Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90sil	4,217,316	1.94	1.95	985,621	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90sil	63.9%	1.92	69.2%	108,532	2,200

- 90sil .. `/ngc/projects2/gm/data/archive/2022/variants/snv/90silfrtf-103892128280-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01484-220914_A01176_AHTNVGDSX3-EXT_LAB`
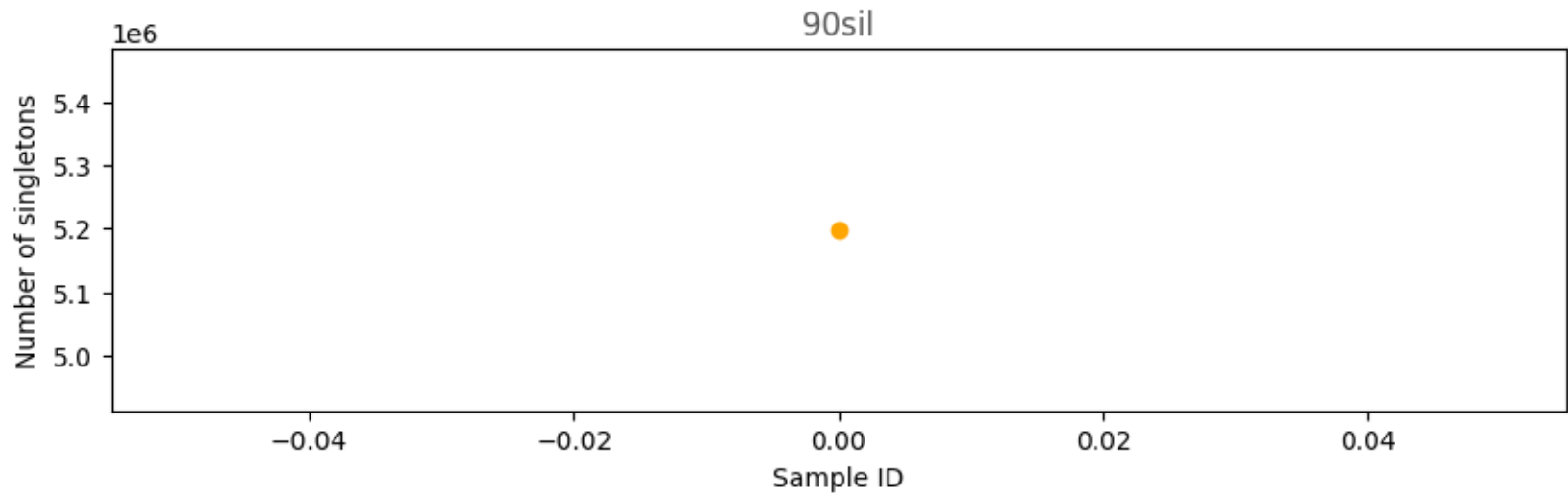 `KA_NGCWGS-NGCWGS05258_22RKG017533_snv_germline_raw.haplotype_caller.vcf.gz`
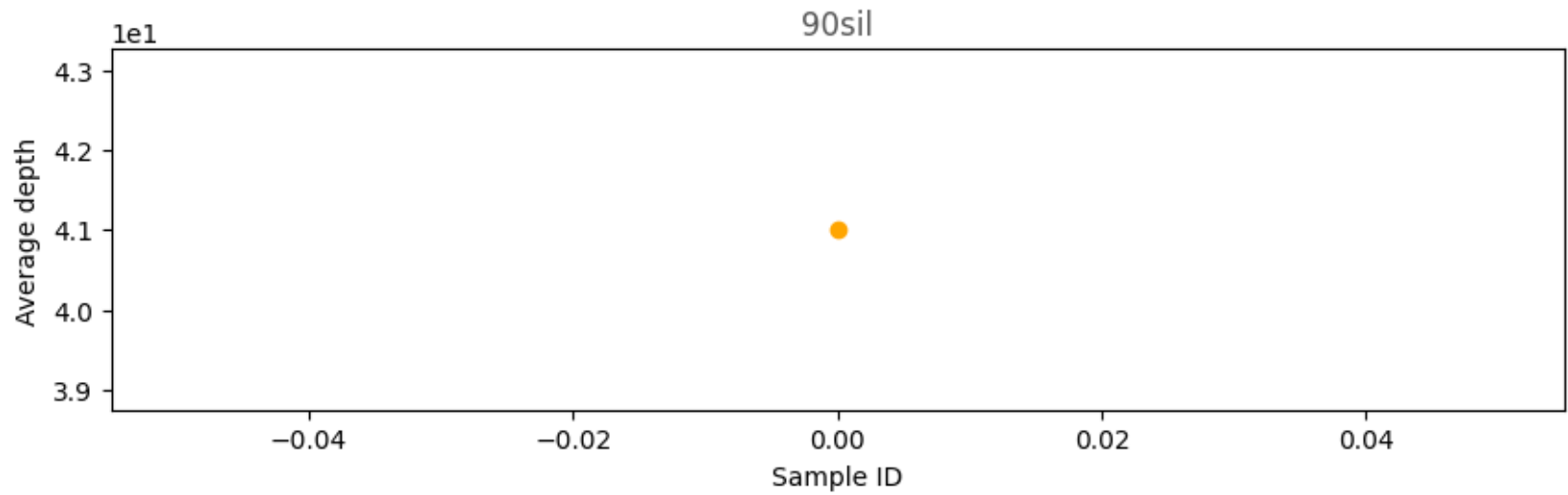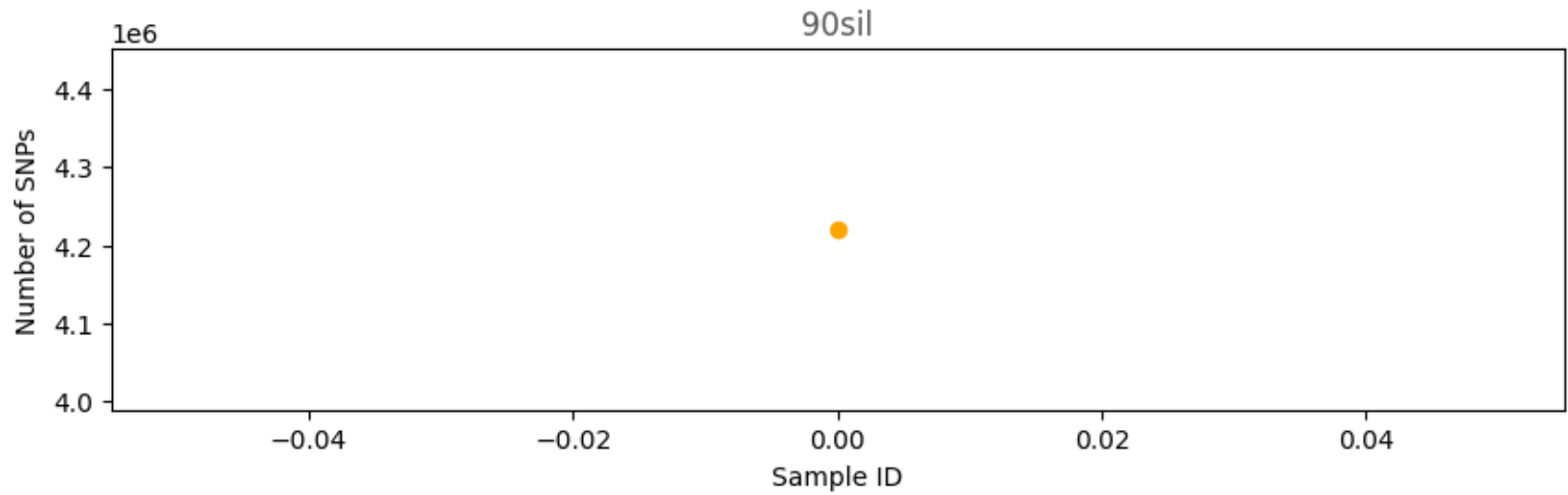
Ts/Tv by sample
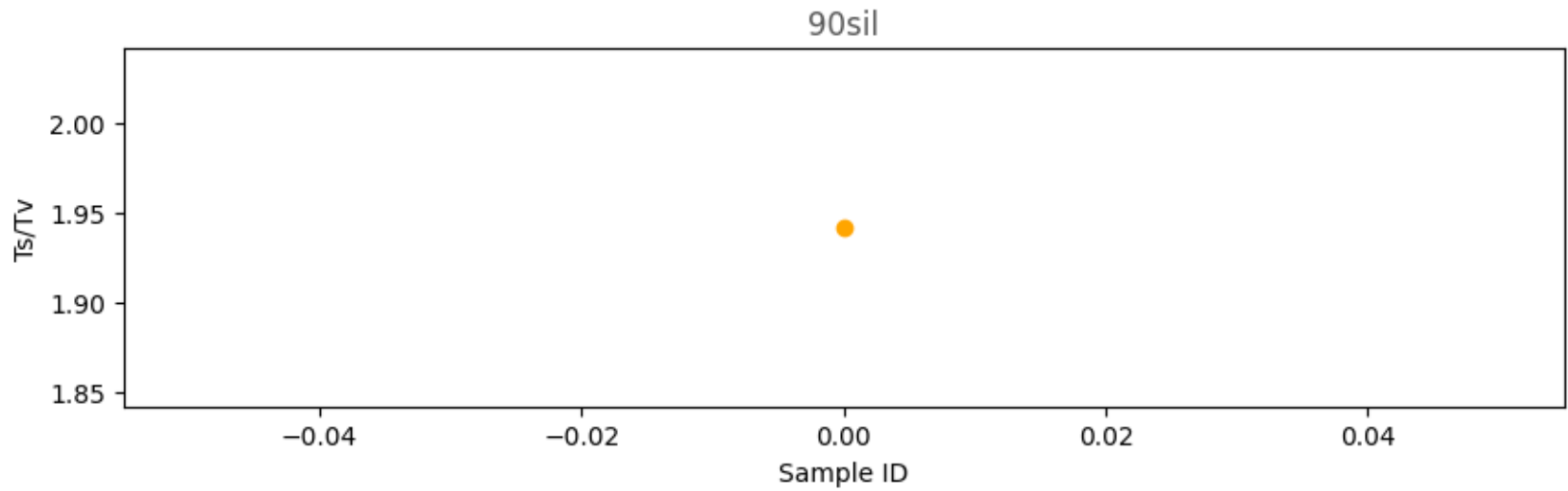
Hets vs non-ref Homs by sample
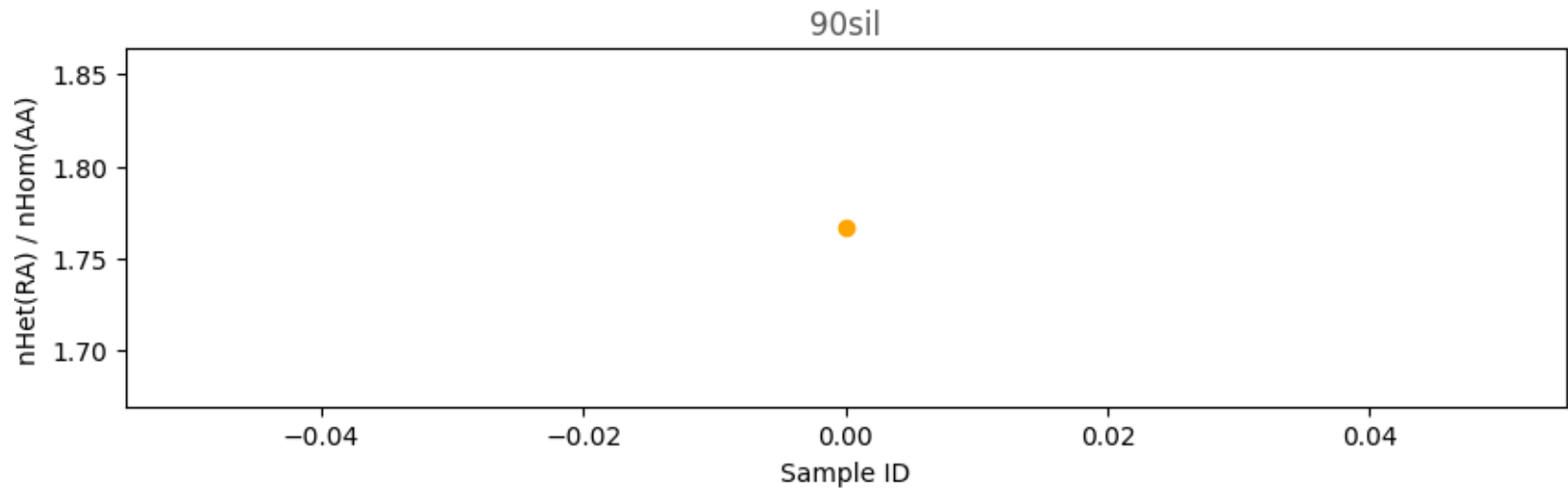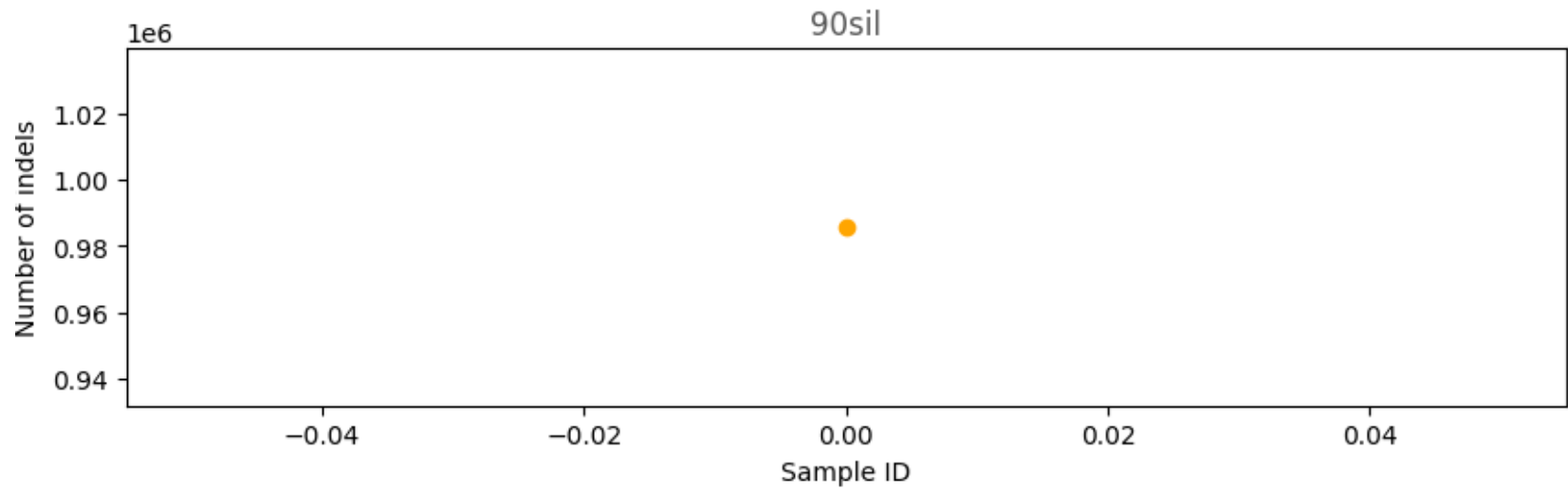
Singletons by sample (hets and homs)
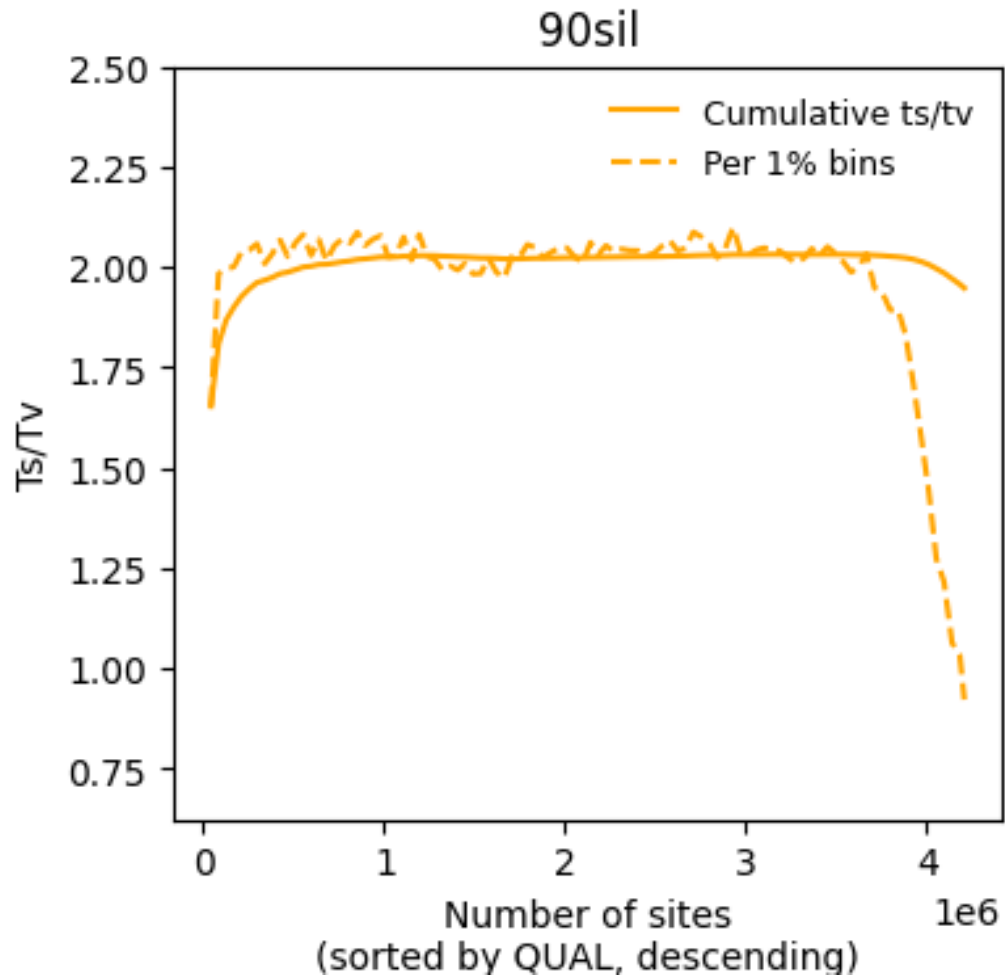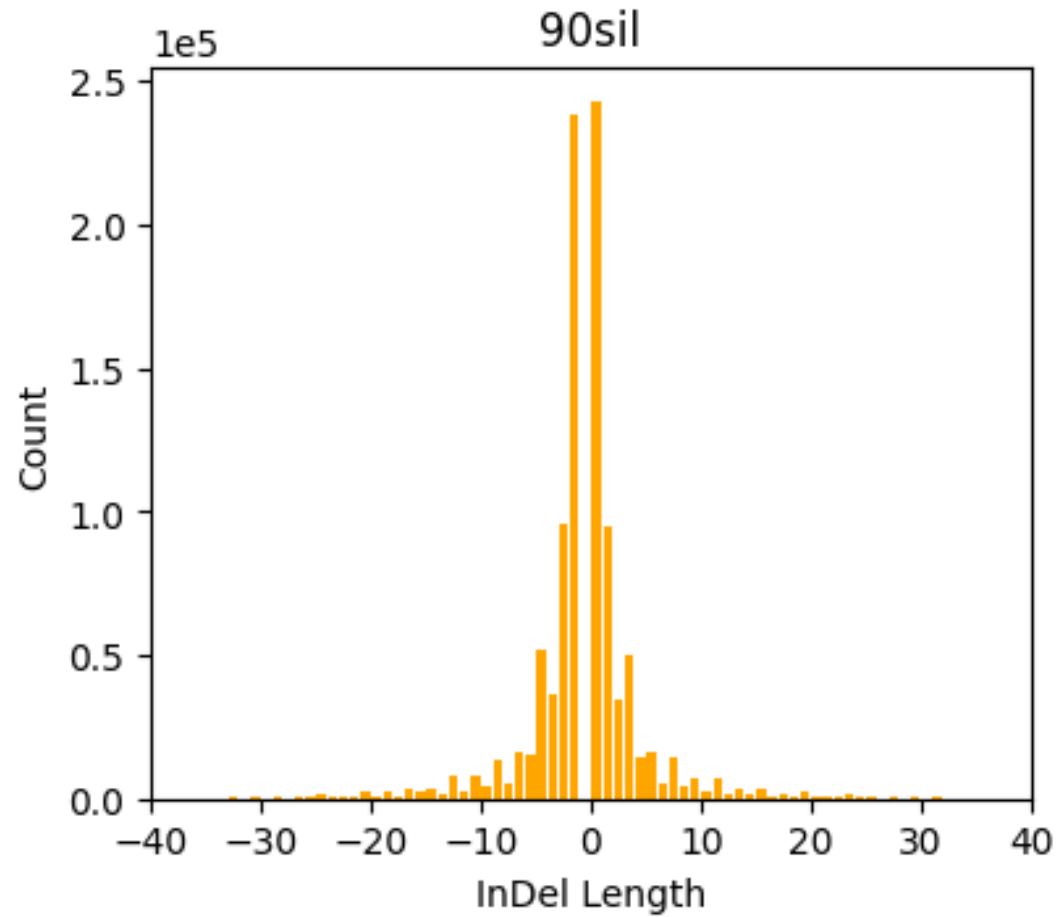
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

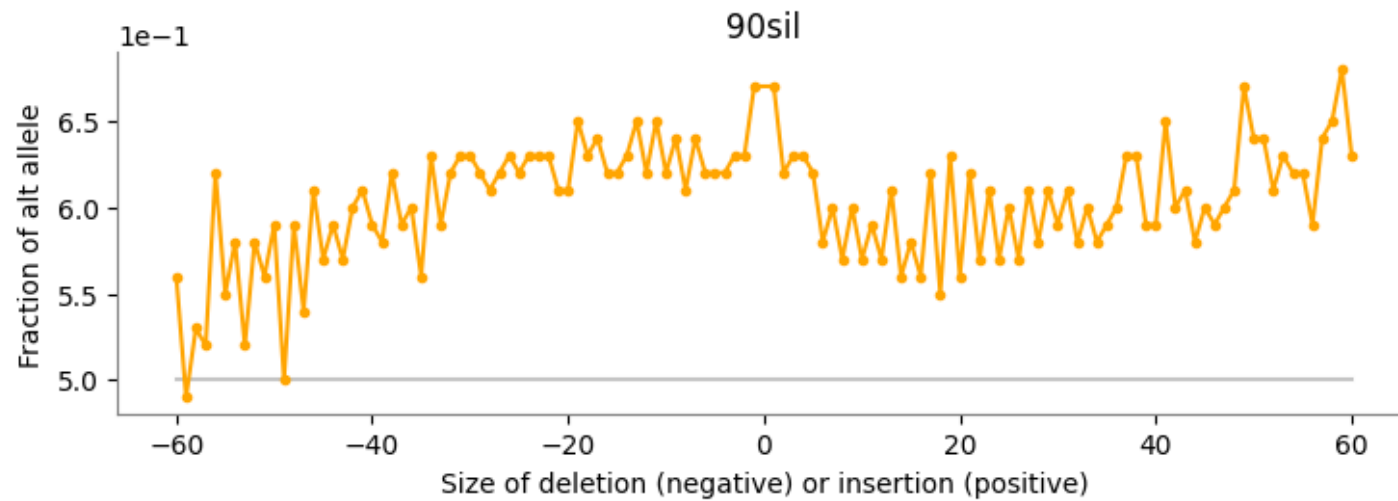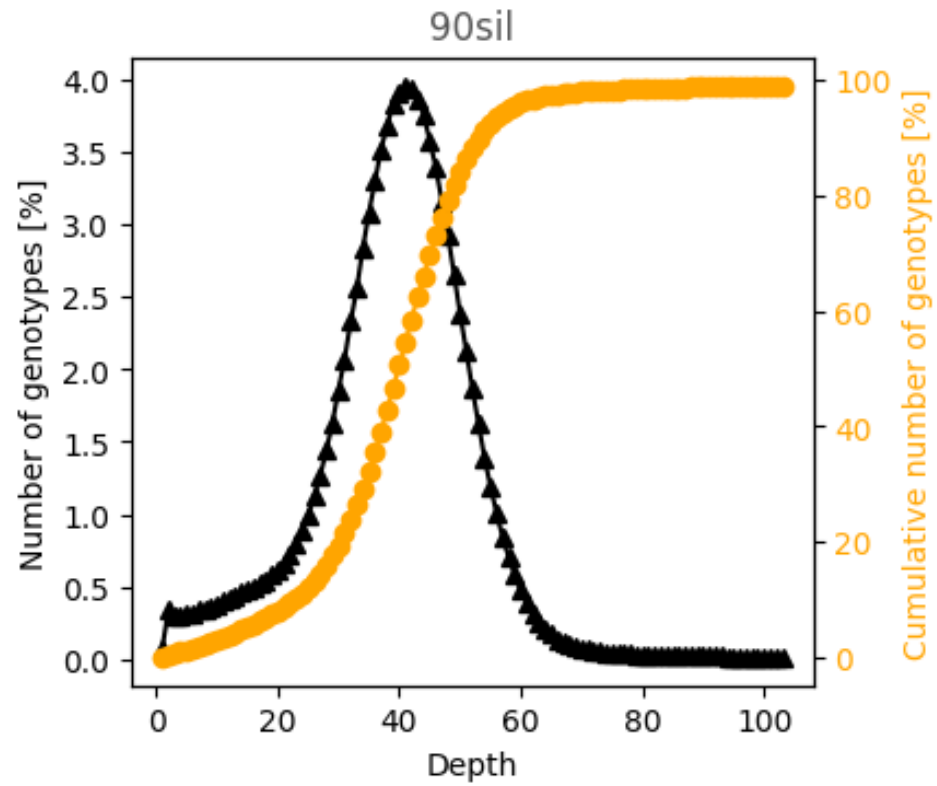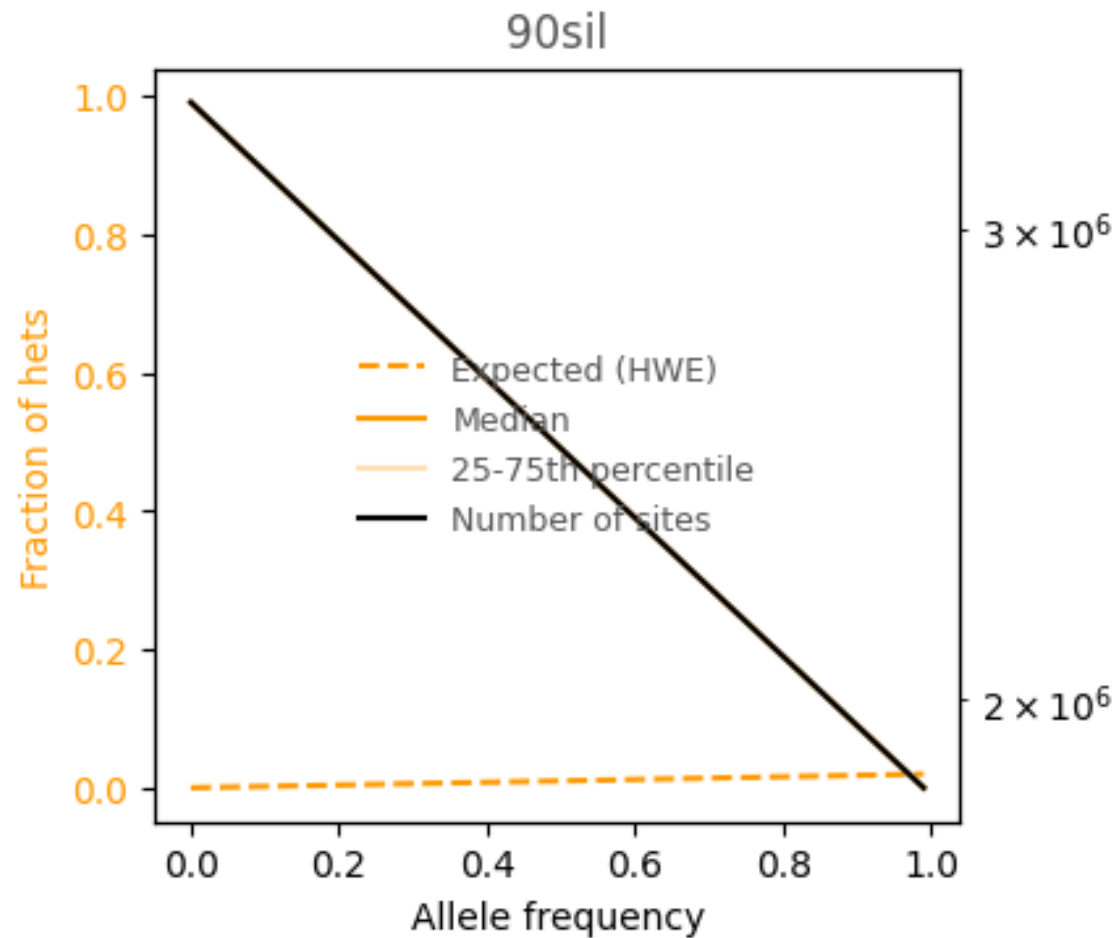# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types