Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
43tro	4,040,370	1.93	1.93	910,774	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
43tro	63.1%	1.89	66.4%	84,188	2,054

- 43tro .. `/ngc/projects2/gm/data/archive/2022/variants/snv/43trovexf-103878803073-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220615_A00559_BHKC2YDSX3-RHGM_LABKA_WGS-`
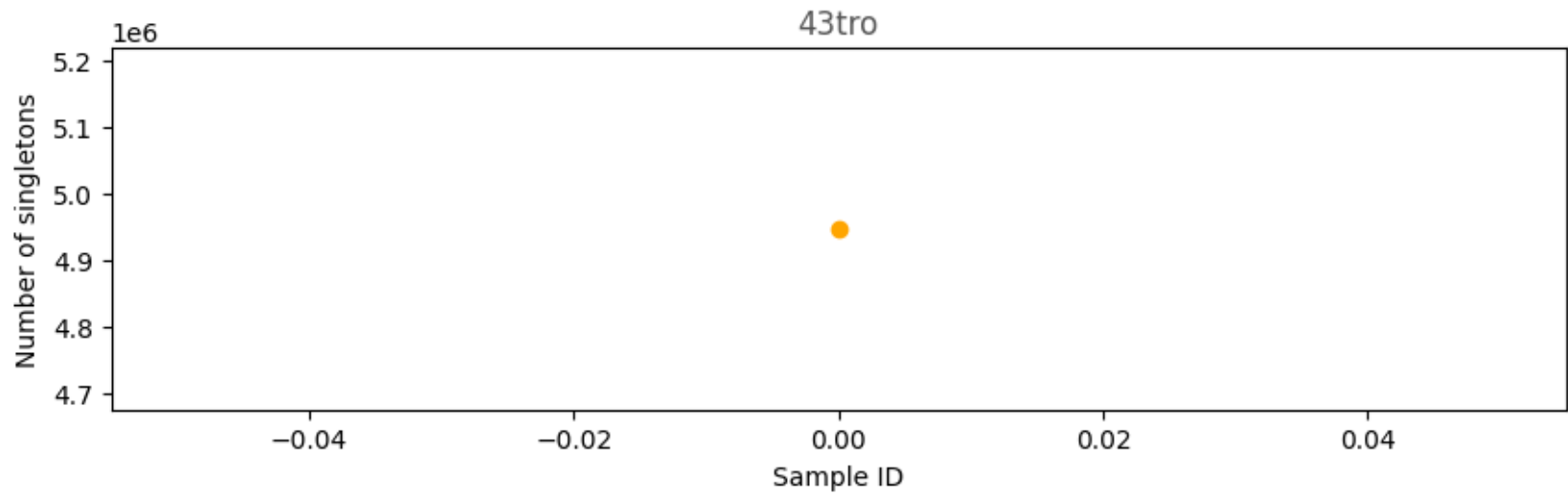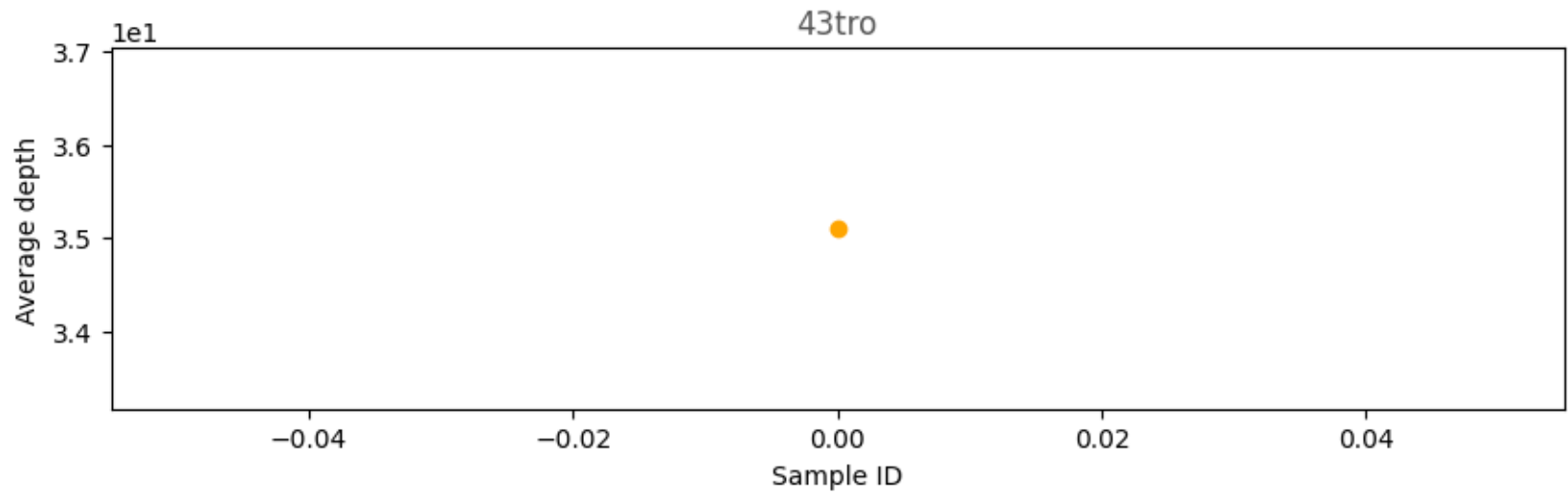 `WGS04447_snv_germline_raw.haplotype_caller.vcf.gz`
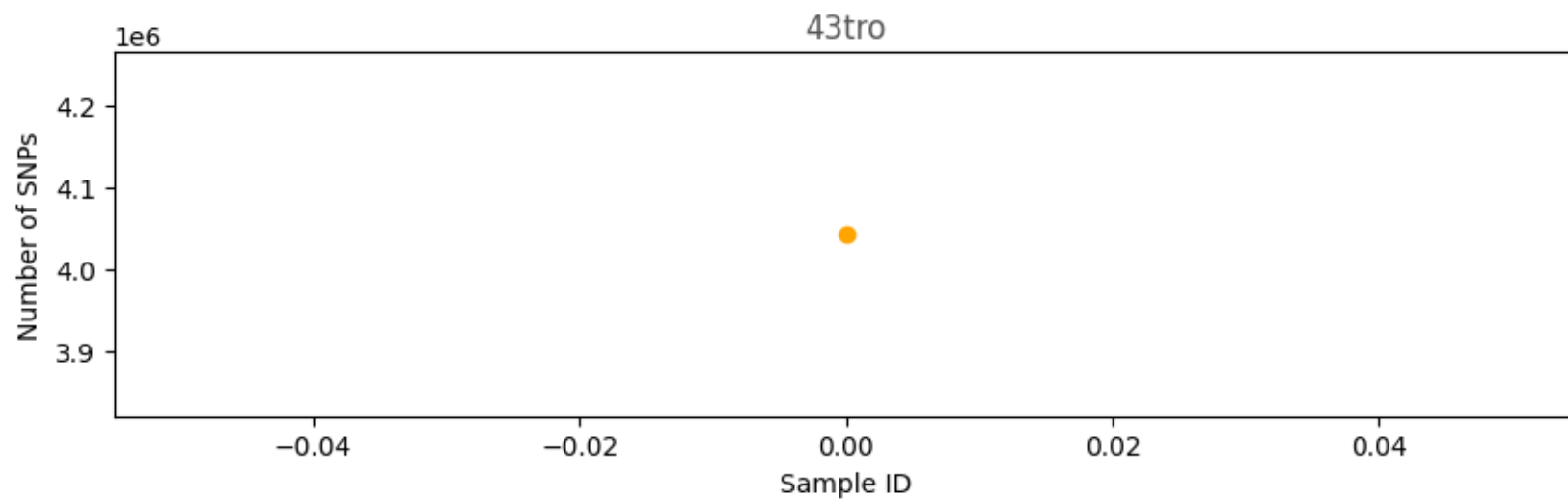
Ts/Tv by sample
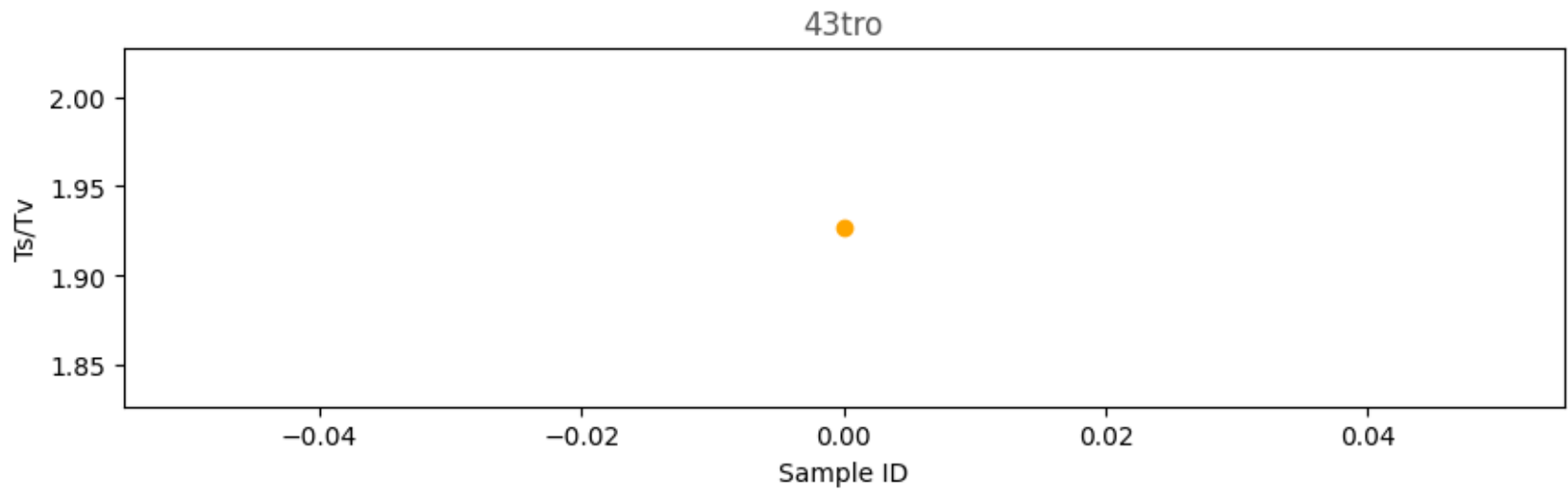
Hets vs non-ref Homs by sample
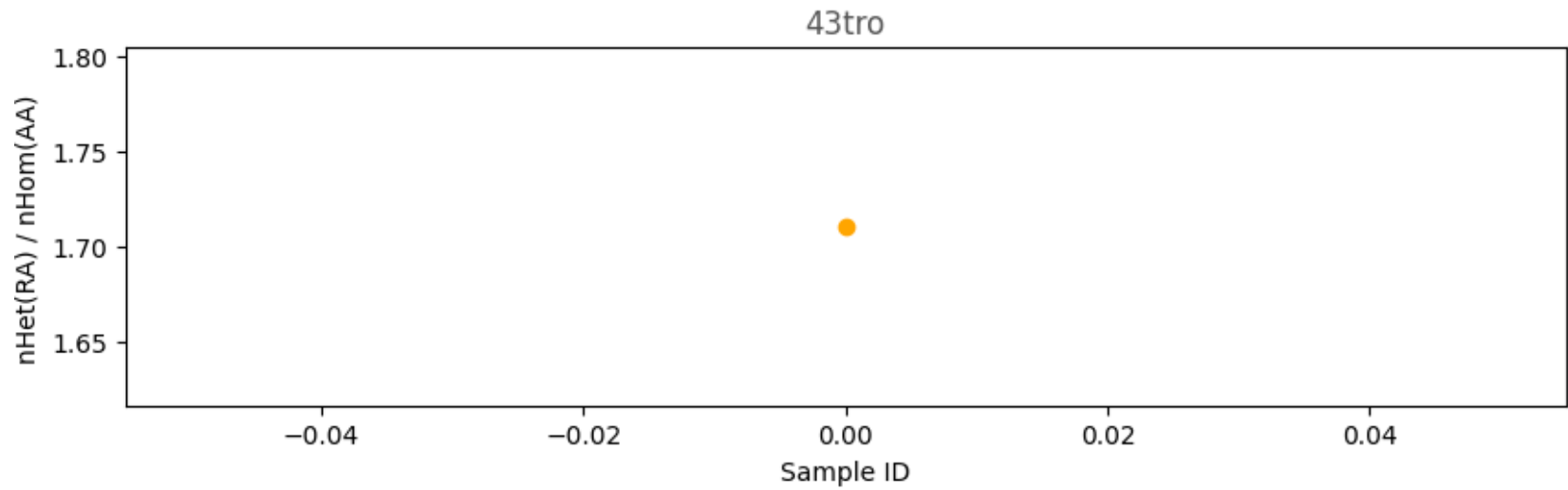
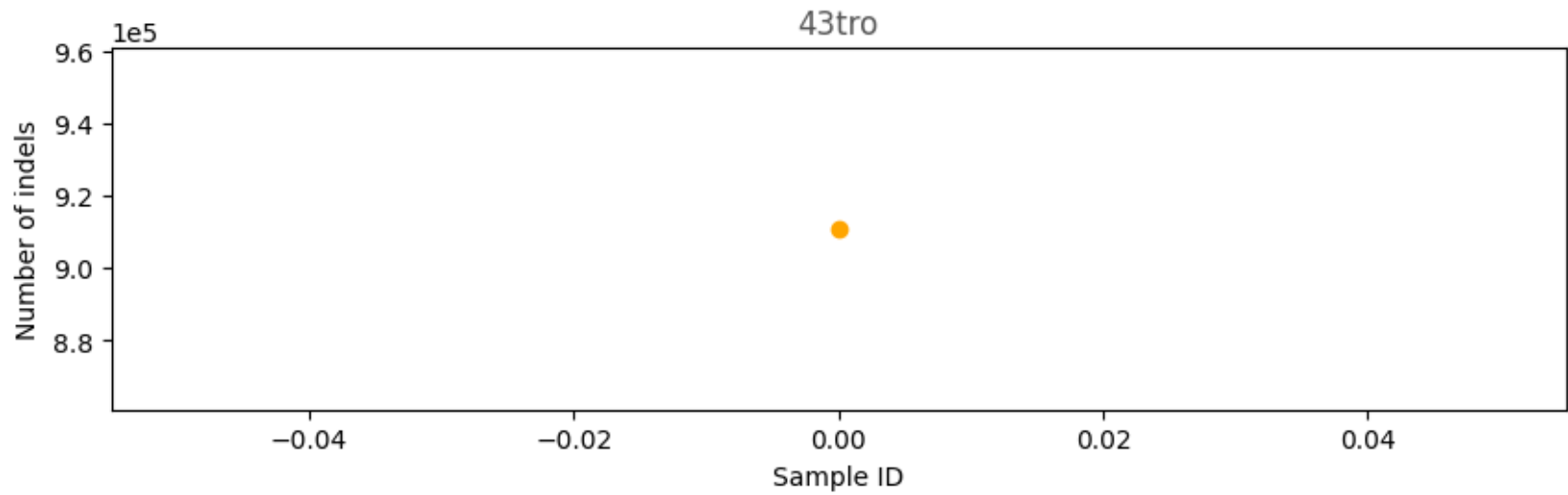# Singletons by sample (hets and homs)
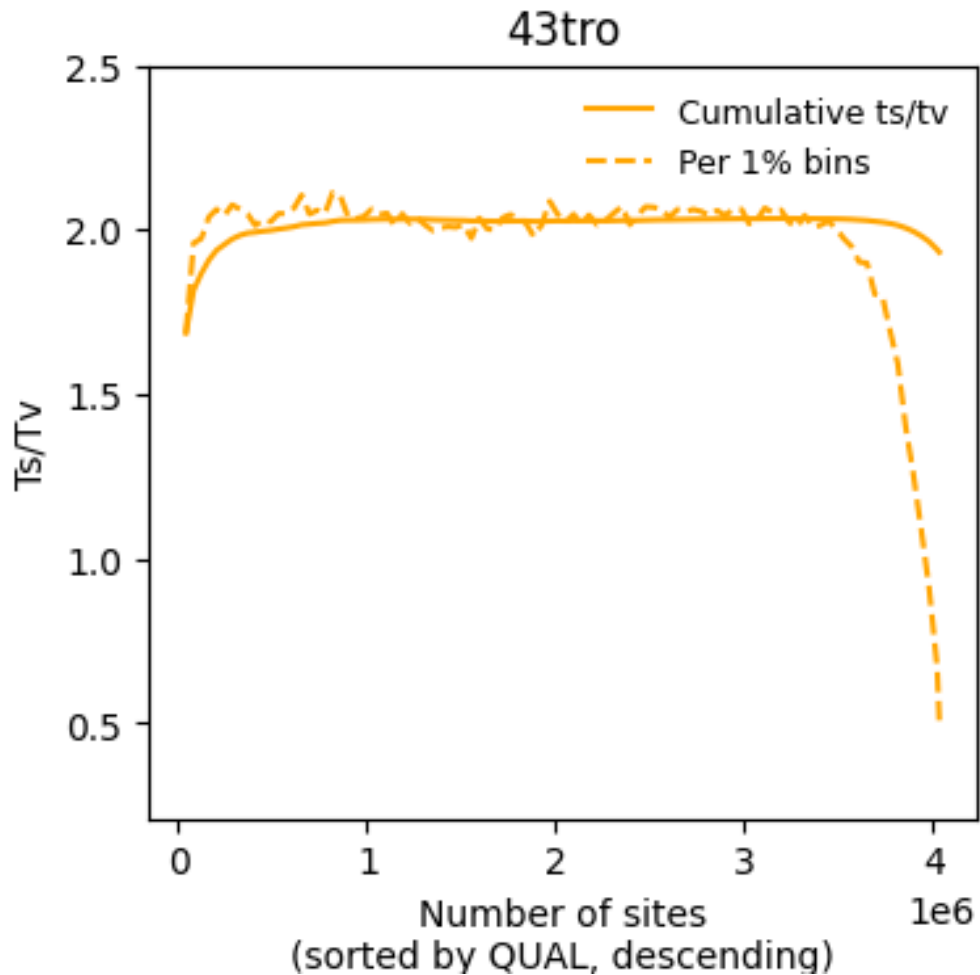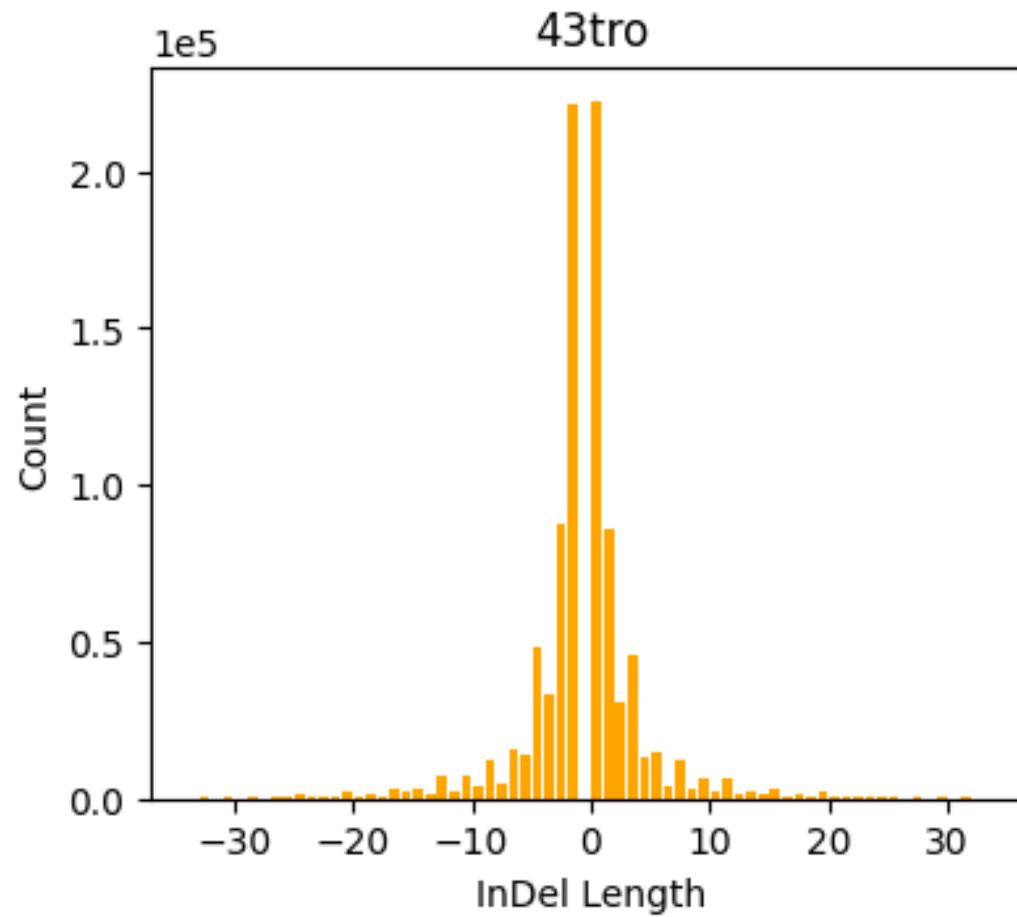
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

43tro

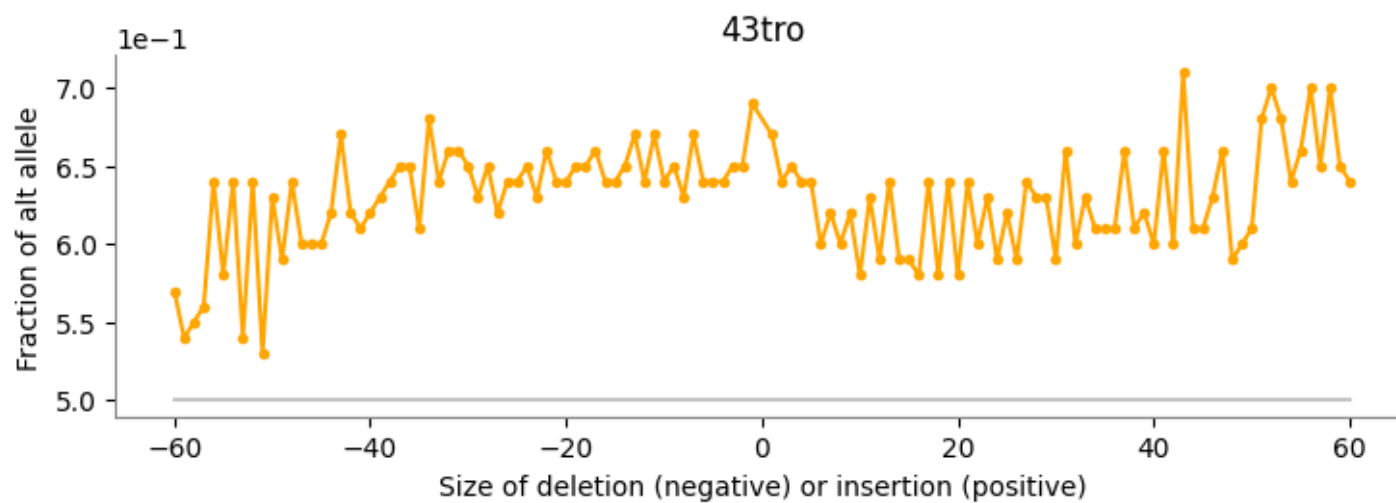# Indel distribution

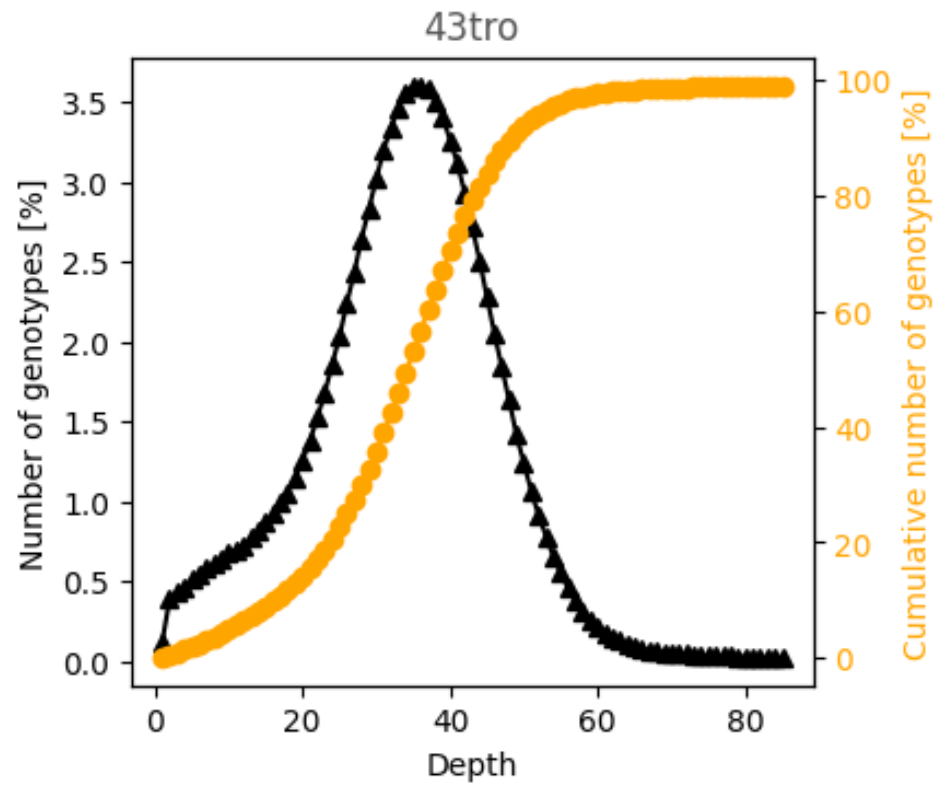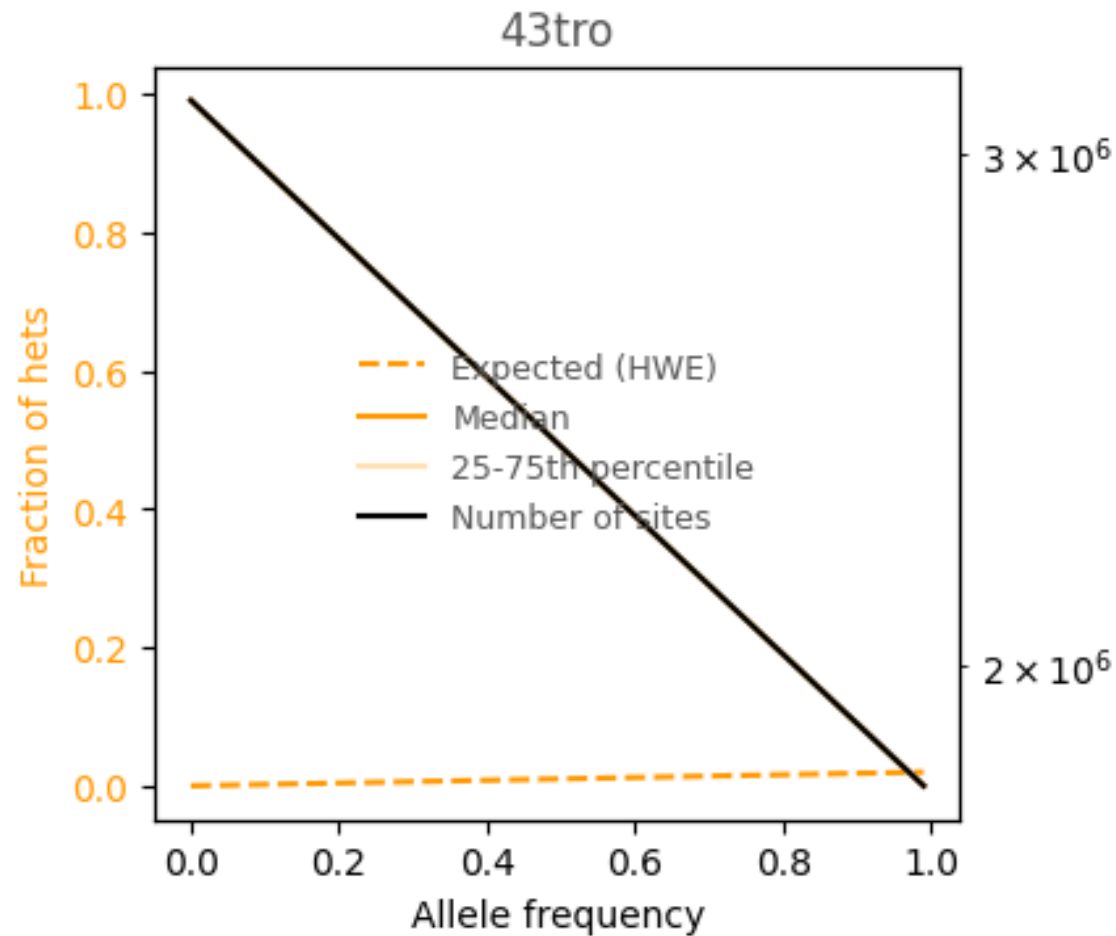# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types