Summary Numbers

	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
59wam	4,120,966	1.93	1.93	958,275	–	0	0
					* frameshift ratio: out/(out+in)		

	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
59wam	63.1%	1.90	68.2%	102,078	2,258

- 59wam .. `/ngc/projects2/gm/data/archive/2022/variants/snv/59wamonnf-110304104833-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02281-221109_A01961_AH3KN7DSX5-EXT_LAB KA_NGCWGS-NGCWGS06166_snv_germline_raw.haplotype_caller.vcf.gz`
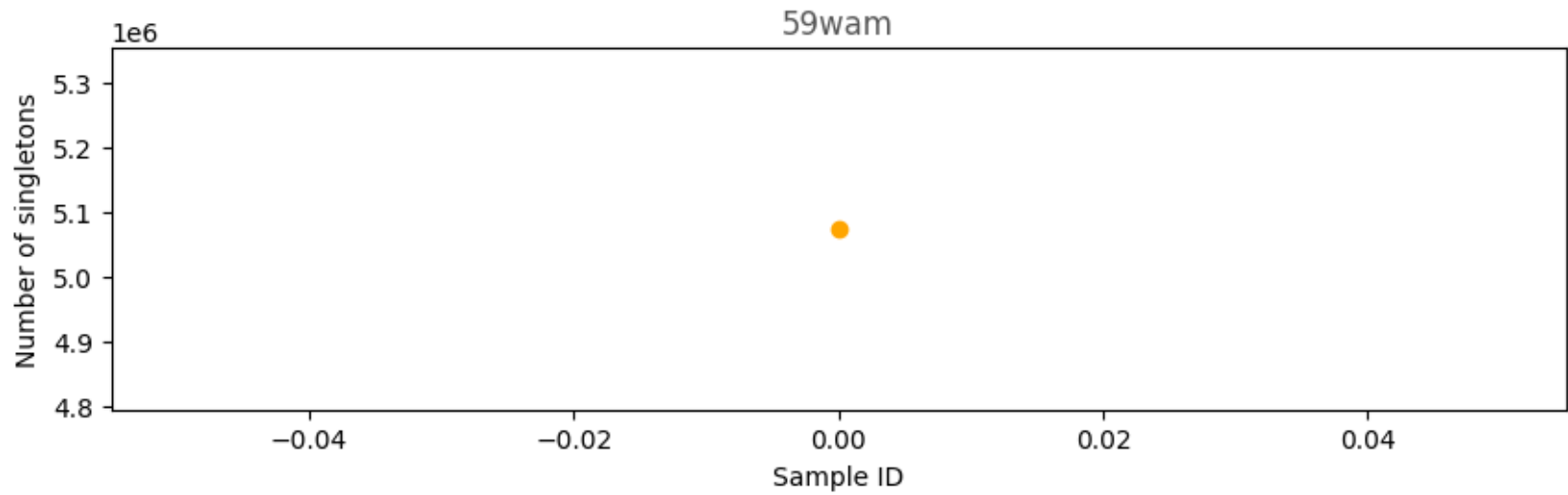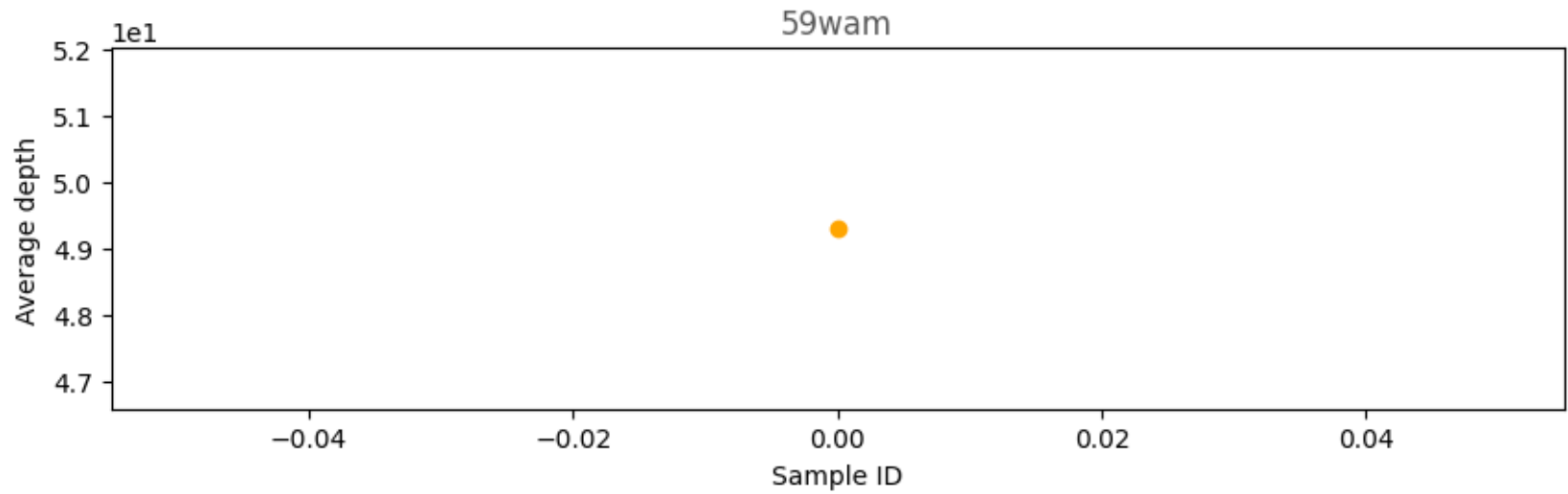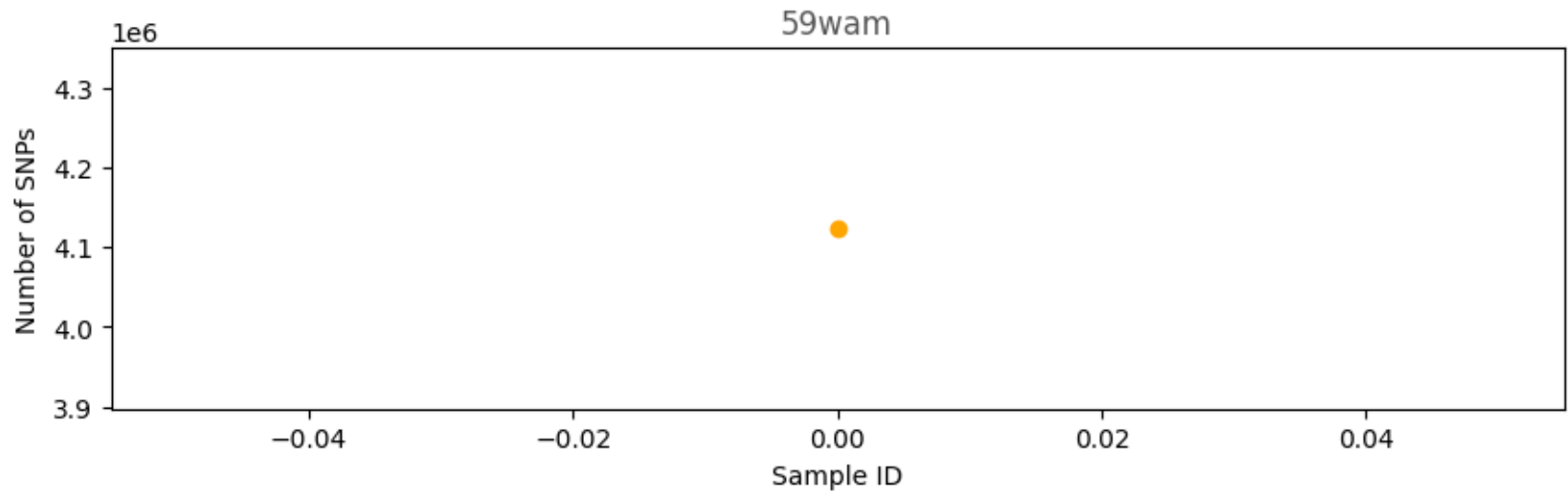
Ts/Tv by sample
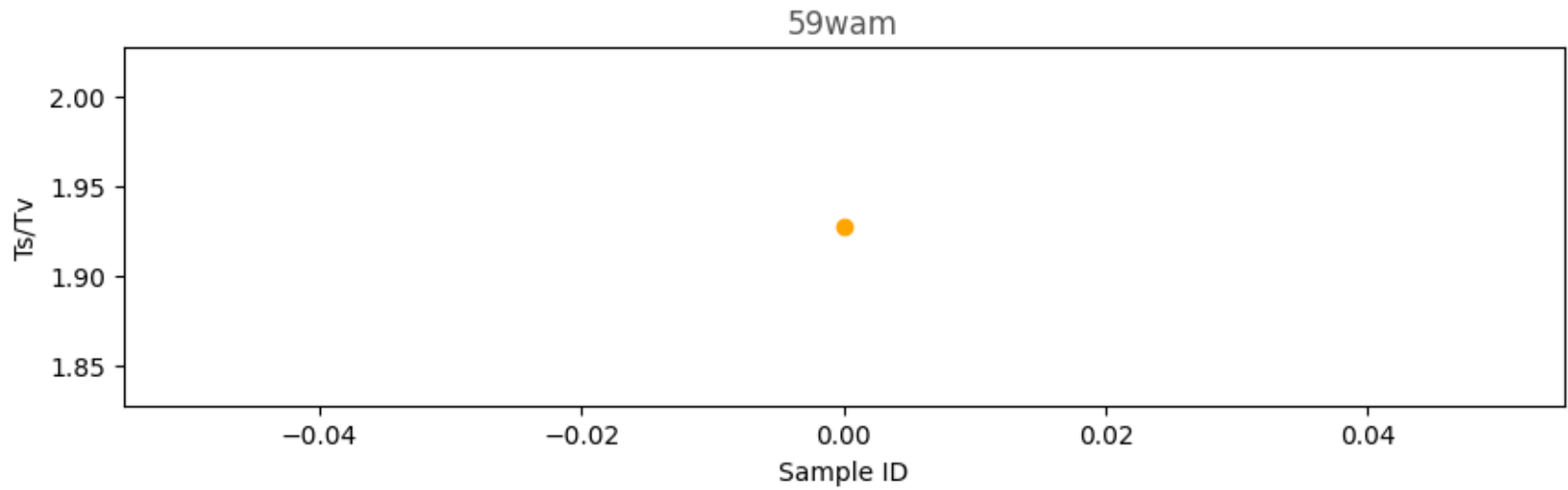
Hets vs non-ref Homs by sample
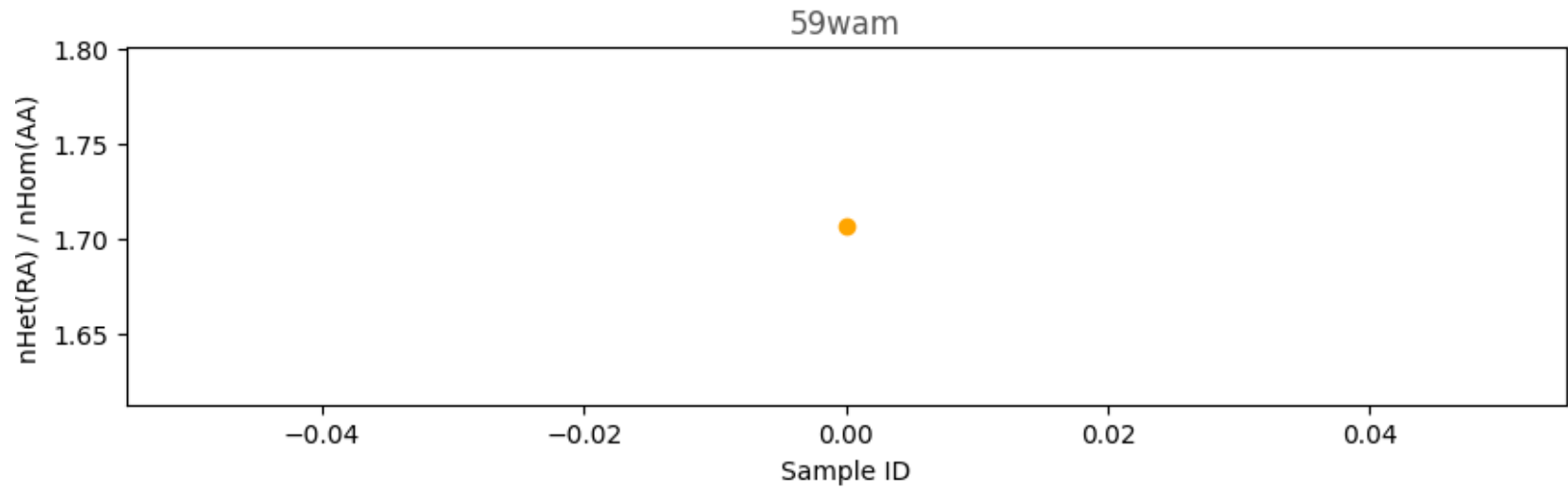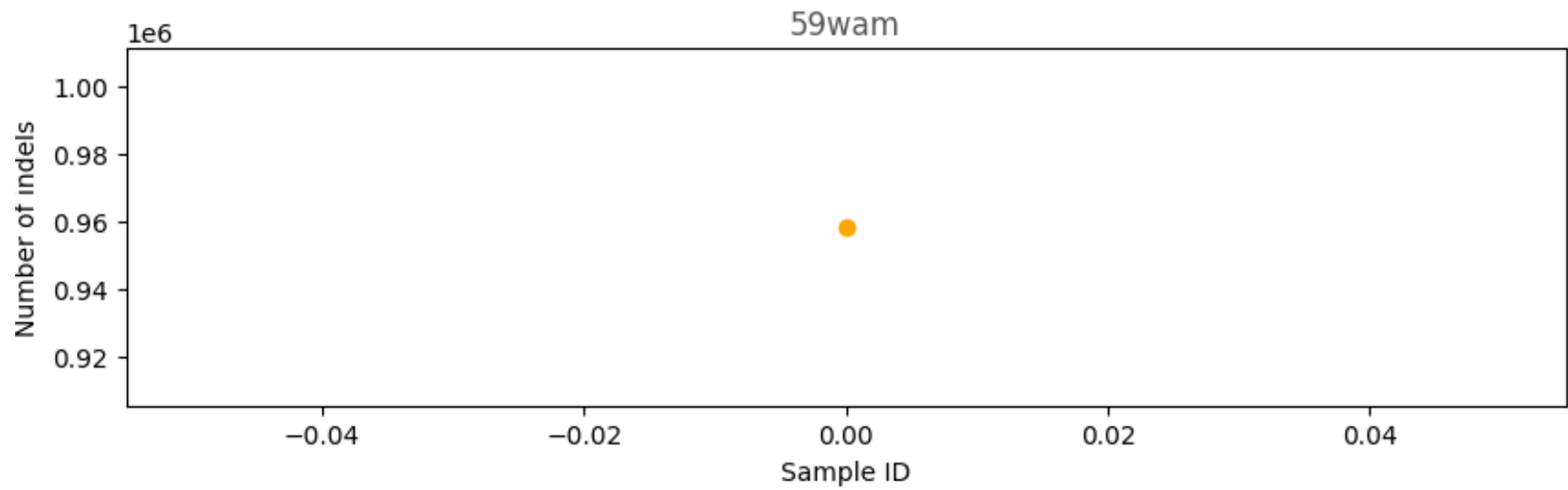
Singletons by sample (hets and homs)
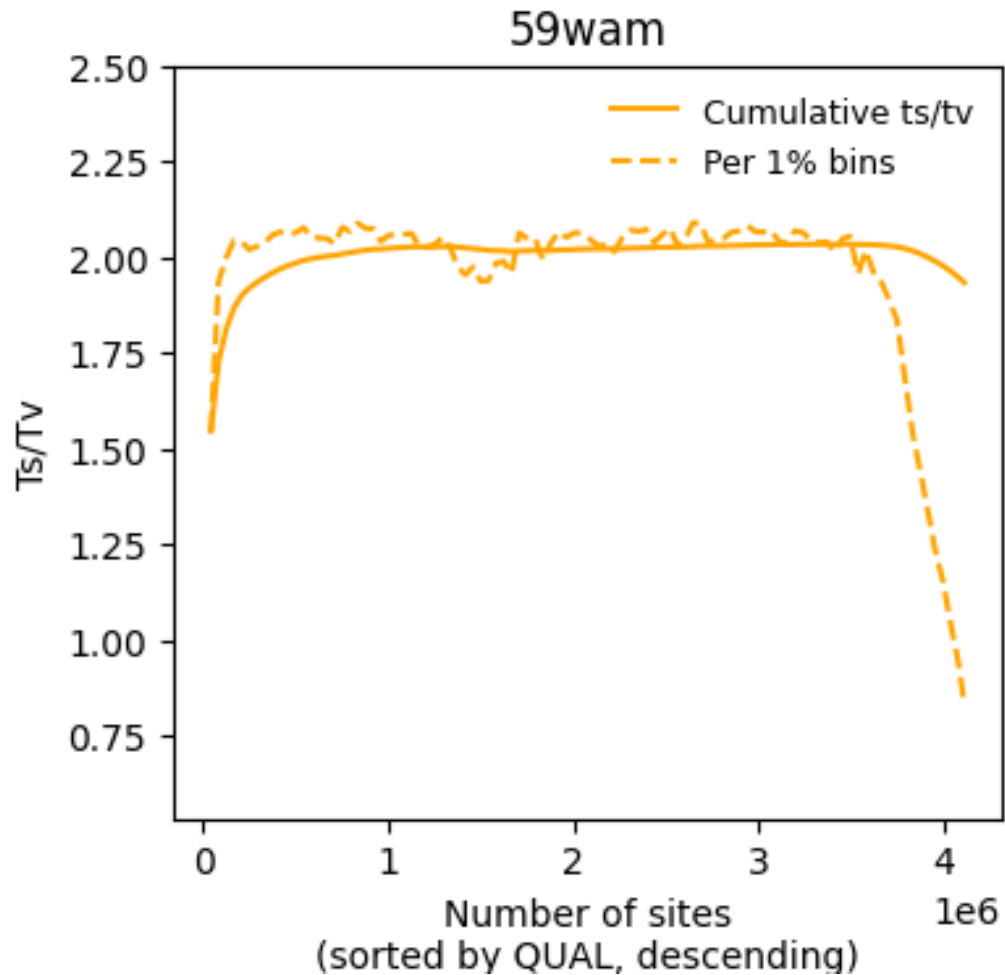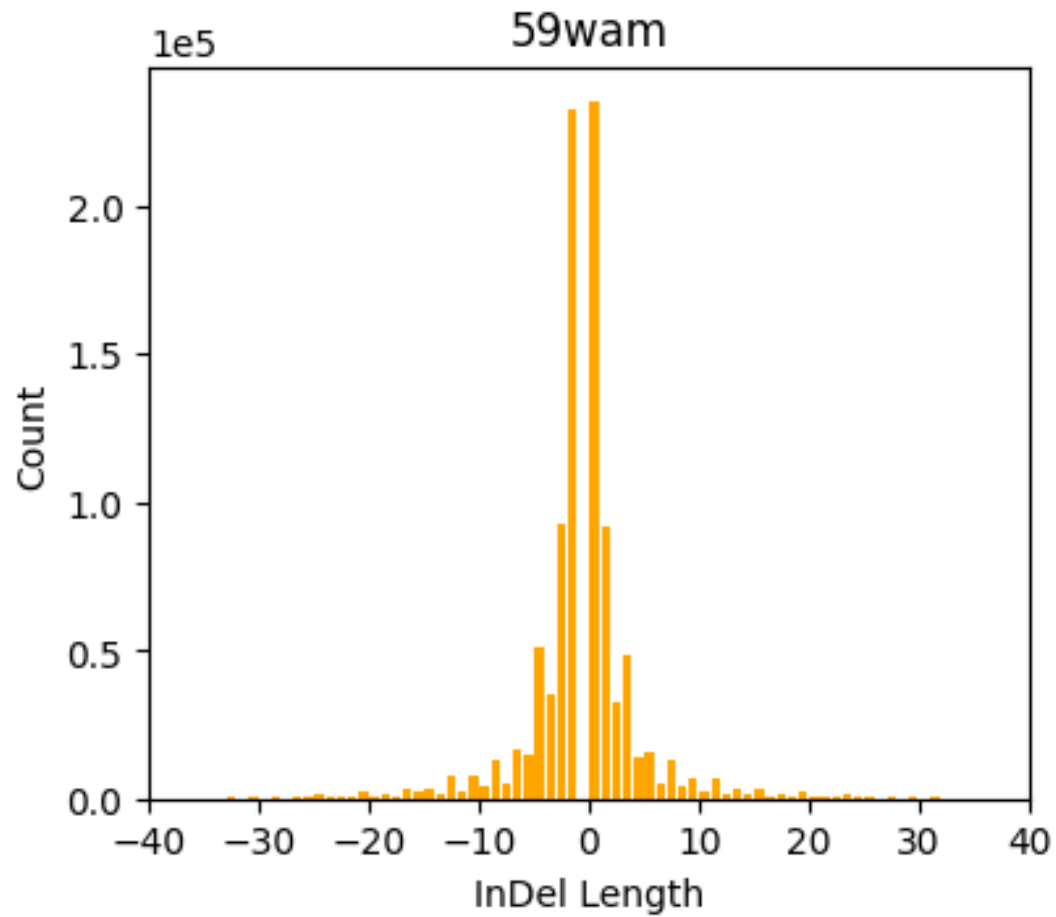
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

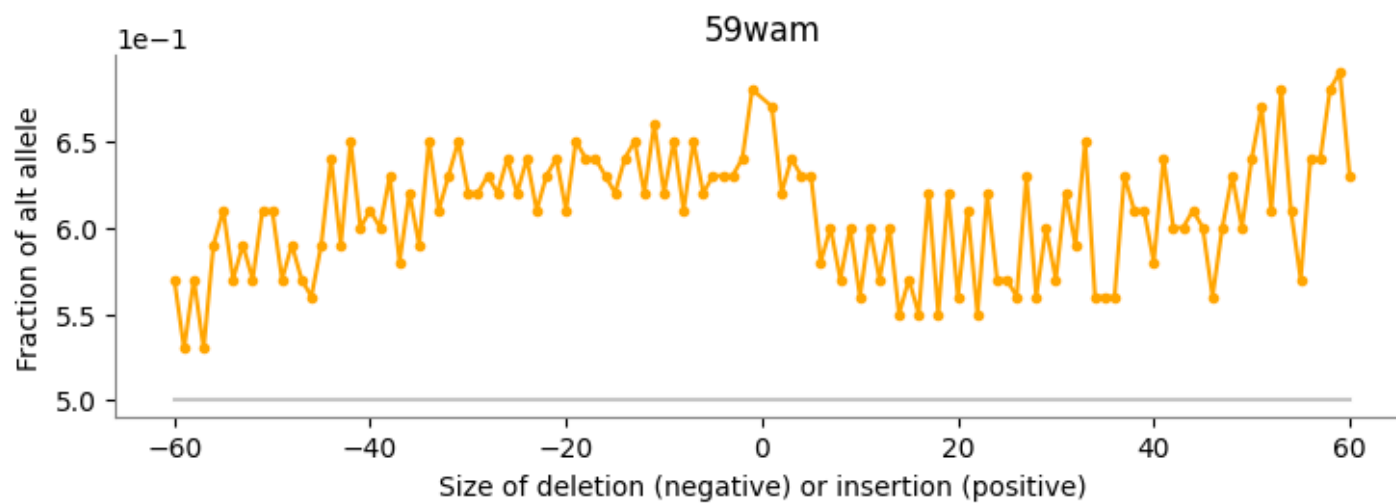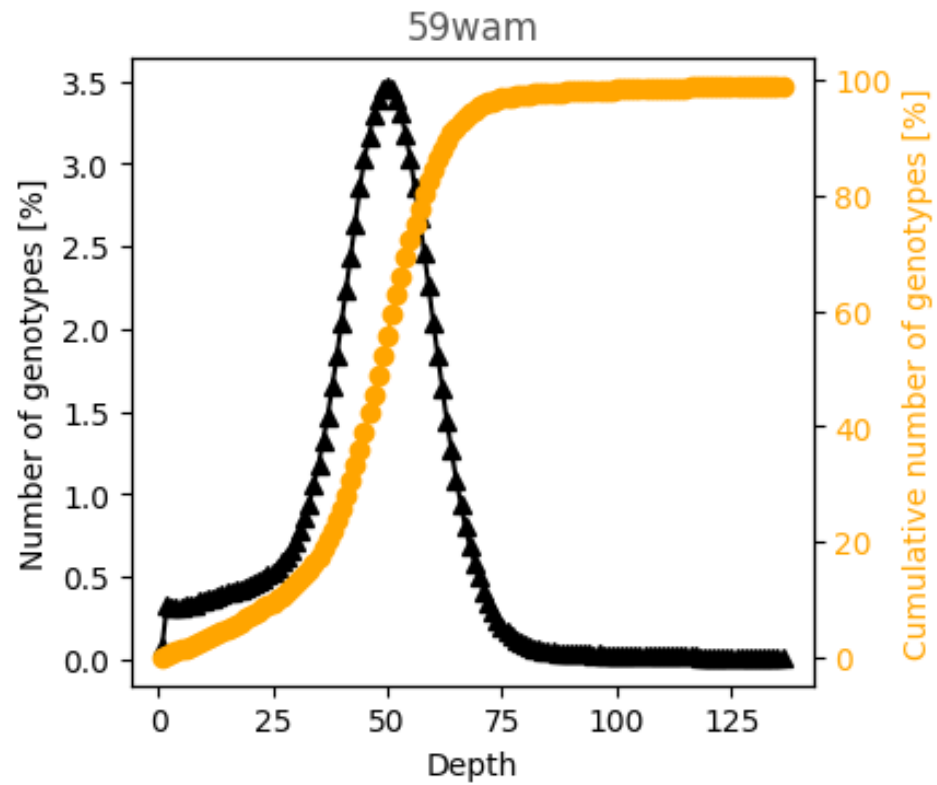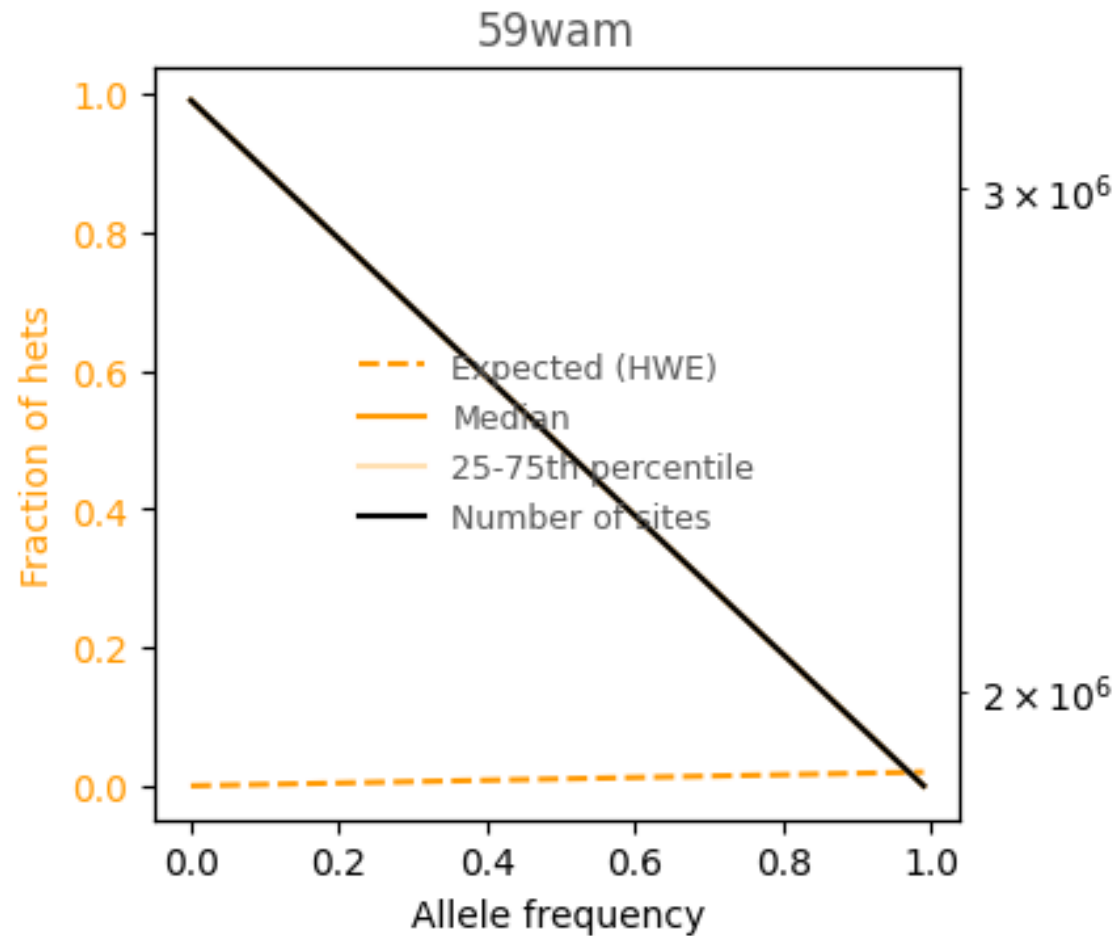# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types