Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
48oma	4,133,795	1.92	1.92	967,656	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
48oma	62.9%	1.89	68.6%	109,097	2,387

- 48oma .. `/ngc/projects2/gm/data/archive/2022/variants/snv/48omathhf-103848682260-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220324_A01176_AHCFCNDSX3-RHGM_LABKA_WGS- WGS04013_snv_germline_raw.haplotype_caller.vcf.gz`
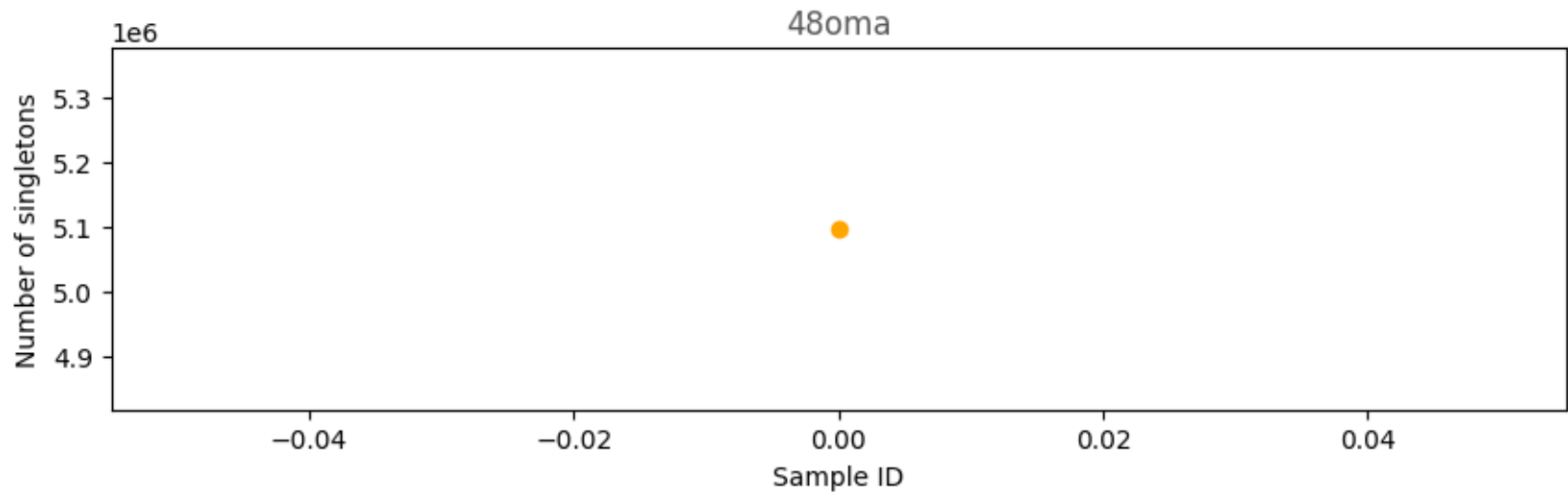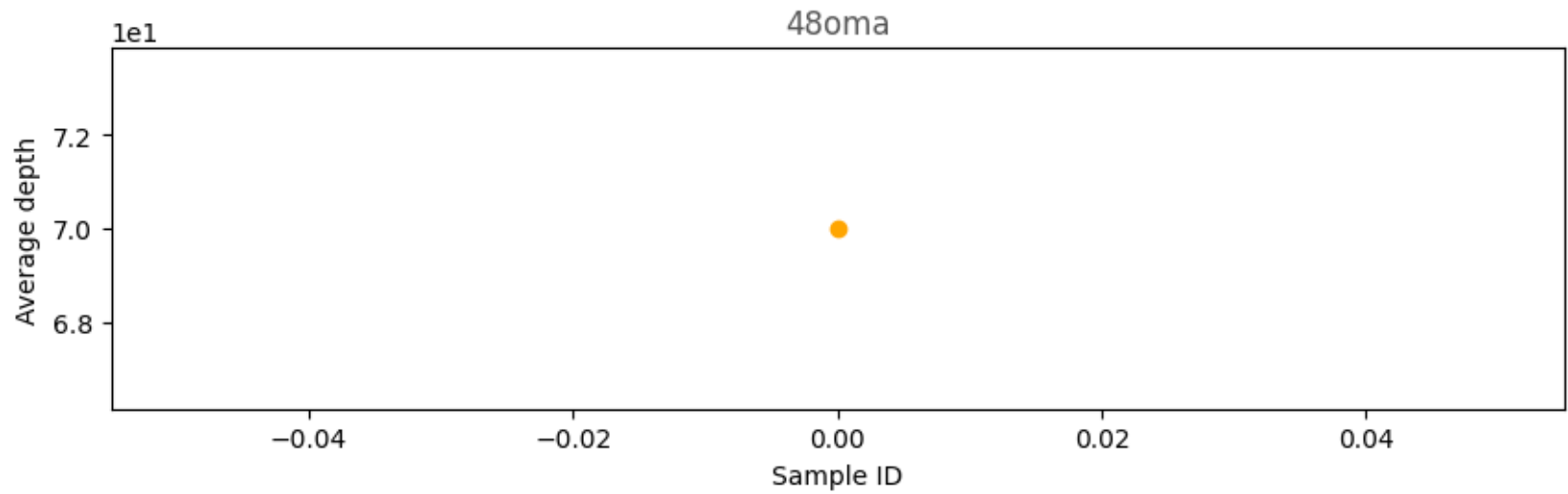
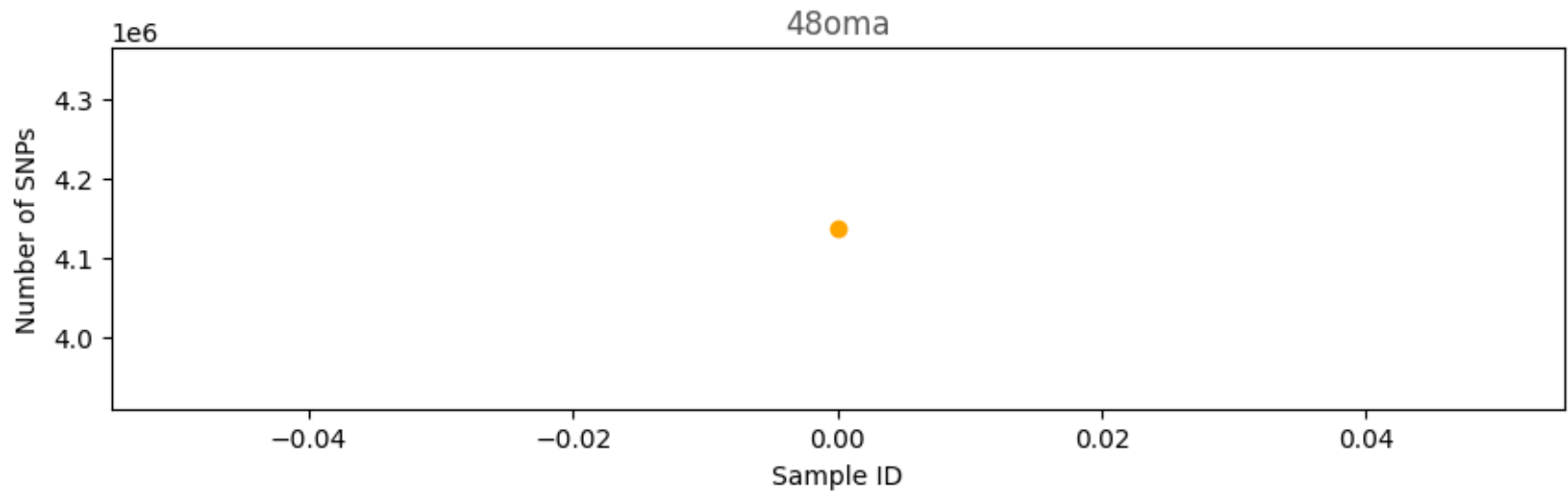# Ts/Tv by sample
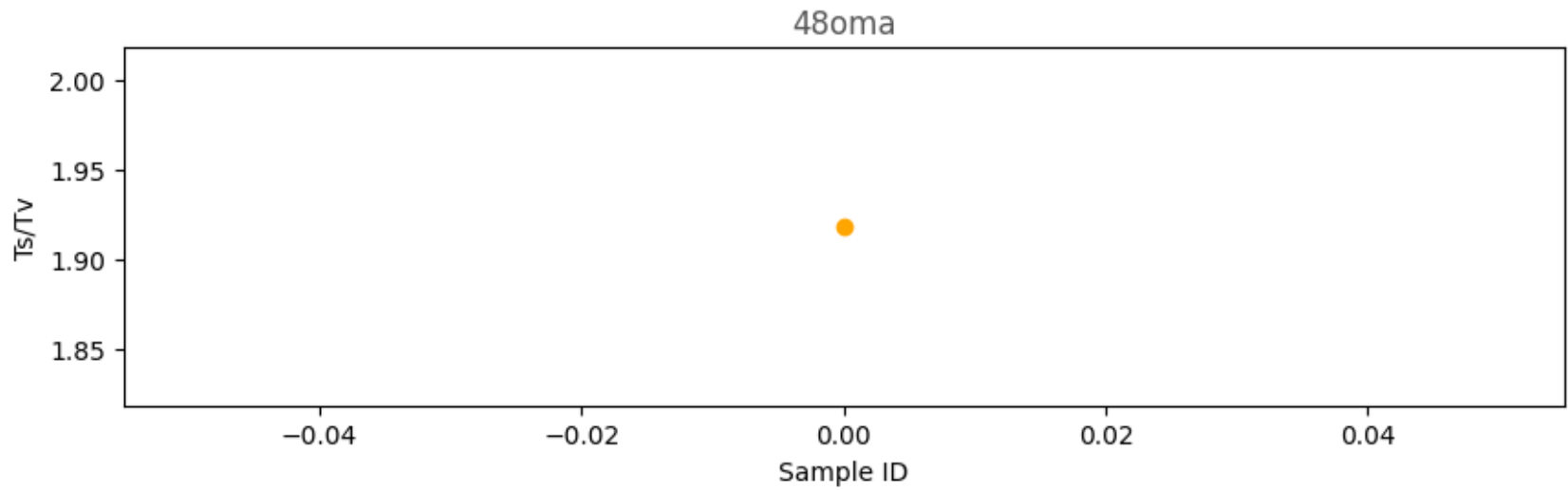
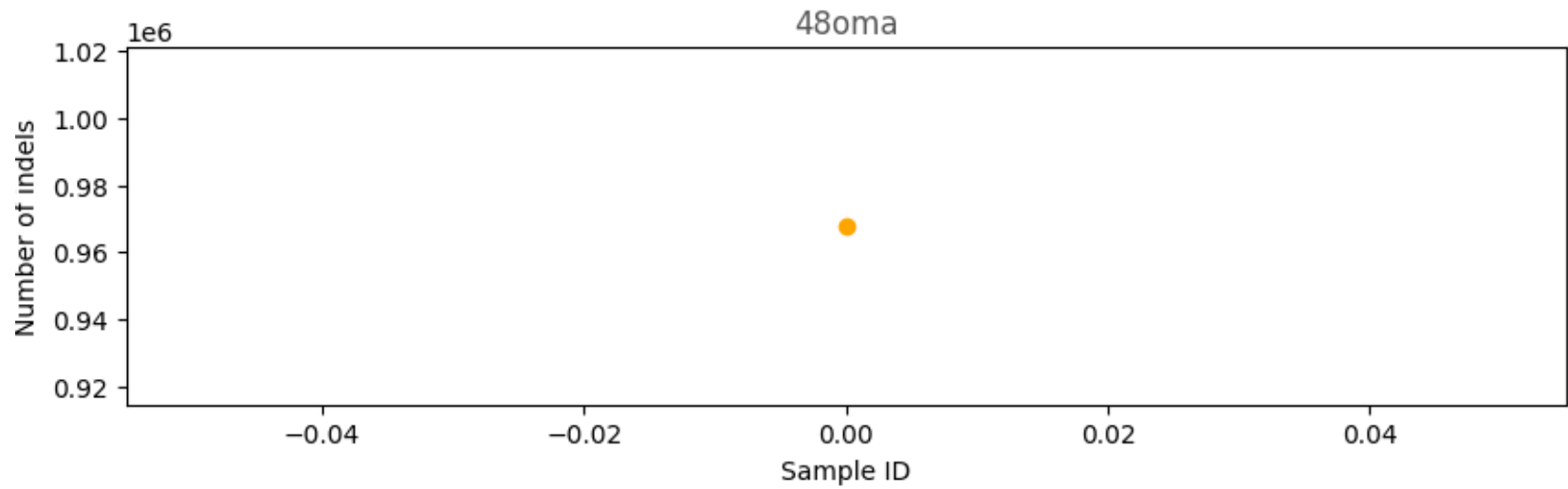# Hets vs non-ref Homs by sample
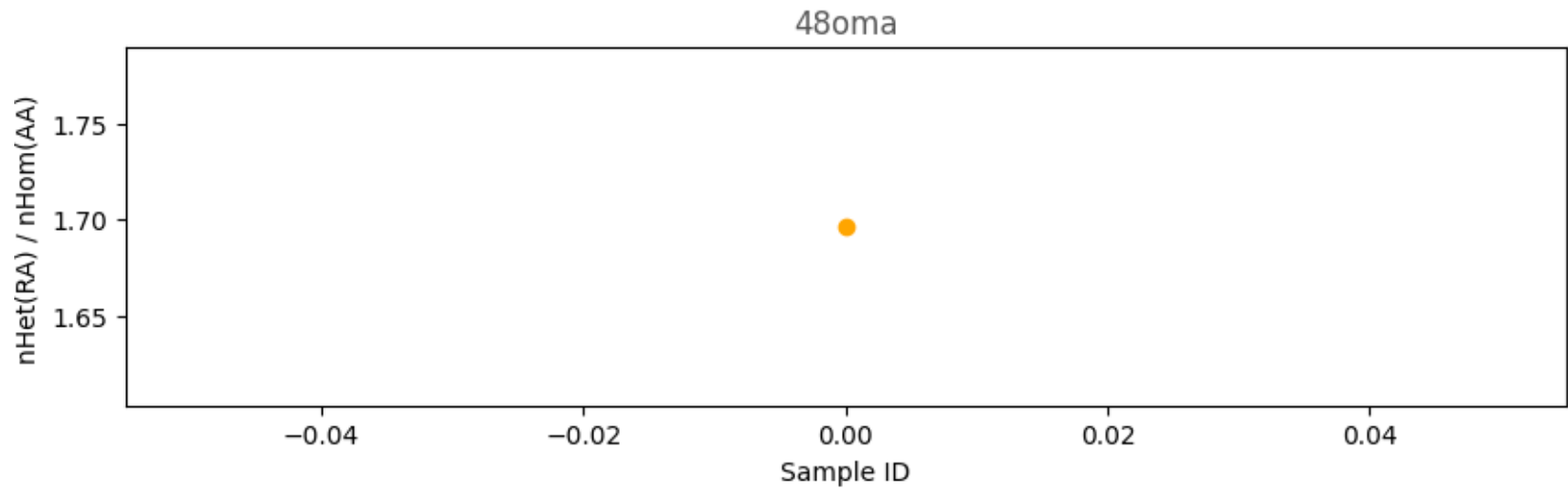
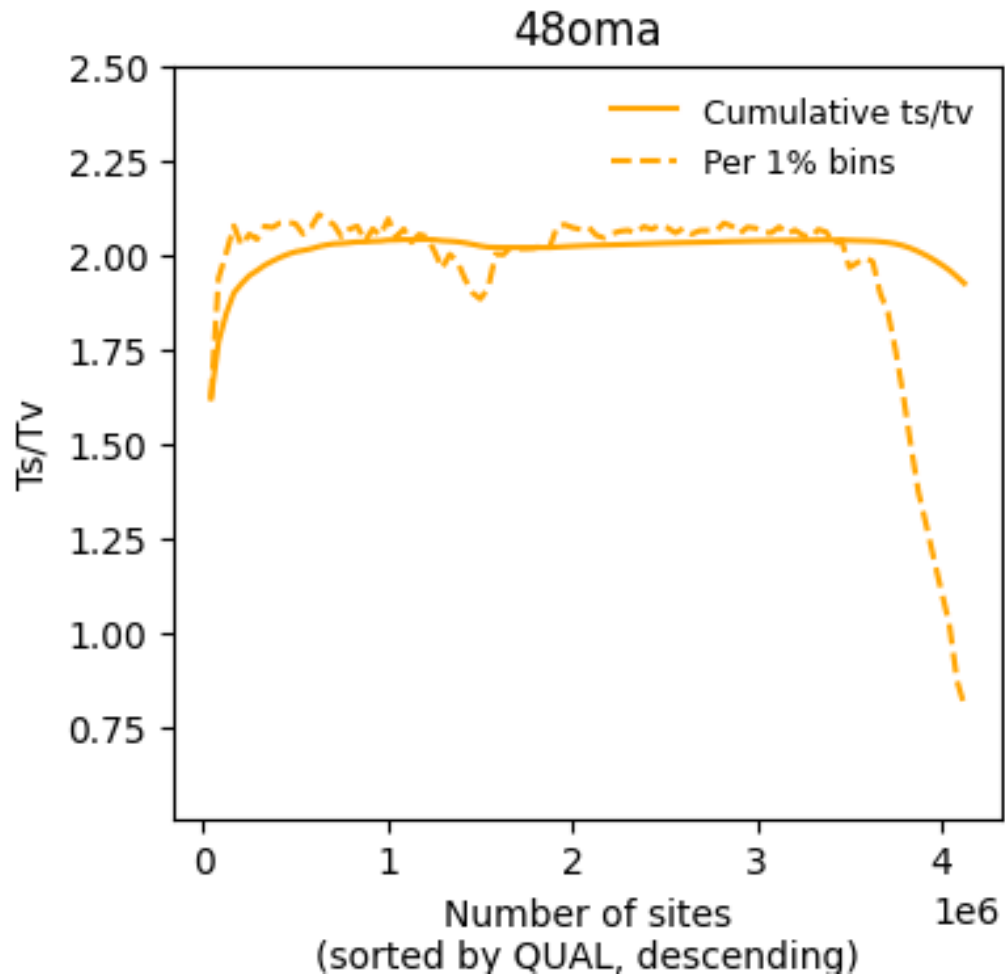# Singletons by sample (hets and homs)
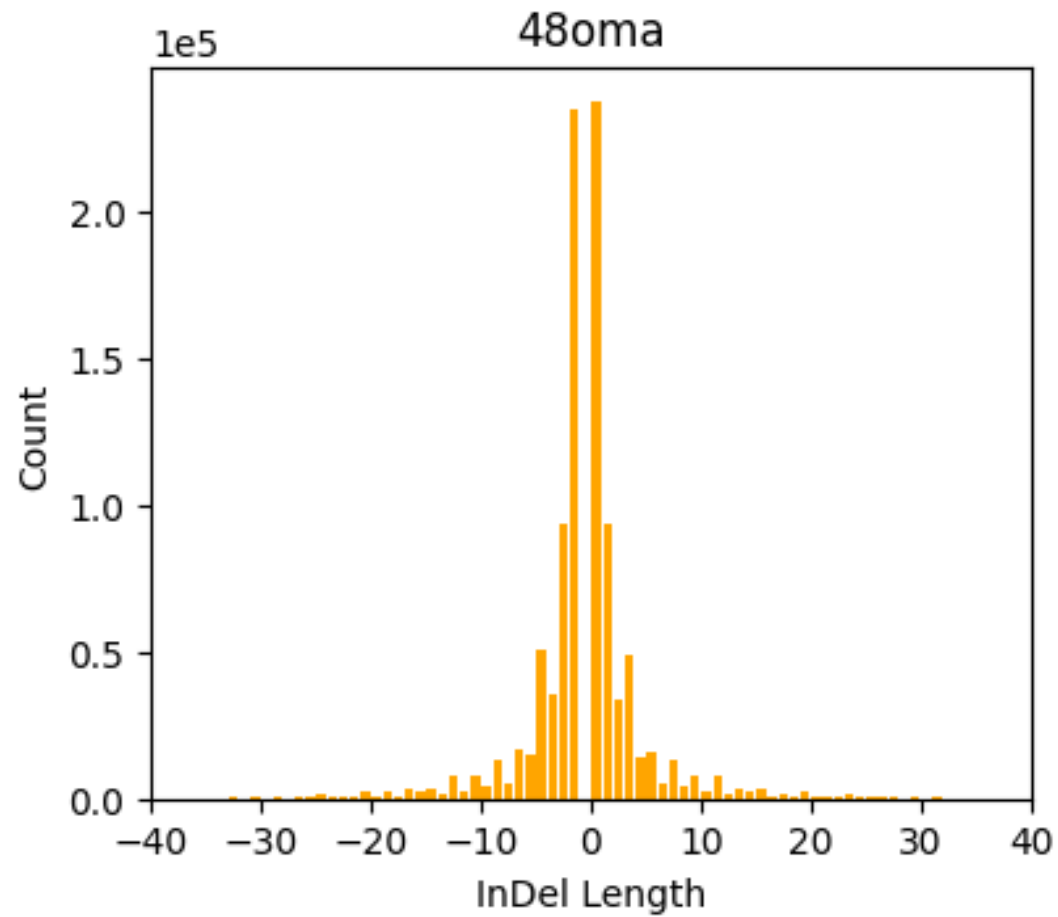
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

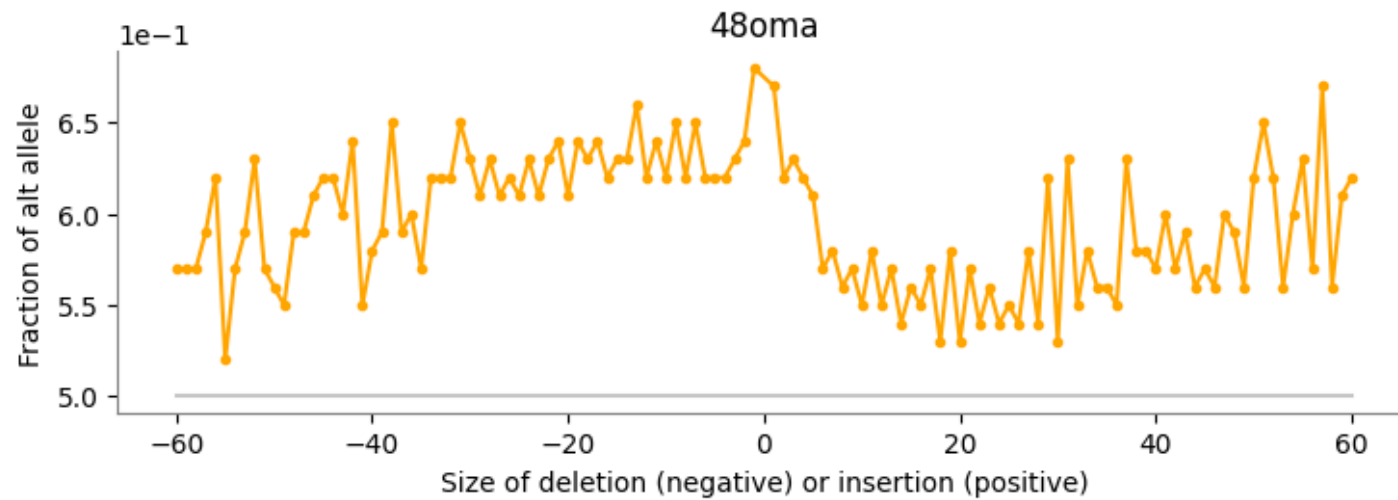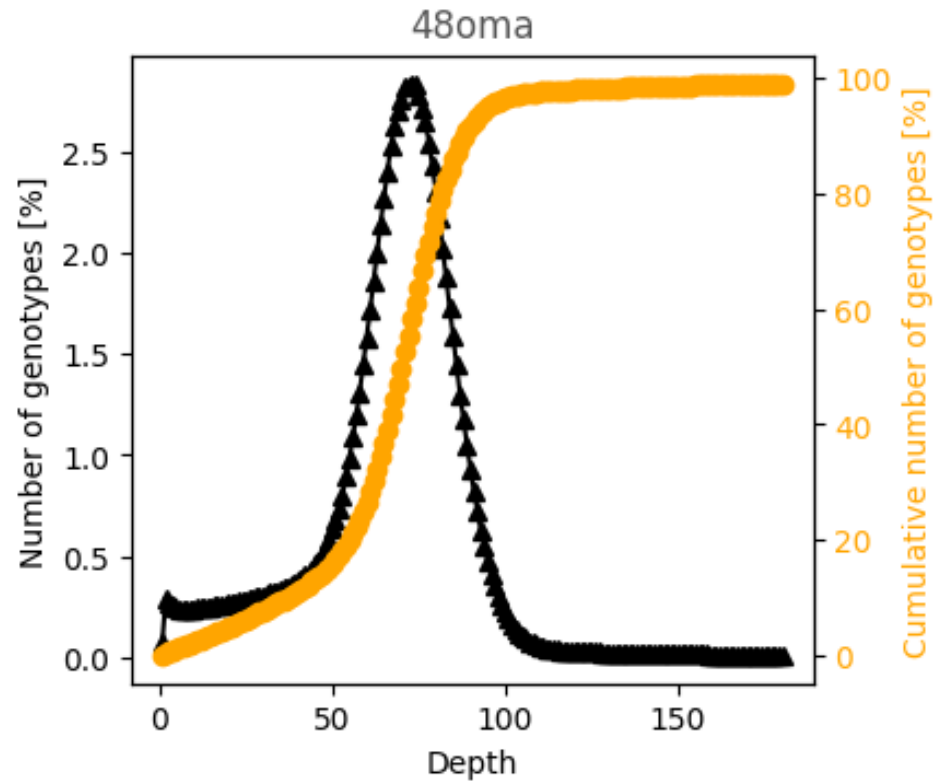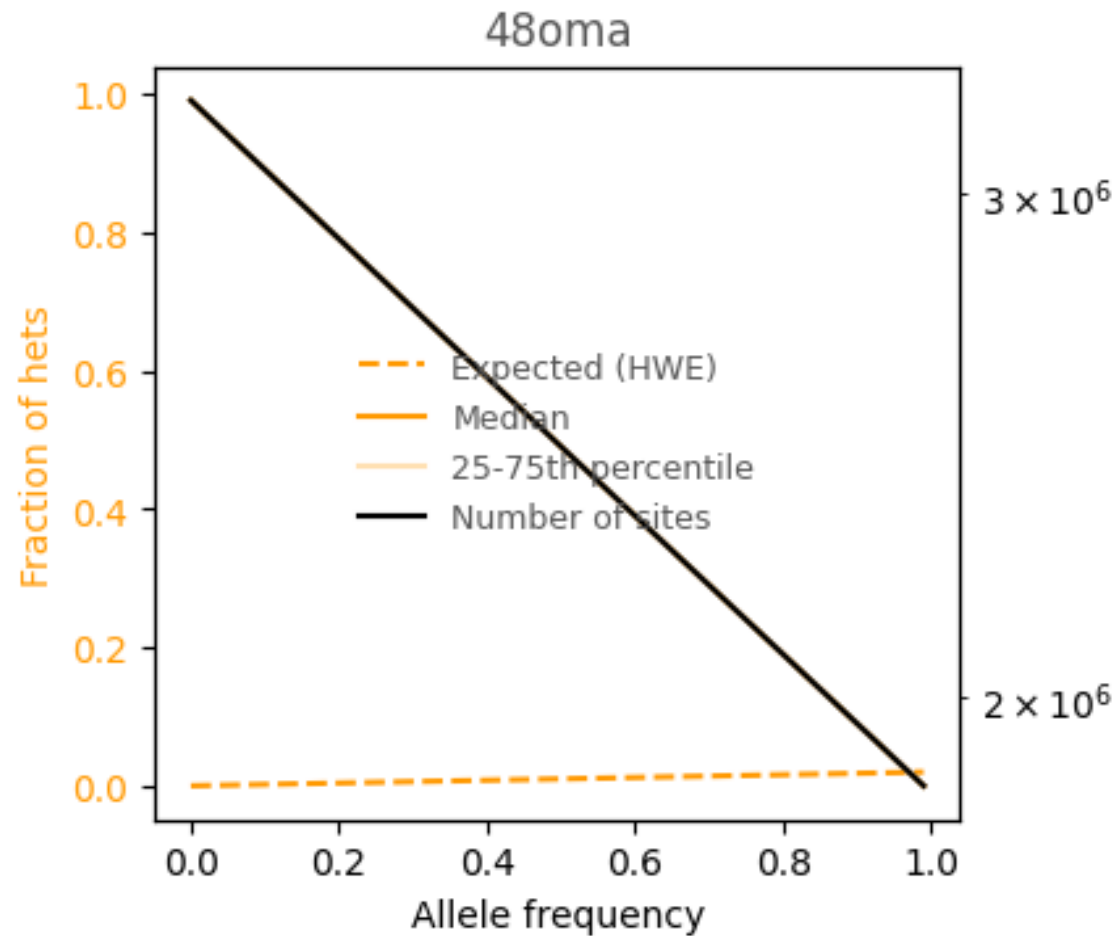# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types