Summary Numbers
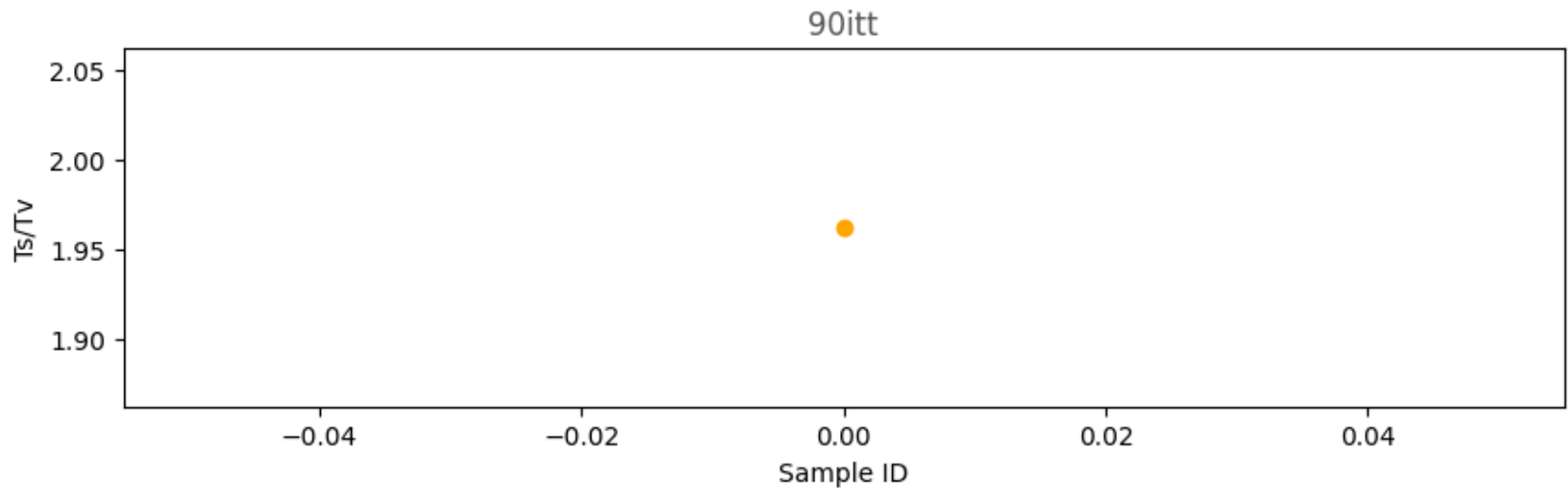
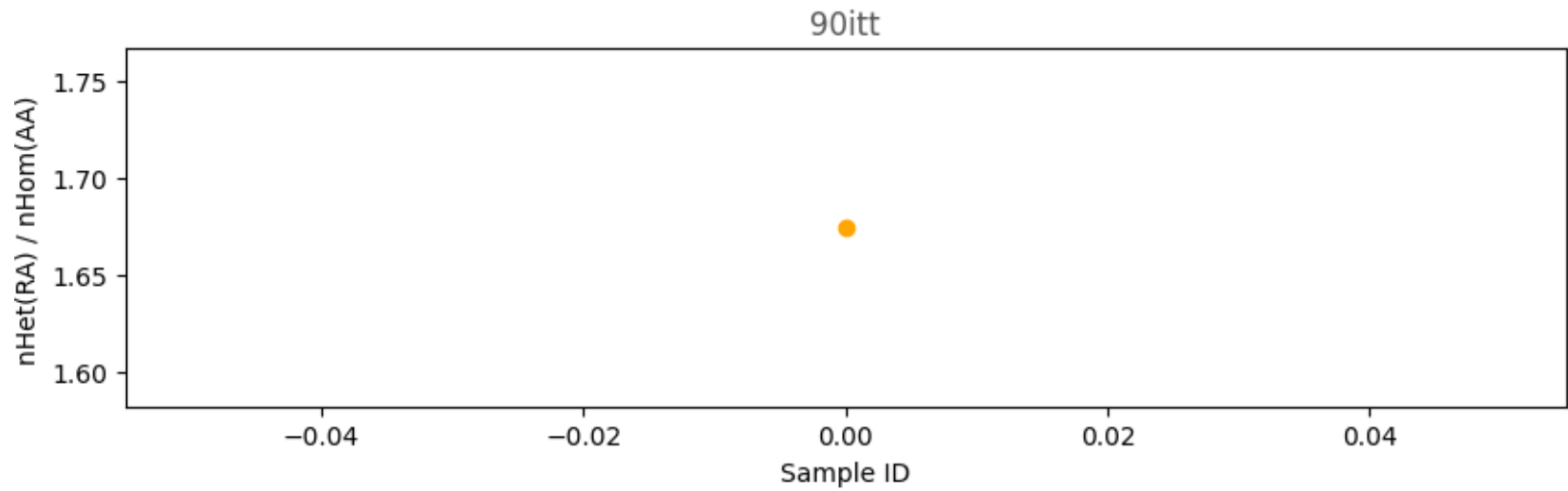| Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90itt	4,031,659	1.96	1.97	937,590	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90itt	62.6%	1.94	67.0%	92,466	1,907

- 90itt .. `/ngc/projects2/gm/data/archive/2022/variants/snv/90ittheef-103890968613-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220720_A01176_BHMWVVDSX3-RHGM_LABKA_WGS-` `WGS04690_snv_germline_raw.haplotype_caller.vcf.gz`
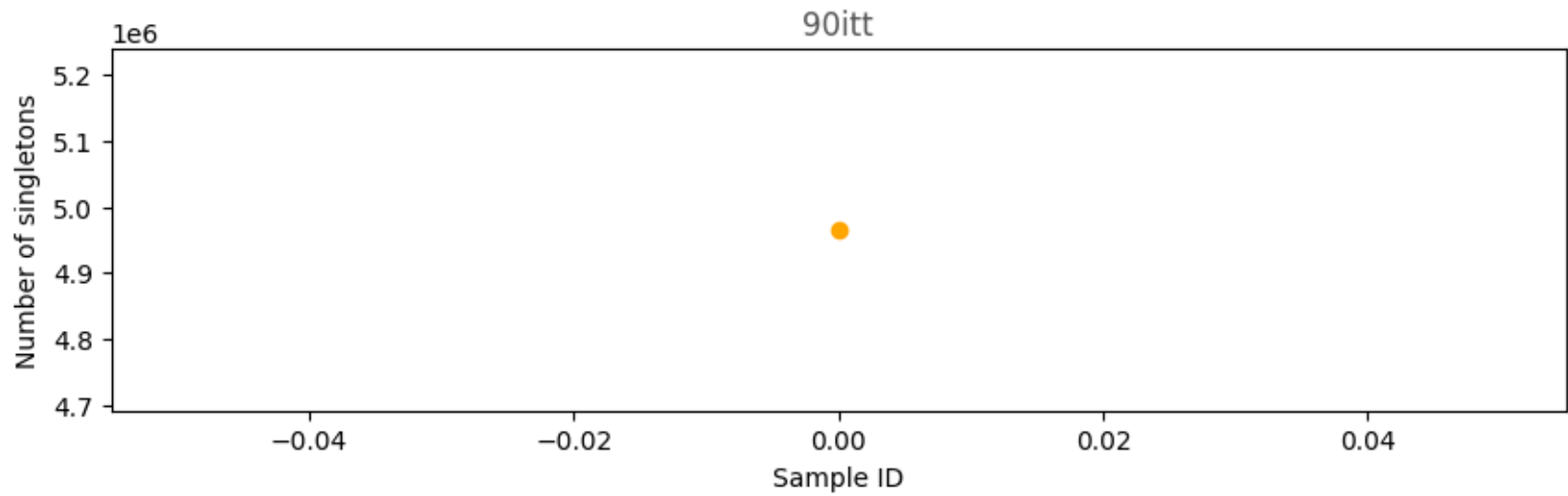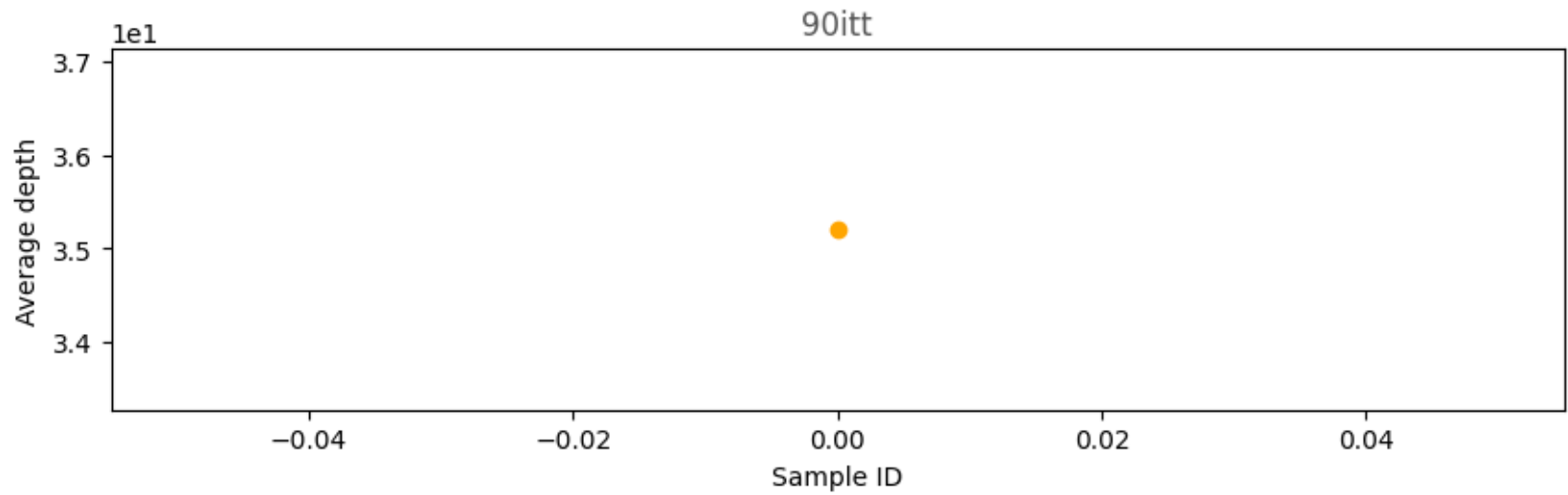
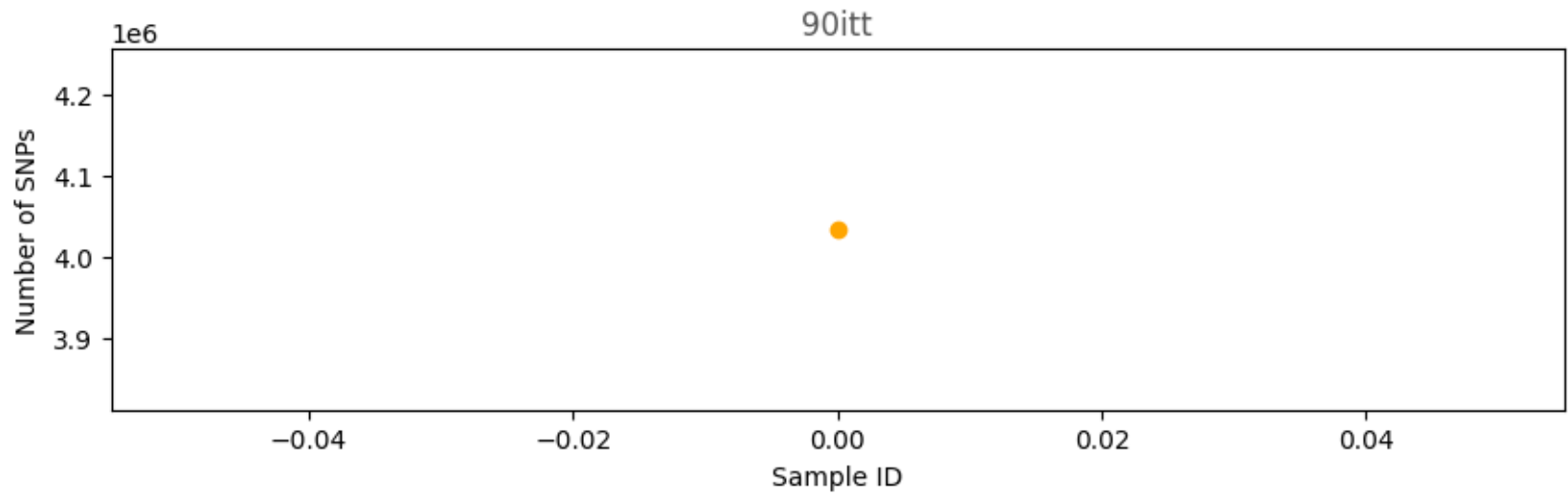# Ts/Tv by sample

Hets vs non-ref Homs by sample

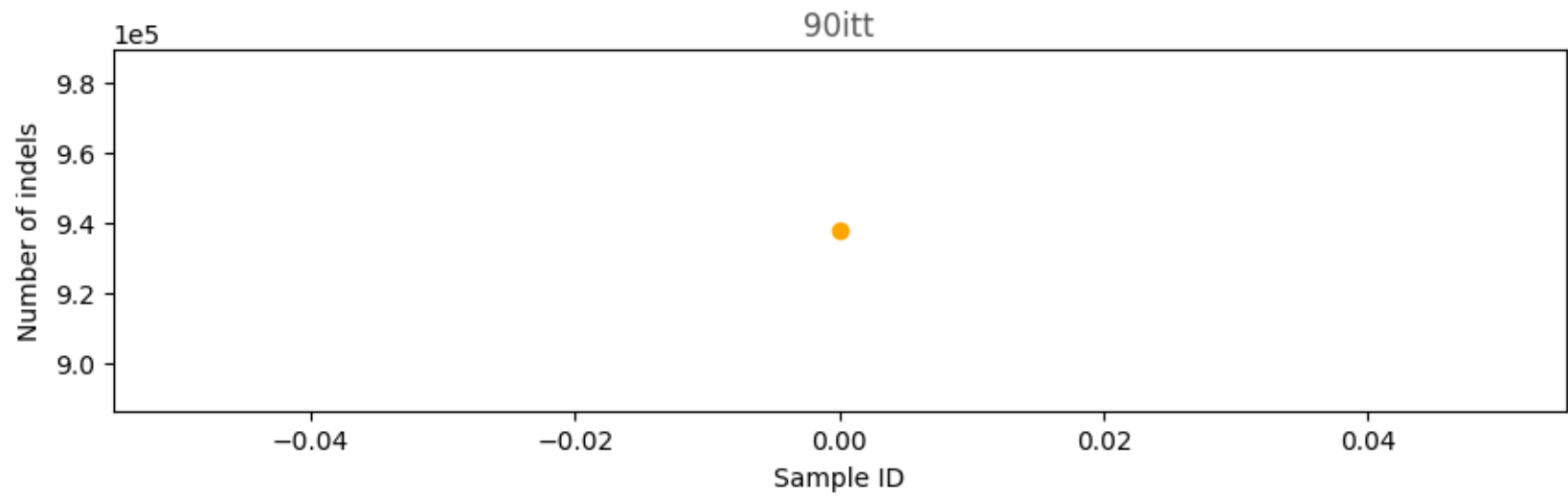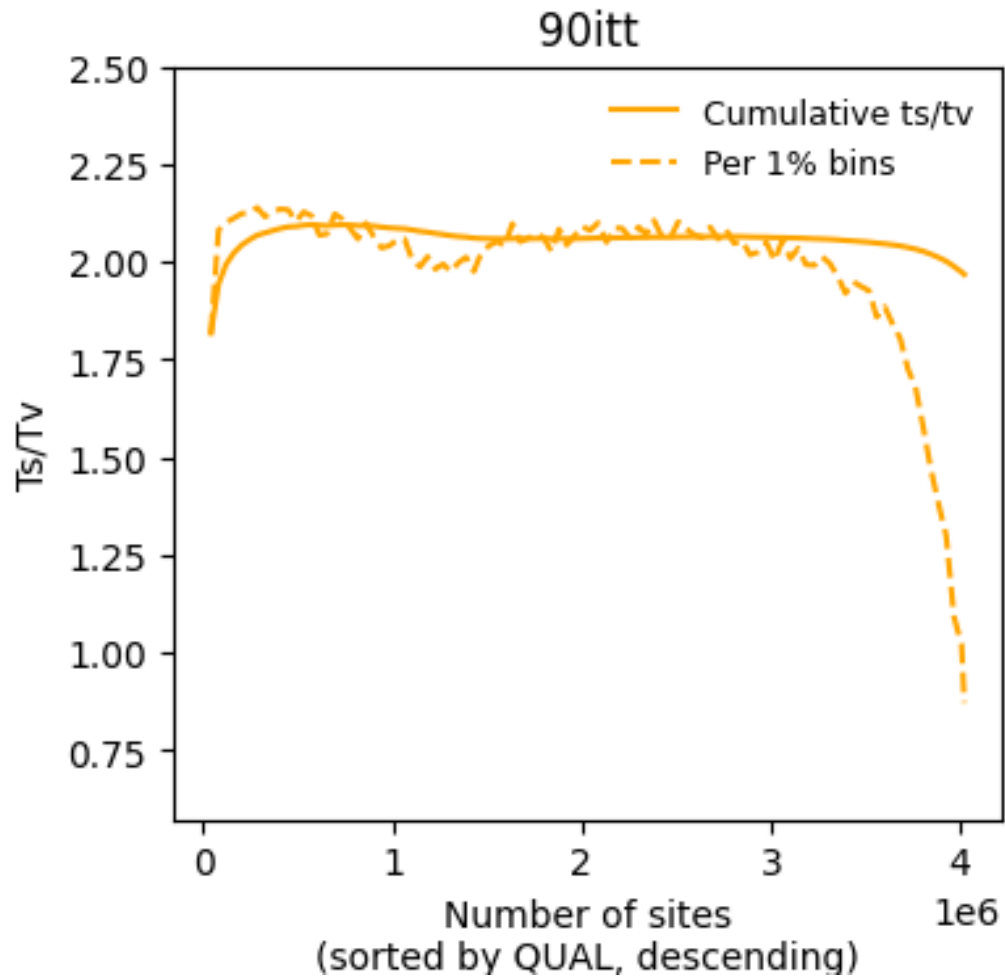# Singletons by sample (hets and homs)
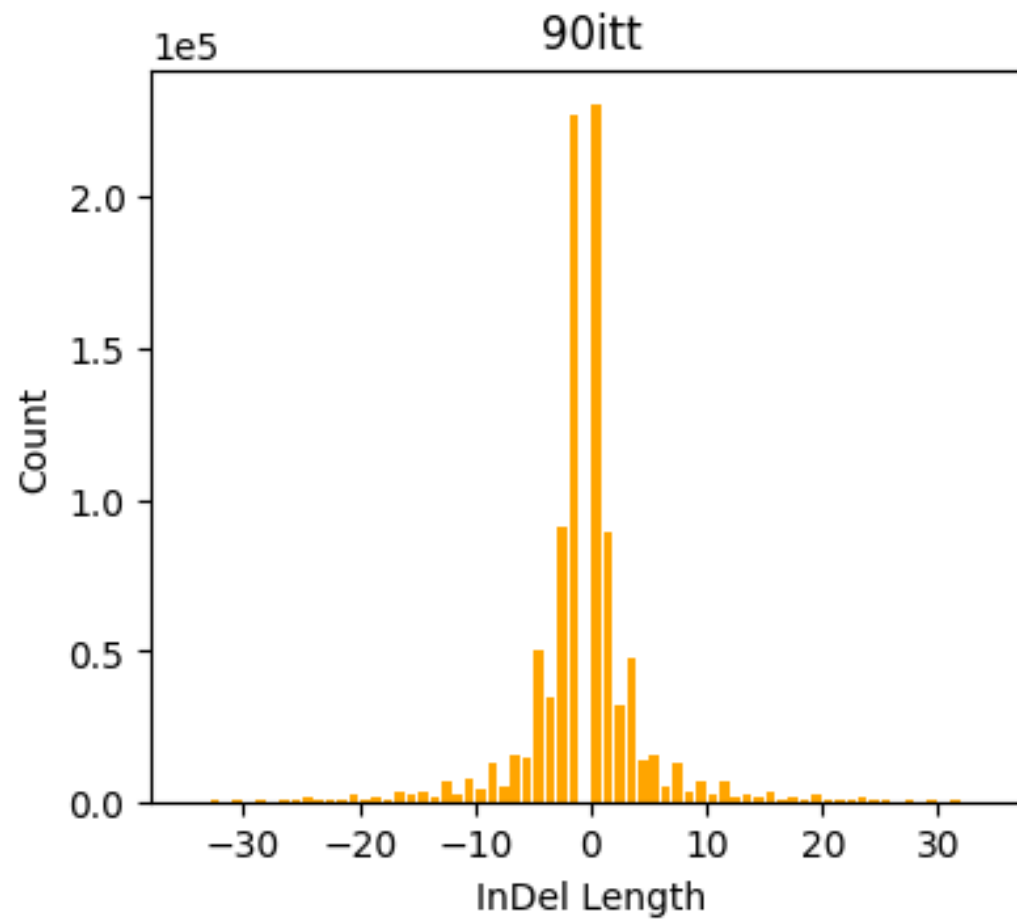
Average depth by sample

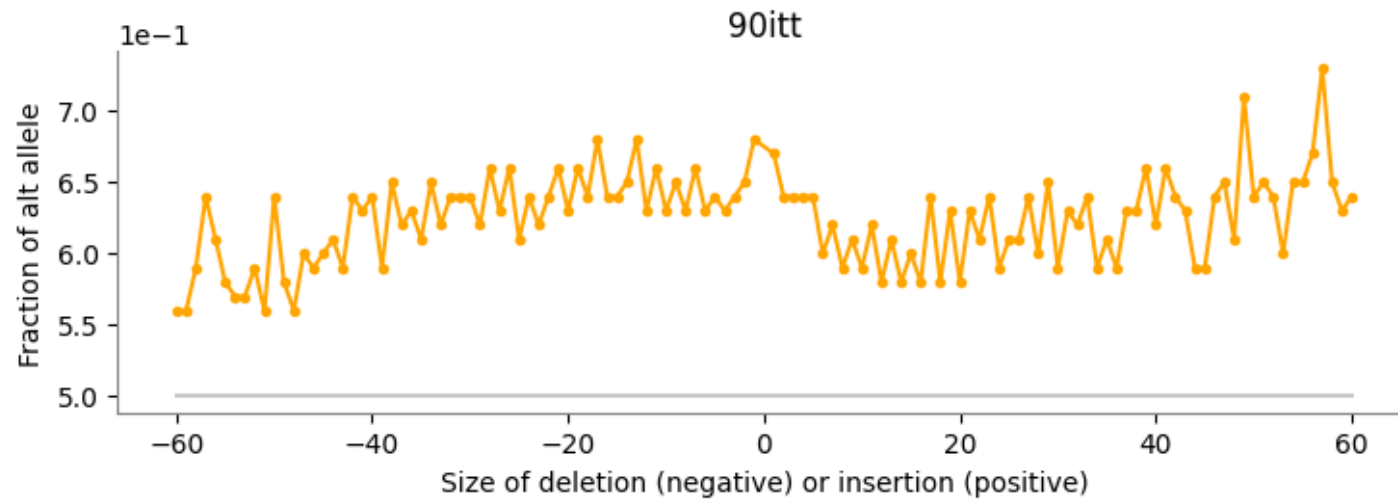# Number of SNPs by sample

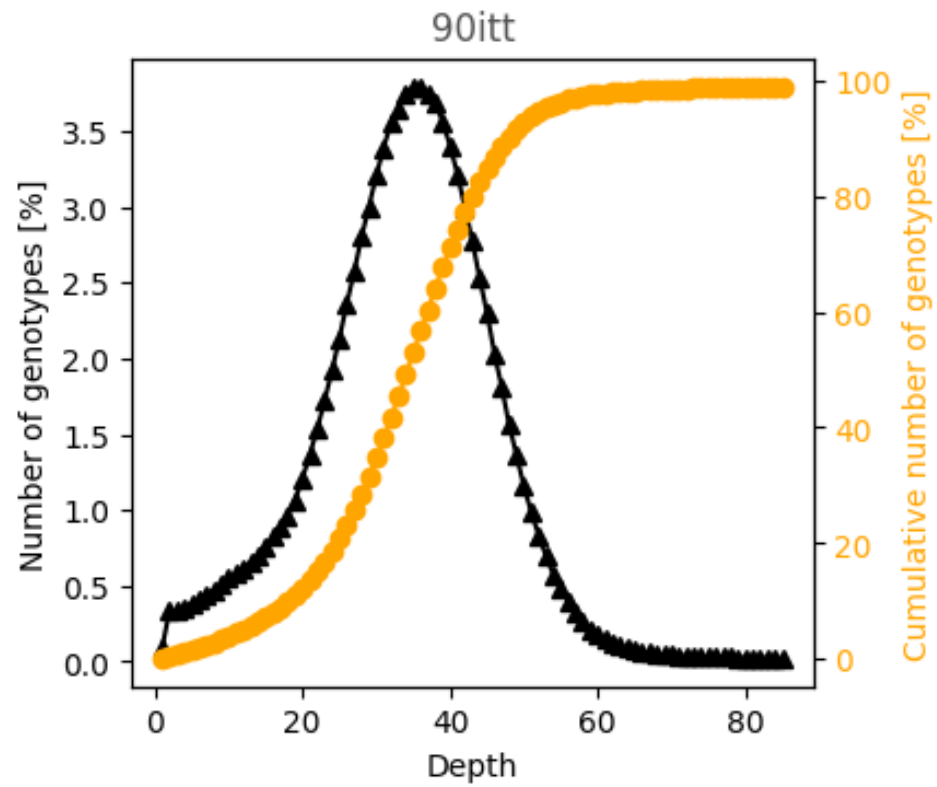# Number of indels by sample
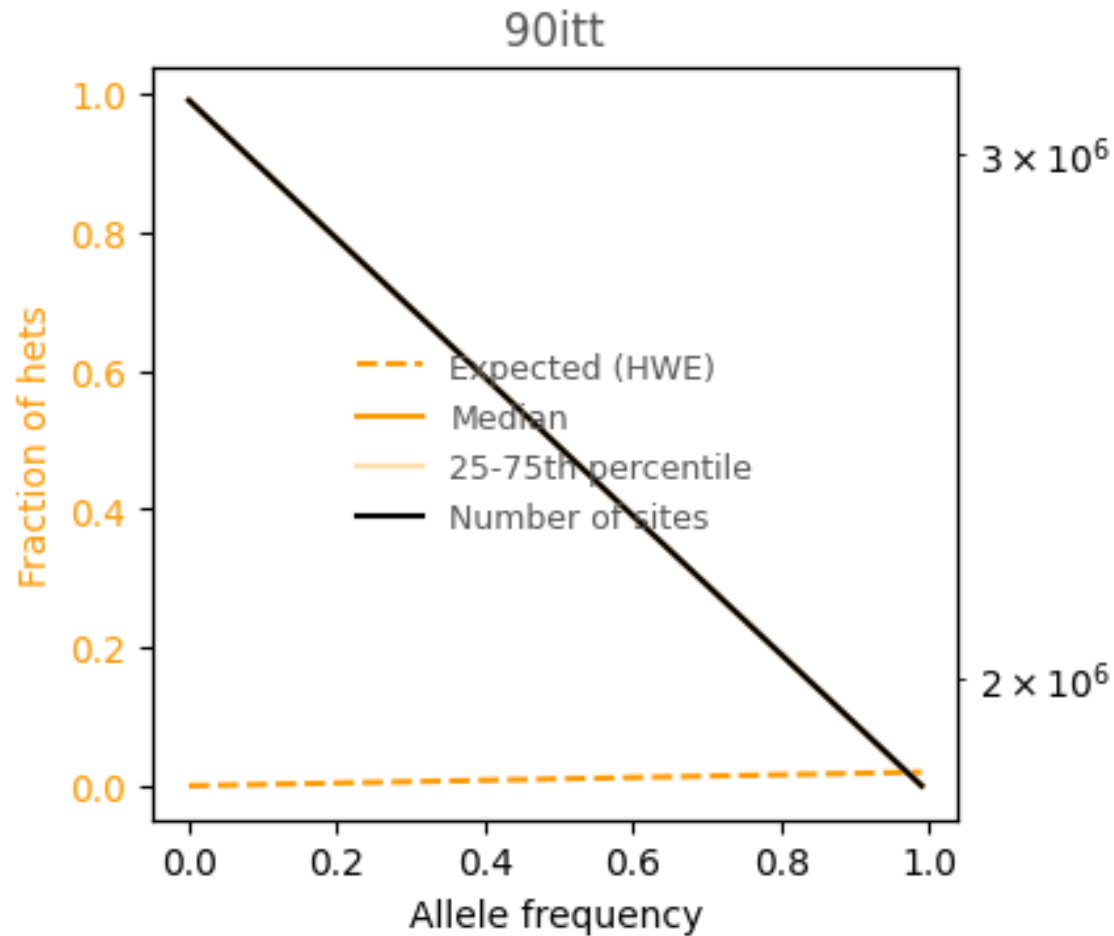
Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types

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