Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90ott	4,060,838	1.93	1.93	941,801	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90ott	62.6%	1.89	66.9%	96,393	1,866

- 90ott .. `/ngc/projects2/gm/data/archive/2022/variants/snv/90ottherm-103818126520-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211217_A01411_BHHJ3KDSX2-RHGM_LABKA_WGSA`
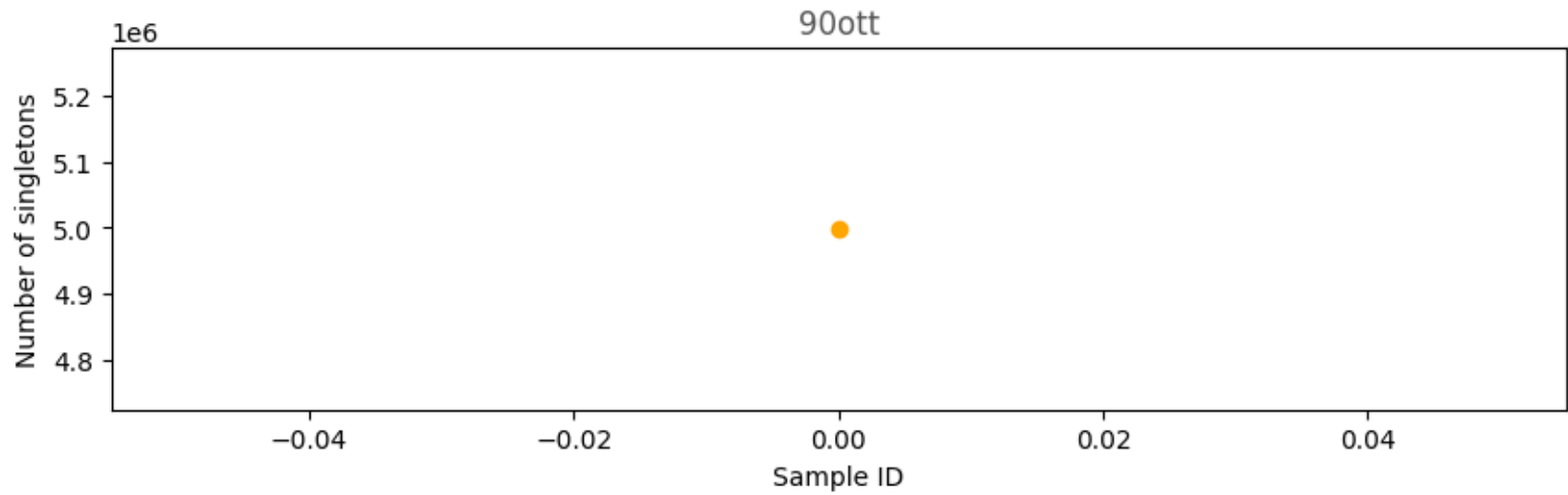 `KUT-WGSAKUT03540_21RKG030757x01_snv_germline_raw.haplotype_caller.vcf.gz`
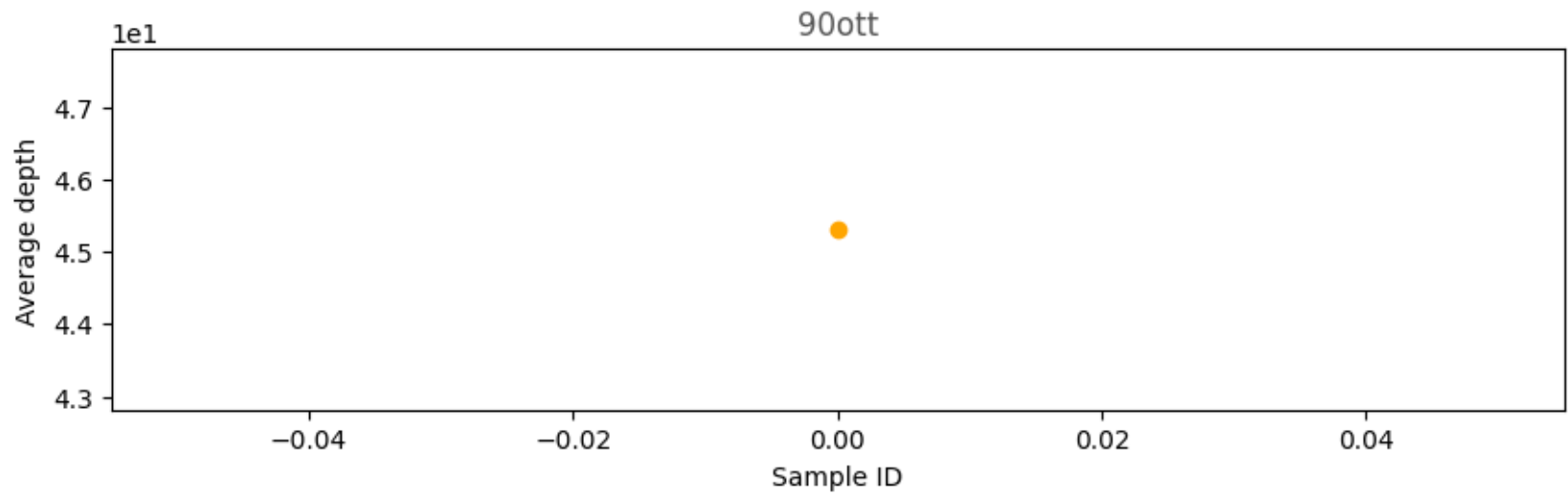
Ts/Tv by sample

90ott
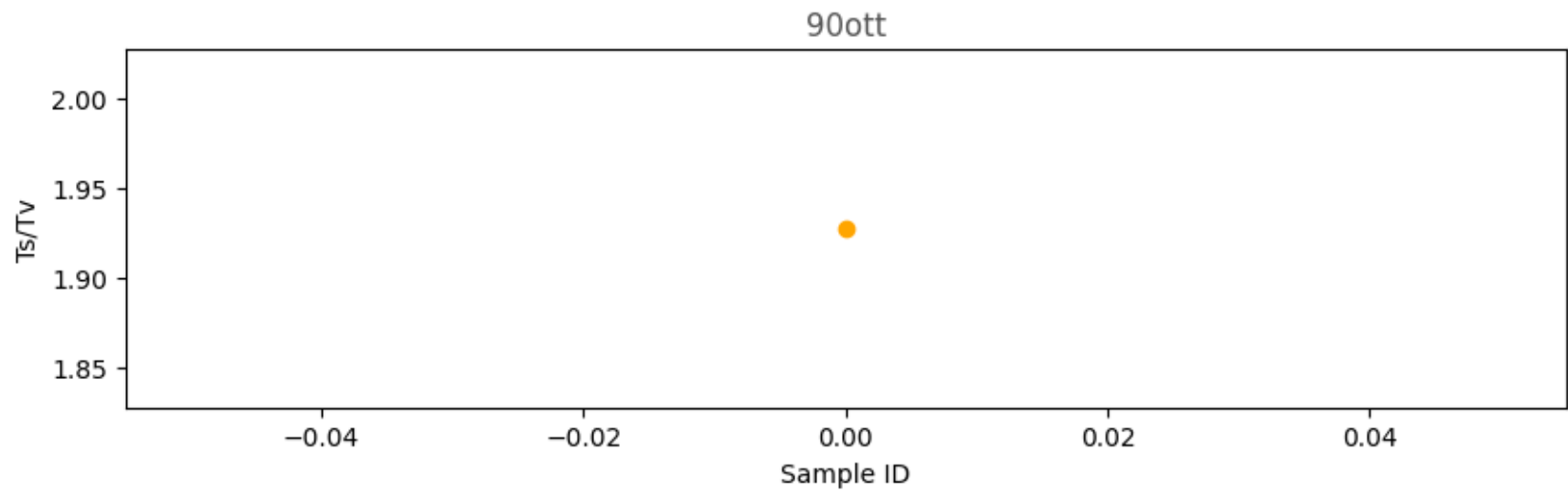
Hets vs non-ref Homs by sample
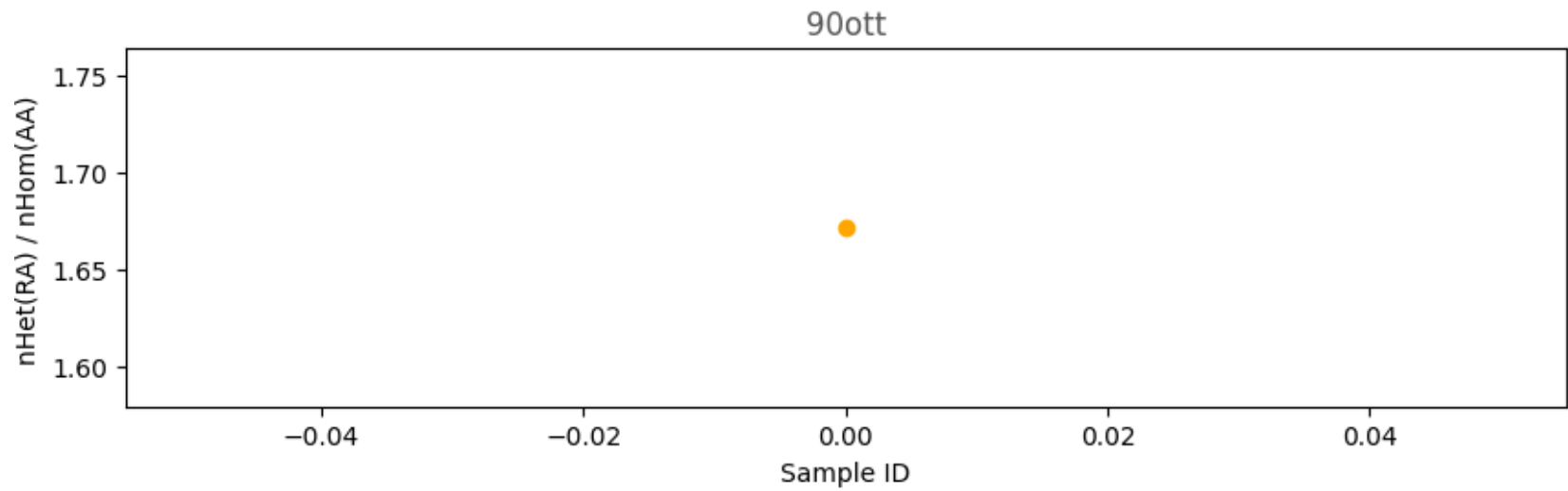
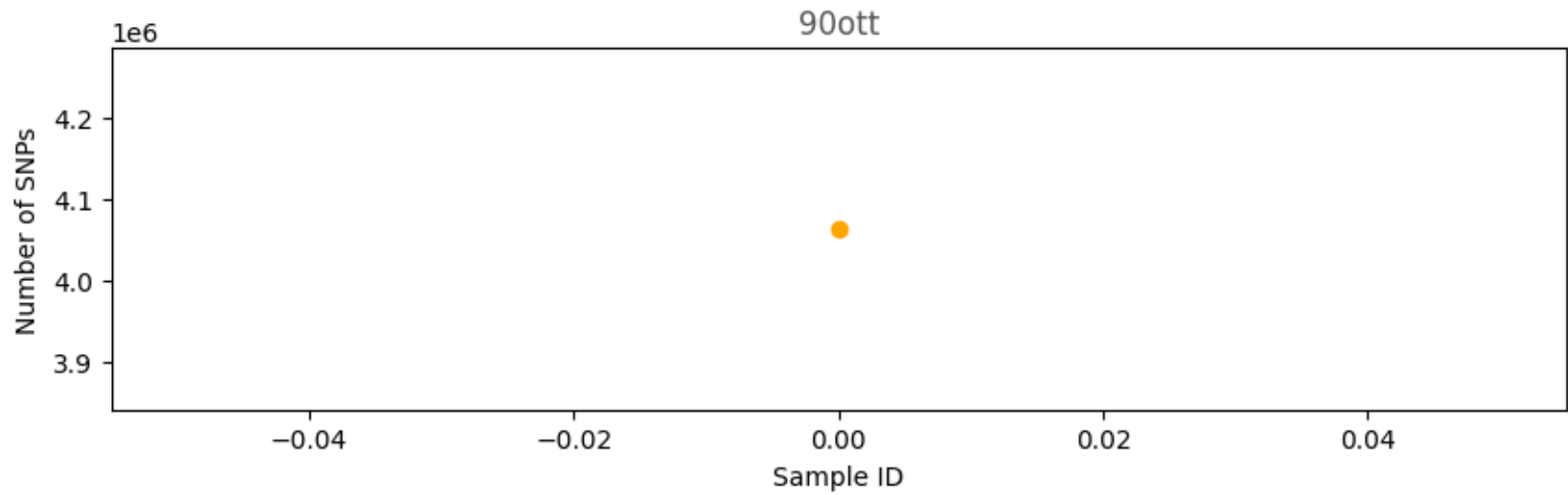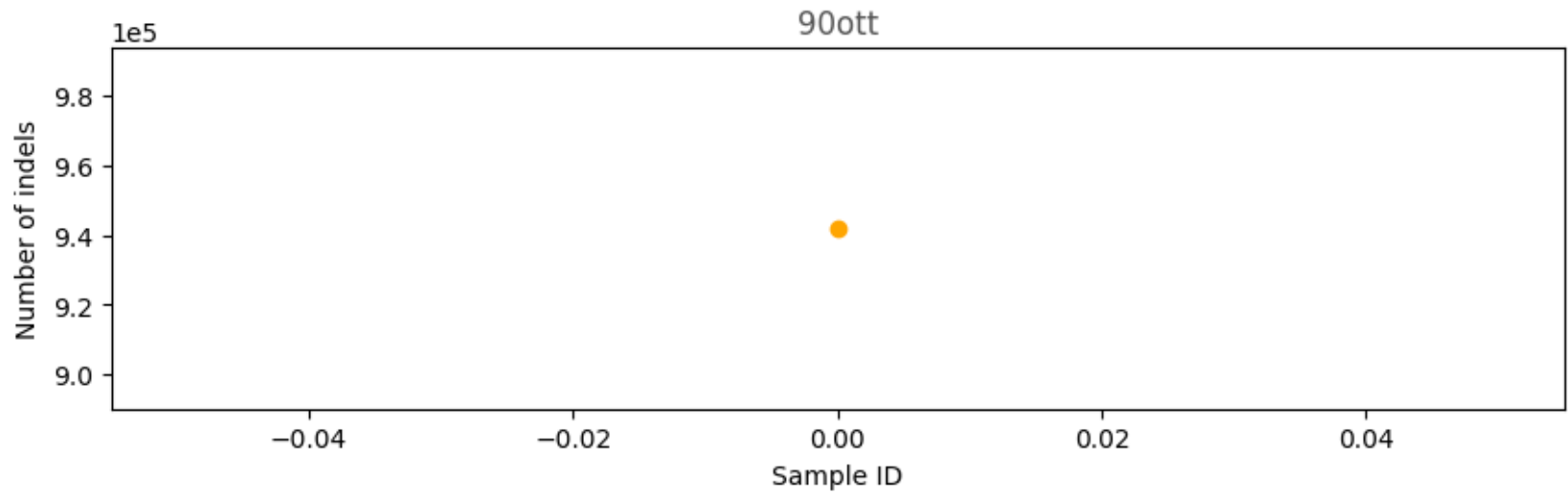# Singletons by sample (hets and homs)
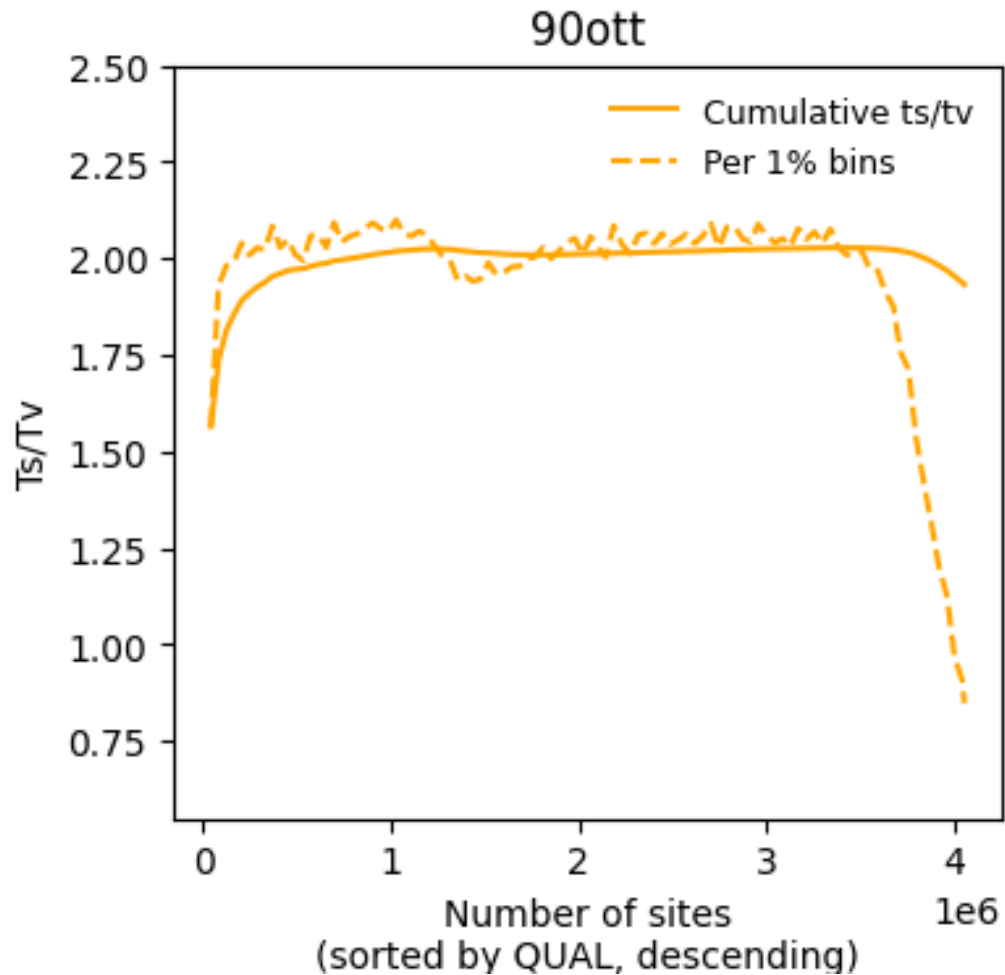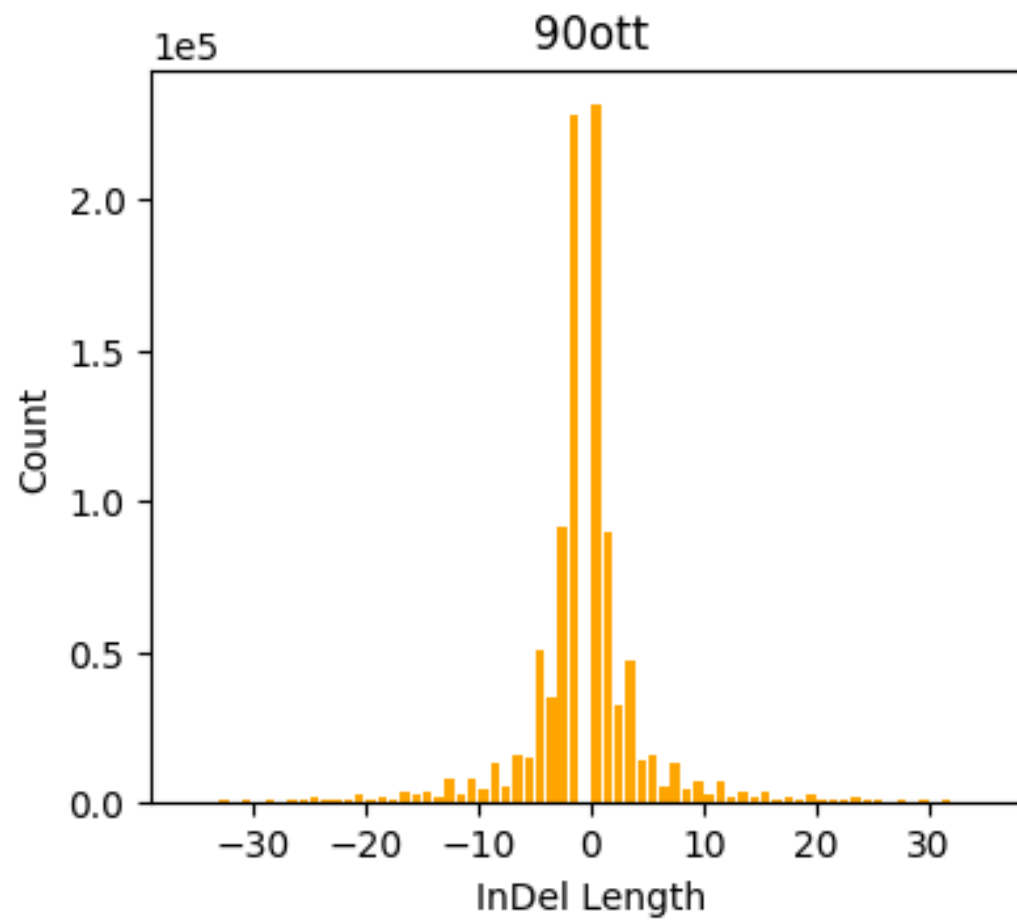
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

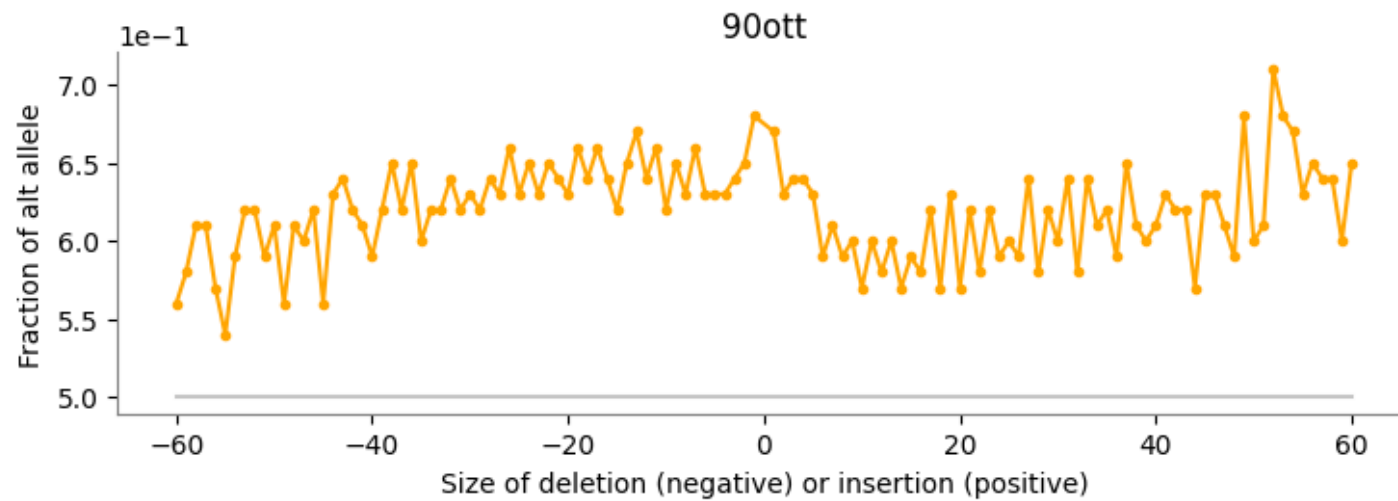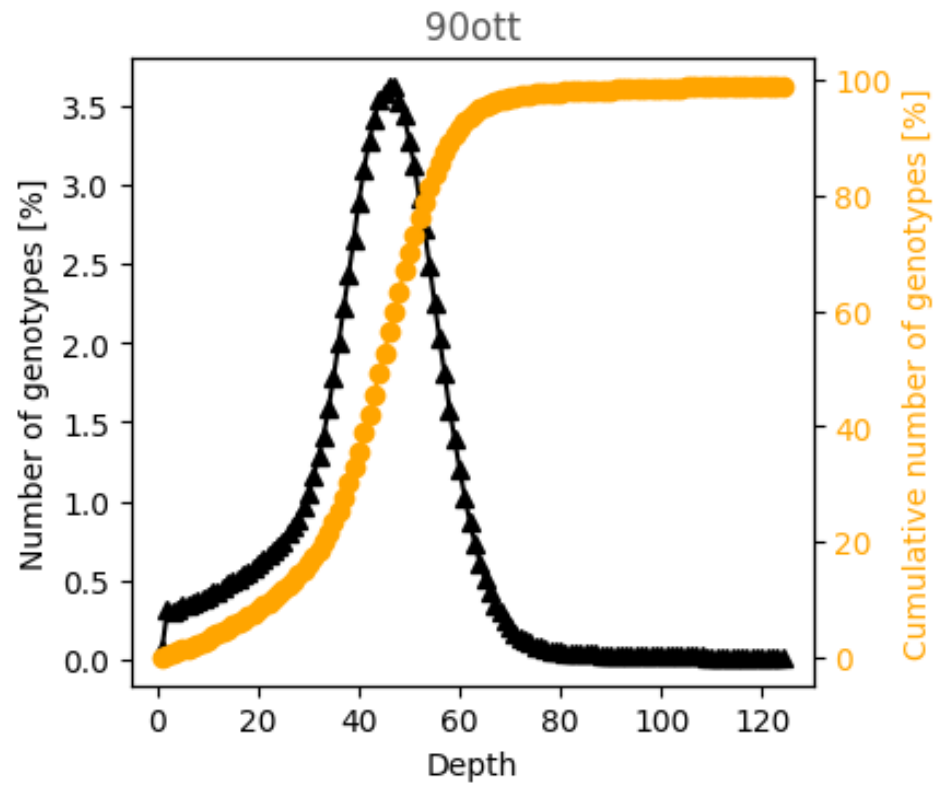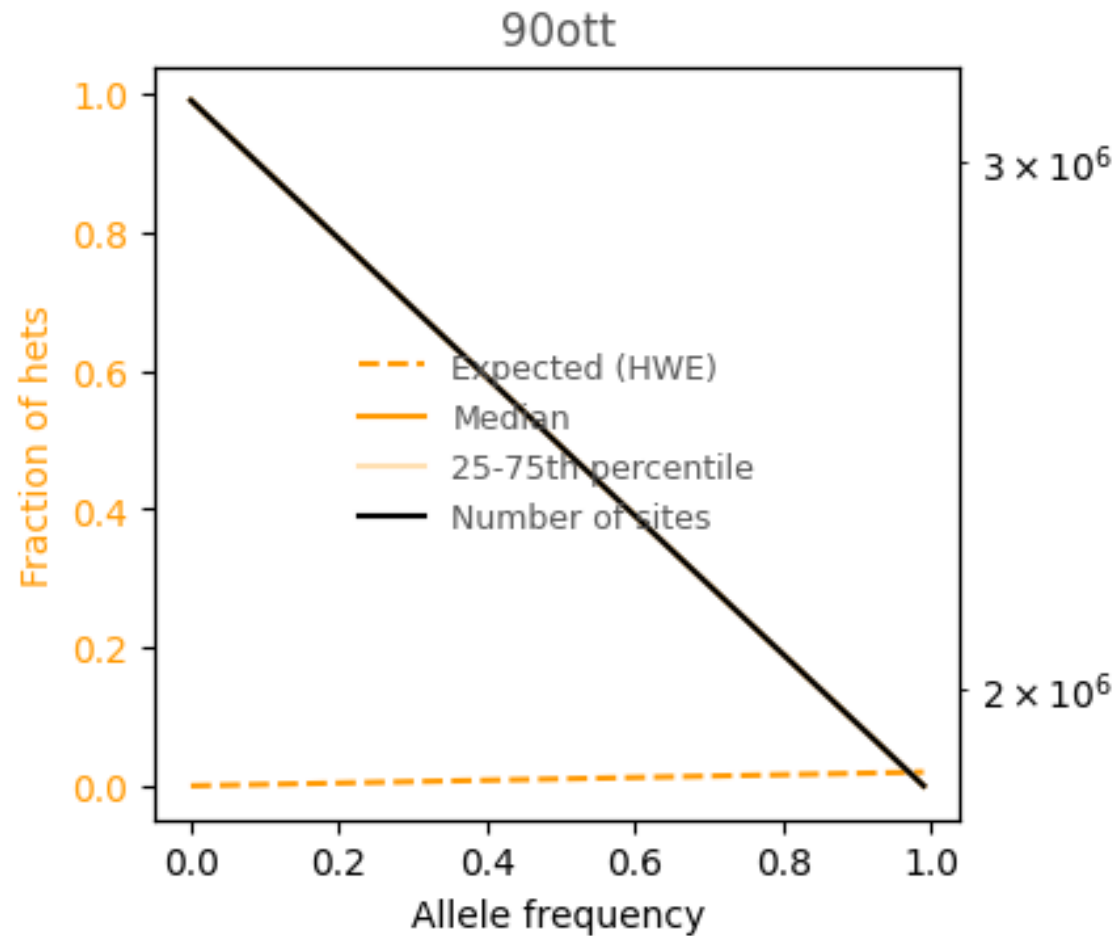# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types