Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
83tar	4,113,592	1.93	1.94	964,841	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
83tar	59.7%	1.89	65.1%	104,519	2,109

- 83tar .. `/ngc/projects2/gm/data/archive/2022/variants/snv/83tarrdvf-103909047689-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01357-220909_A01961_BHY2CGDSX3-EXT_LAB`
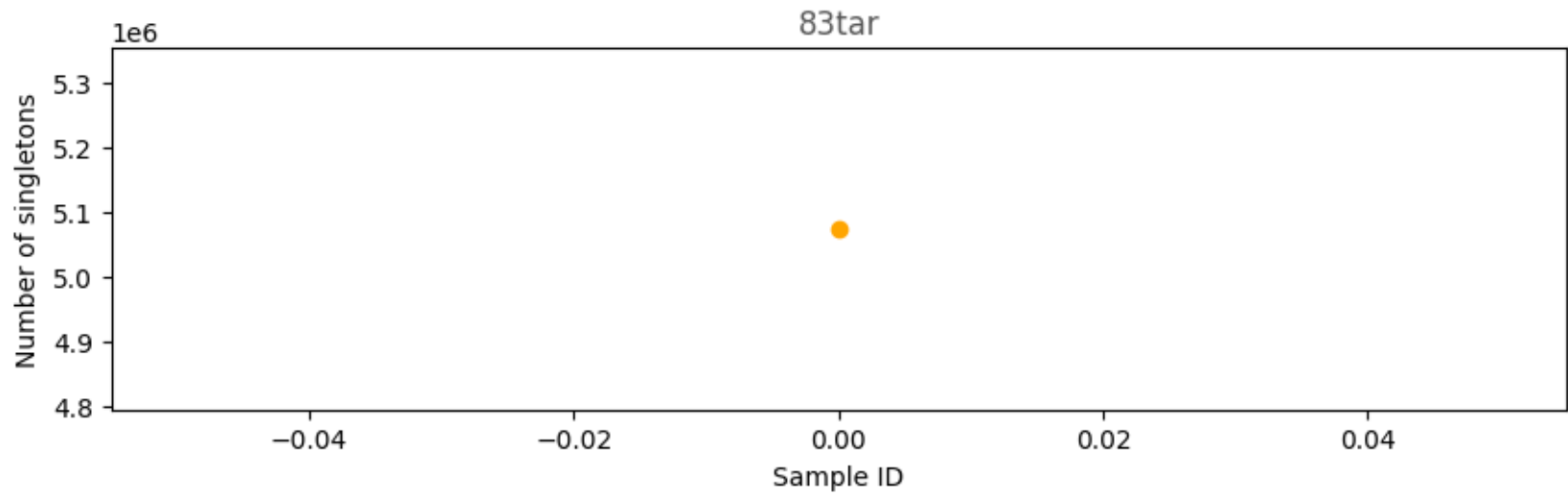 `KA_NGCWGS-NGCWGS05135_22RKG015211_snv_germline_raw.haplotype_caller.vcf.gz`
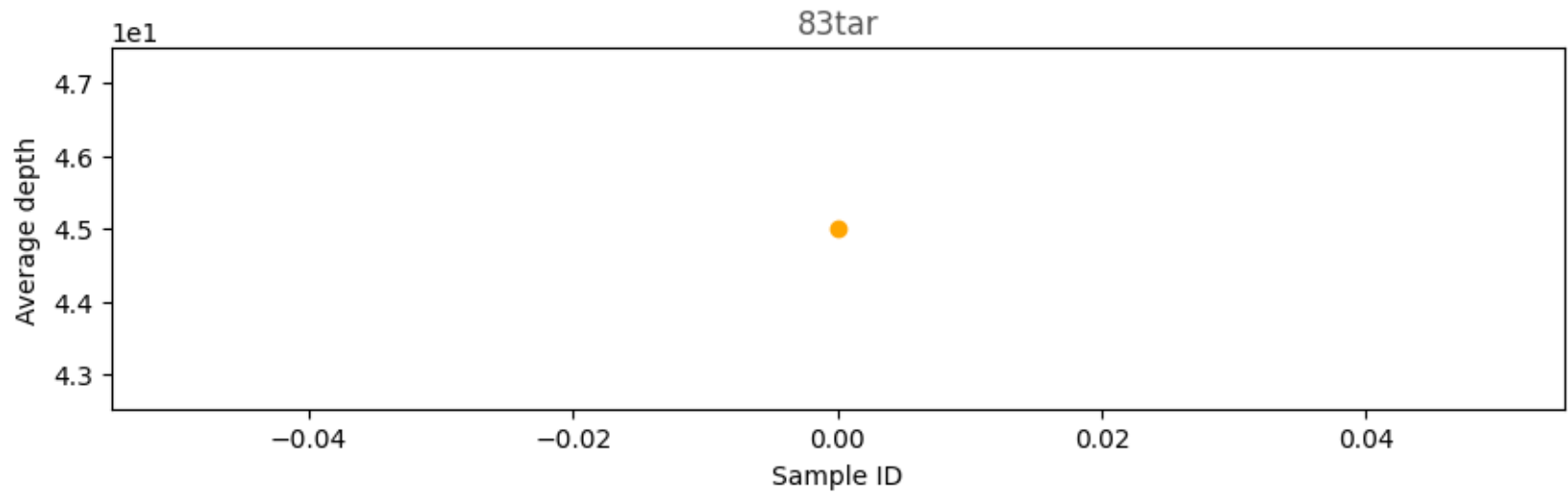
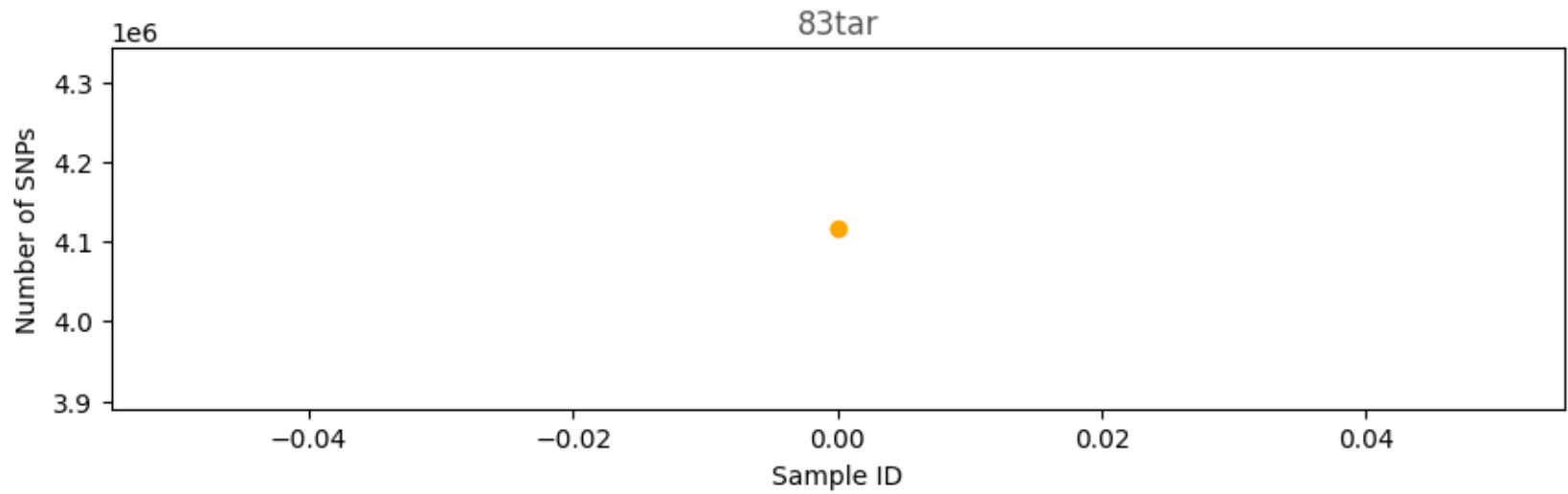# Ts/Tv by sample
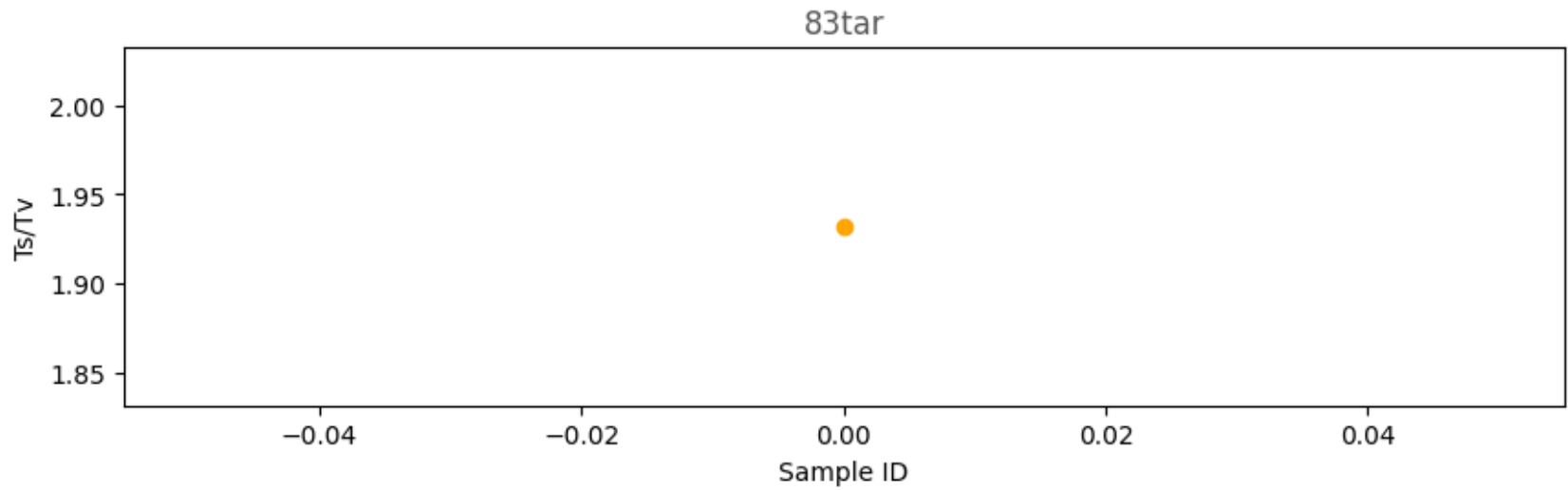
Hets vs non-ref Homs by sample
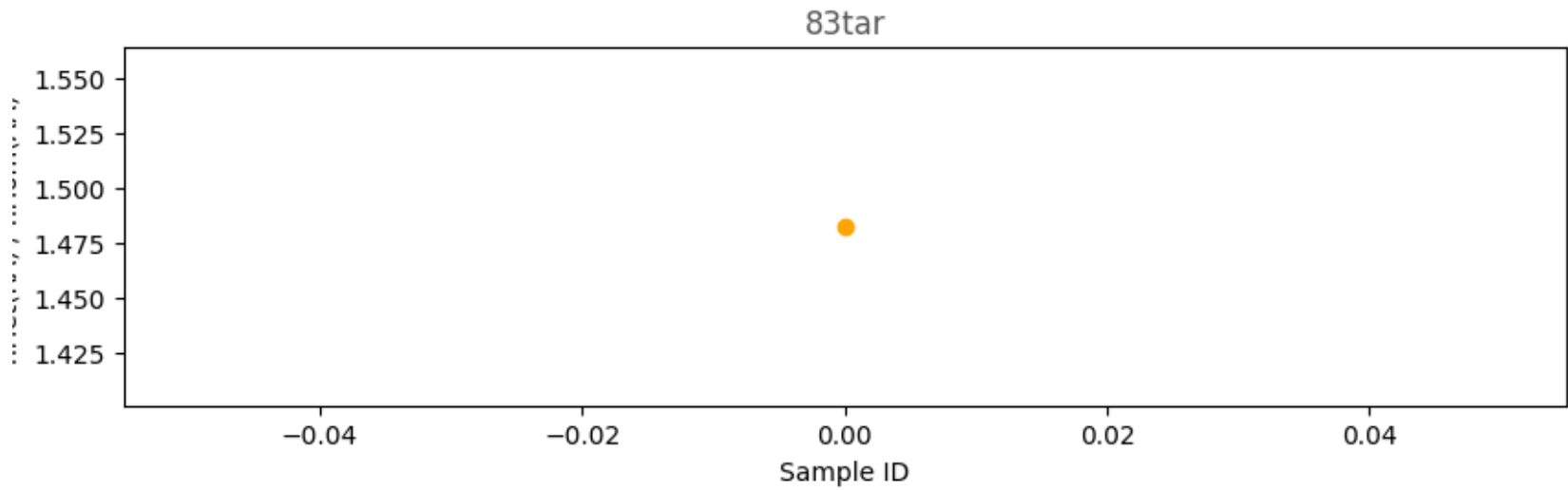
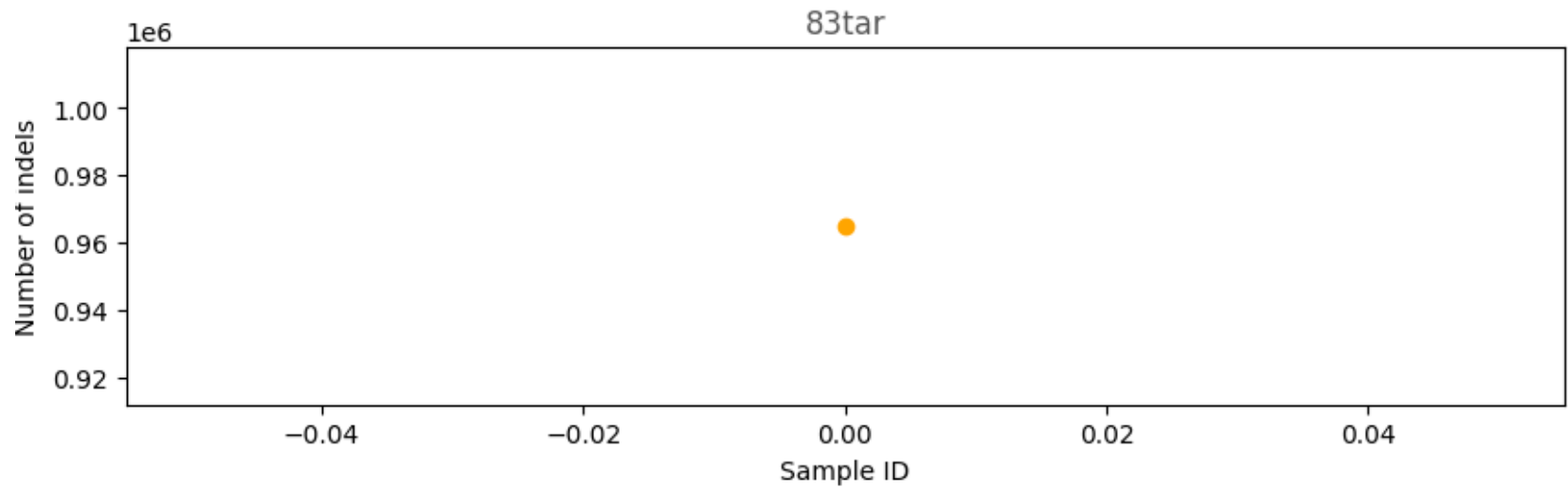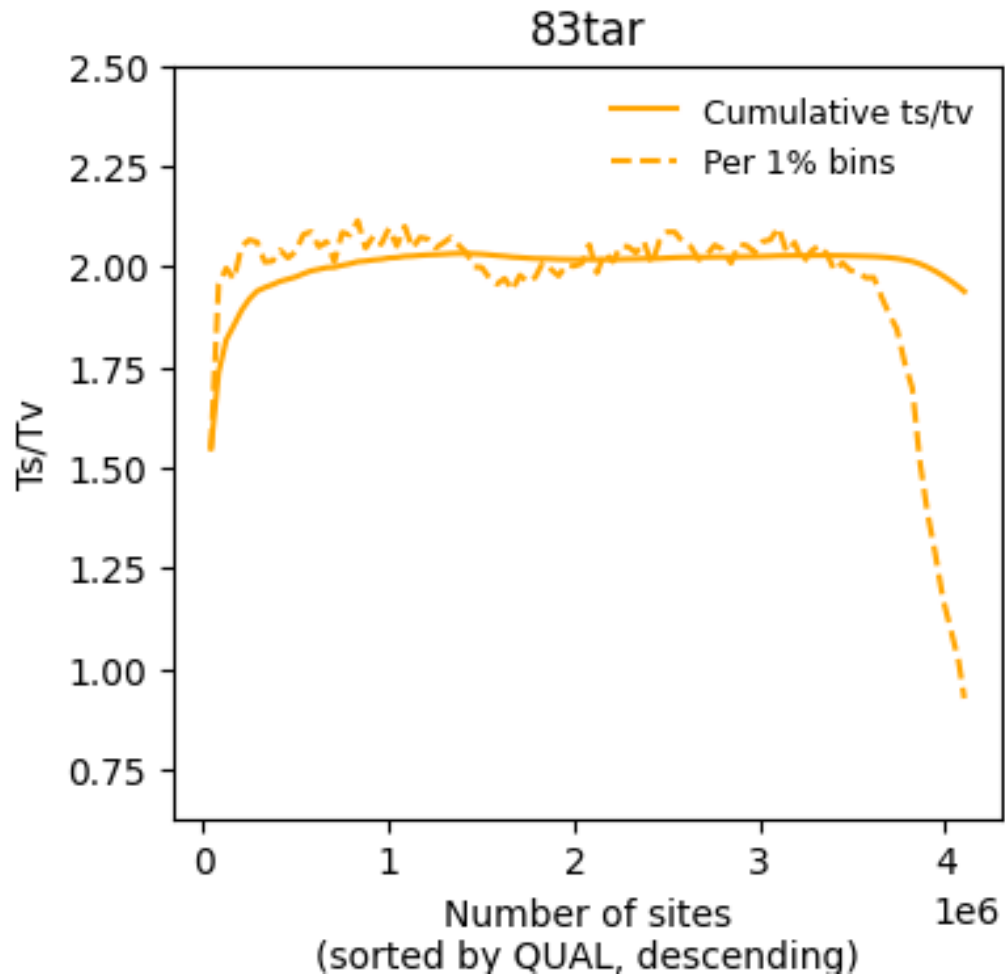# Singletons by sample (hets and homs)
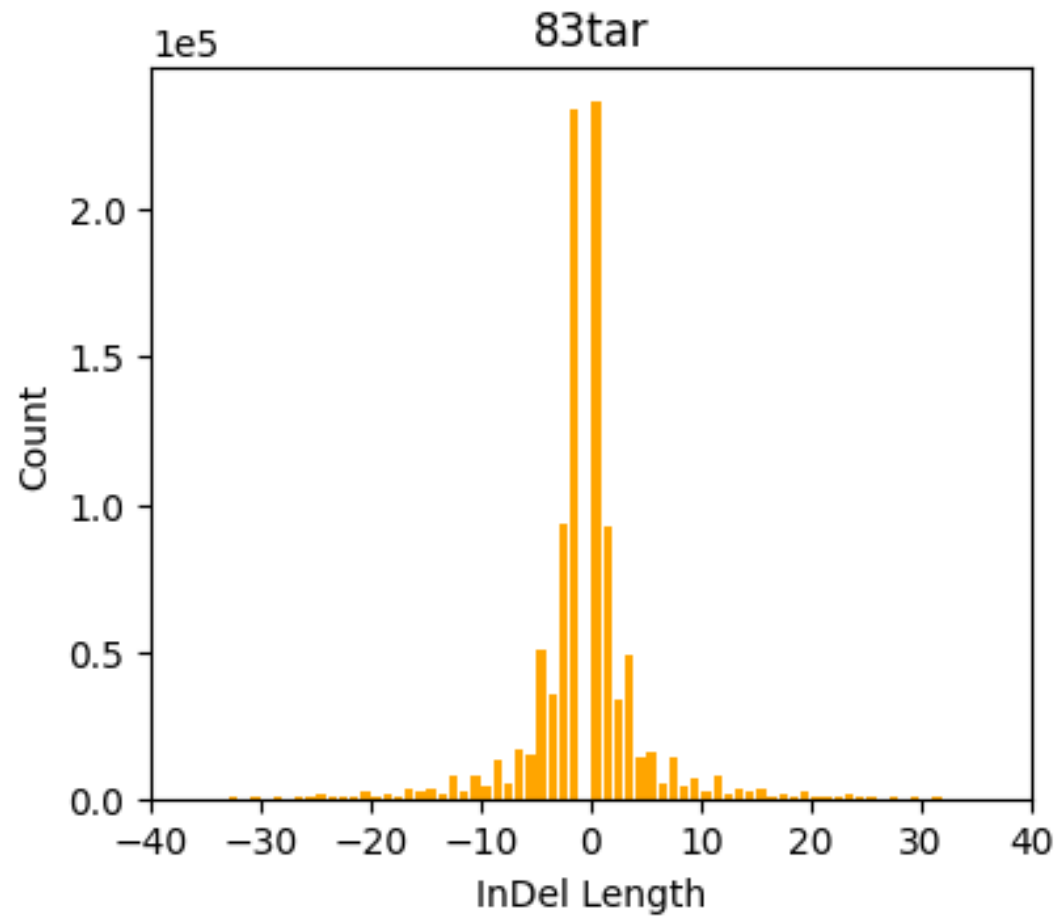
Average depth by sample
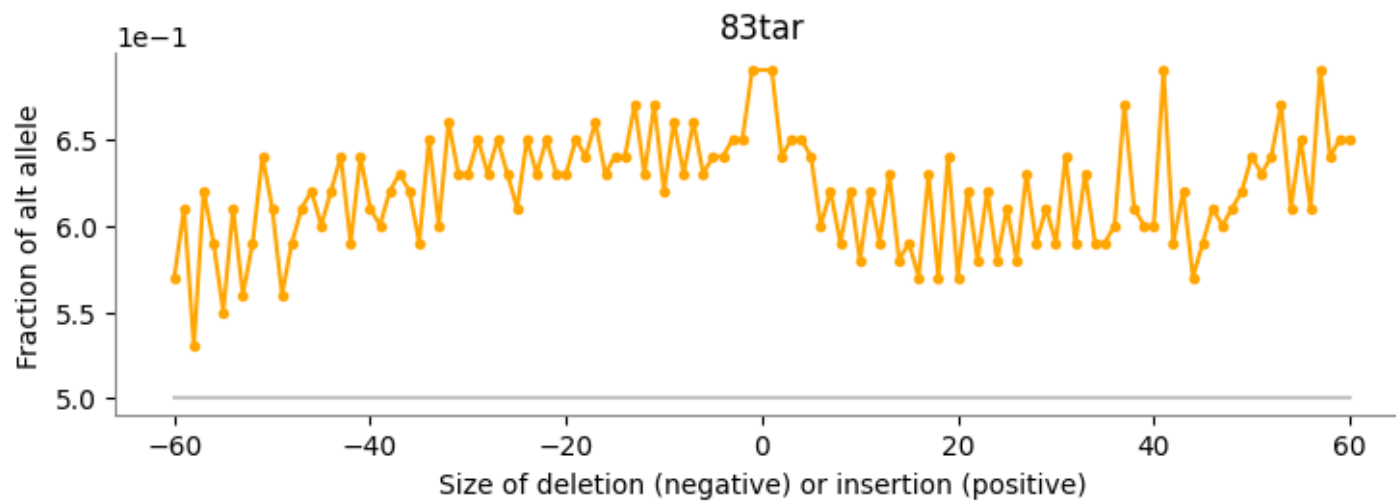
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

Indel distribution

83tar

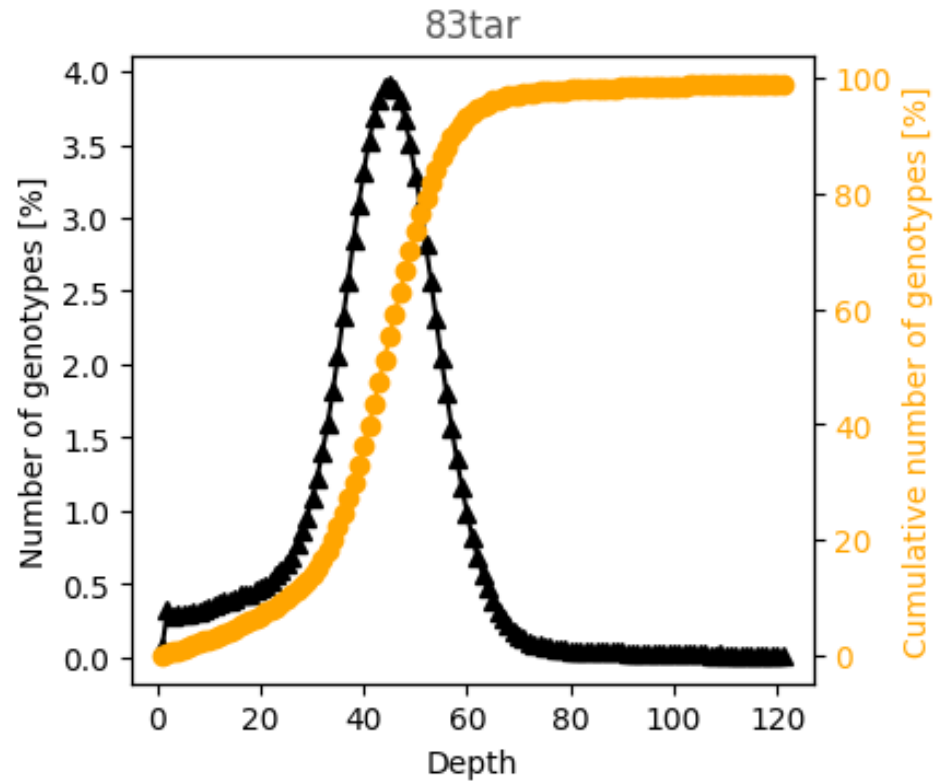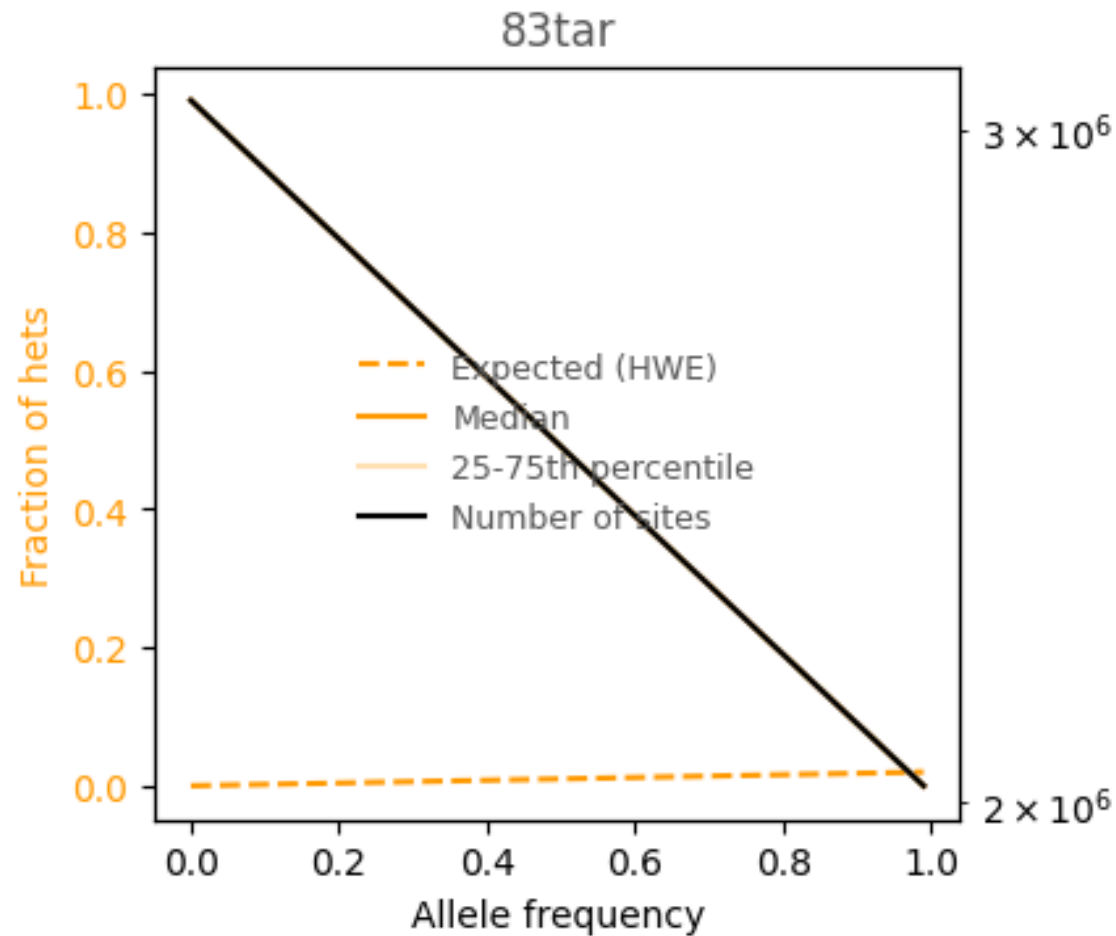# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types