Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
75tpo	4,074,662	1.92	1.93	951,013	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
75tpo	62.1%	1.89	67.0%	100,915	2,162

- 75tpo .. `/ngc/projects2/gm/data/archive/2022/variants/snv/75tpoform-103818401688-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211222_A01466_AH2CJCDSX3-RHGM_LABKA_WGSA`
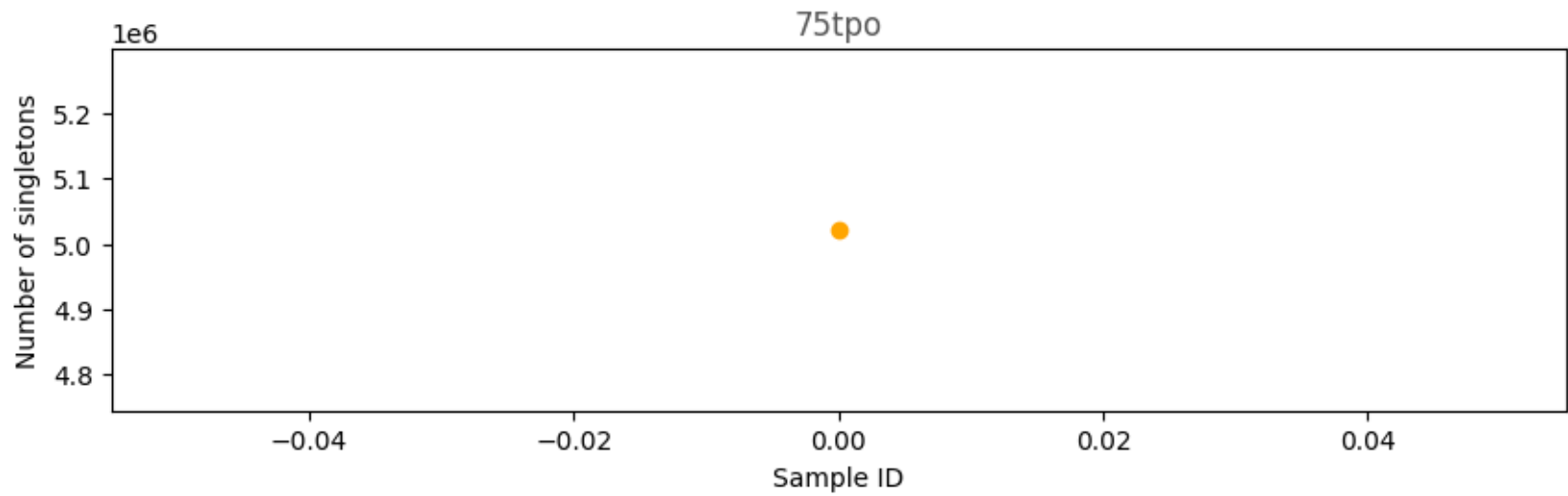 `KUT-WGSAKUT03548_21RKG029747x01_snv_germline_raw.haplotype_caller.vcf.gz`
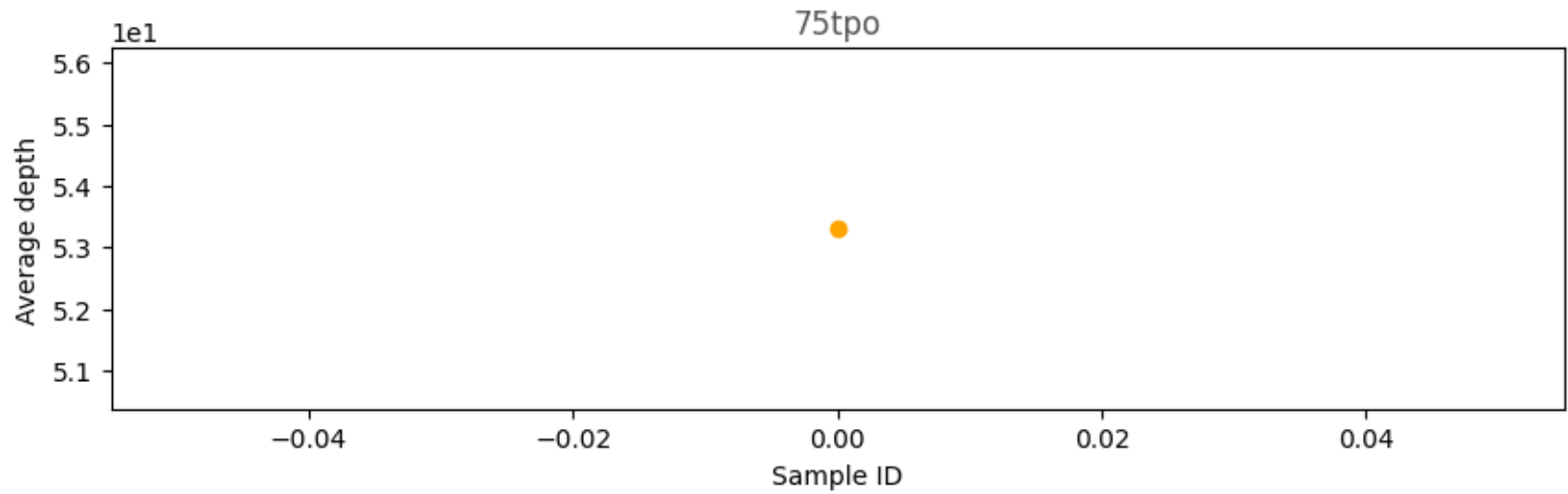
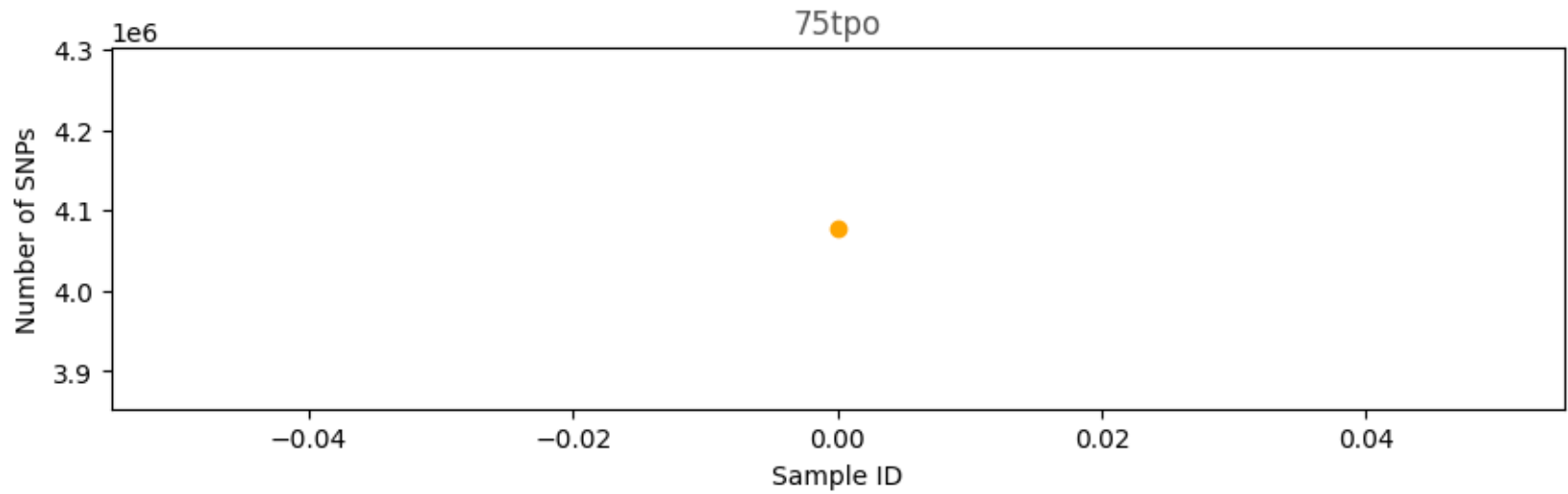# Ts/Tv by sample
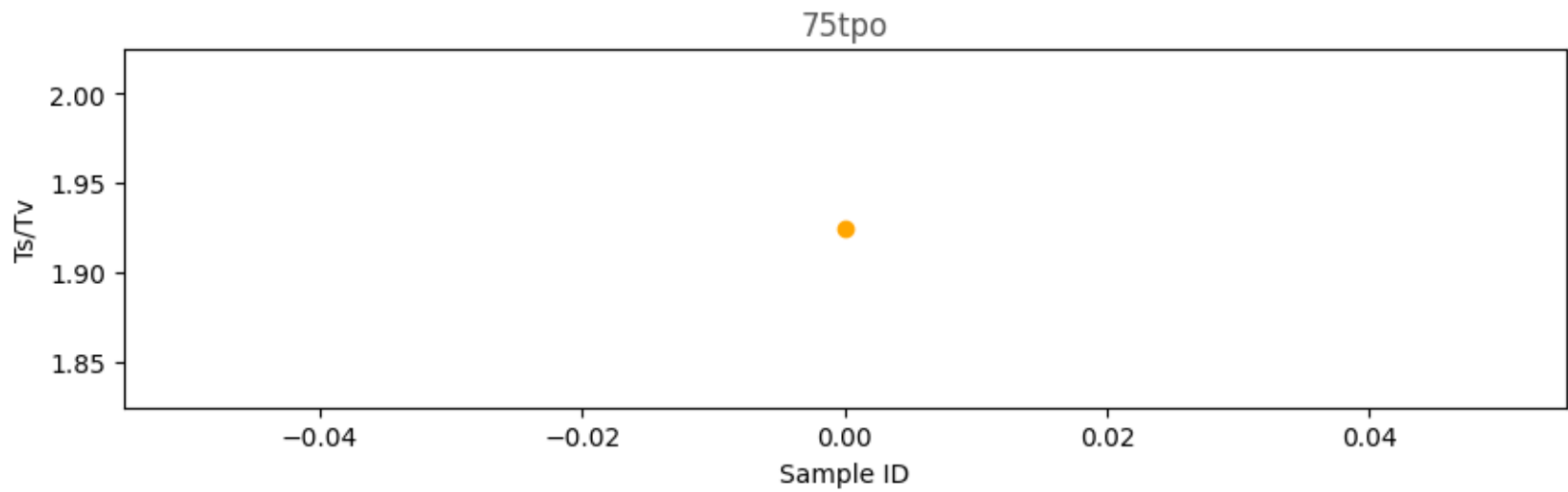
Hets vs non-ref Homs by sample
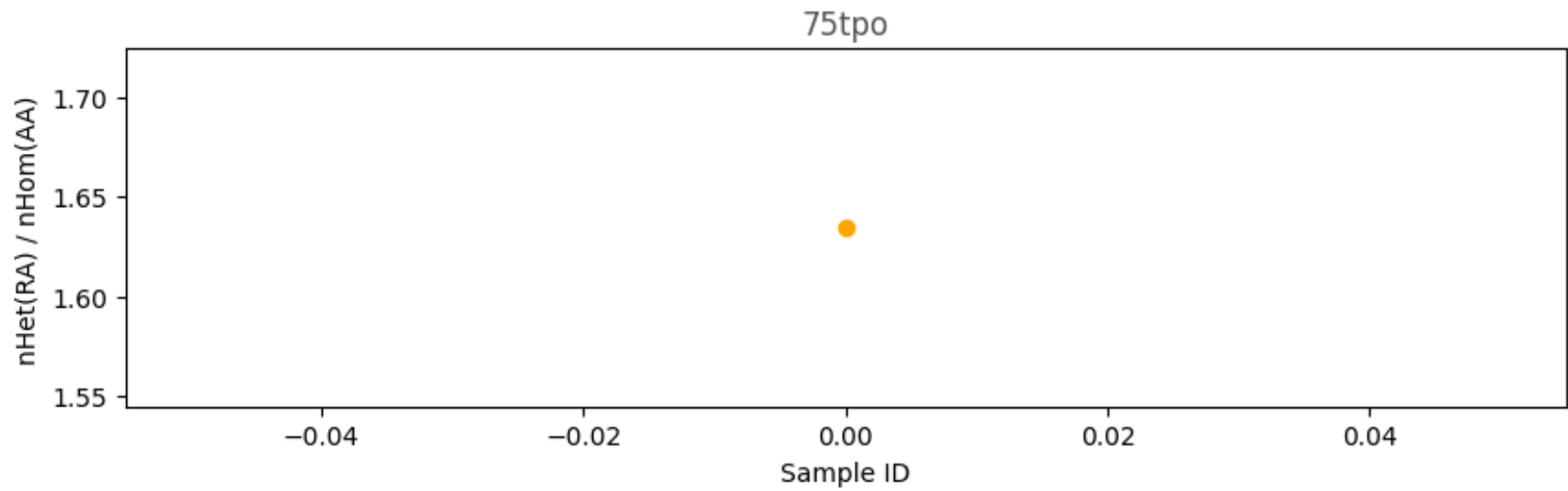
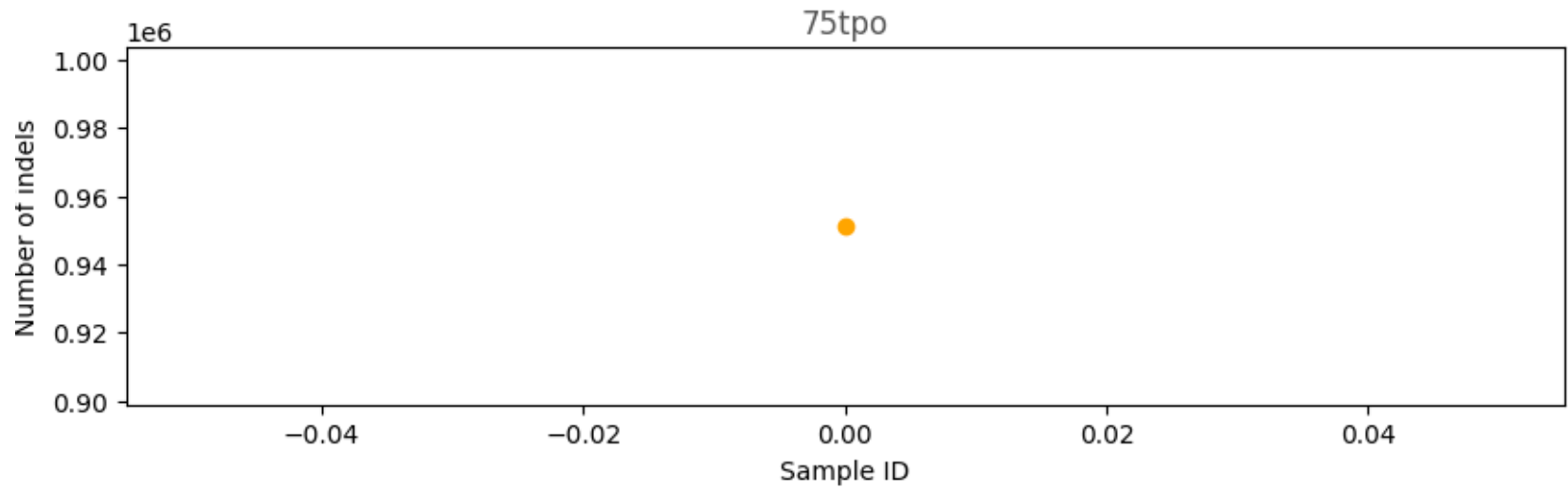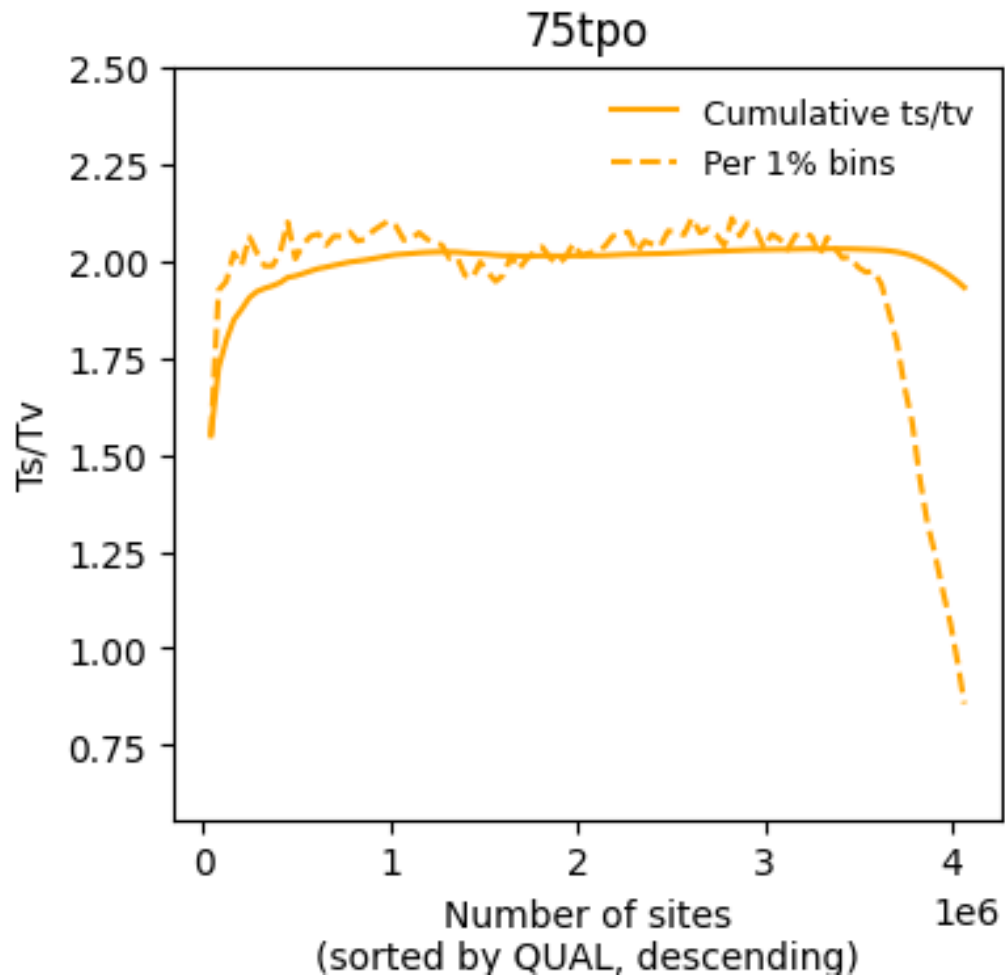# Singletons by sample (hets and homs)
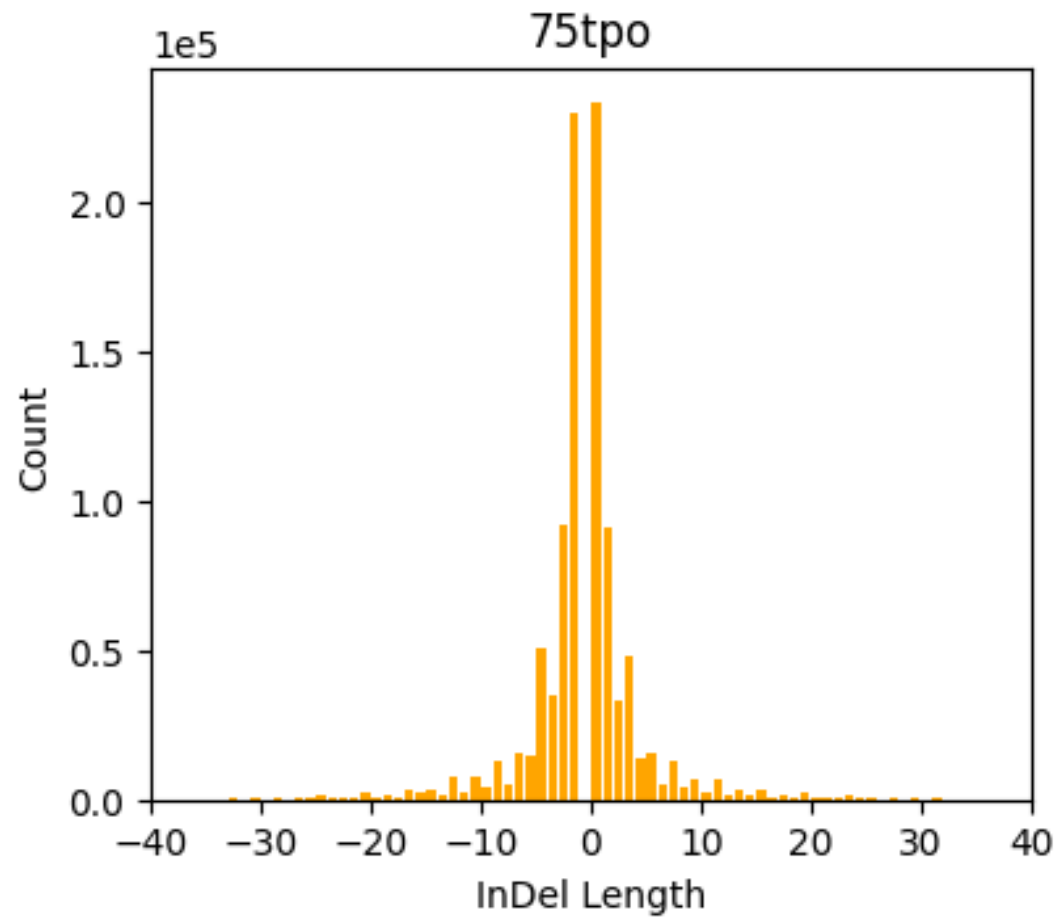
Average depth by sample

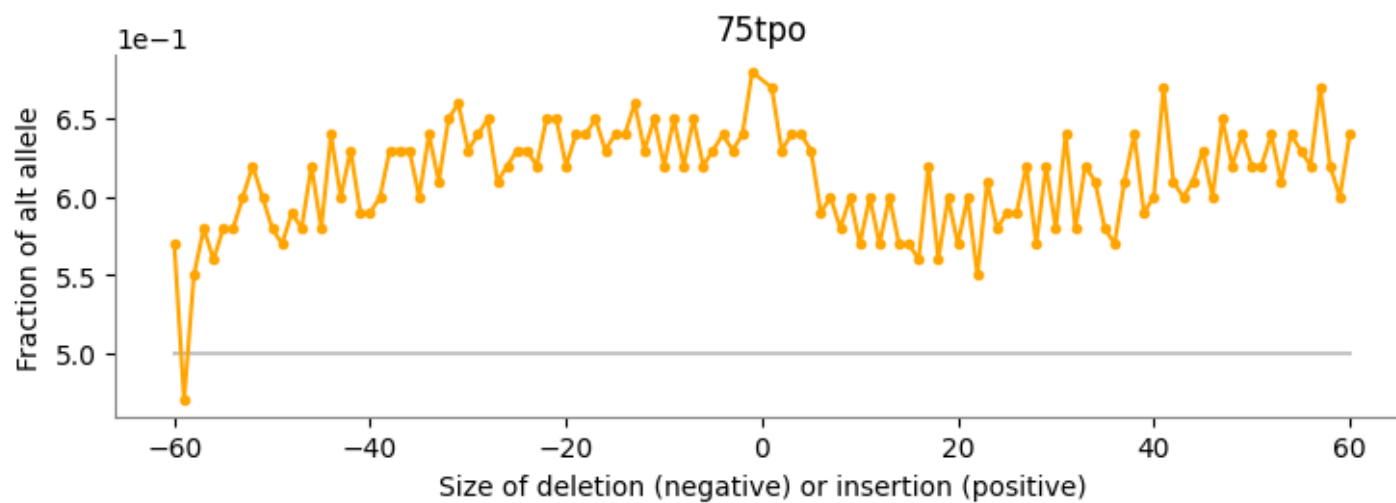# Number of SNPs by sample

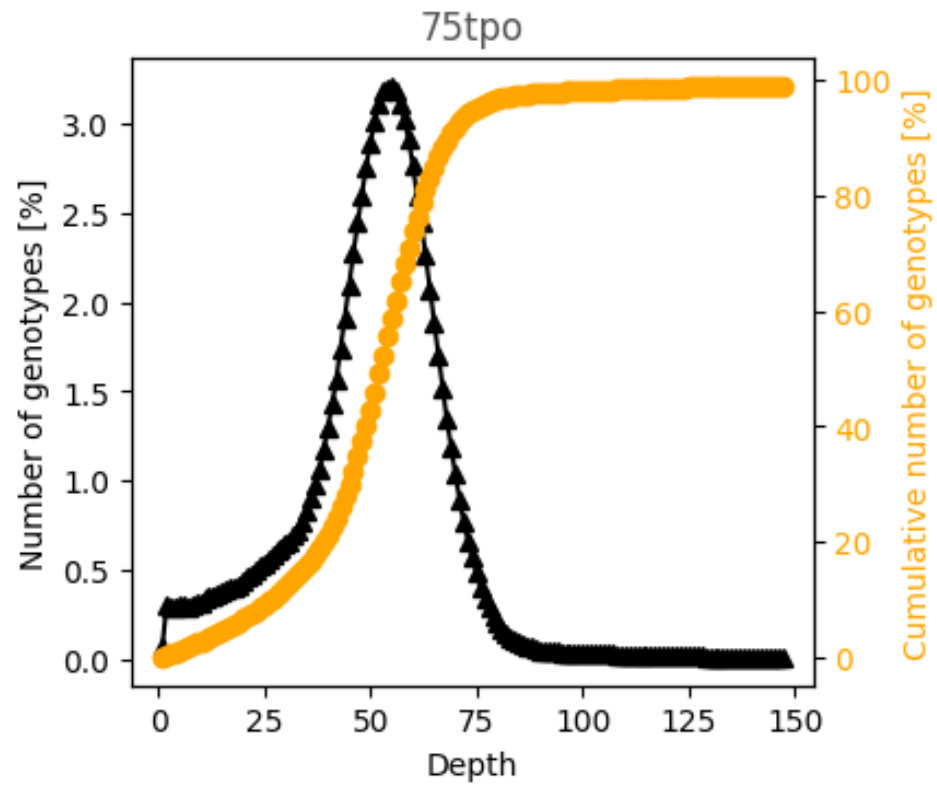# Number of indels by sample

Ts/Tv stratified by QUAL
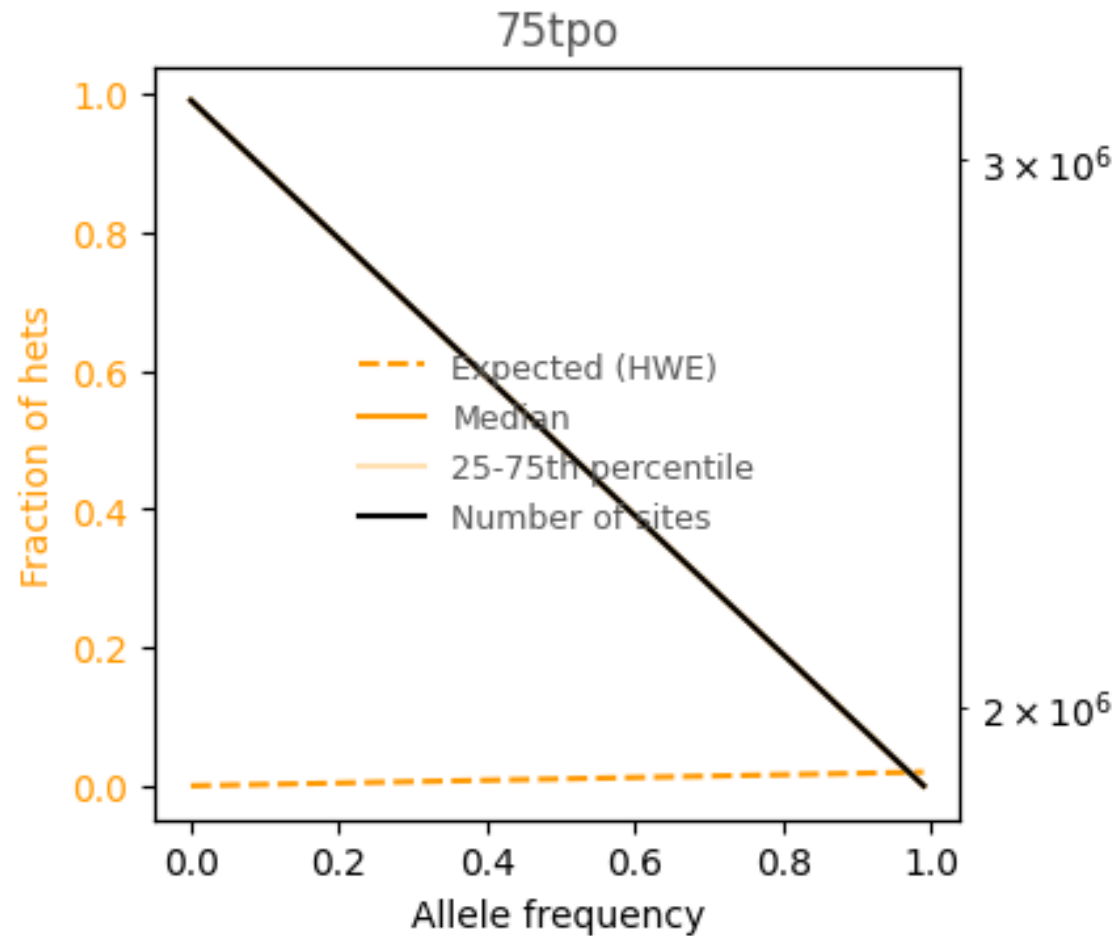

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Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types

75tpo

1e5

Count

A>C A>G A>T C>A C>G C>T G>A G>C G>T T>A T>C T>G