Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78hon	4,083,139	1.92	1.93	958,326	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78hon	62.5%	1.89	67.5%	103,661	2,044

- 78hon .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78hondinm-103807550792-Normal_B`
    `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00412-211203_A01466_AHN25LDSX2-EXT_LAB`
    `KA_NGCWGS-NGCWGS03434_21RKG024662x01_snv_germline_raw.haplotype_caller.vcf.gz`
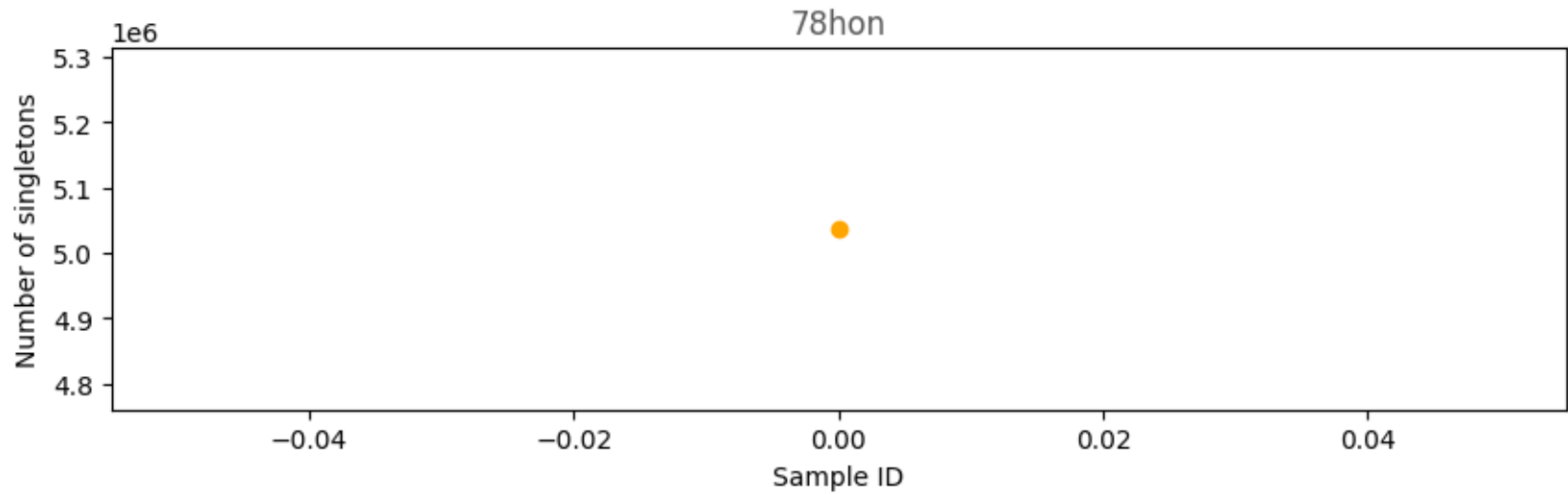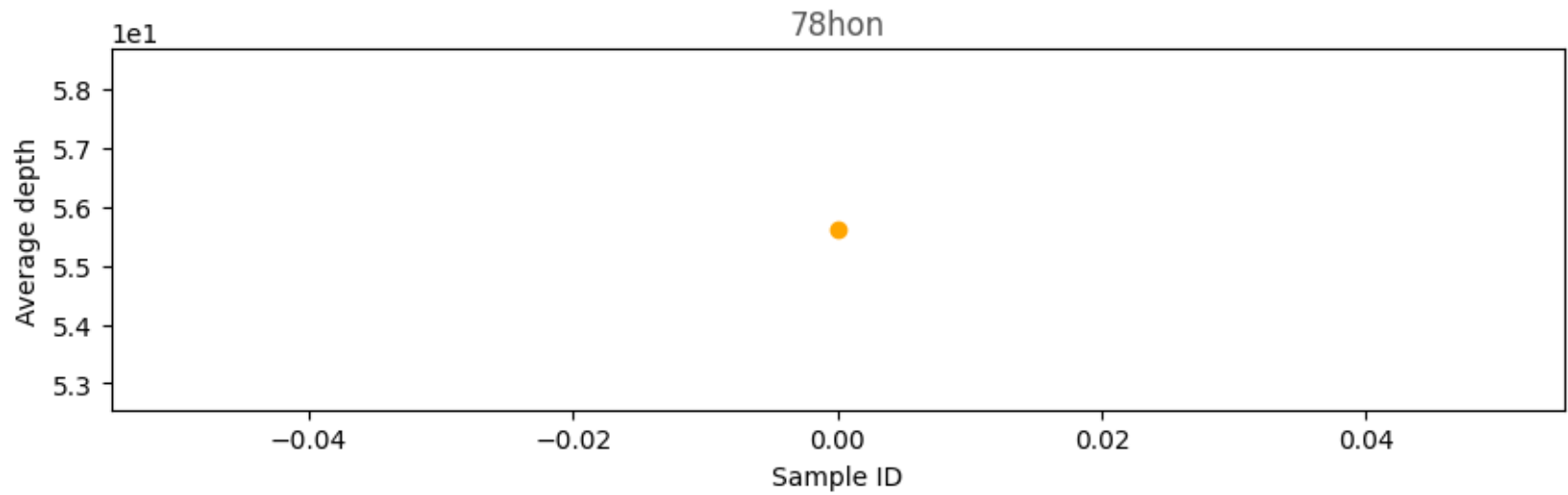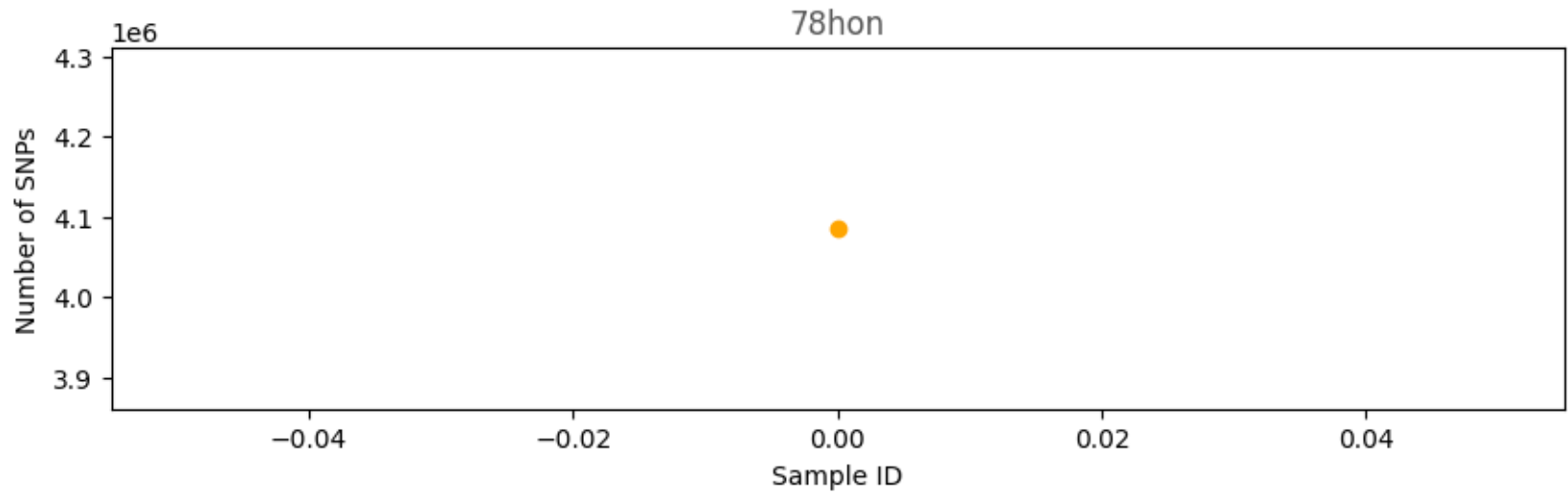
Ts/Tv by sample
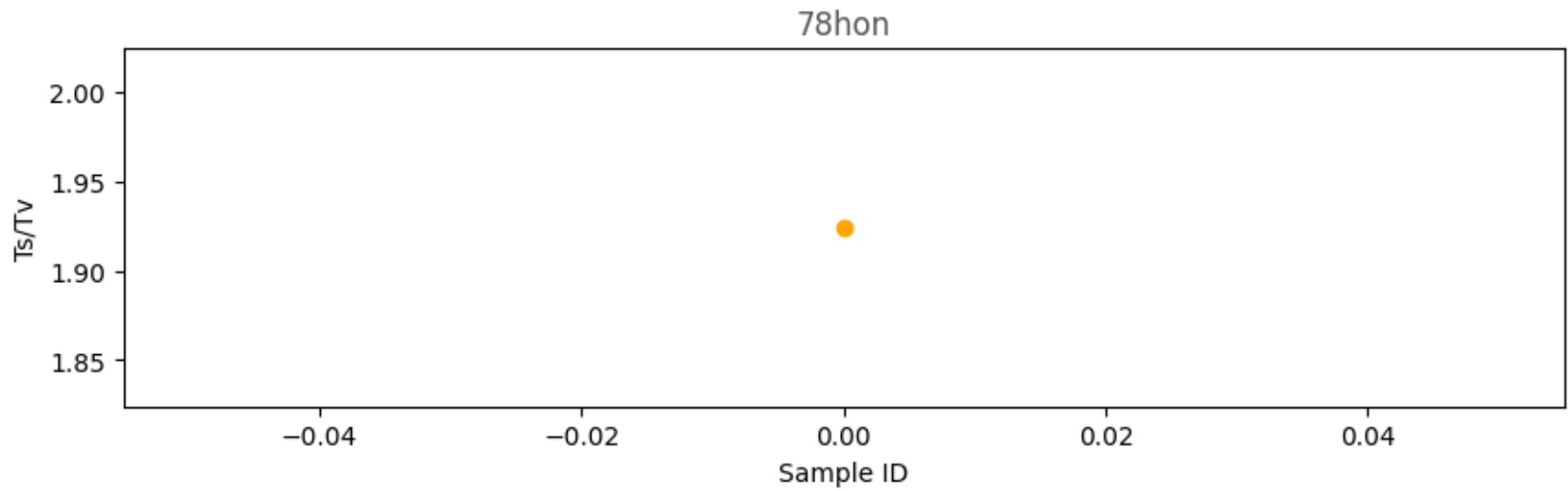
Hets vs non-ref Homs by sample
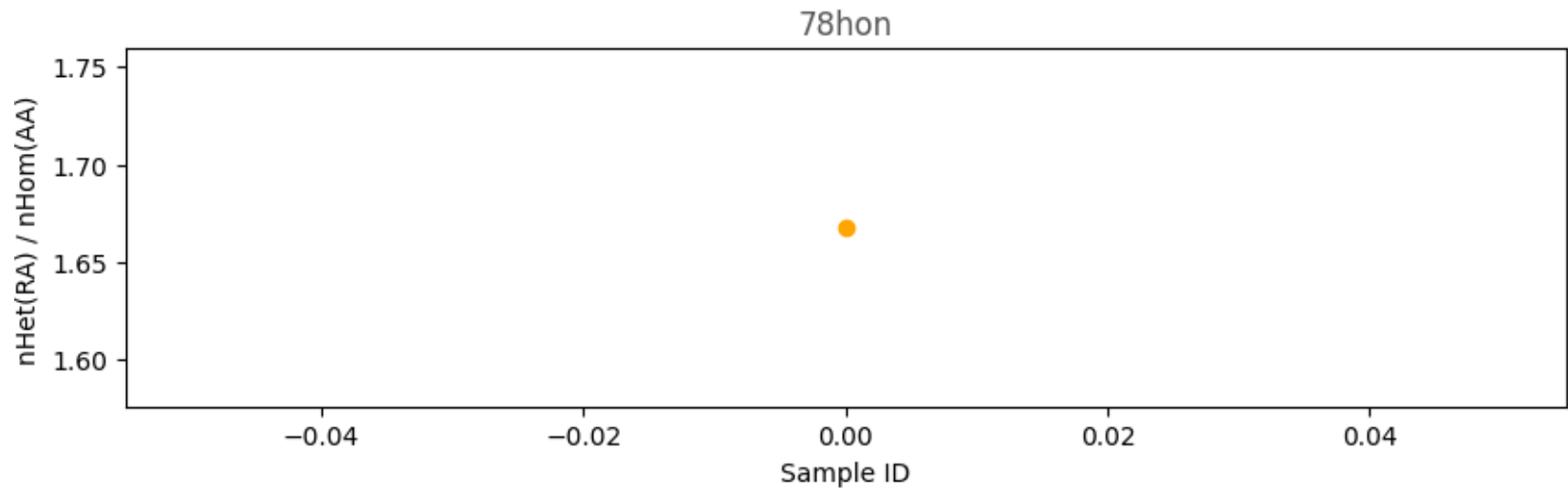
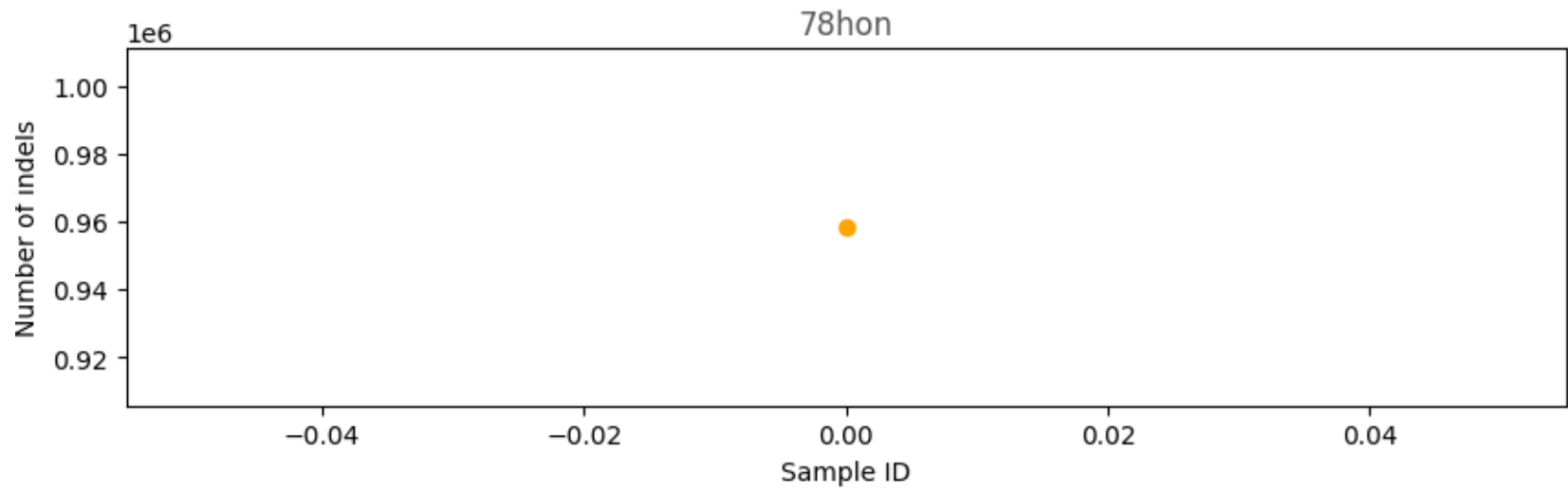# Singletons by sample (hets and homs)
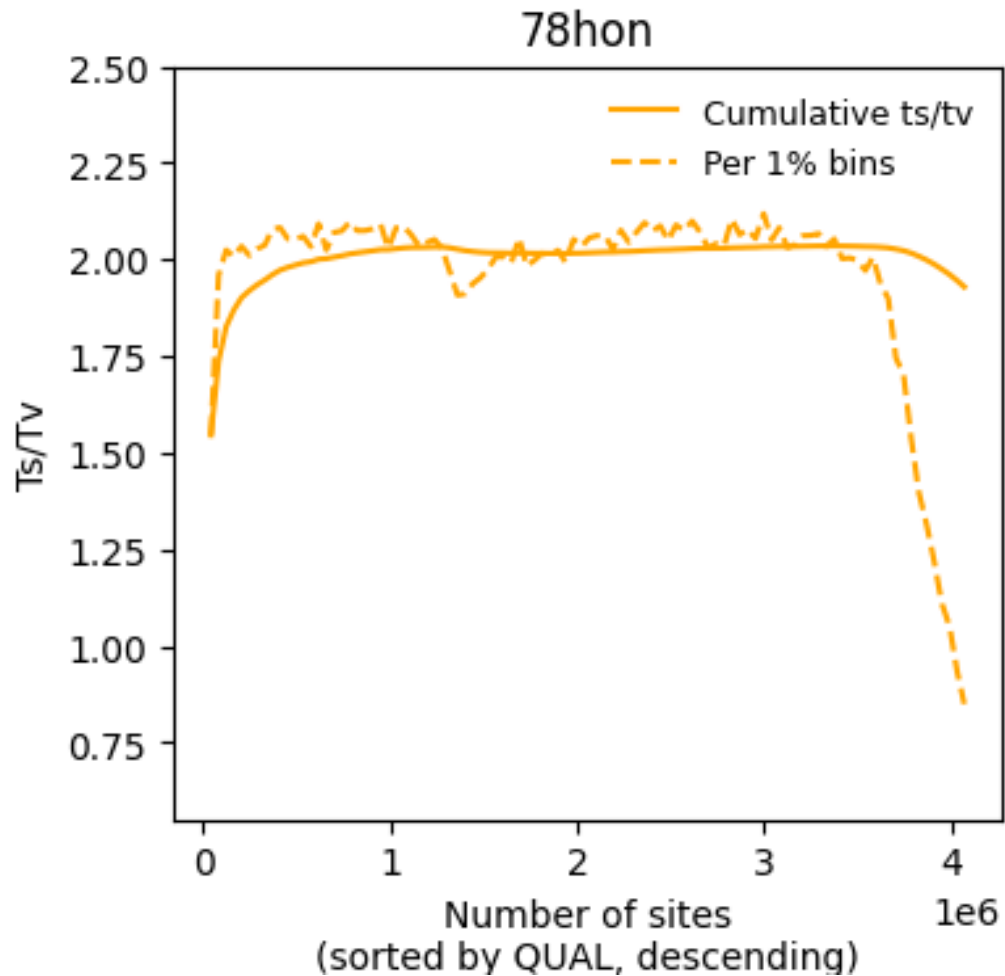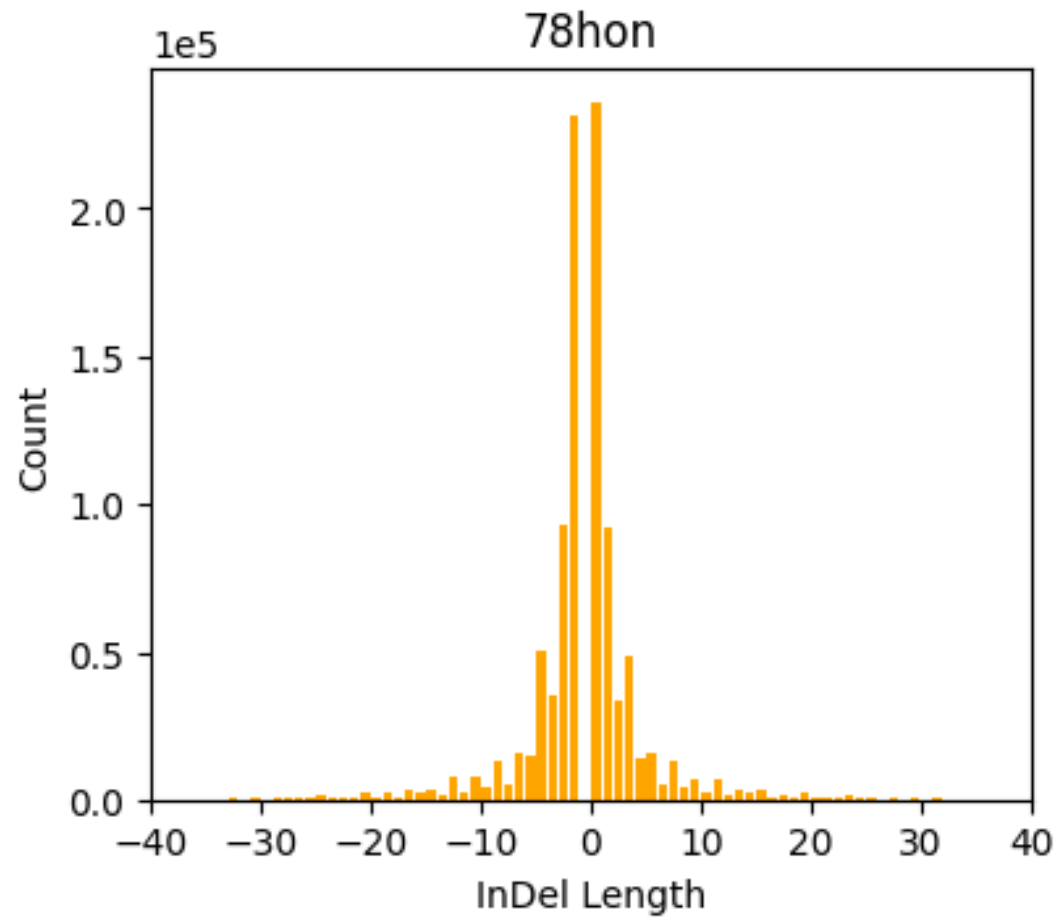
Average depth by sample
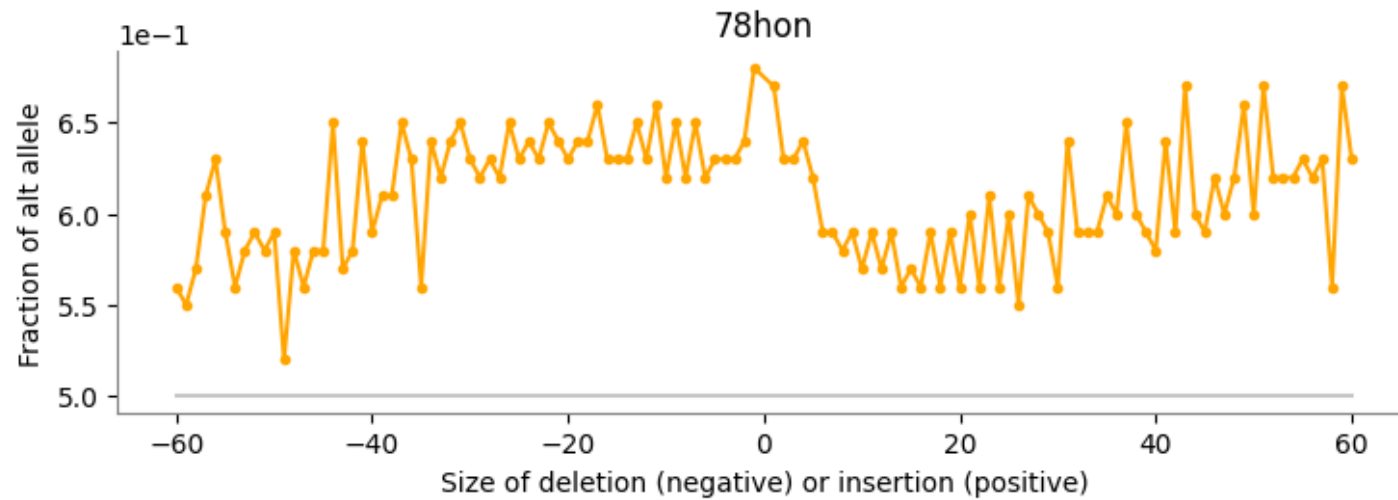
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

78hon

Indel distribution

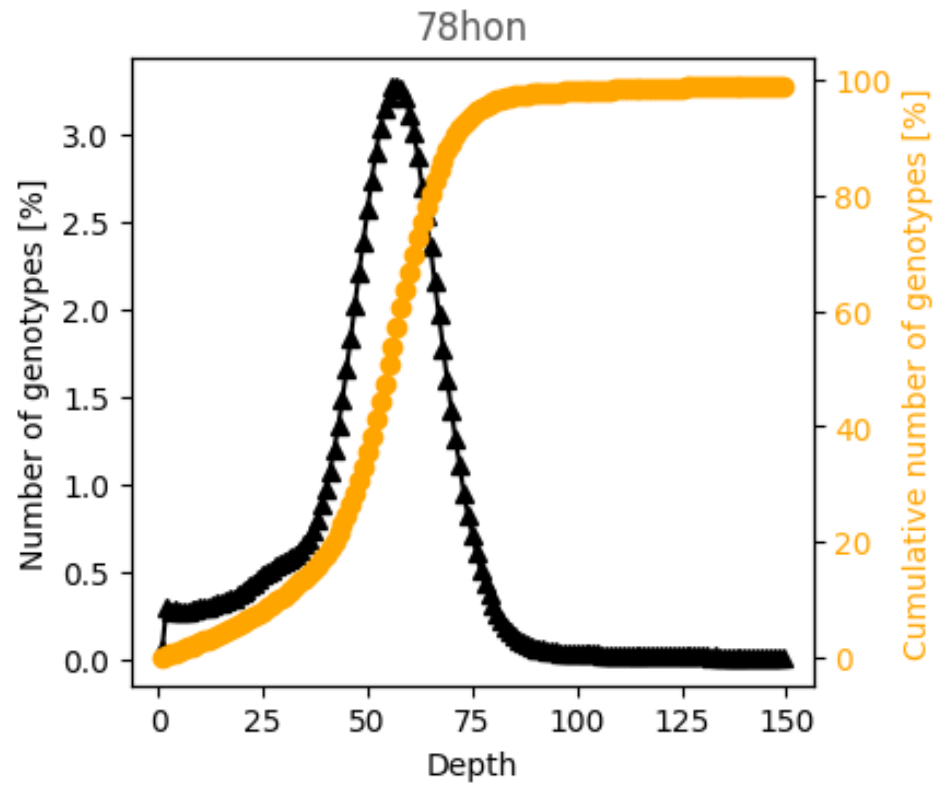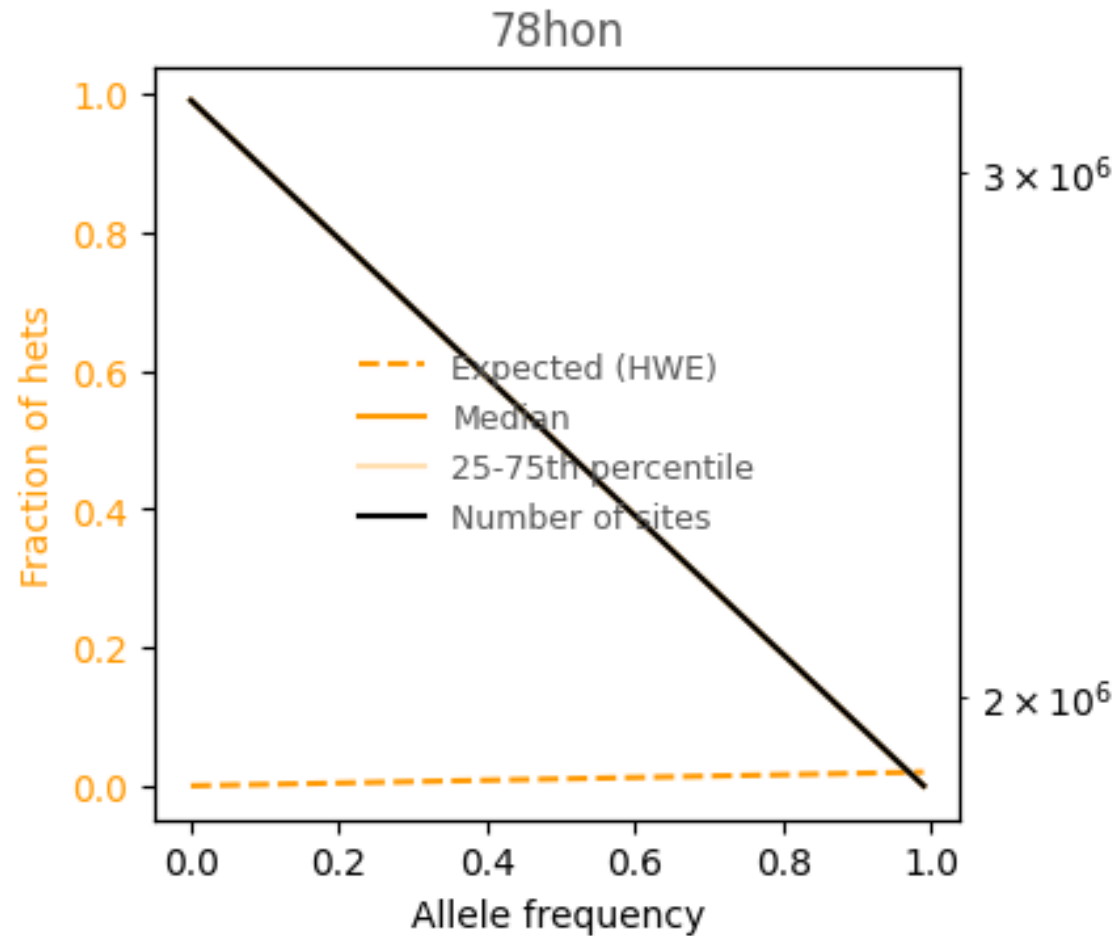# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types