Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
50she	4,103,451	1.93	1.93	960,094	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
50she	62.6%	1.90	68.0%	103,874	2,133

- 50she .. `/ngc/projects2/gm/data/archive/2022/variants/snv/50sheheef-103910868763-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01538-220916_A01411_AHTJ7JDSX3-EXT_LAB`
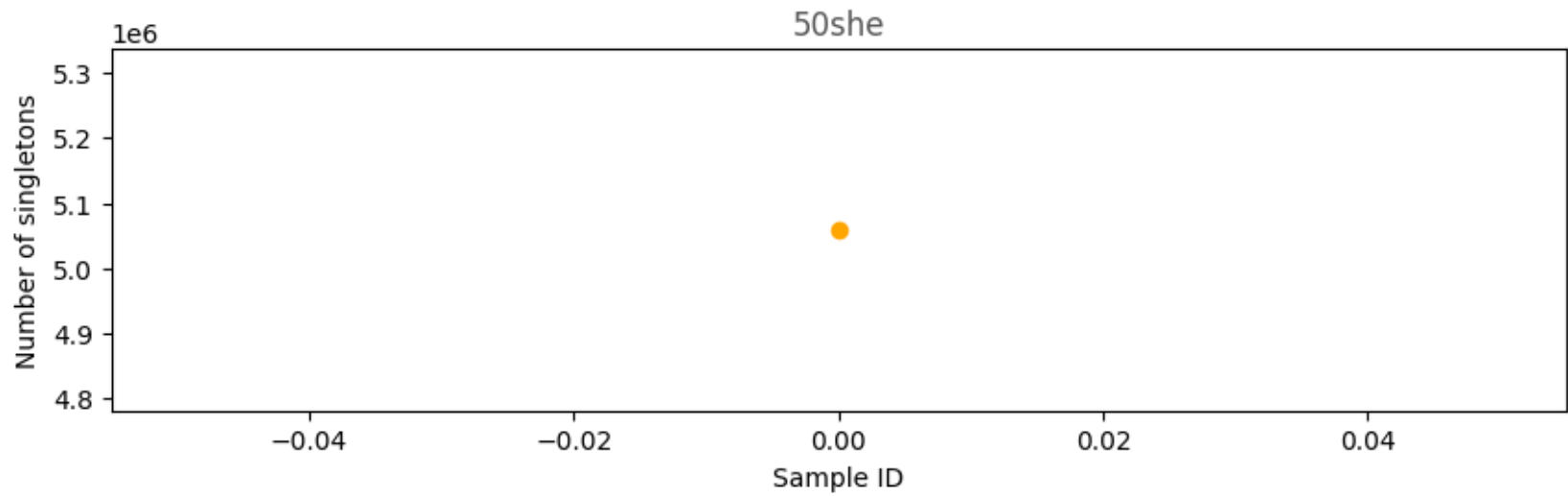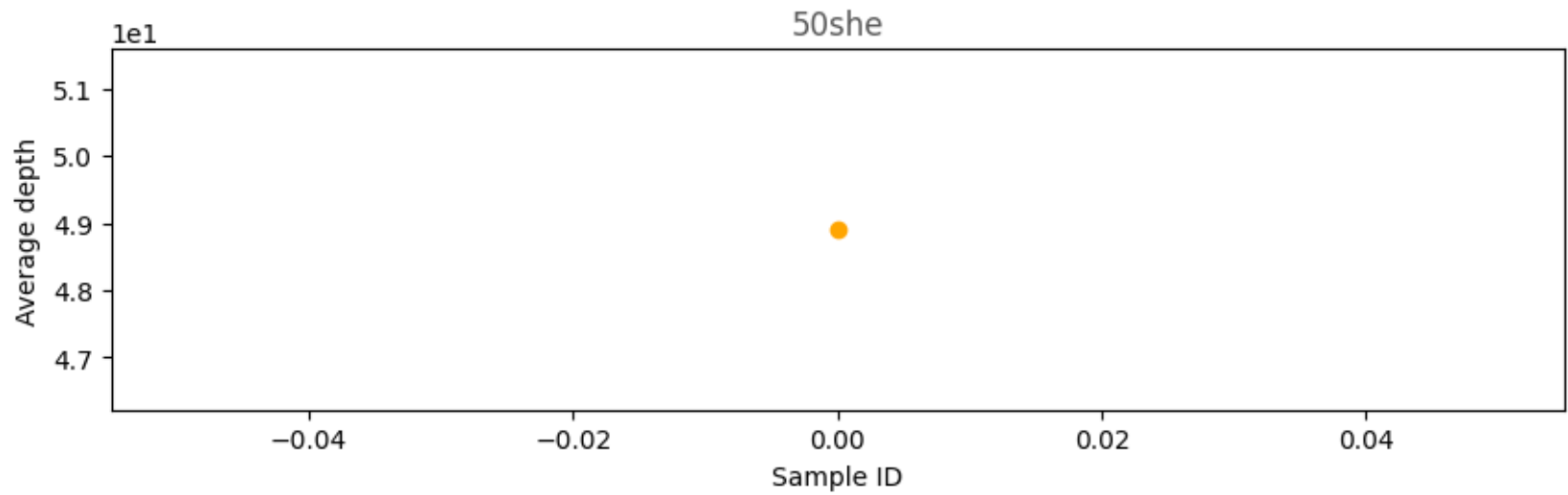 `KA_NGCWGS-NGCWGS05380_snv_germline_raw.haplotype_caller.vcf.gz`

Ts/Tv by sample
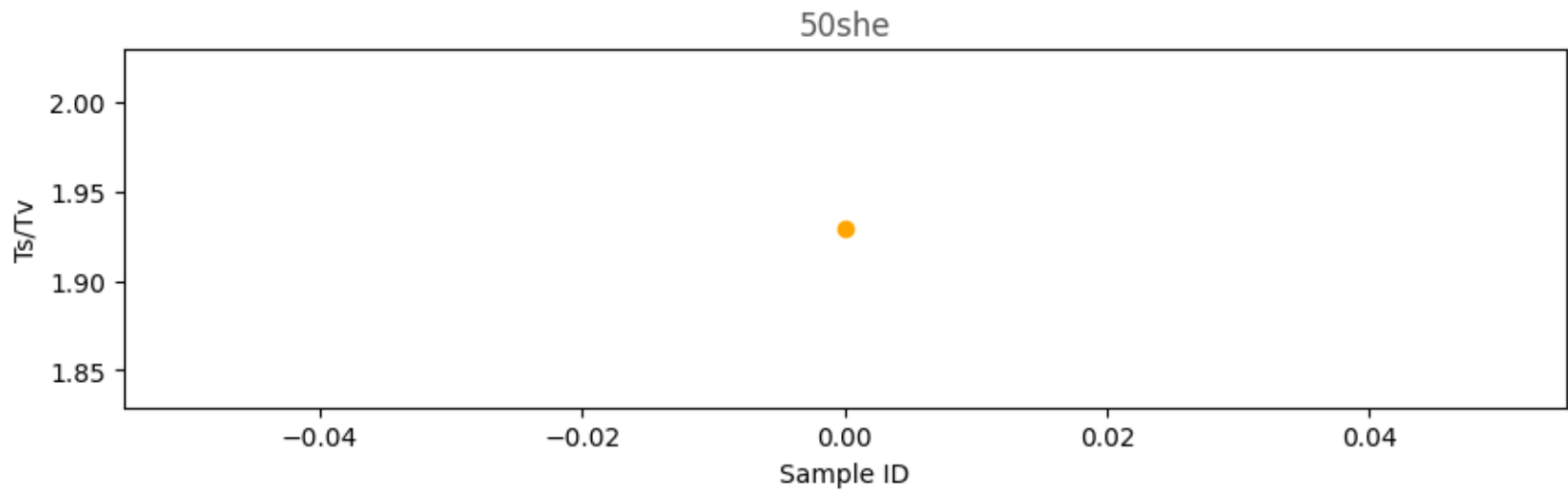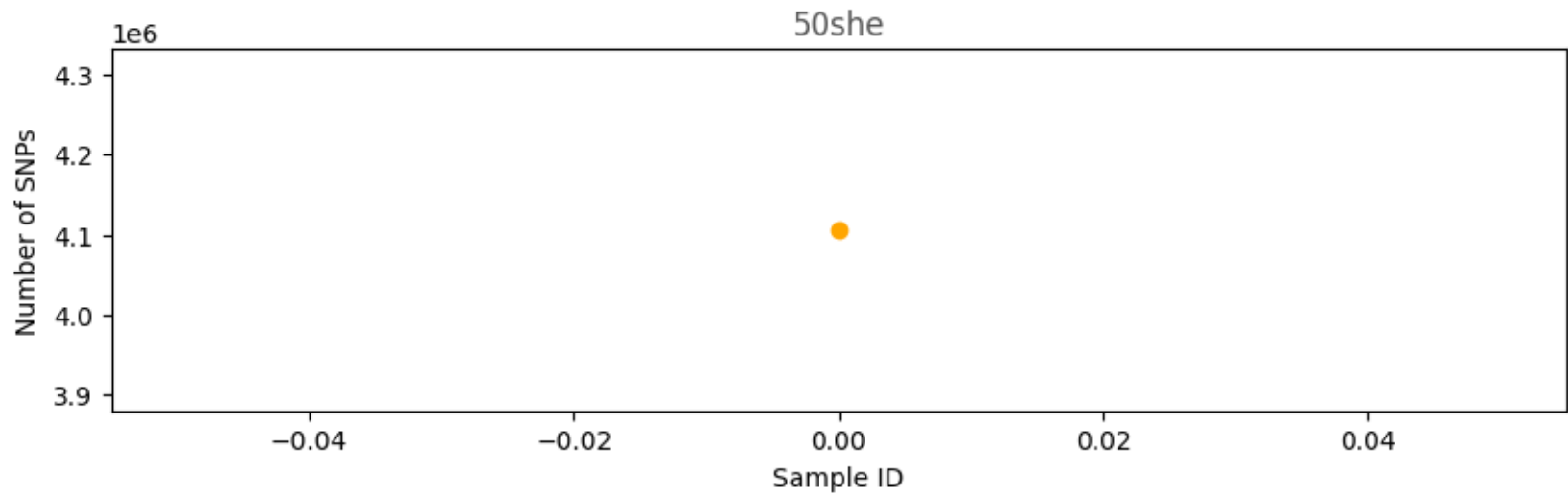
Hets vs non-ref Homs by sample
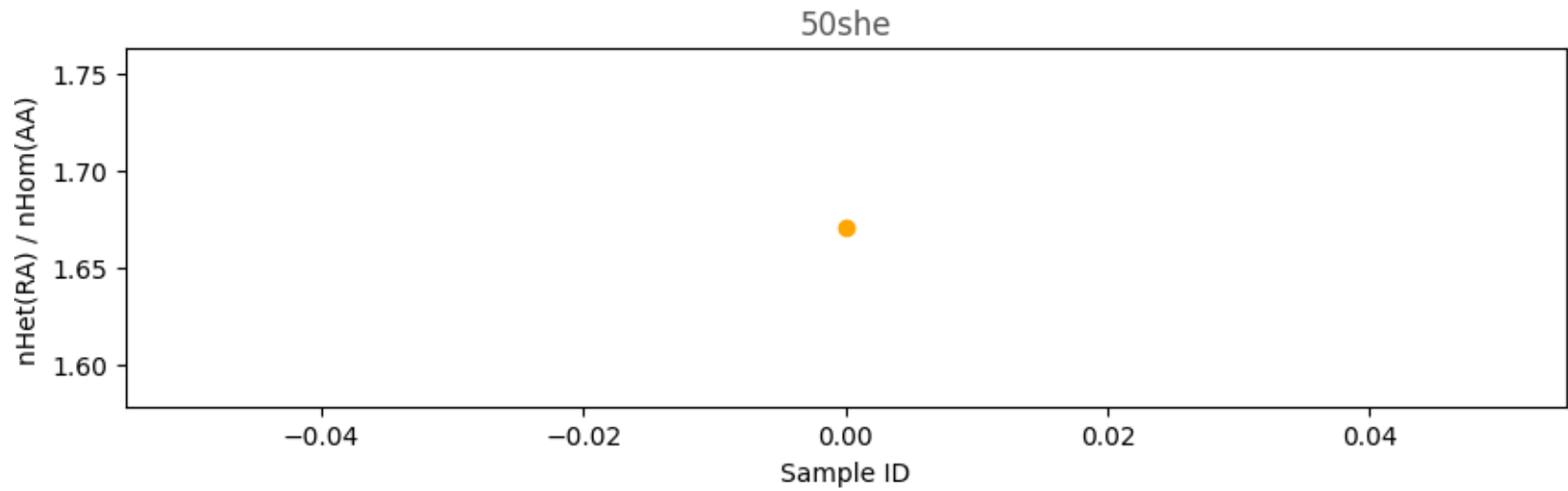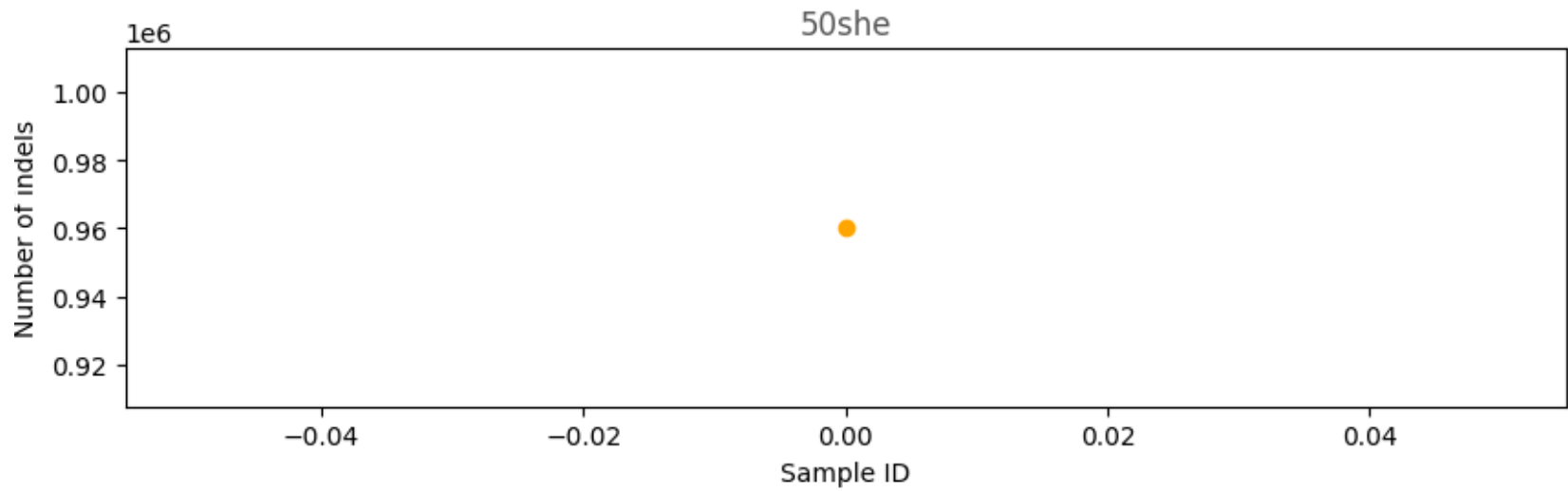
Singletons by sample (hets and homs)
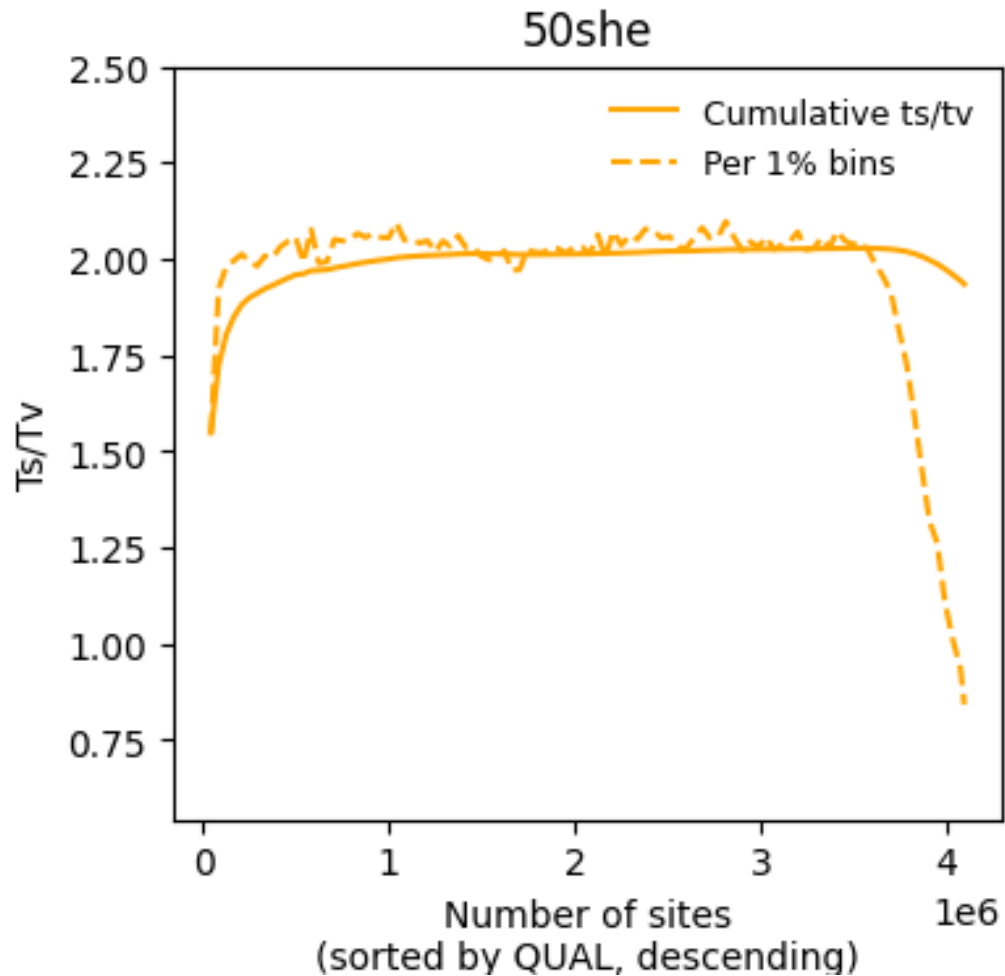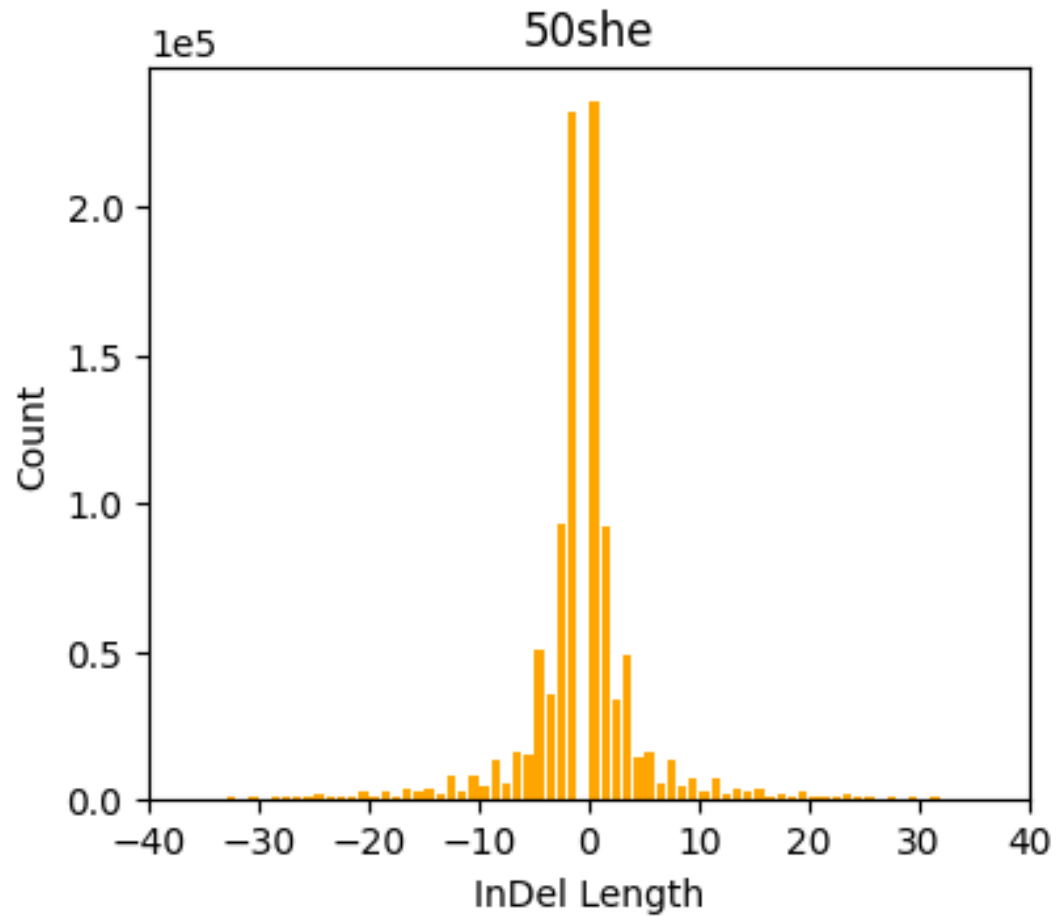
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

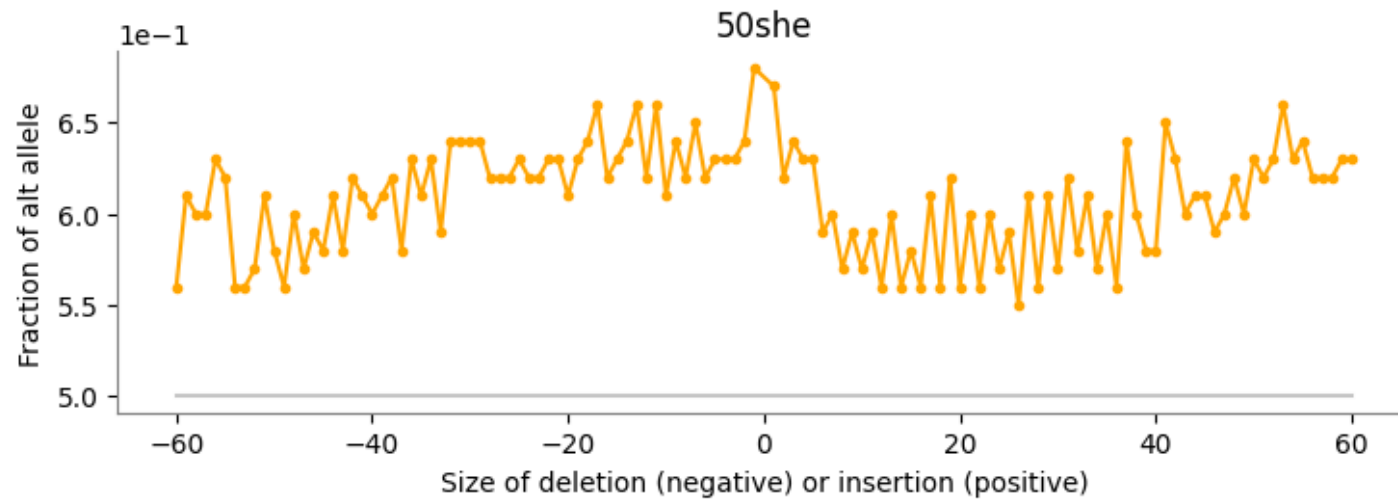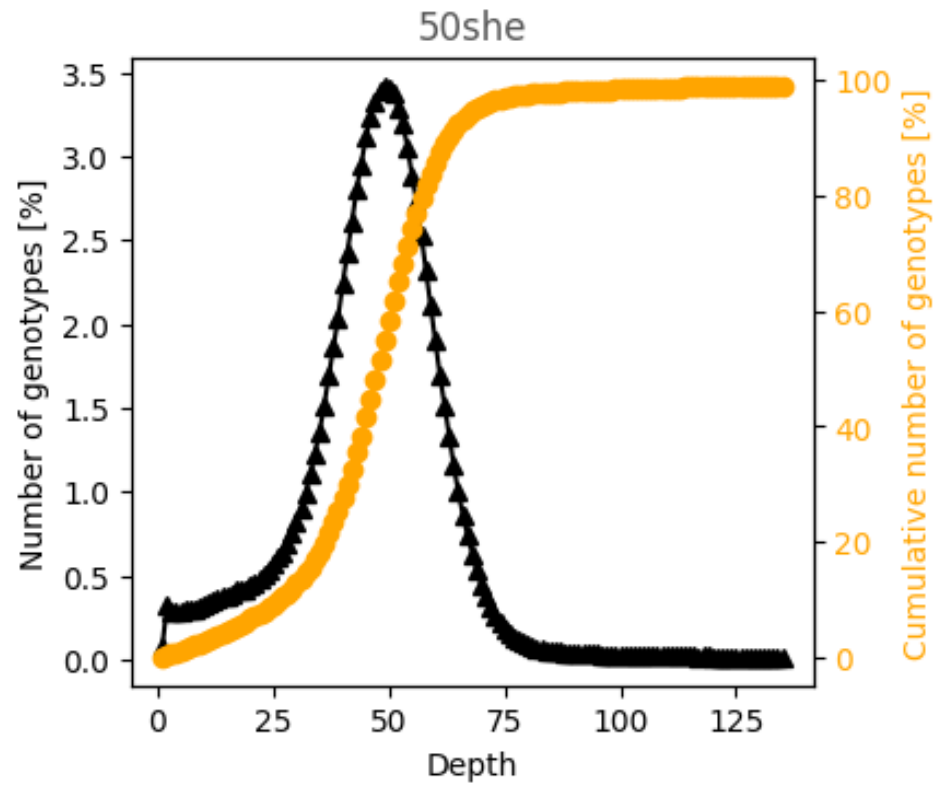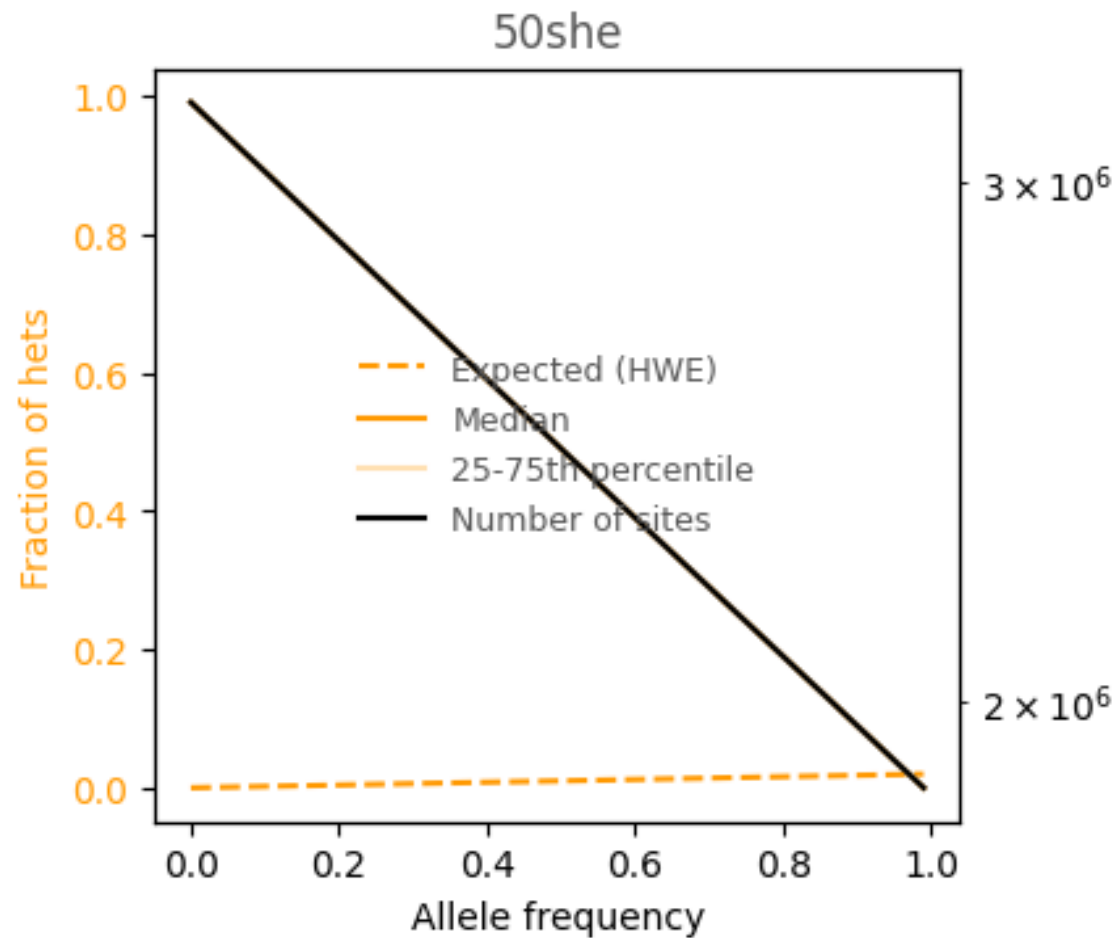# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types