Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
83ros	4,042,409	1.94	1.94	922,135	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
83ros	61.8%	1.91	65.3%	87,390	2,020

- 83ros .. `/ngc/projects2/gm/data/archive/2022/variants/snv/83rosoldm-103846323505-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220309_A00559_BHCGGNDSX3-RHGM_LABKA_WGSA`
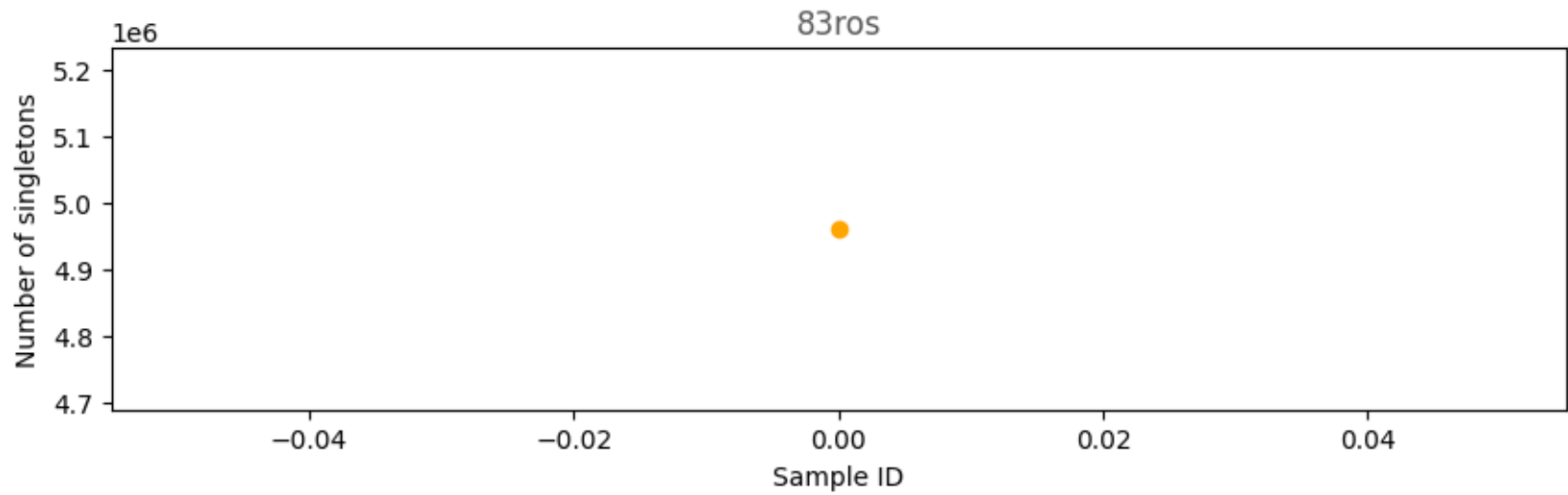 `KUT-WGSAKUT03955_22RKG005264x01_snv_germline_raw.haplotype_caller.vcf.gz`
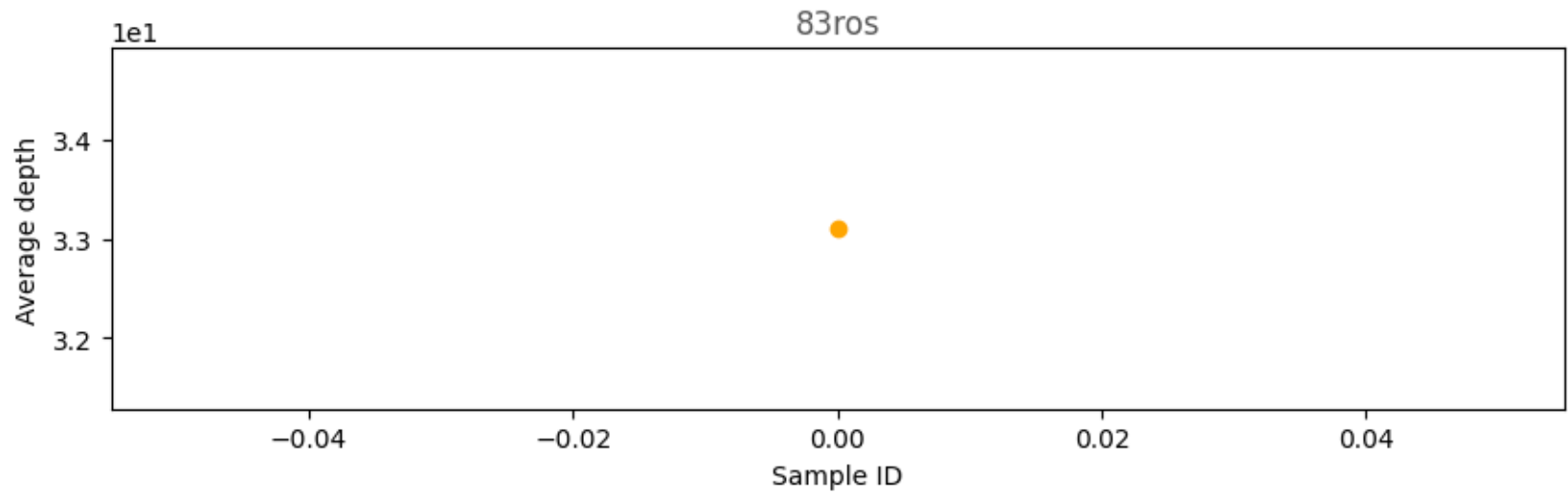
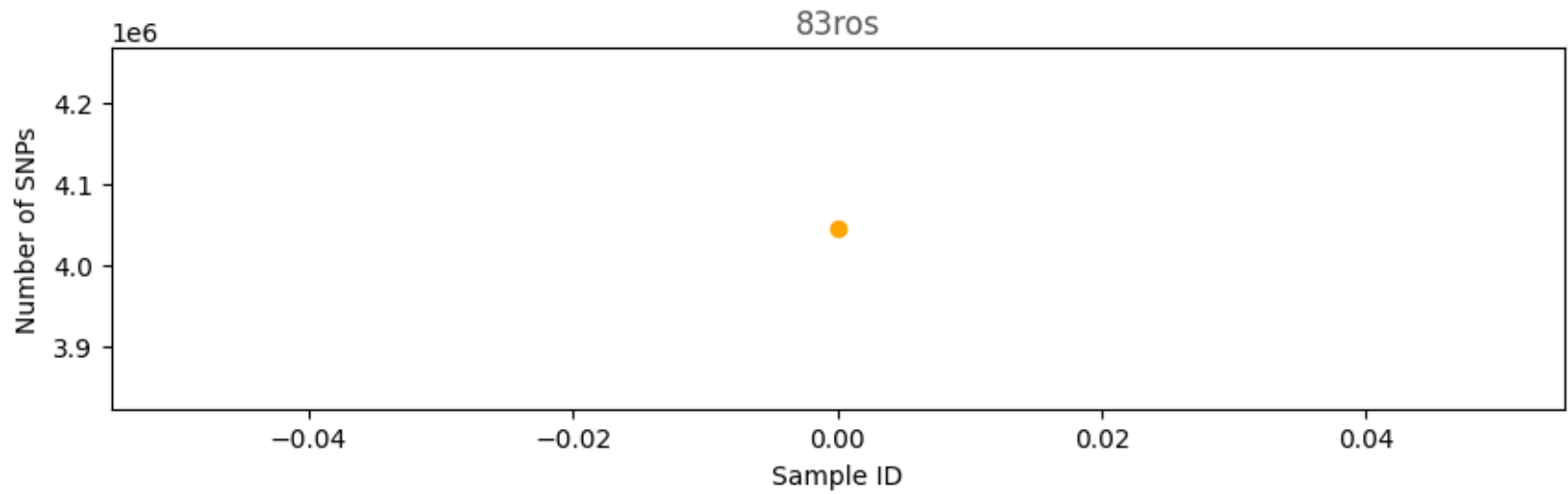# Ts/Tv by sample
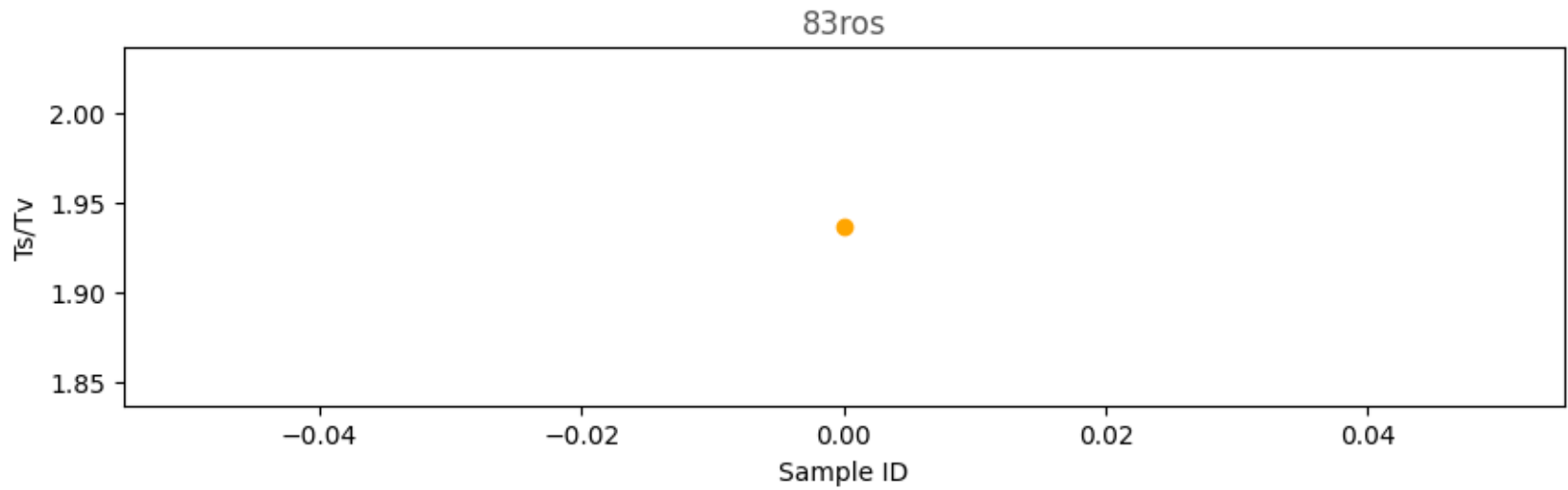
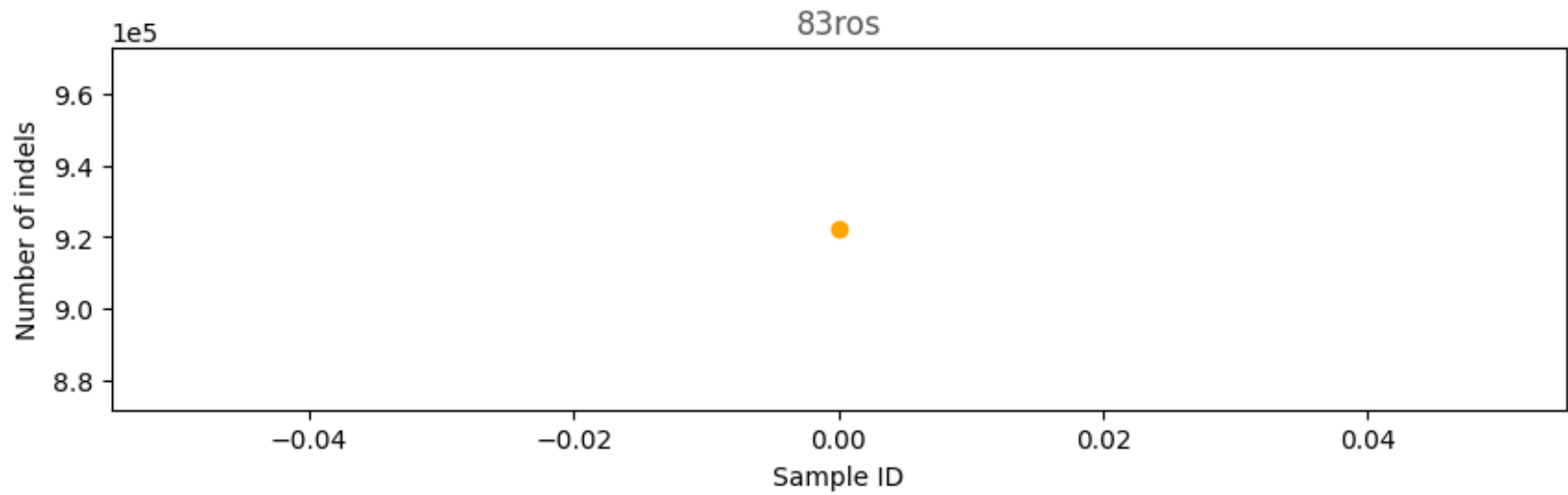# Hets vs non-ref Homs by sample
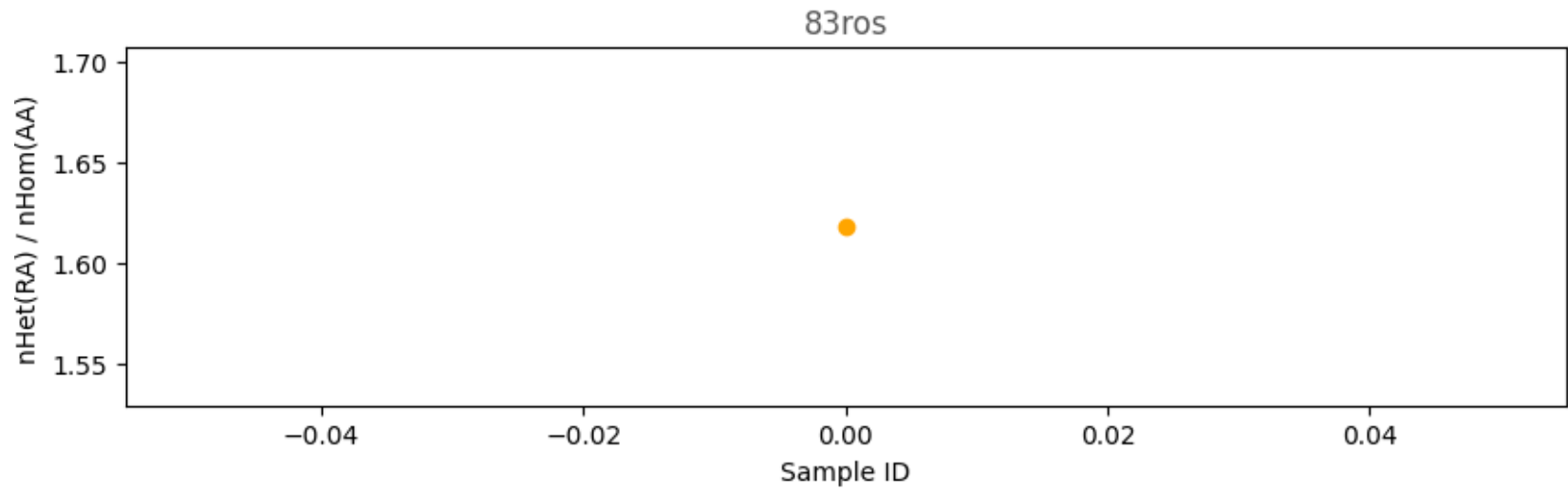
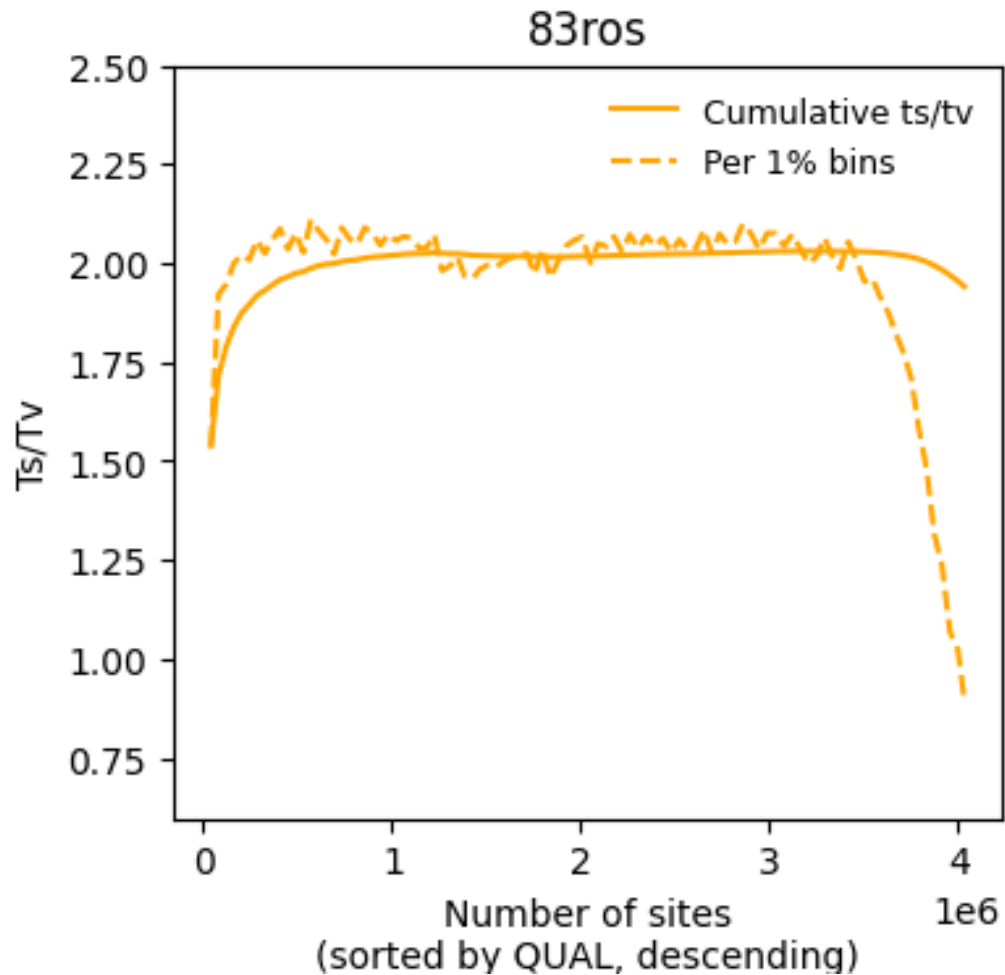# Singletons by sample (hets and homs)
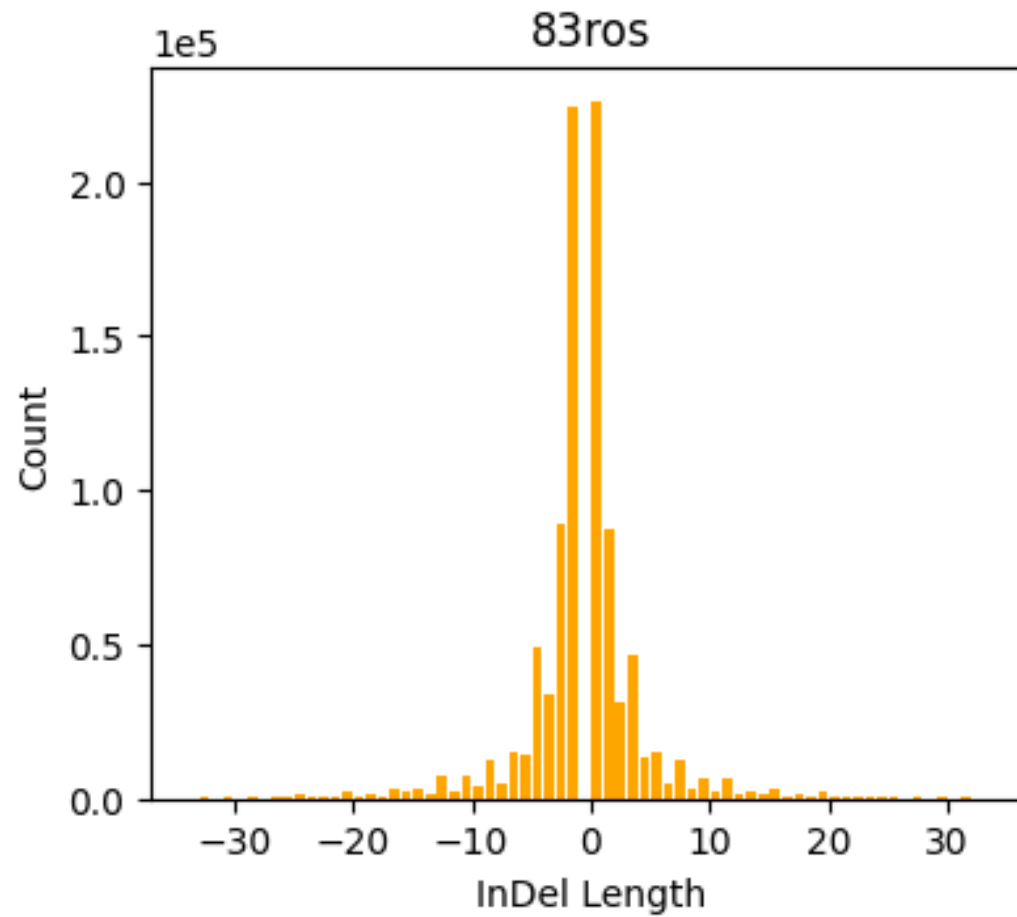
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

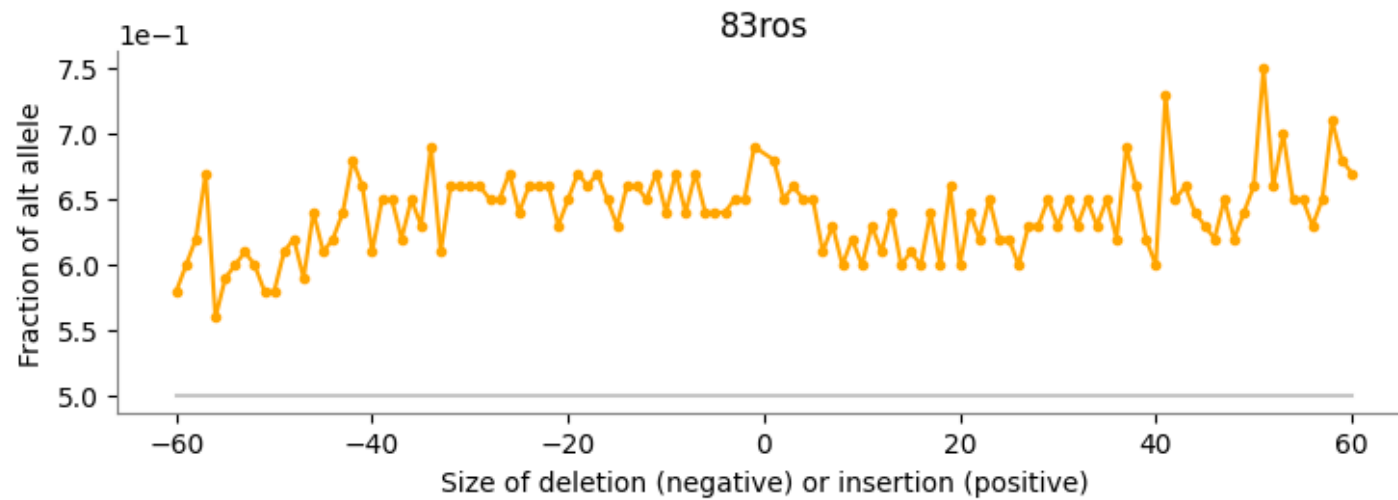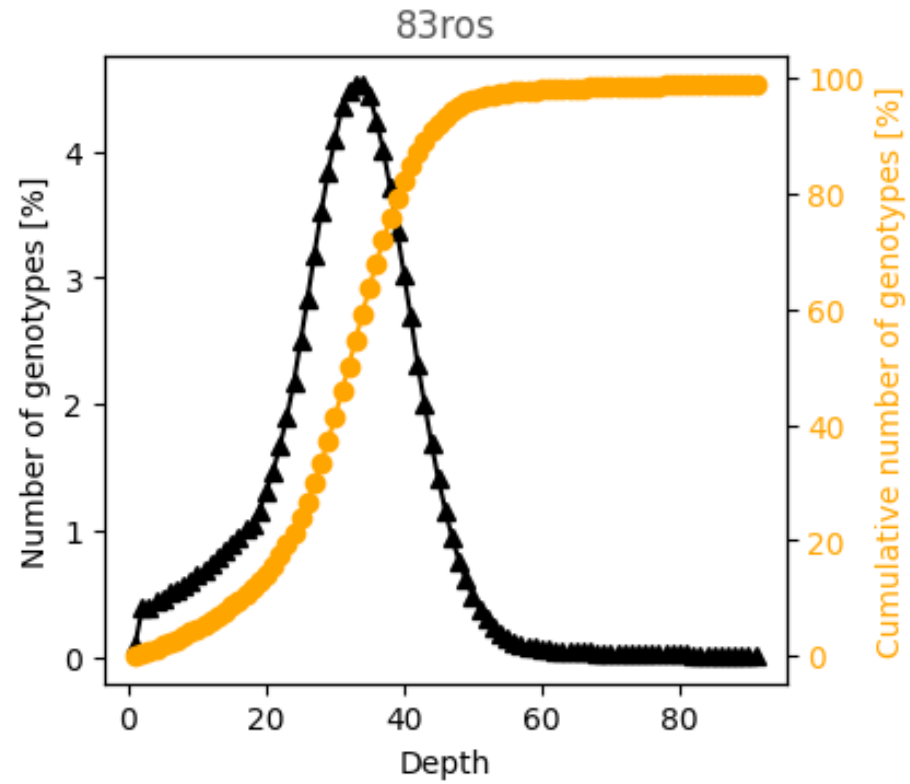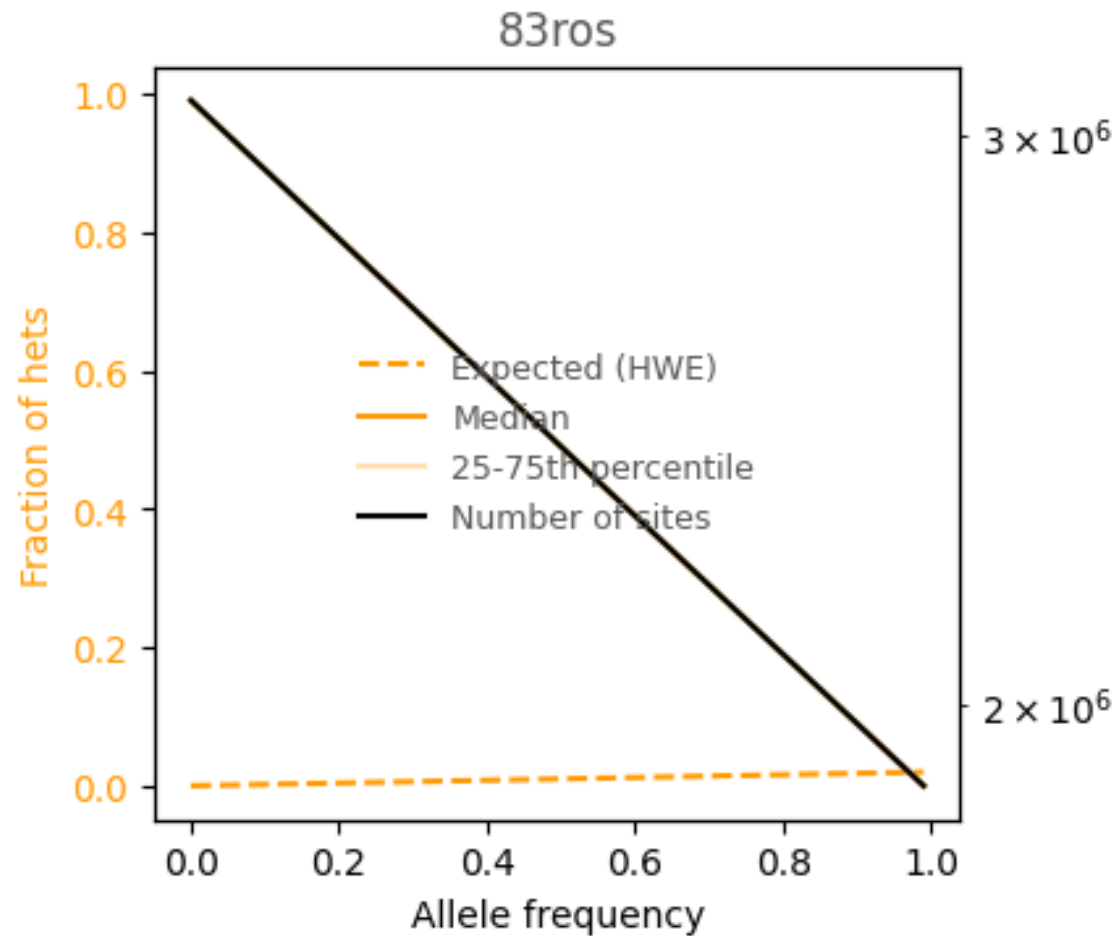# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types