Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
84ome	4,211,506	1.94	1.94	980,546	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
84ome	65.0%	1.91	70.1%	107,769	2,274

- 84ome .. `/ngc/projects2/gm/data/archive/2022/variants/snv/84omermif-103874678533-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220525_A01176_BHJ5LVDSX3-RHGM_LABKA_WGSA`
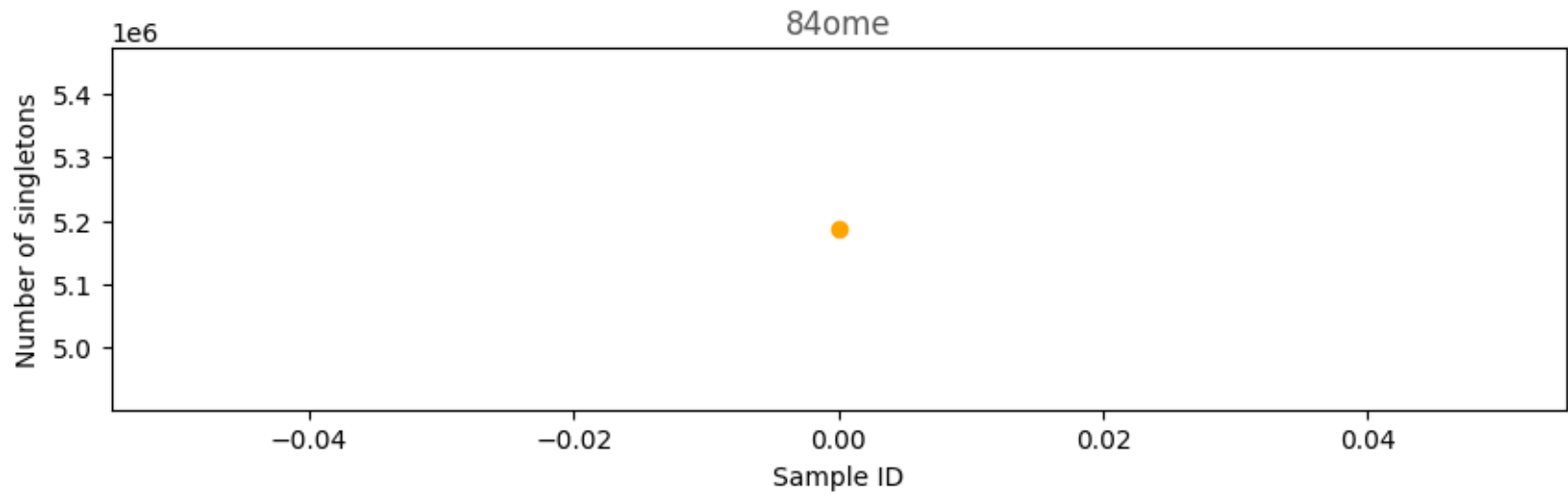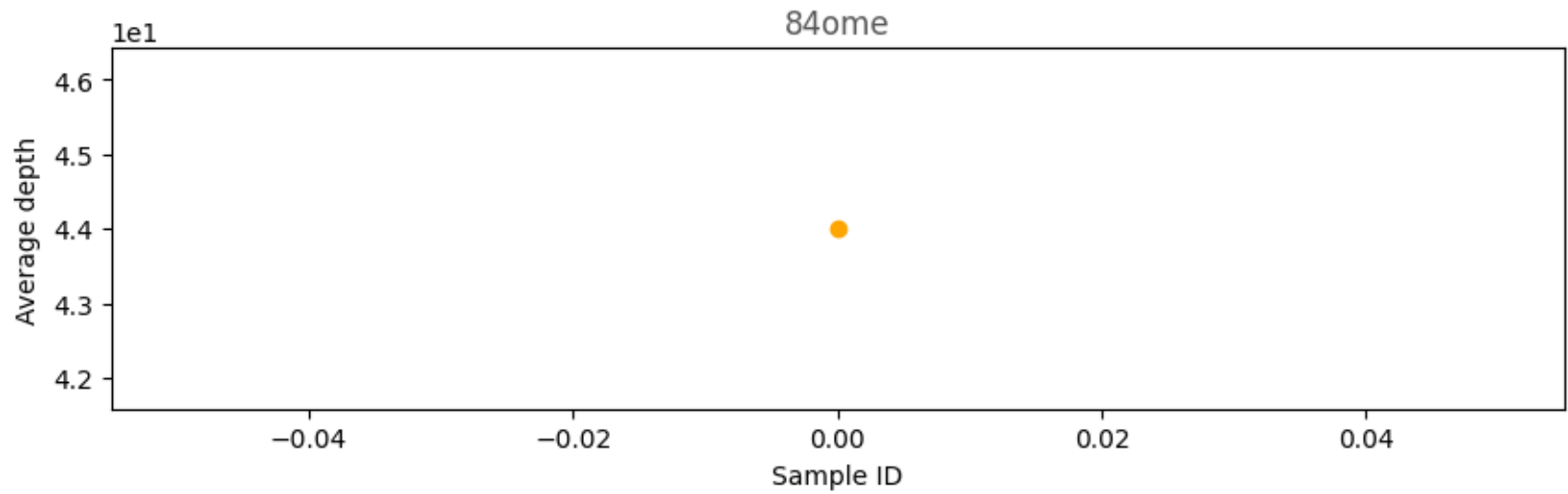 `KUT-WGSAKUT04357_22RKG012687x01_snv_germline_raw.haplotype_caller.vcf.gz`

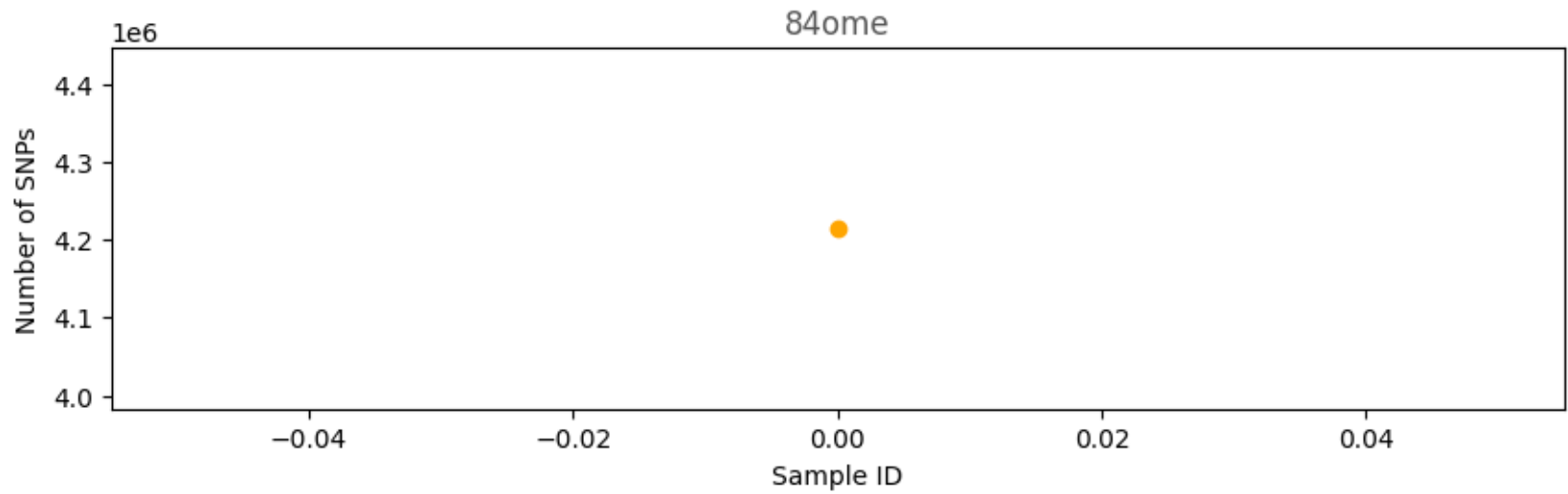# Ts/Tv by sample

84ome
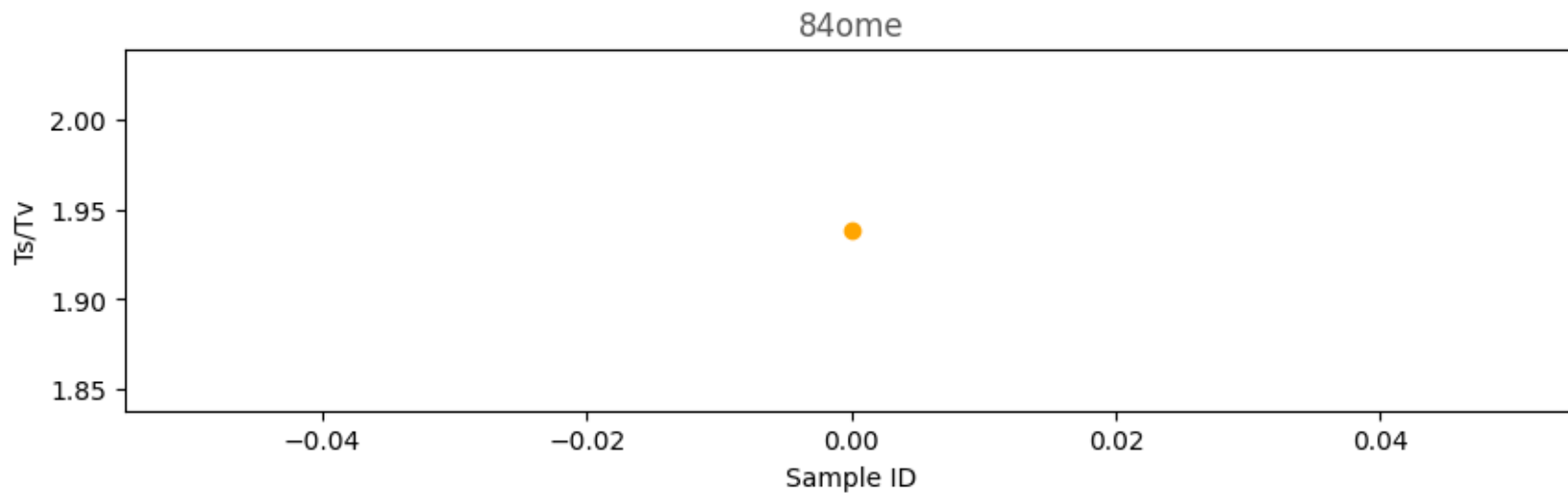
Hets vs non-ref Homs by sample
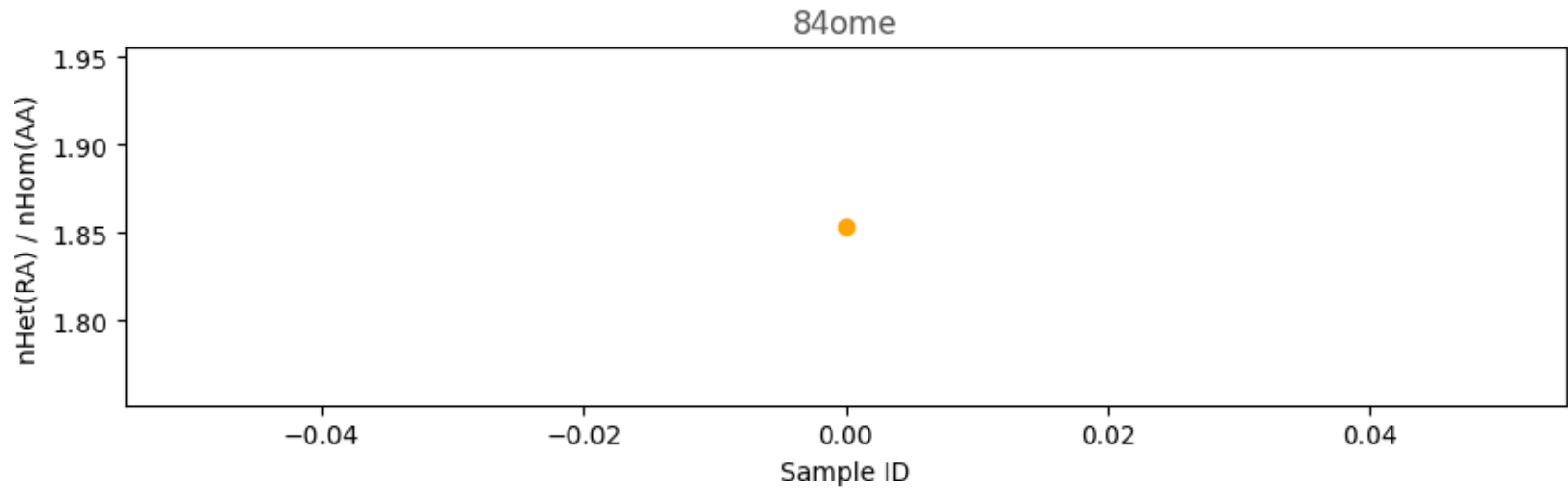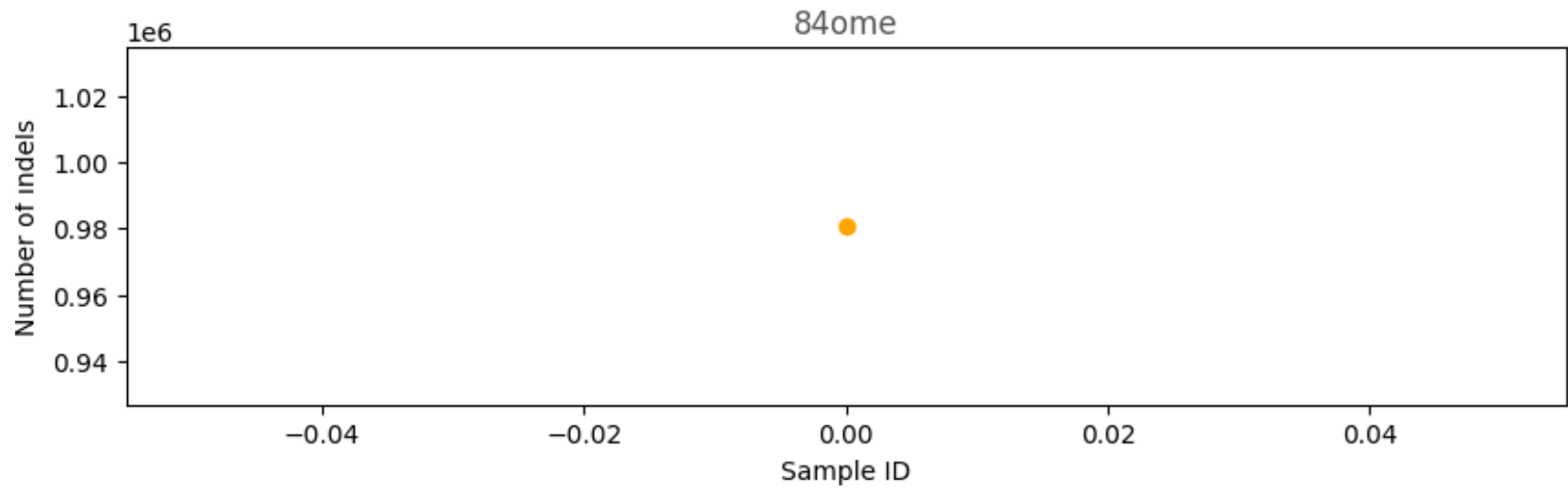
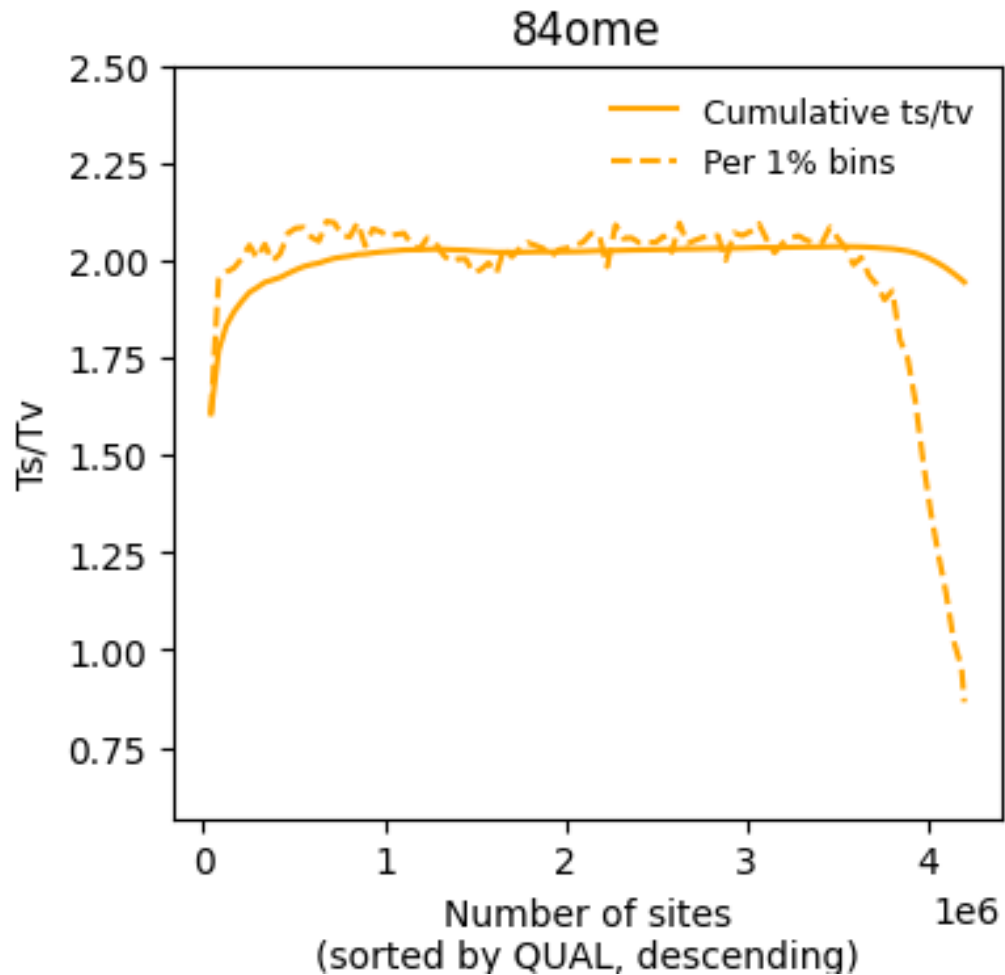# Singletons by sample (hets and homs)
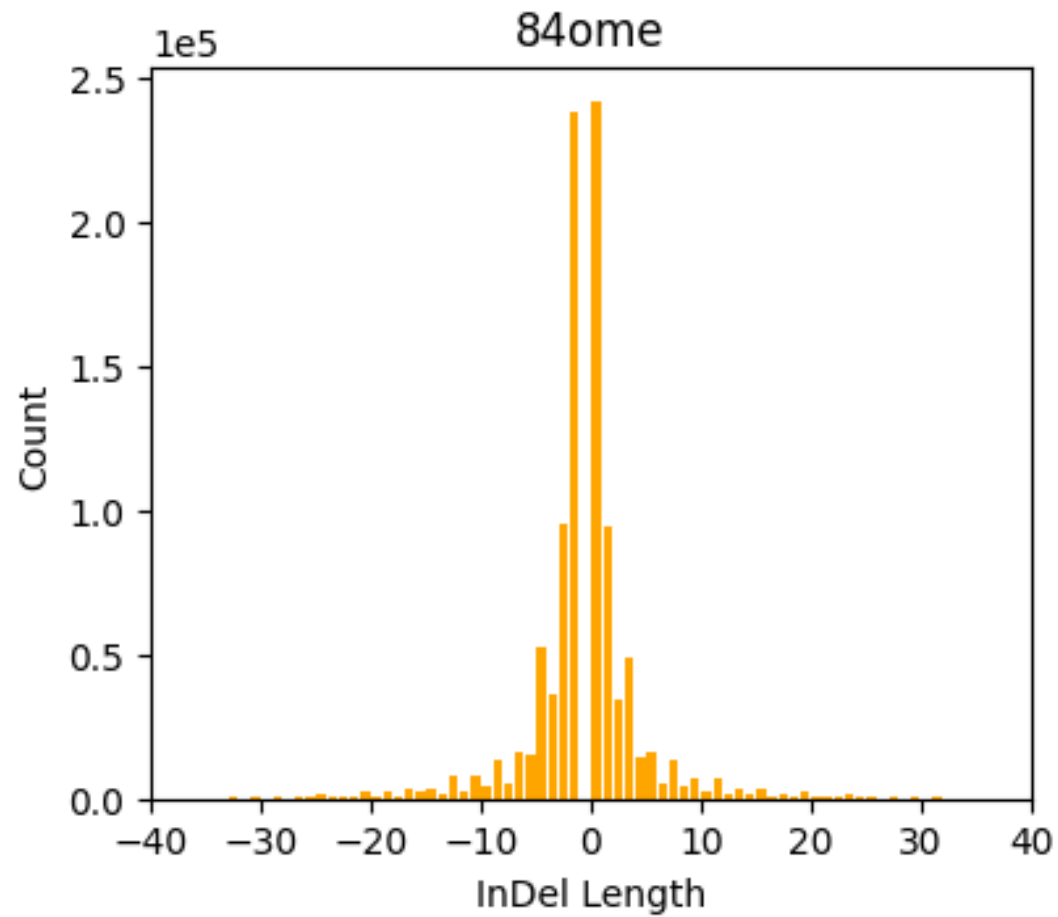
Average depth by sample
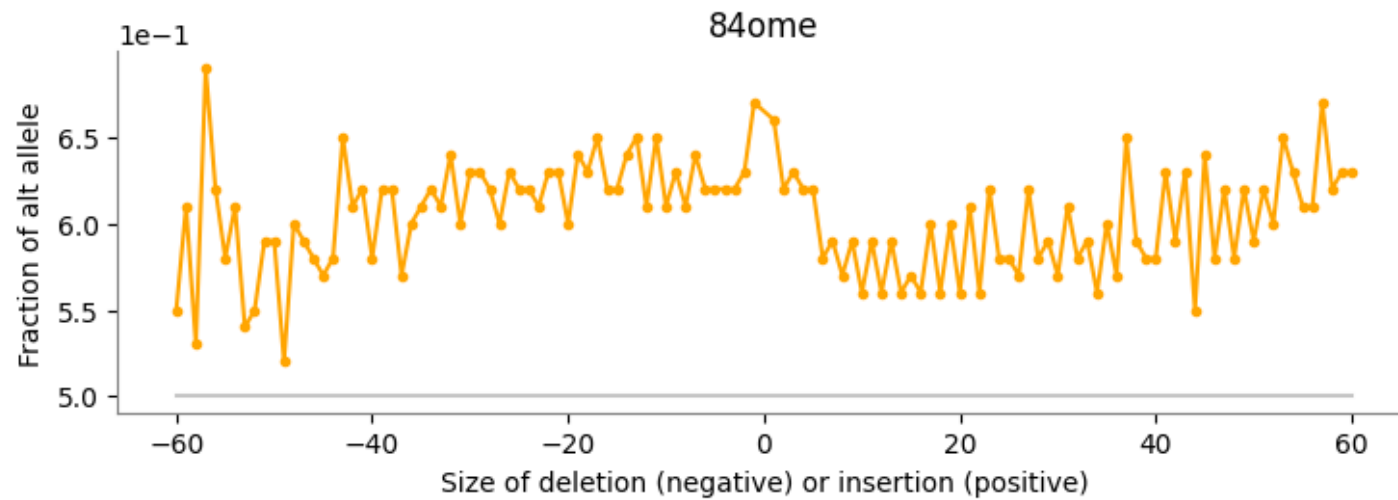
Number of SNPs by sample

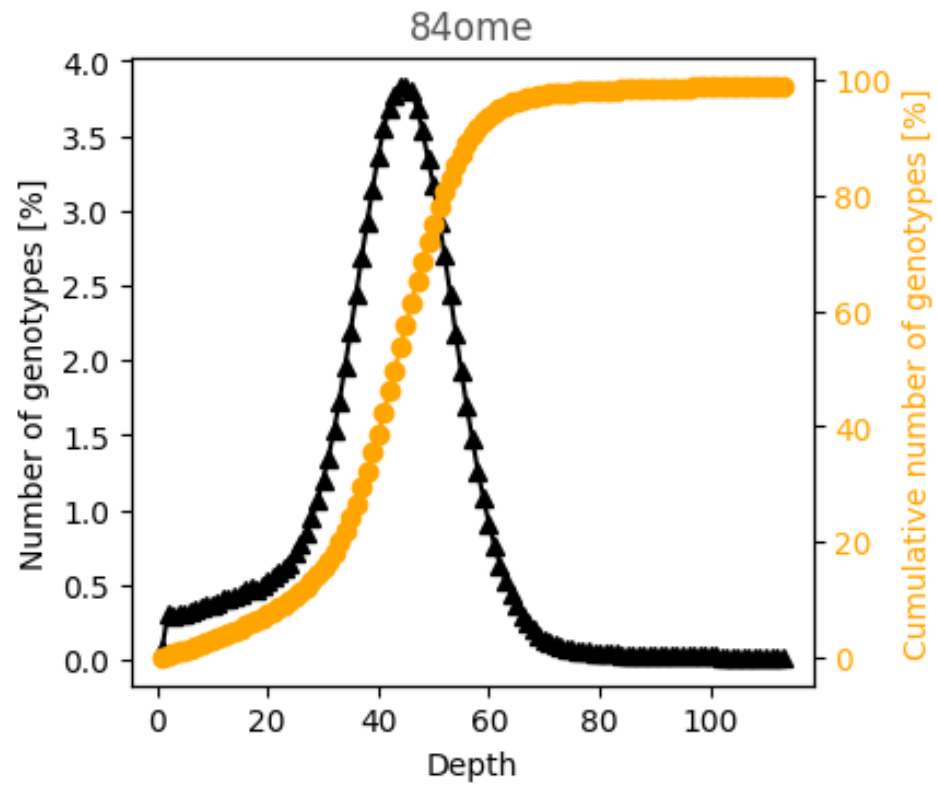# Number of indels by sample

Ts/Tv stratified by QUAL

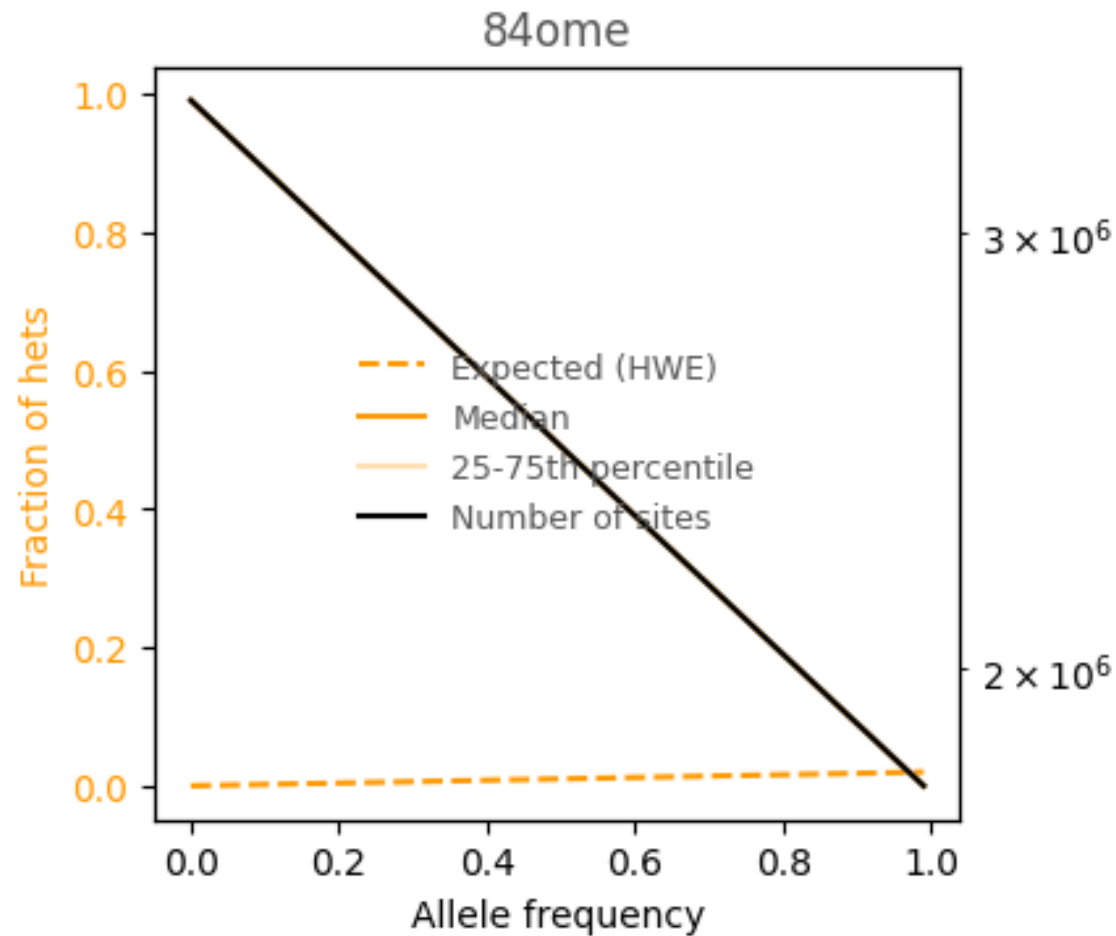

84ome

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types