Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76wia	4,107,723	1.93	1.93	957,877	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76wia	63.6%	1.90	68.7%	103,768	2,077

- 76wia .. `/ngc/projects2/gm/data/archive/2022/variants/snv/76wiajuif-103817926370-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00461-211222_A01176_AHT7KCDSX2-EXT_LAB`
  `KA_NGCWGS-NGCWGS03558_21RKG028419x01_snv_germline_raw.haplotype_caller.vcf.gz`
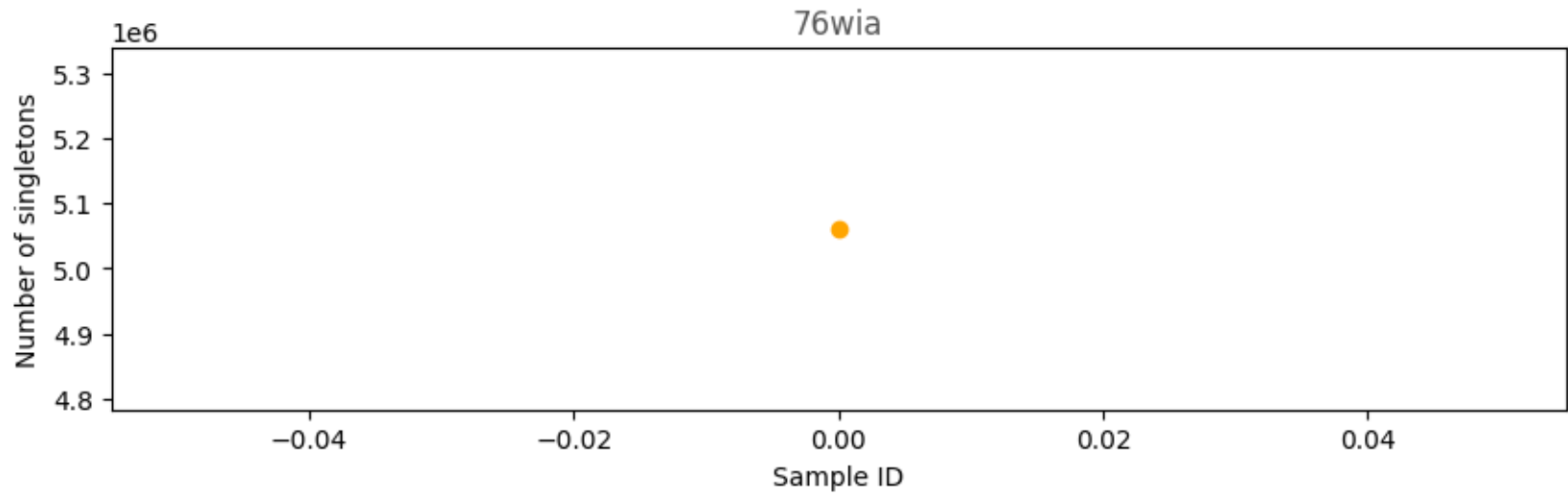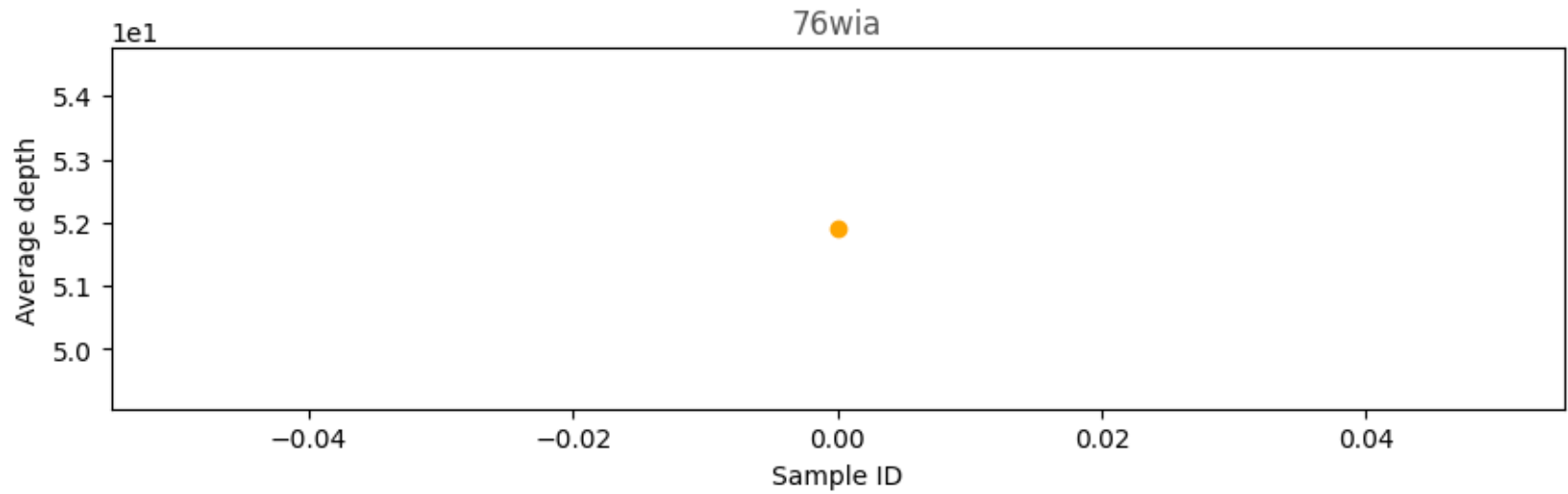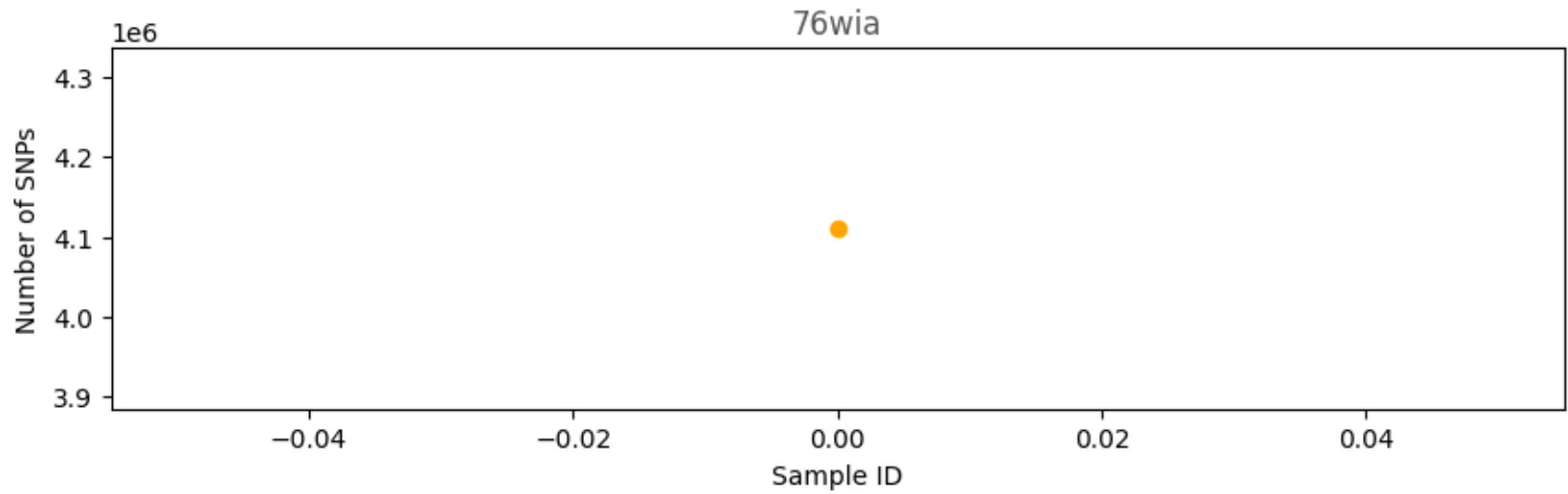
Ts/Tv by sample
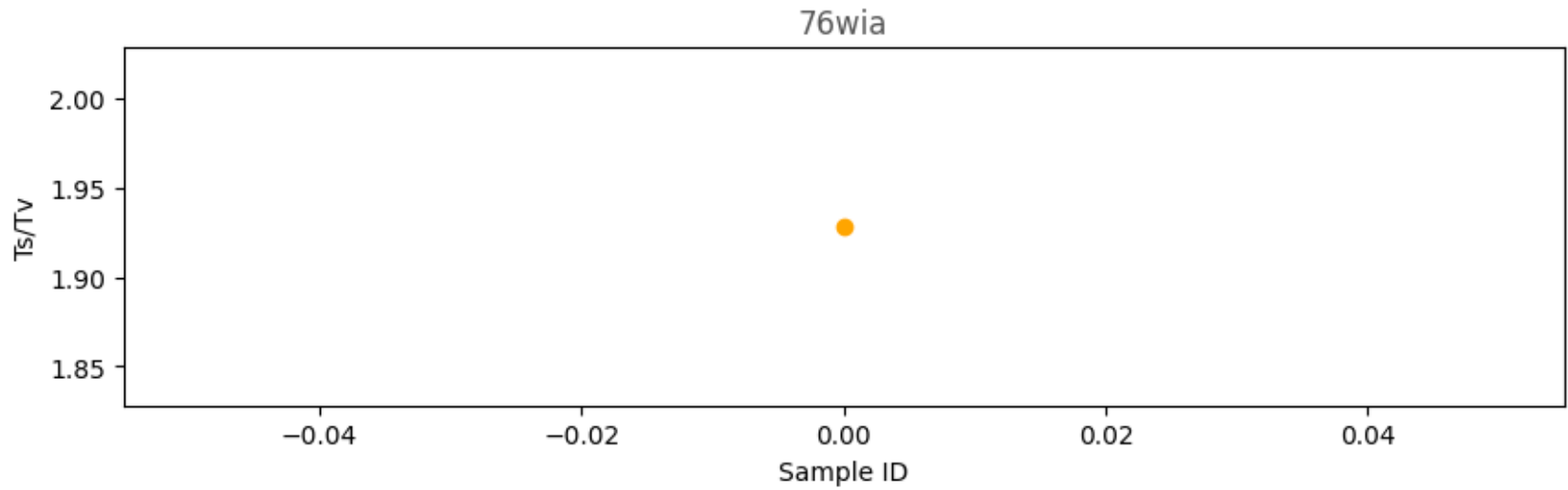
Hets vs non-ref Homs by sample
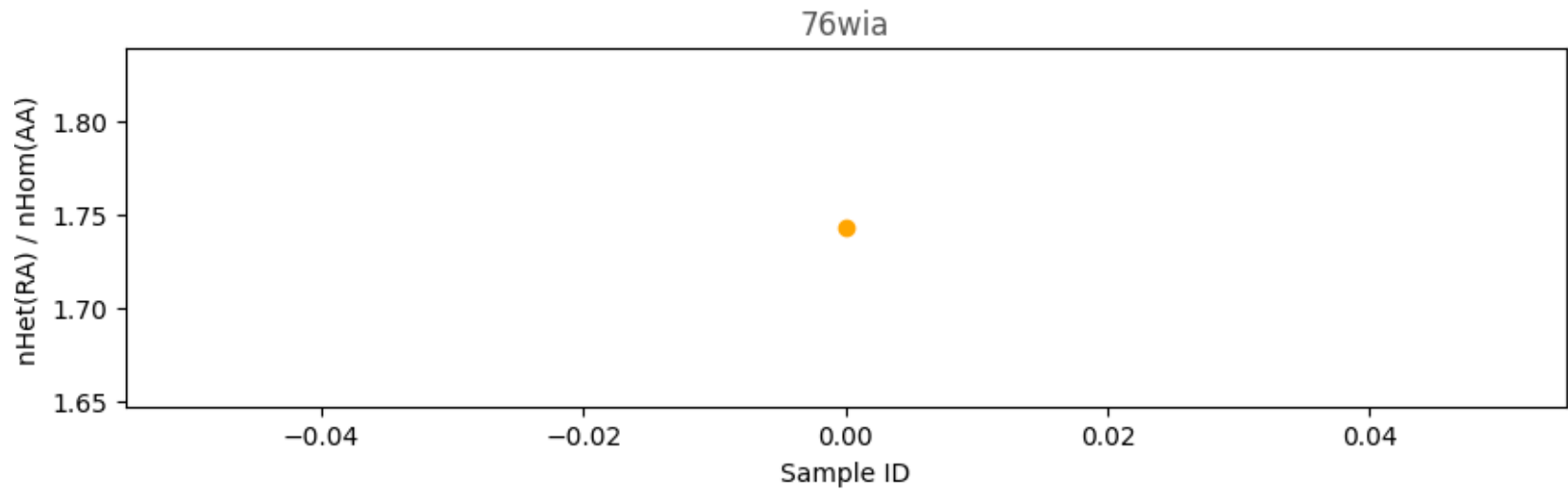
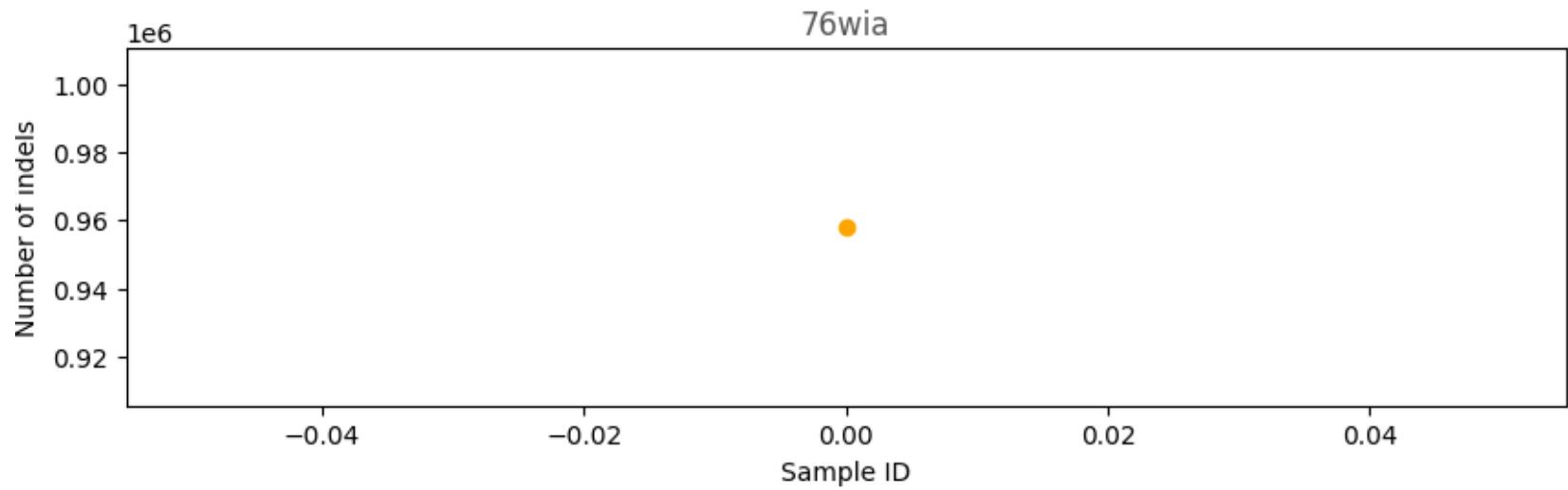# Singletons by sample (hets and homs)

Average depth by sample
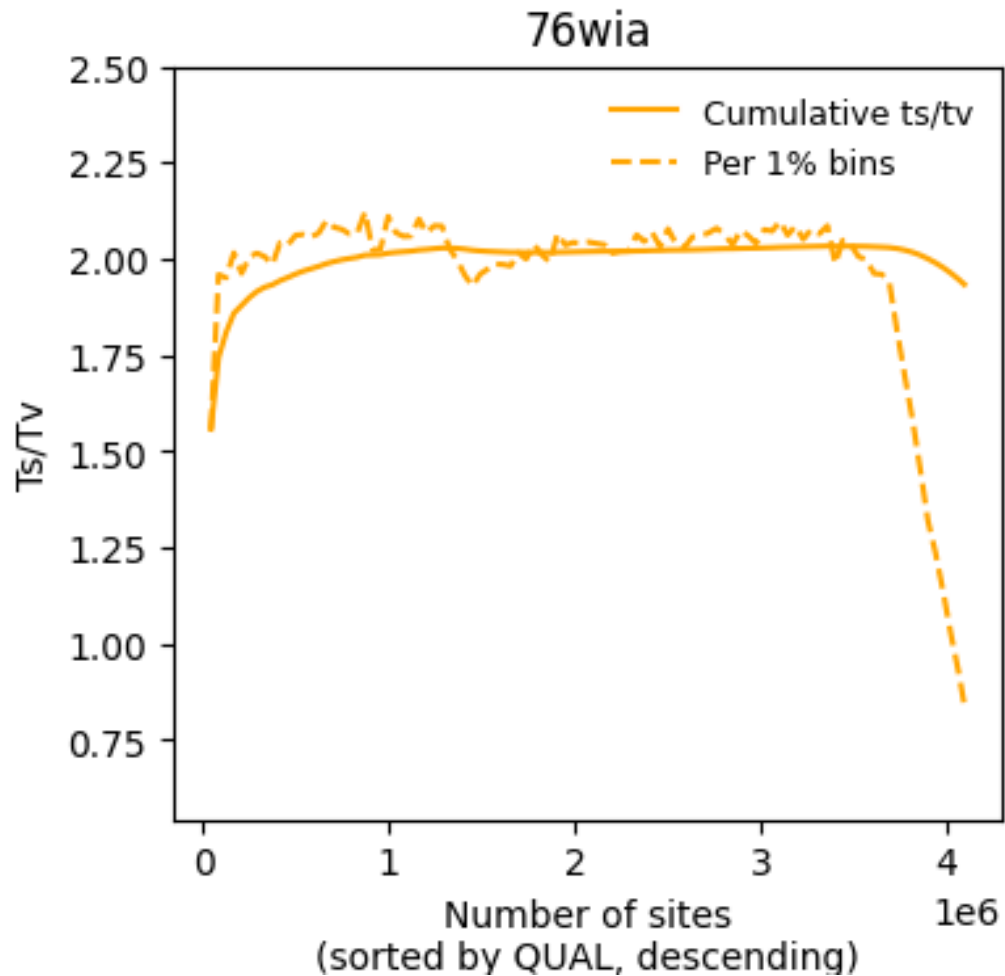
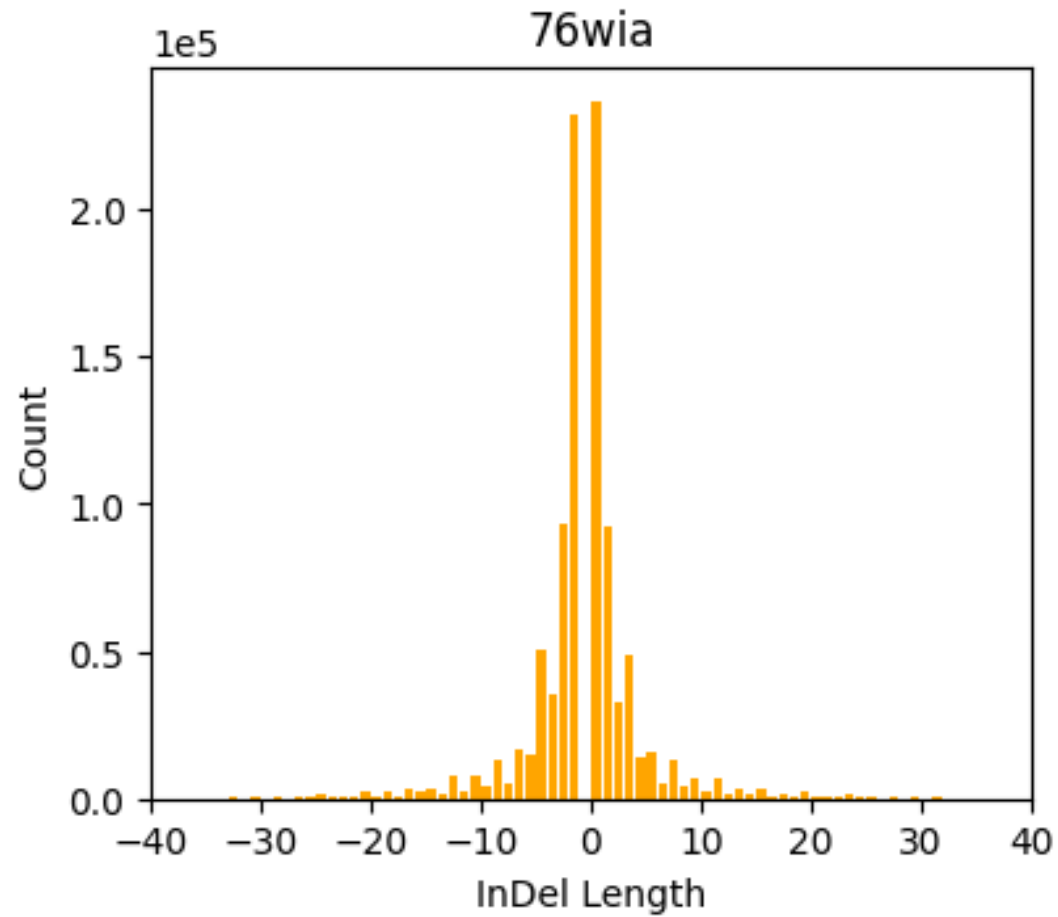# Number of SNPs by sample

Number of indels by sample
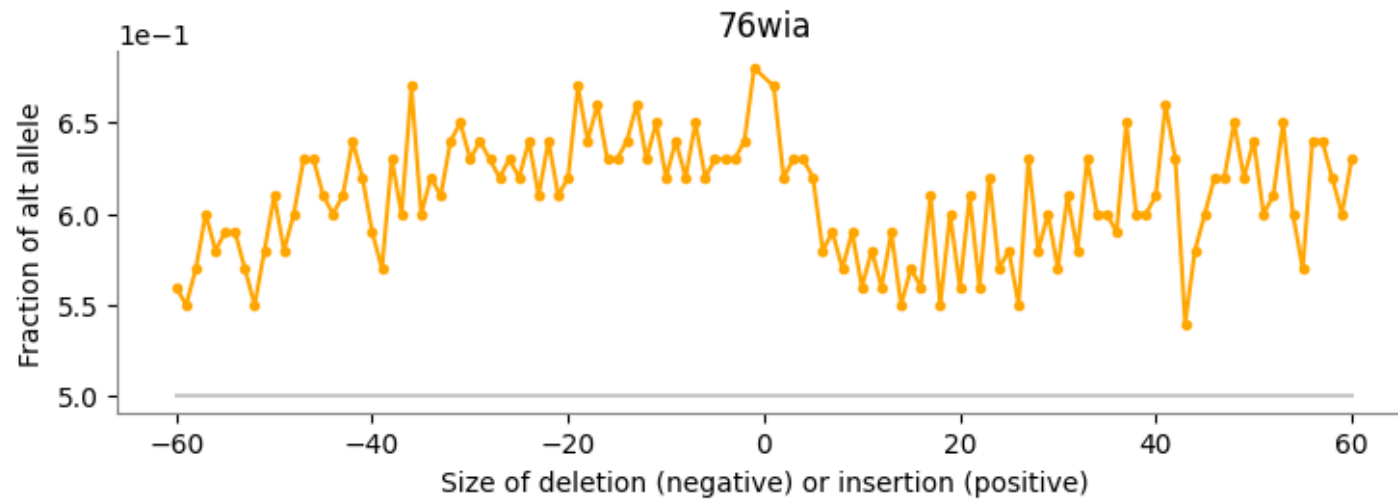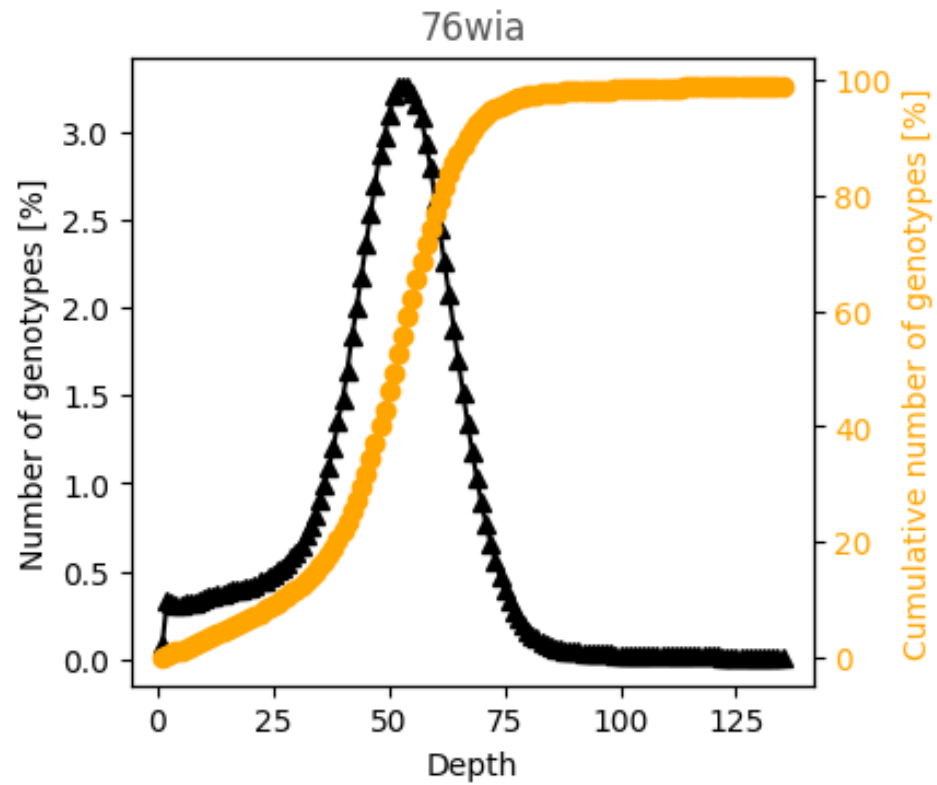
Ts/Tv stratified by QUAL

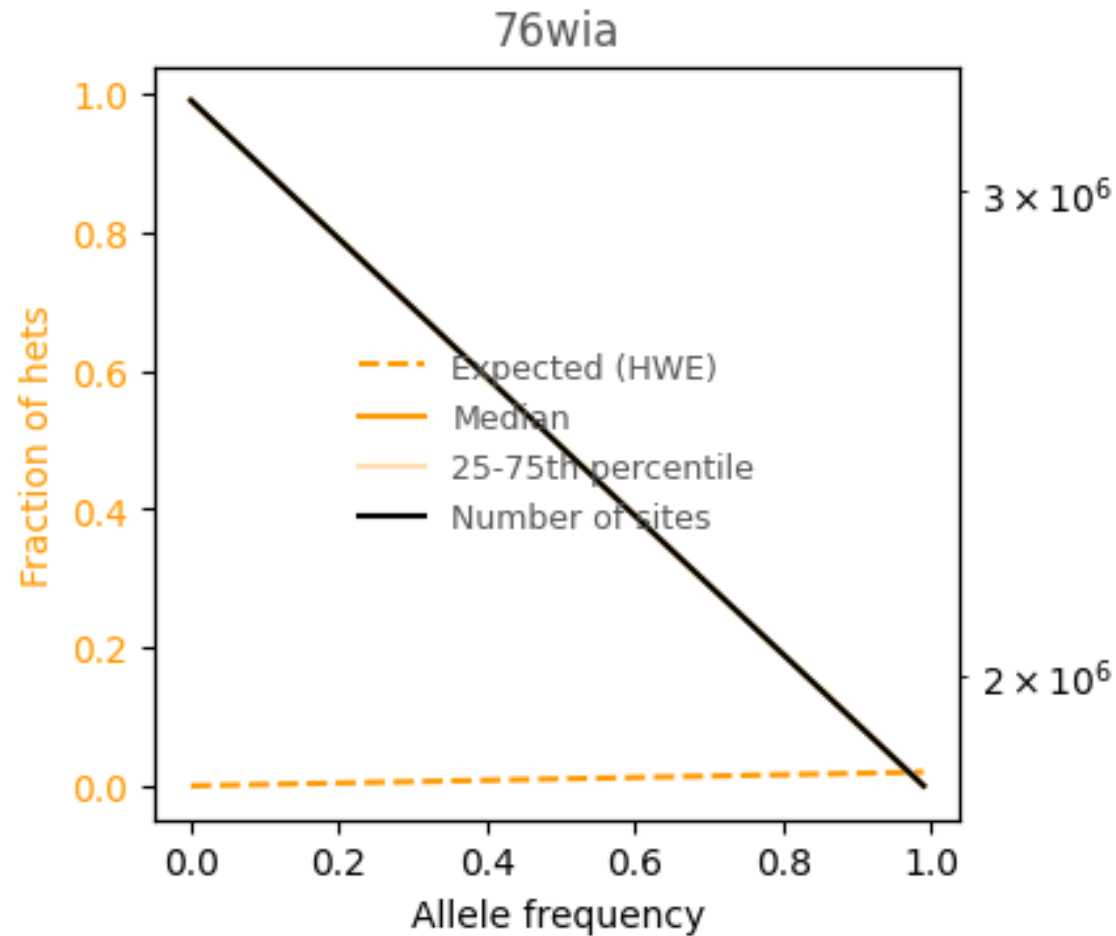

76wia

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types