Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
89ind	4,052,189	1.93	1.94	931,820	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
89ind	61.9%	1.90	65.8%	92,623	2,005

- 89ind .. `/ngc/projects2/gm/data/archive/2022/variants/snv/89indhemm-103872894026-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00889-220615_A00559_AHJTYVDSX3-EXT_LAB`
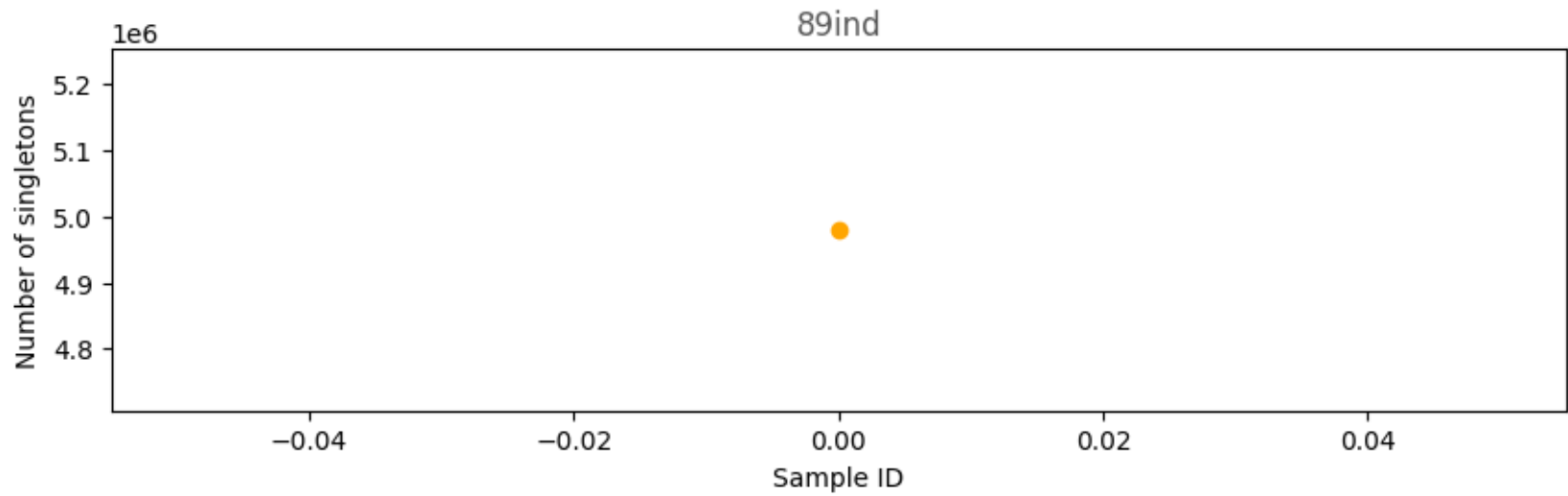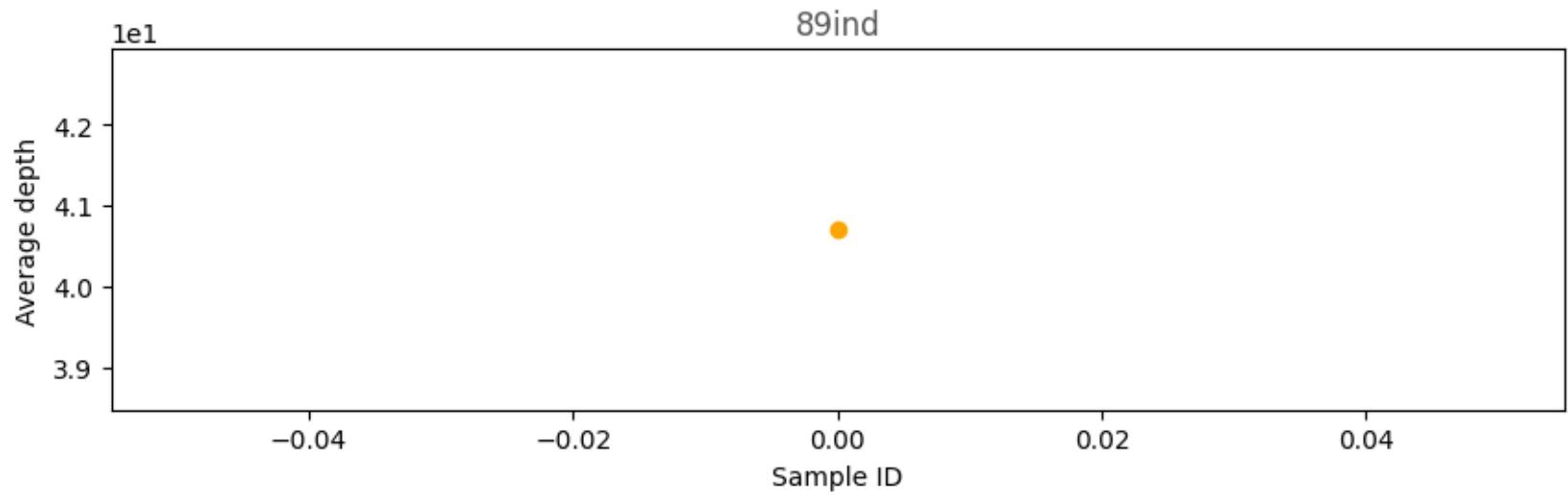 `KA_NGCWGS-NGCWGS04458_snv_germline_raw.haplotype_caller.vcf.gz`
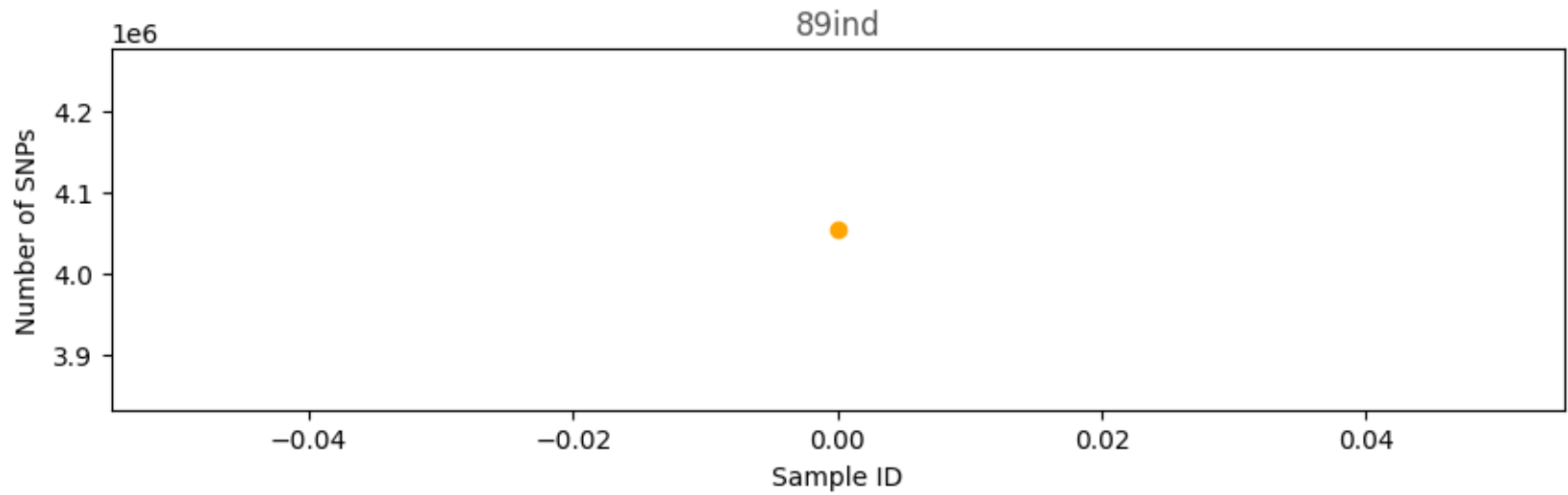
Ts/Tv by sample
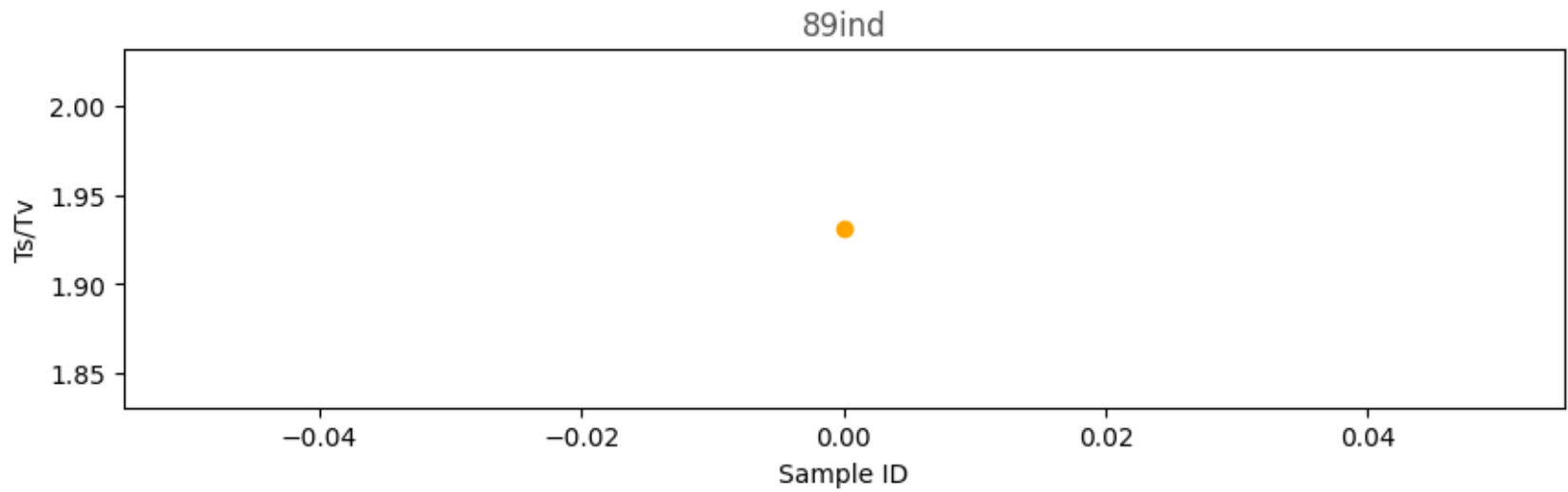
Hets vs non-ref Homs by sample
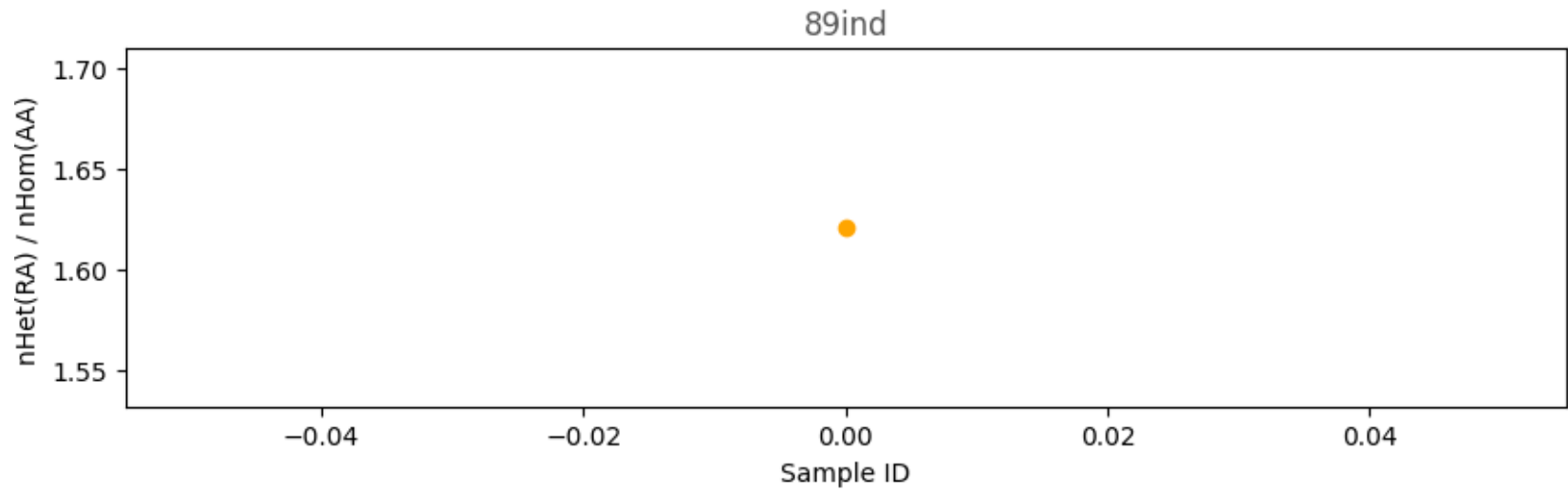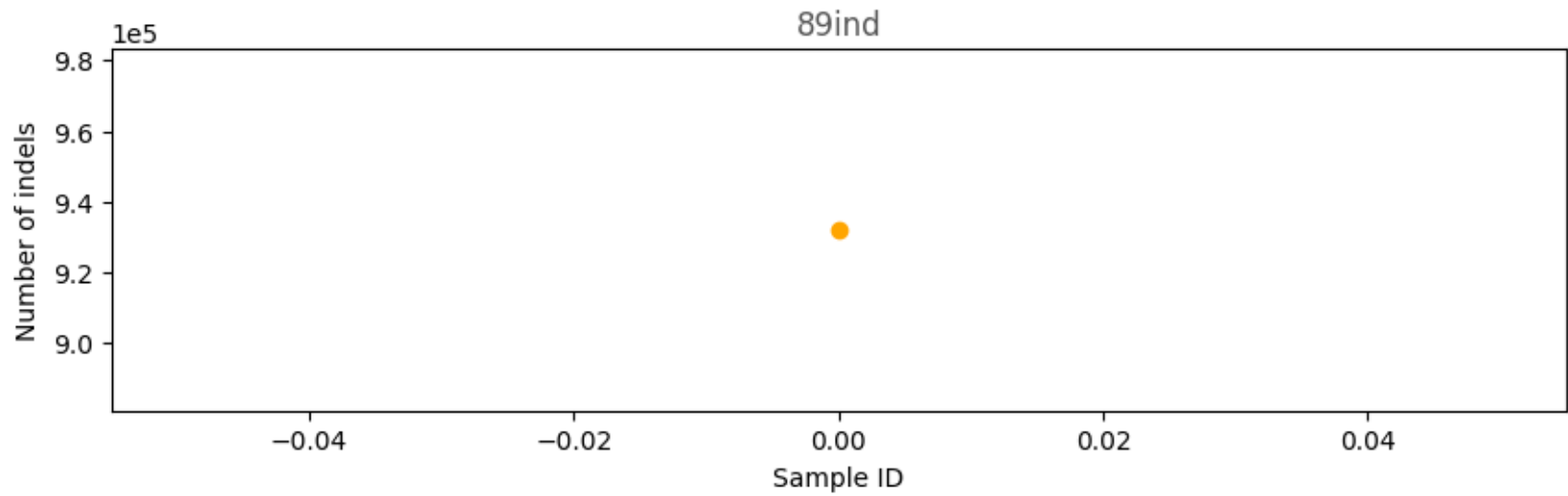
Singletons by sample (hets and homs)
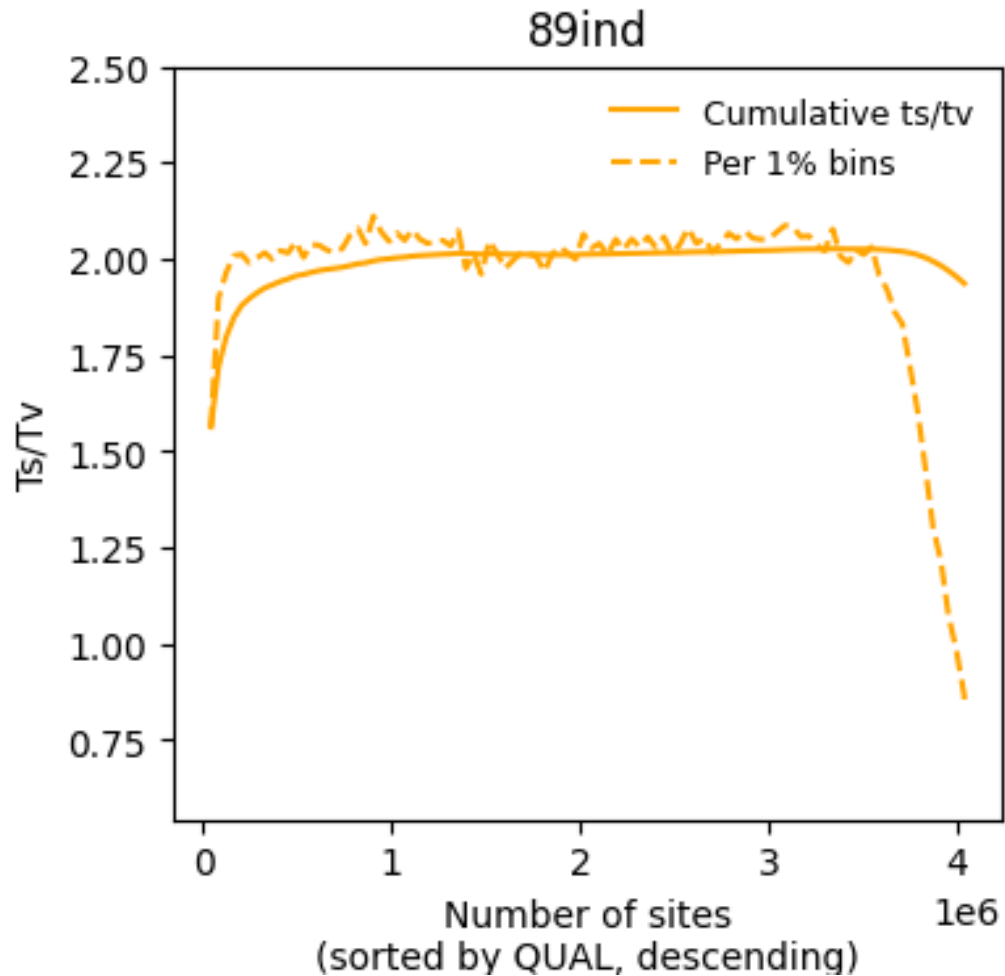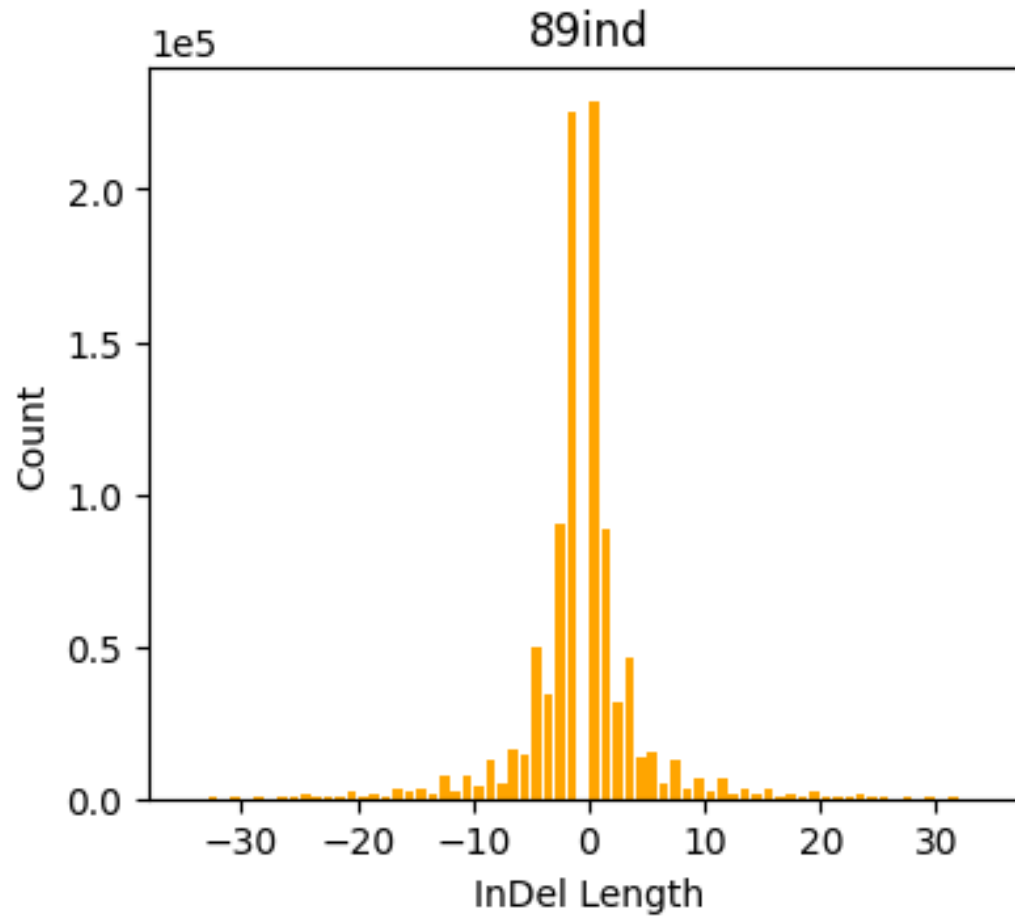
Average depth by sample
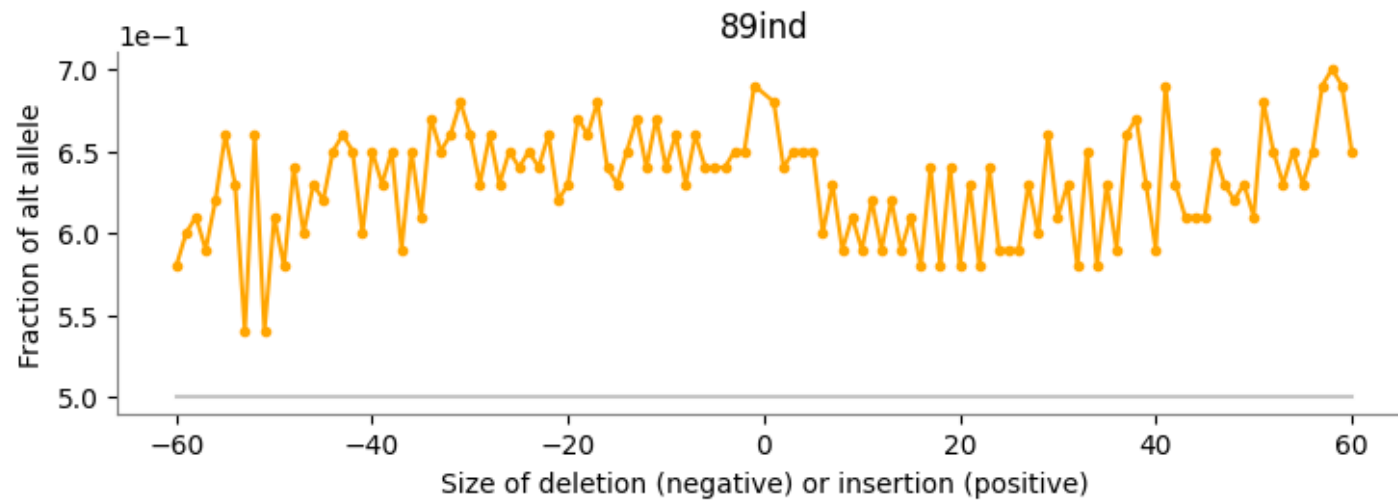
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

89ind

Indel distribution

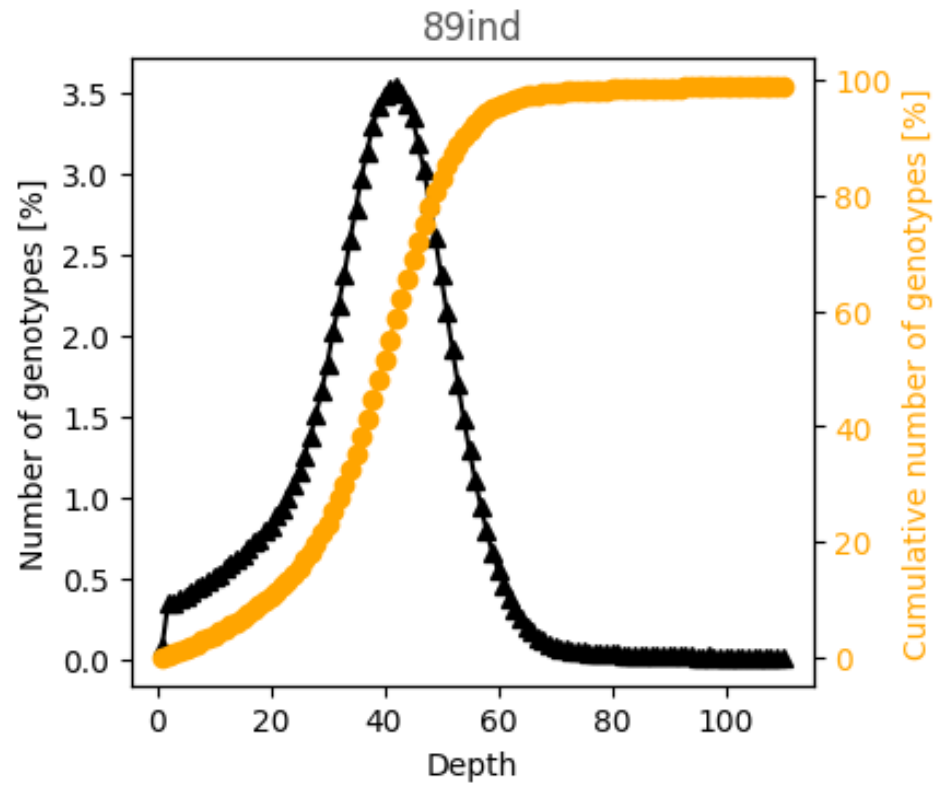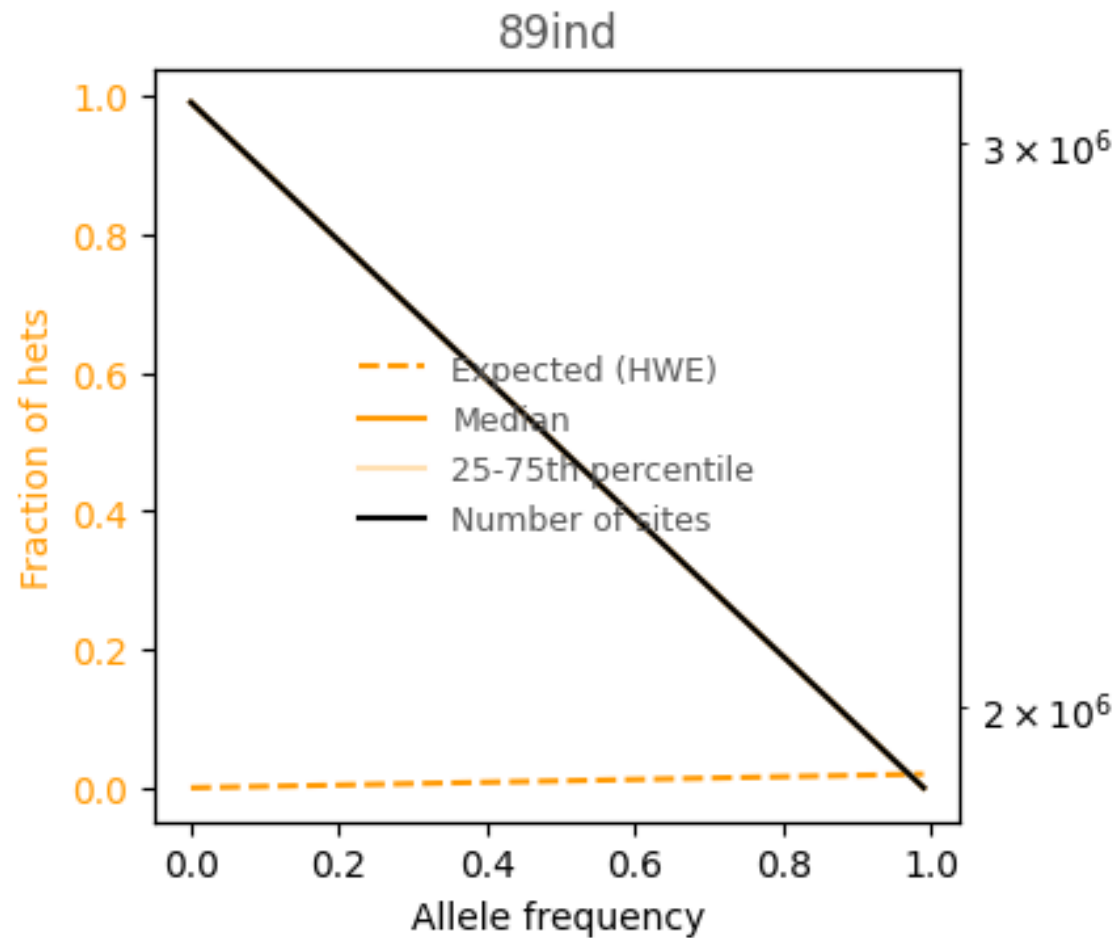# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types