Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
79ola	4,088,382	1.92	1.93	958,970	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
79ola	62.1%	1.89	67.3%	103,229	2,051

- 79ola .. `/ngc/projects2/gm/data/archive/2022/variants/snv/79olarosm-110296262246-Normal_B`
`lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02035-221019_A01176_BH3FN7DSX5-EXT_LAB`
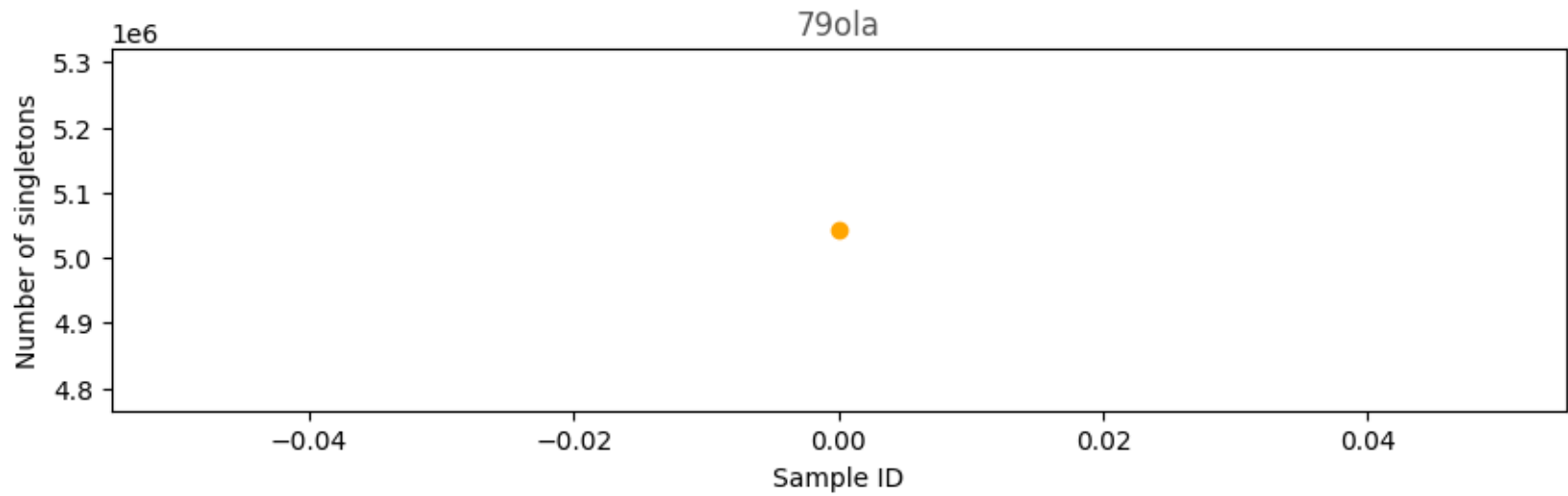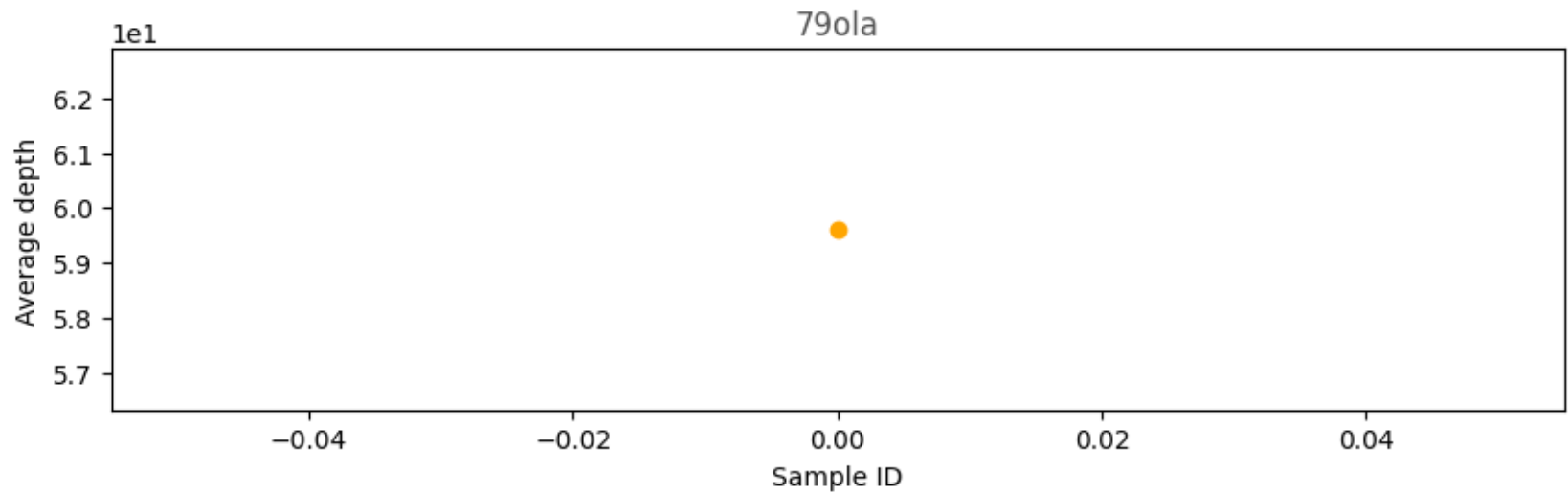`KA_NGCWGS-NGCWGS05909_snv_germline_raw.haplotype_caller.vcf.gz`

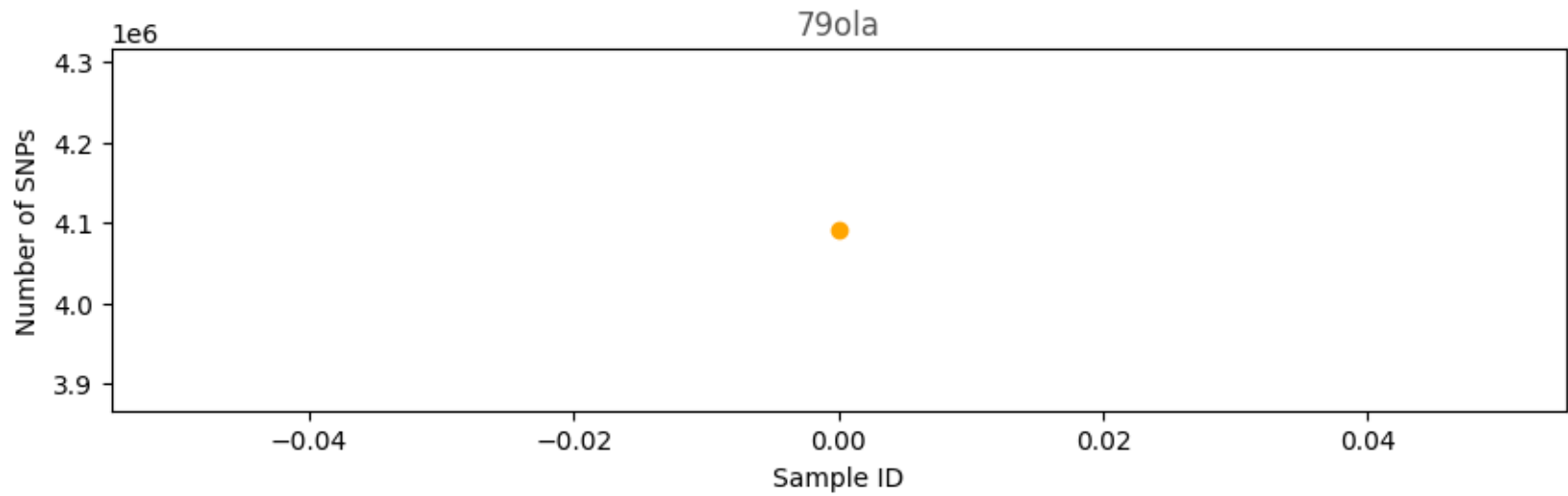# Ts/Tv by sample

79ola
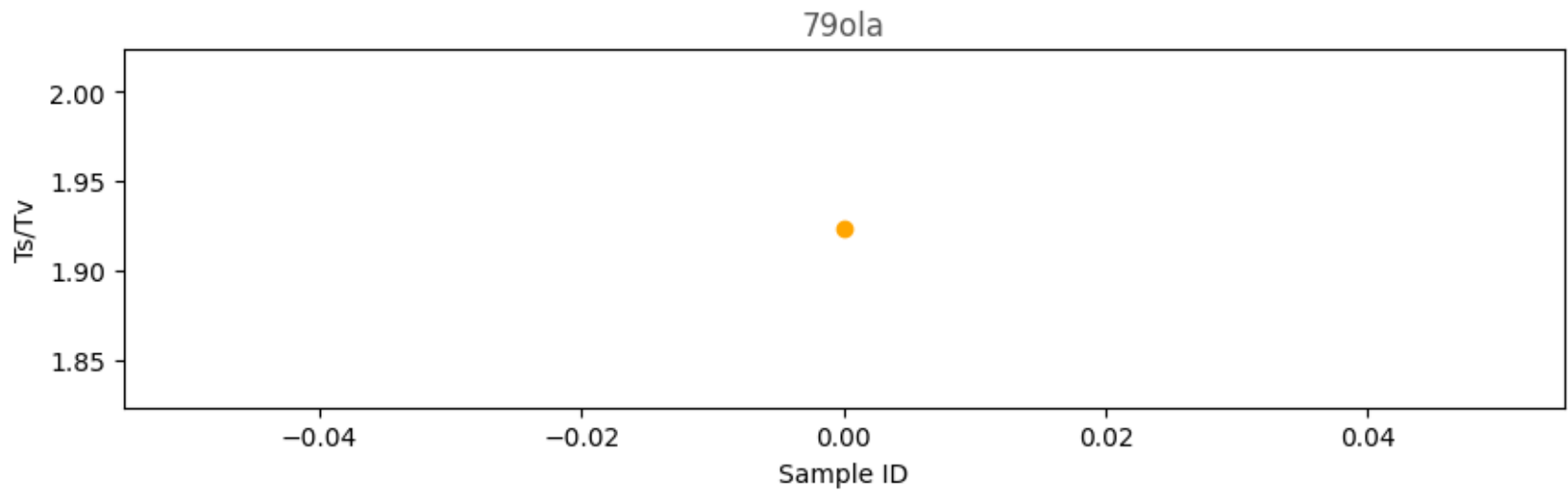
Hets vs non-ref Homs by sample
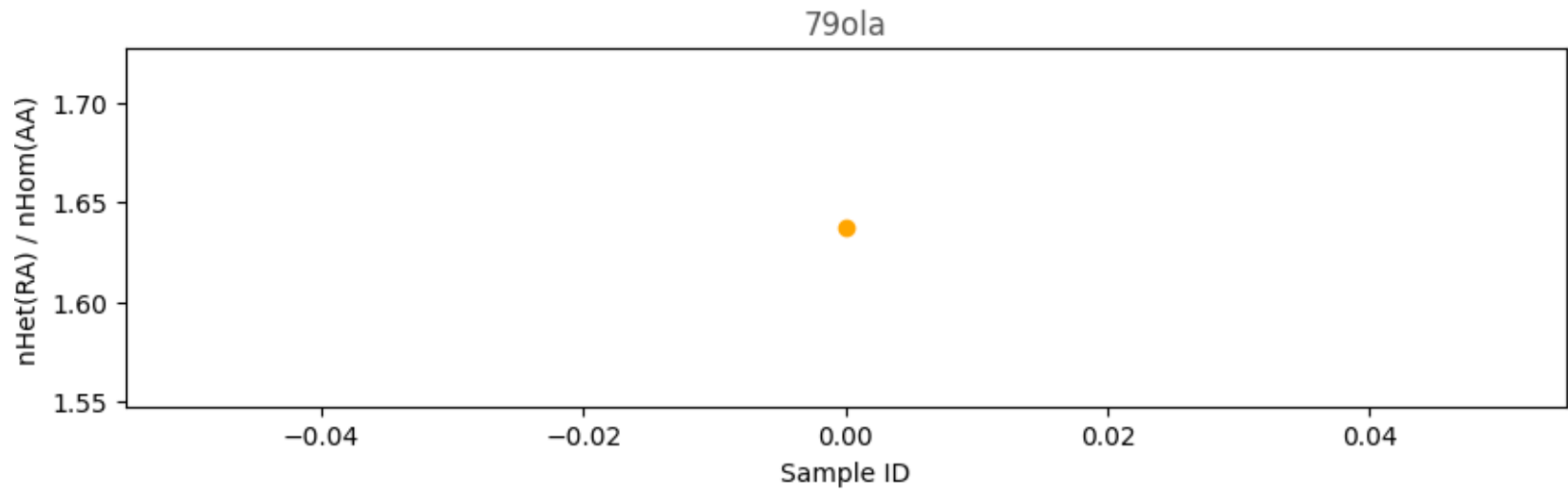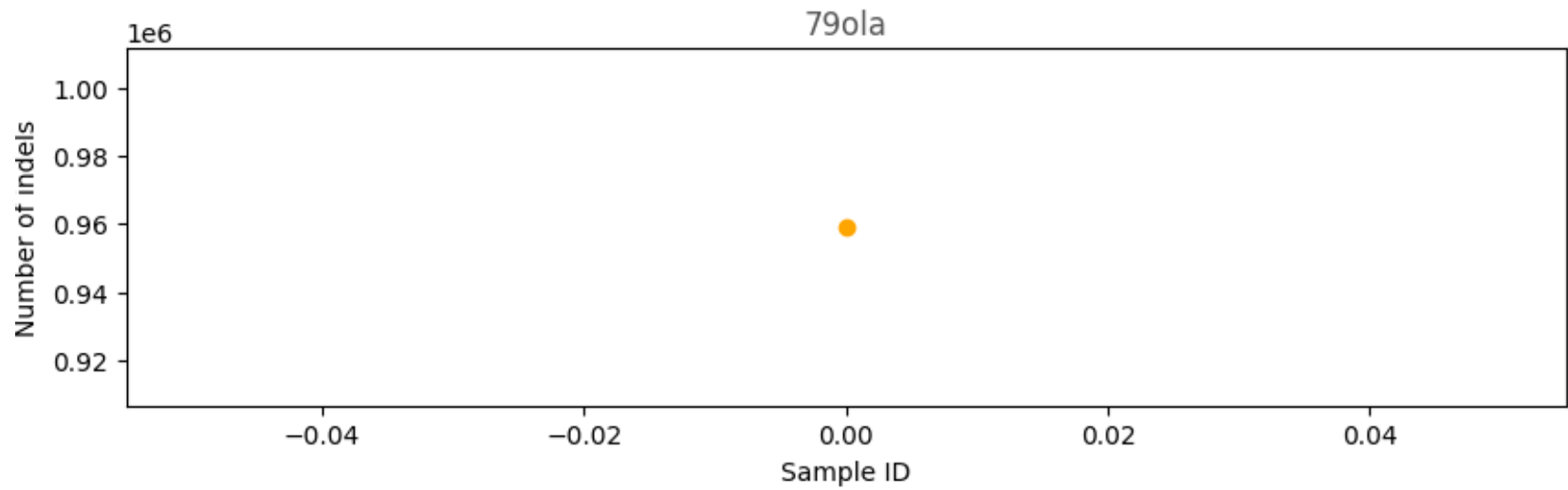
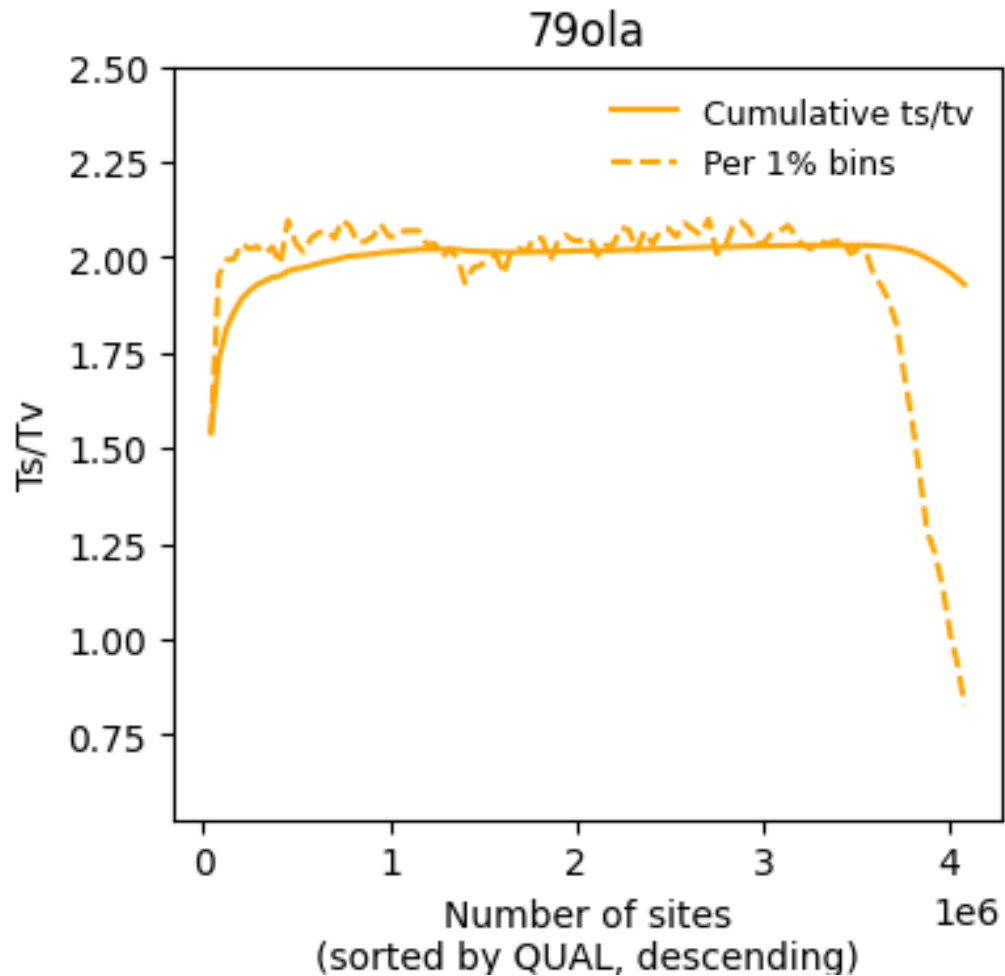# Singletons by sample (hets and homs)
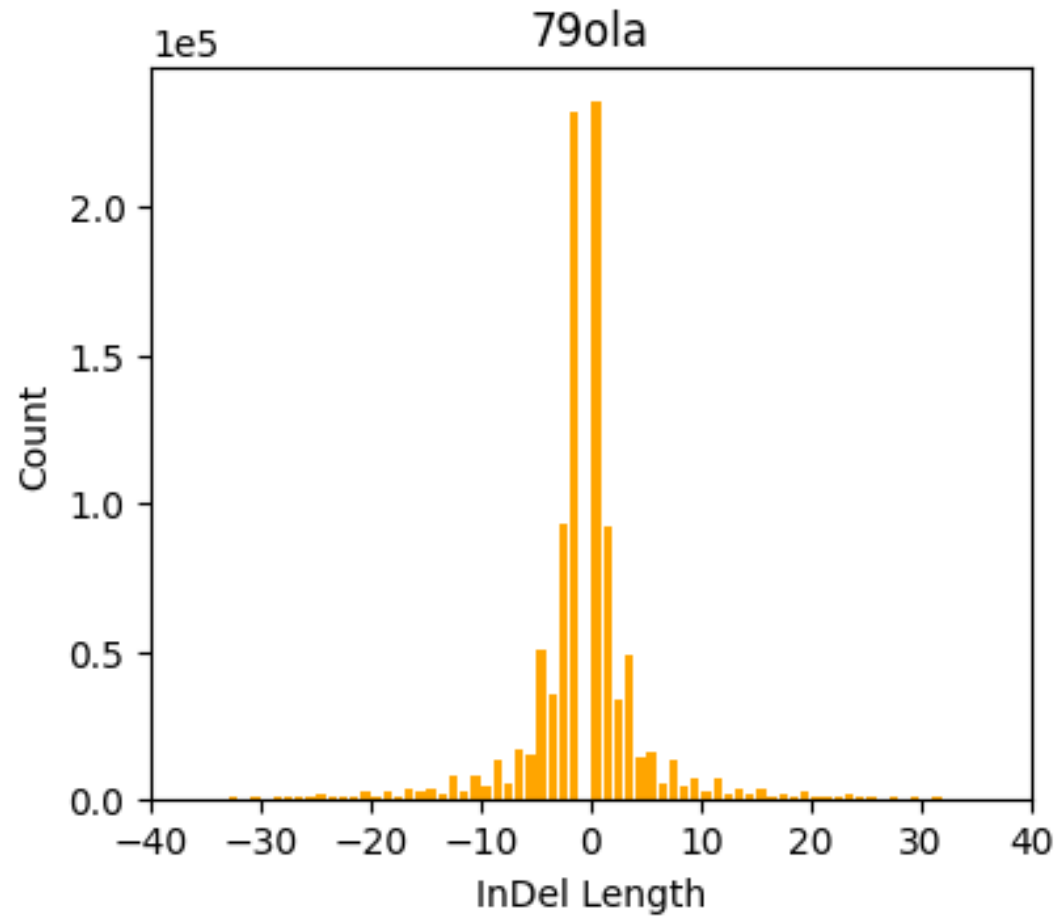
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

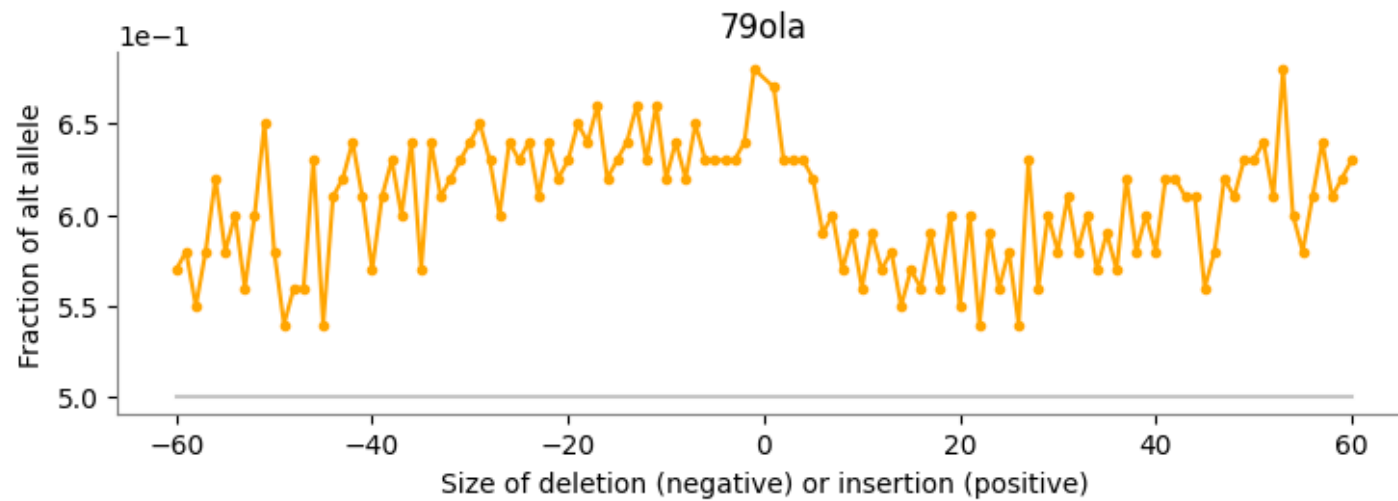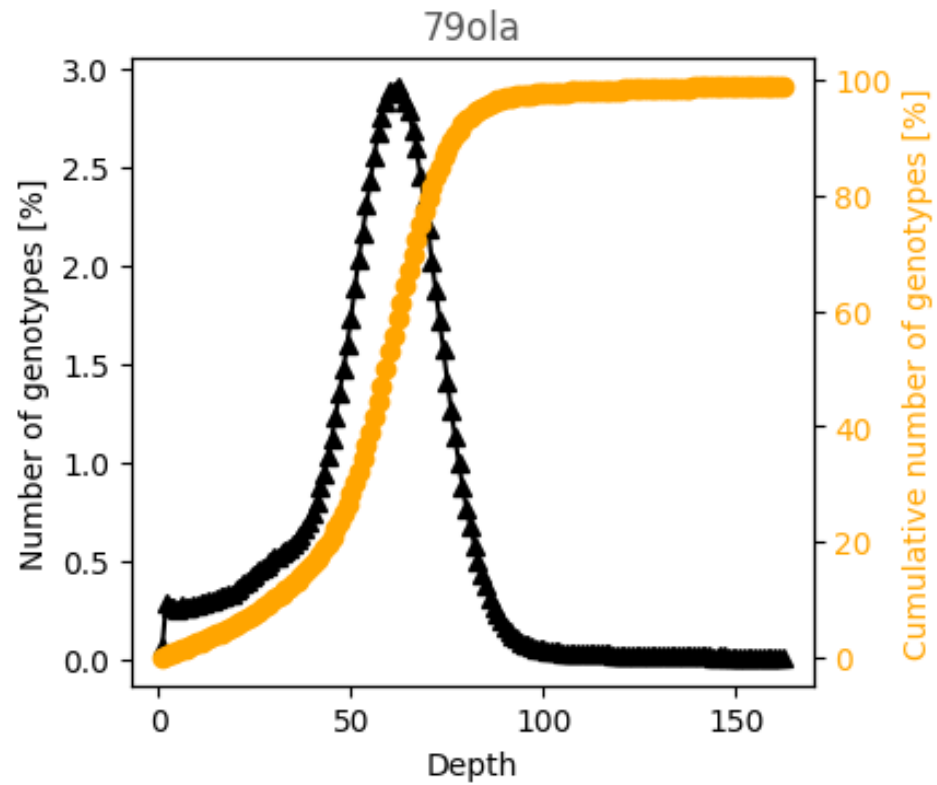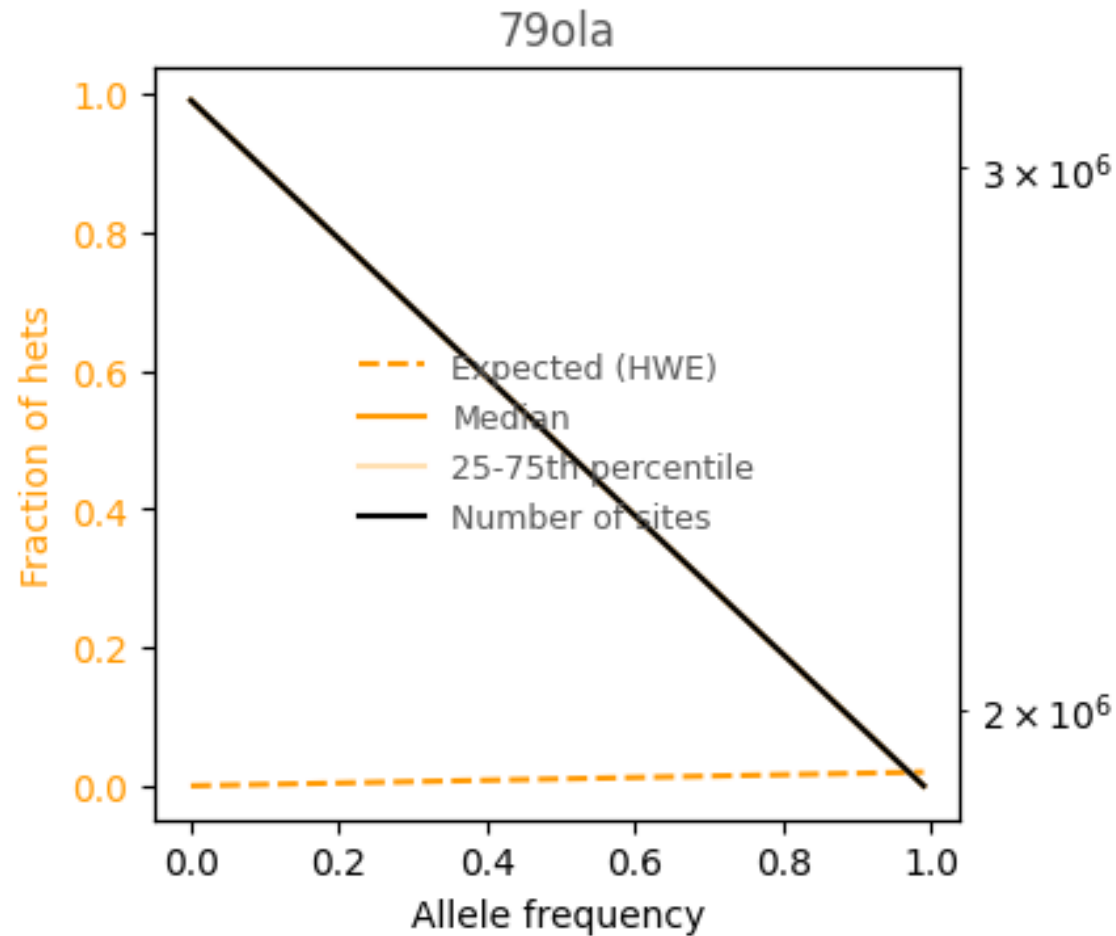# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types