Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
91ved	4,131,309	1.92	1.92	961,214	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
91ved	63.6%	1.88	68.7%	104,509	2,086

- 91ved .. `/ngc/projects2/gm/data/archive/2022/variants/snv/91vedffrf-103857992497-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00717-220413_A00559_AHGTFCDSX3-EXT_LAB`
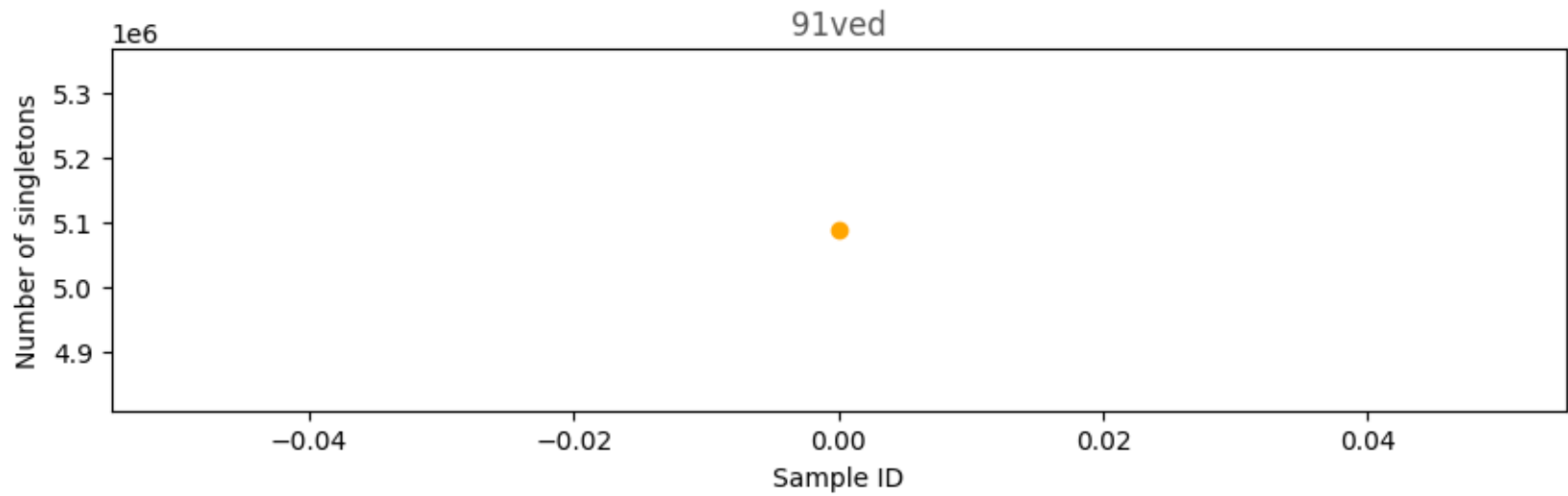 `KA_NGCWGS-NGCWGS04132_snv_germline_raw.haplotype_caller.vcf.gz`
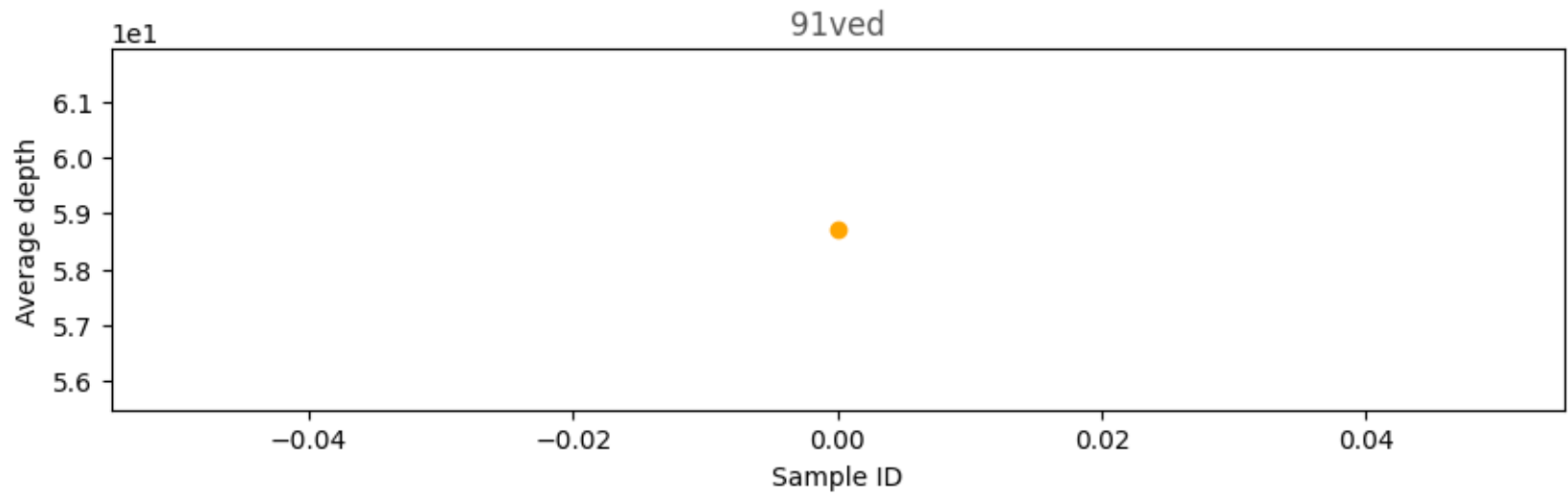
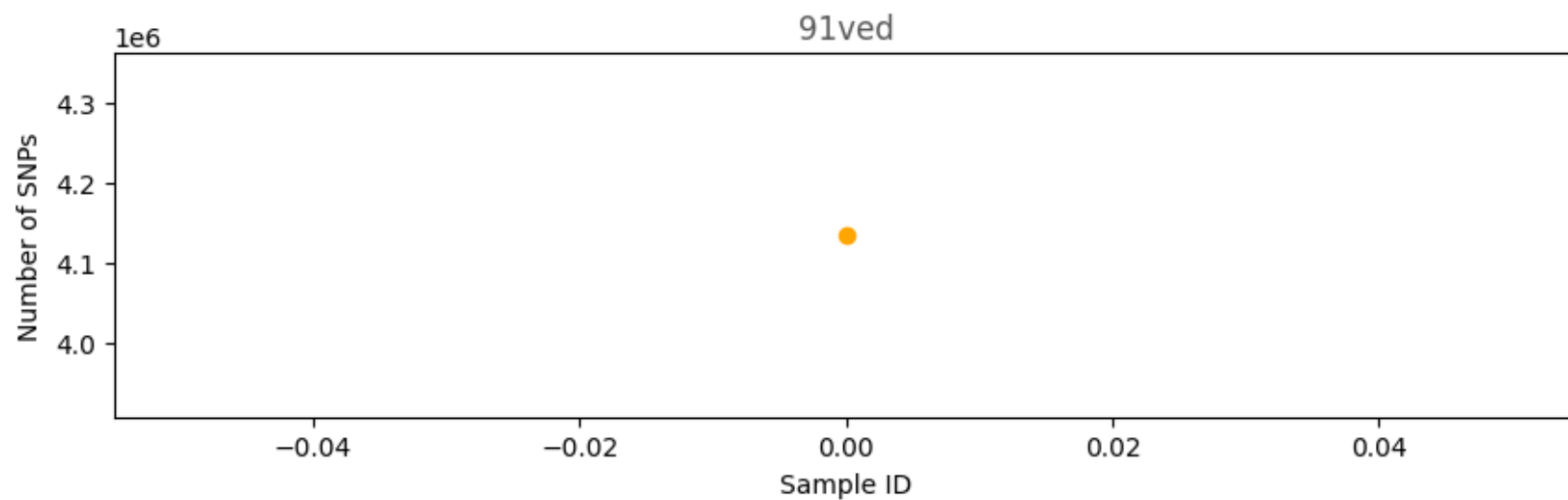# Ts/Tv by sample

91ved
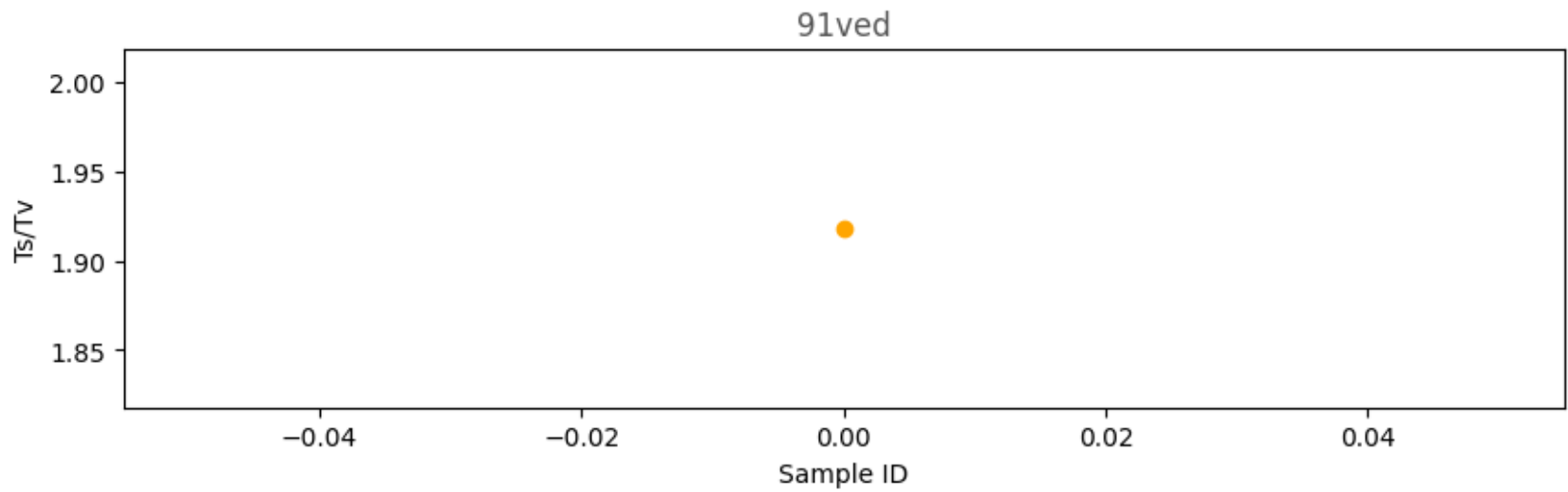
Hets vs non-ref Homs by sample
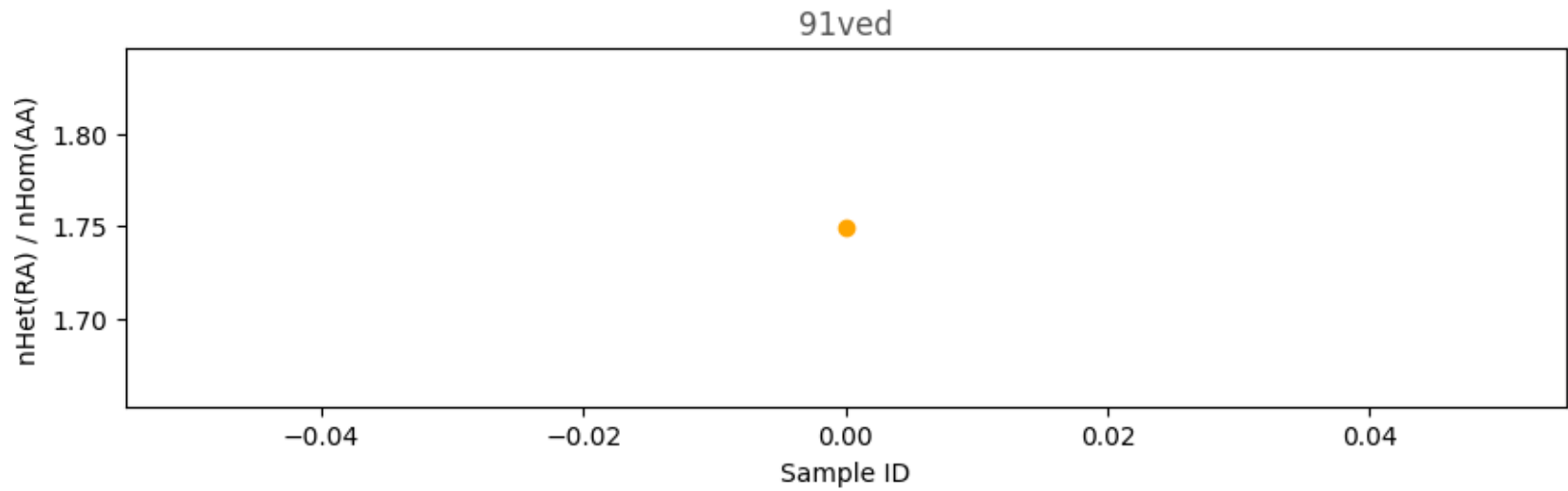
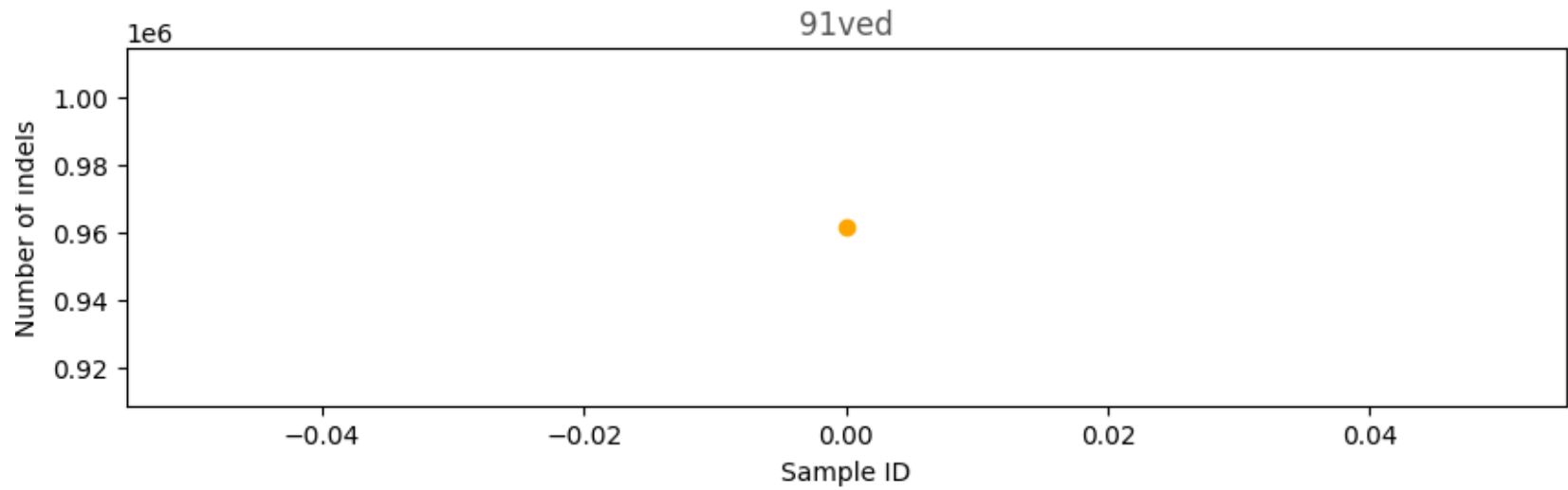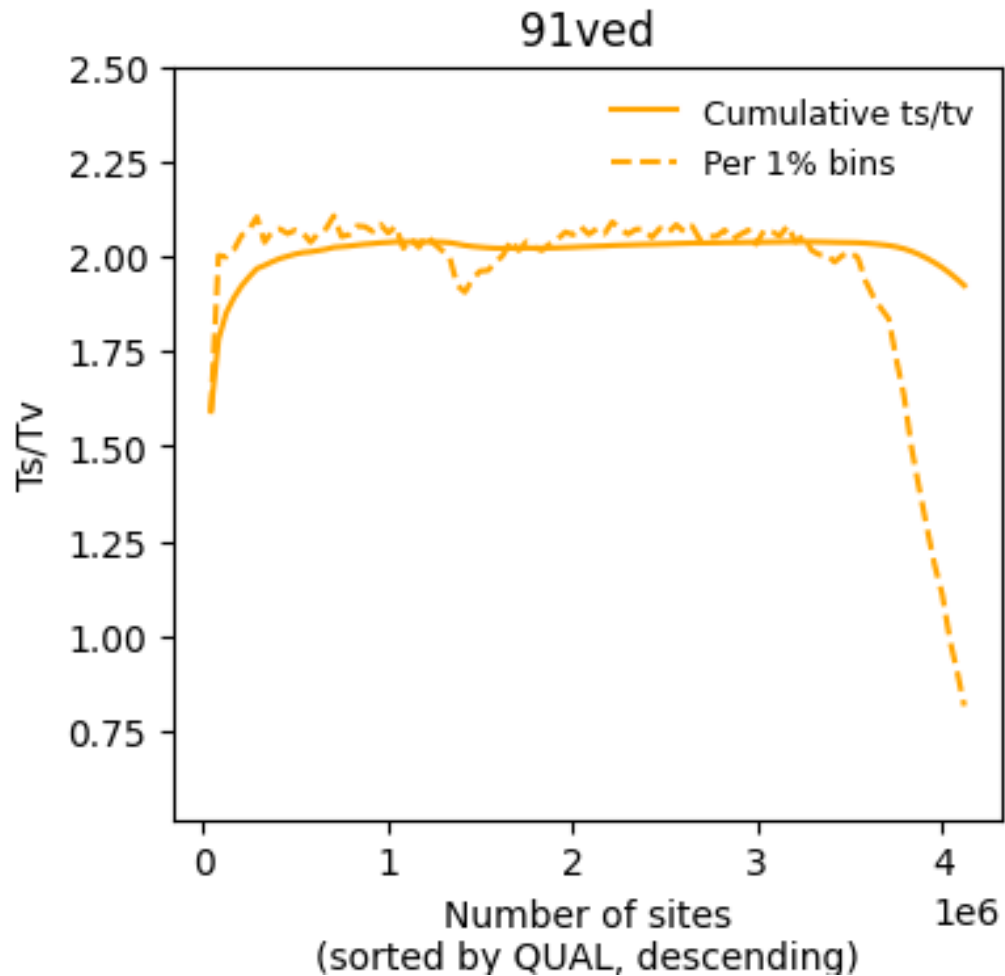# Singletons by sample (hets and homs)
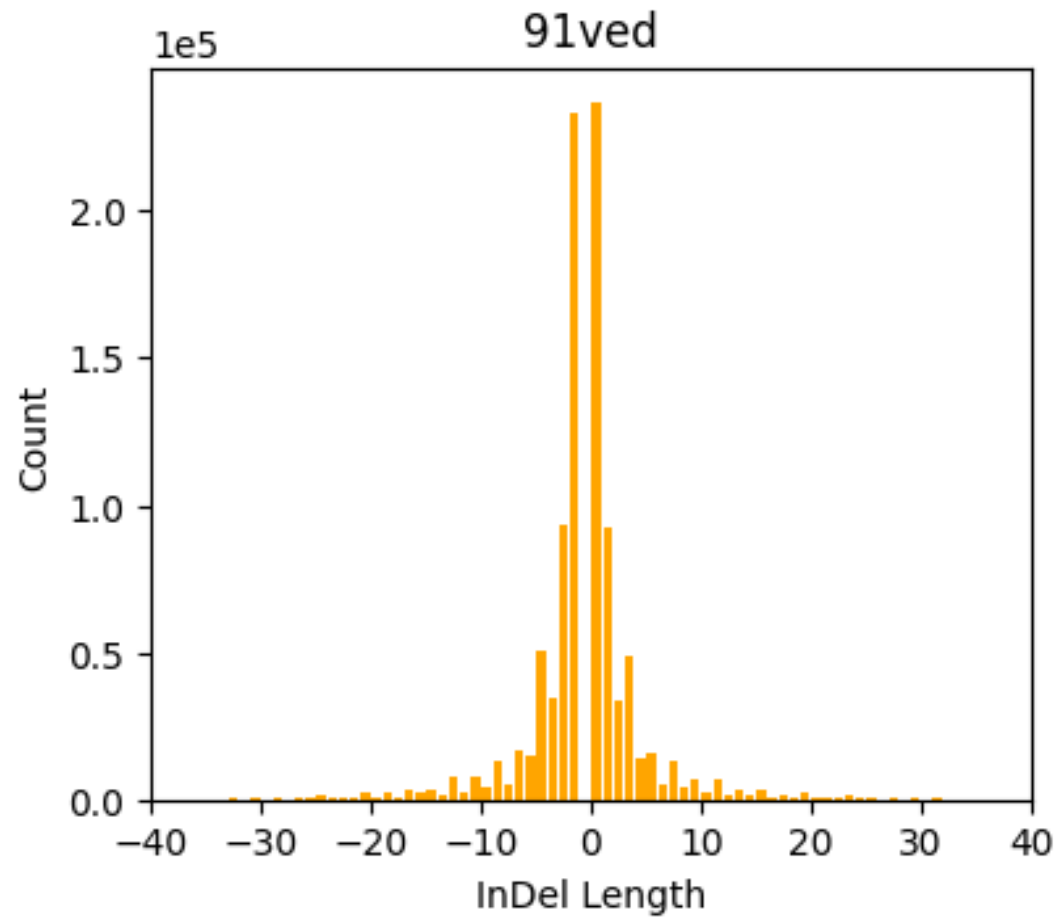
Average depth by sample
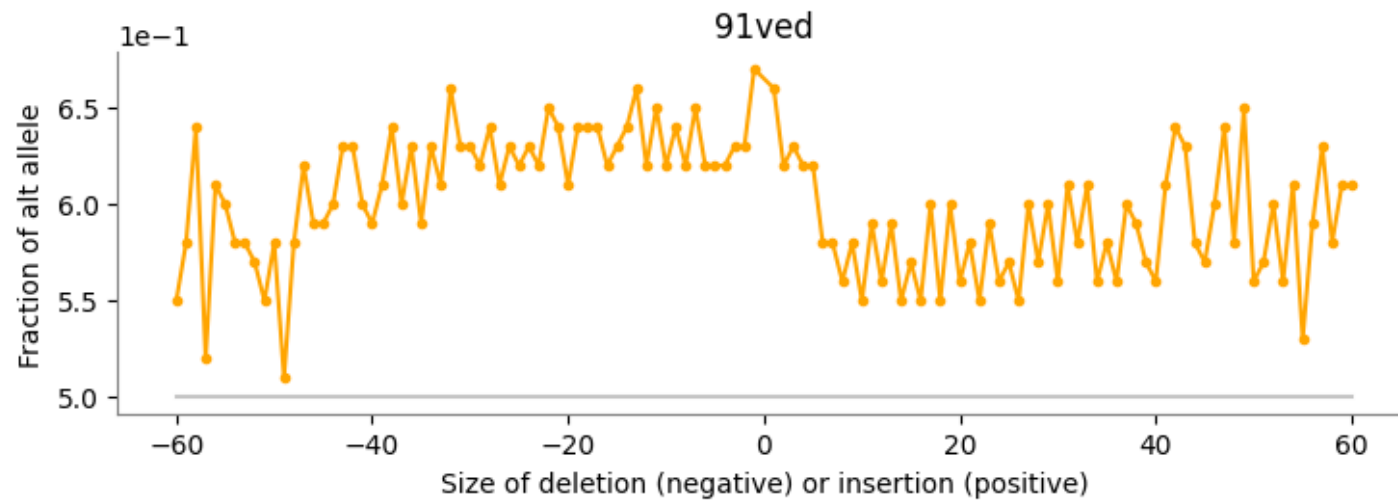
Number of SNPs by sample

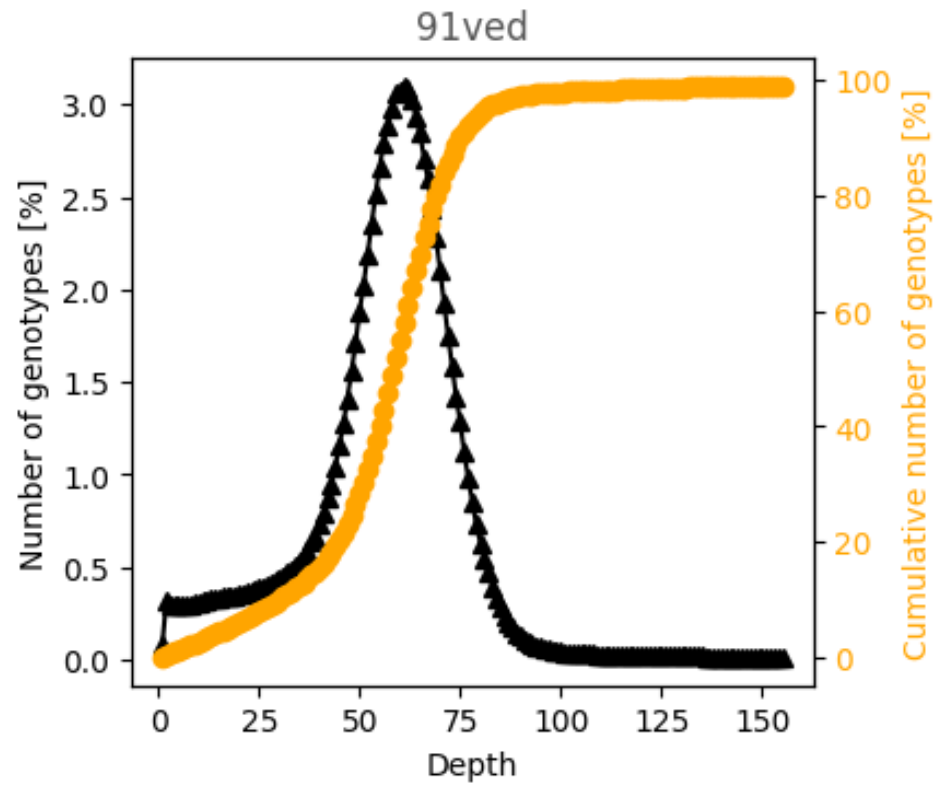# Number of indels by sample
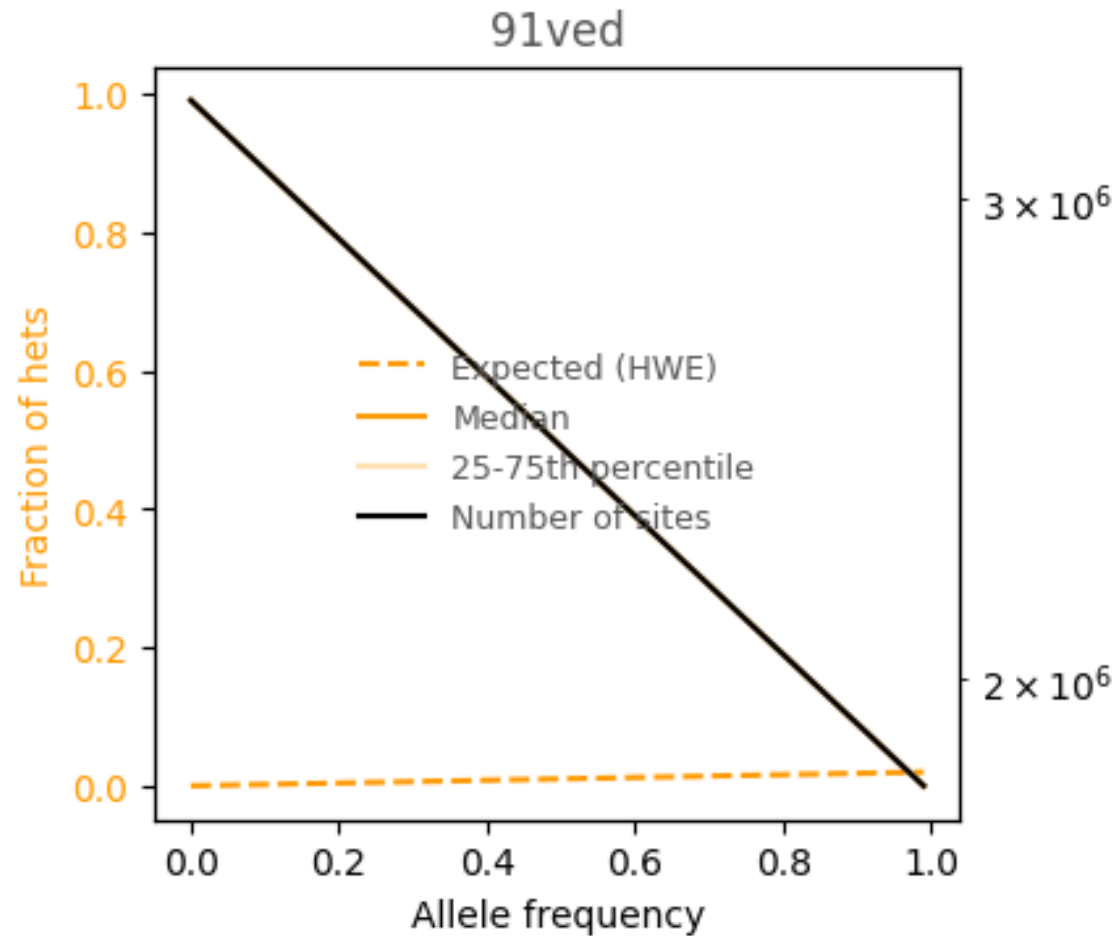
Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types

91ved