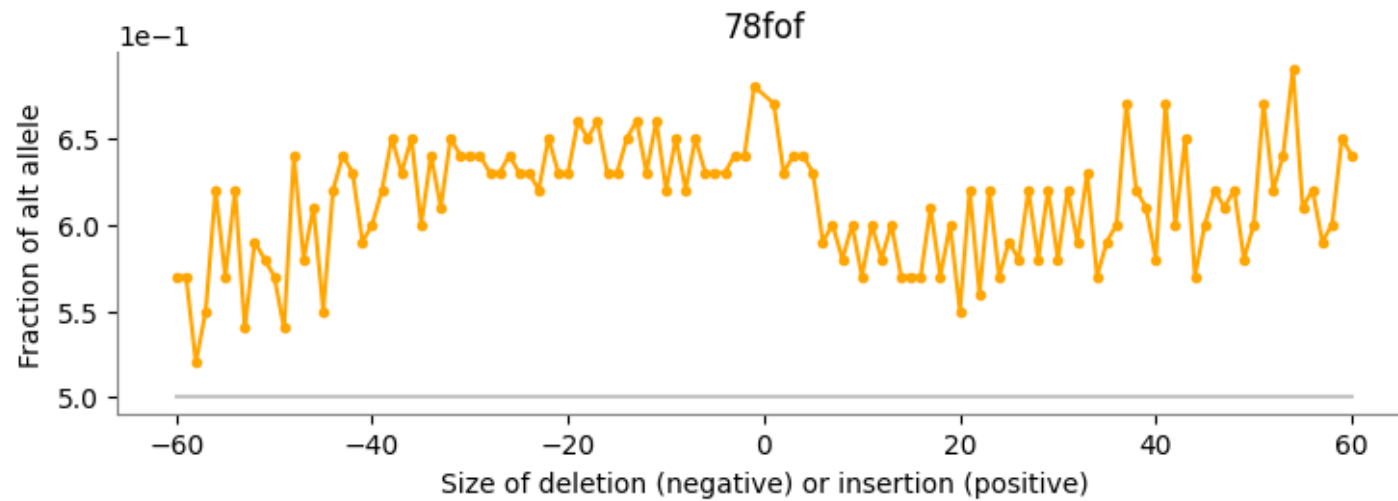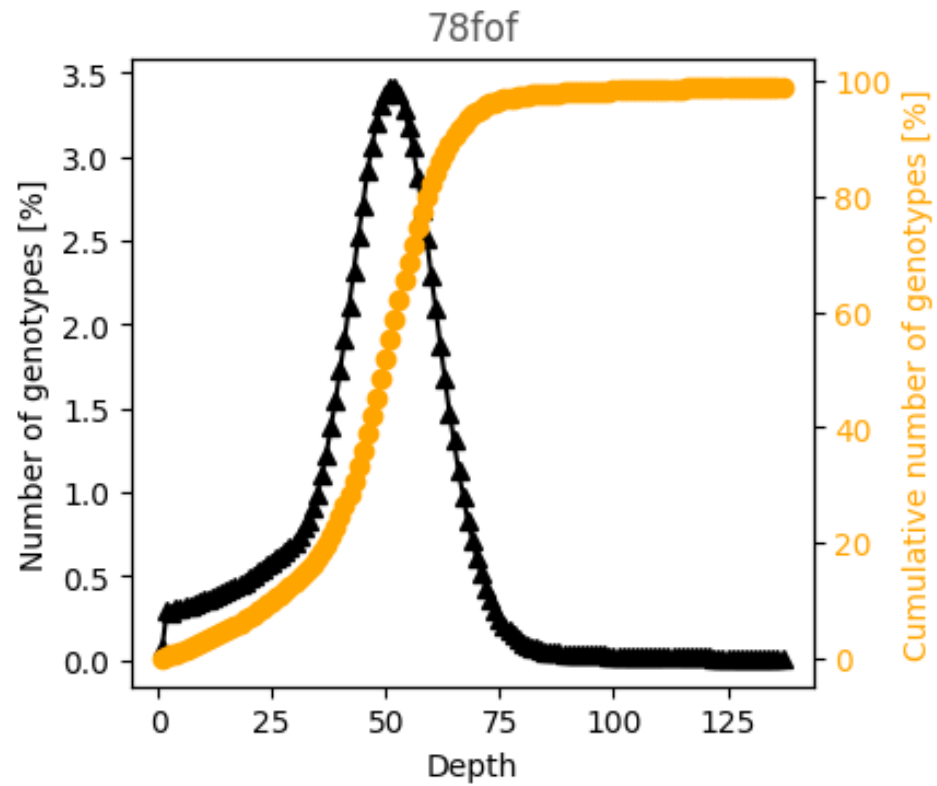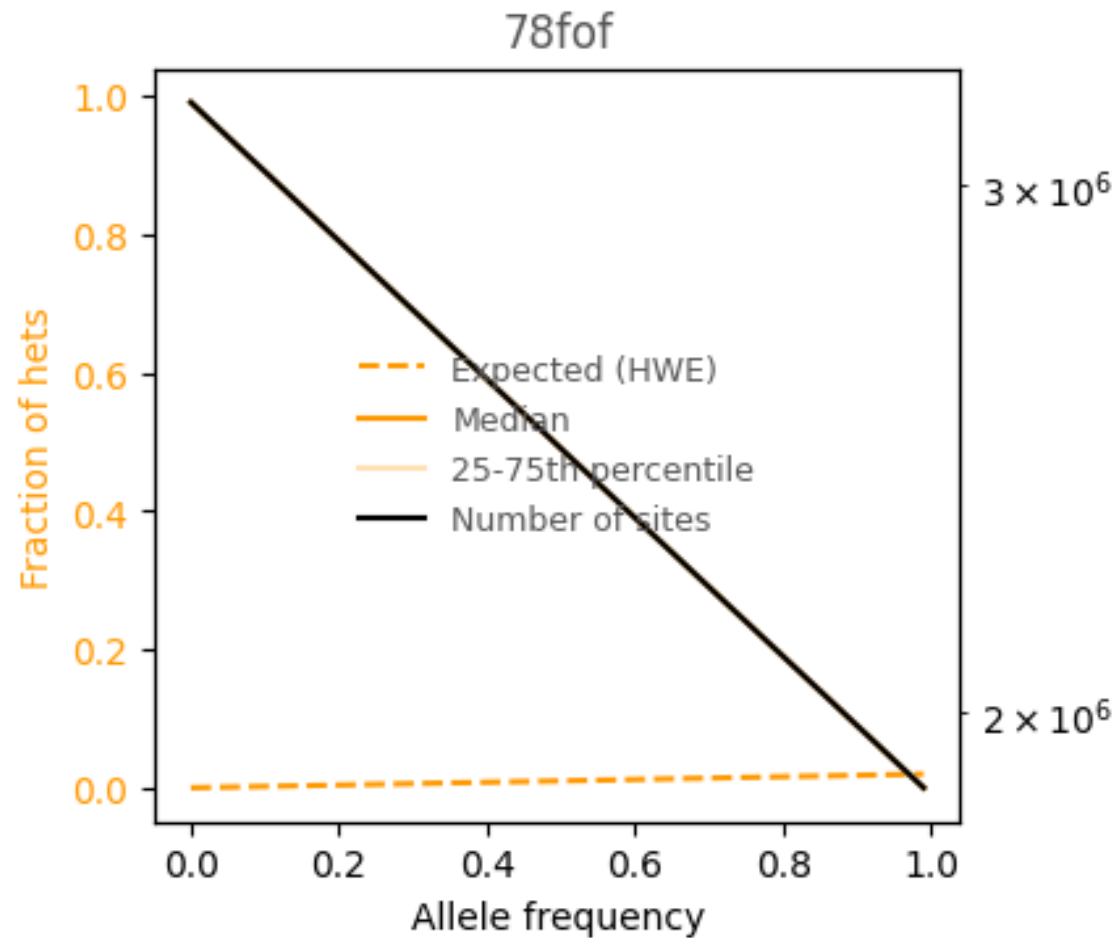# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types