Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
85mam	4,108,435	1.96	1.96	960,751	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
85mam	62.8%	1.93	67.5%	99,722	1,865

- 85mam .. `/ngc/projects2/gm/data/archive/2022/variants/snv/85mamedtm-103838575240-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00602-220225_A01176_AH7CT5DSX3-EXT_LAB`
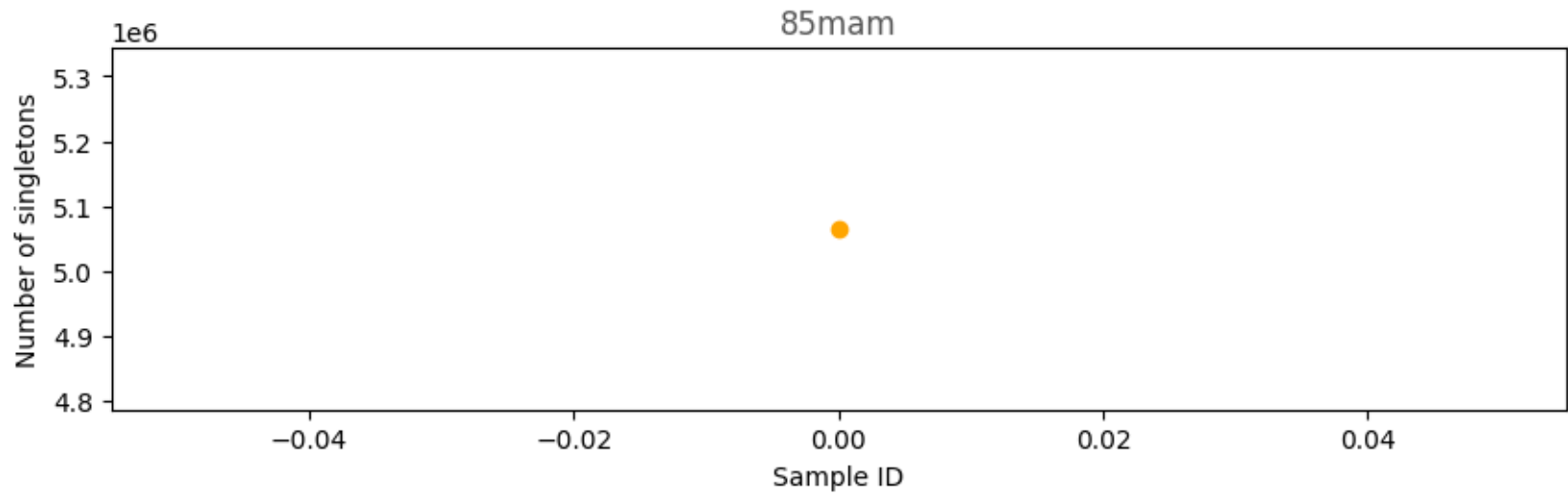 `KA_NGCWGS-NGCWGS03879_22RKG000473x01_snv_germline_raw.haplotype_caller.vcf.gz`
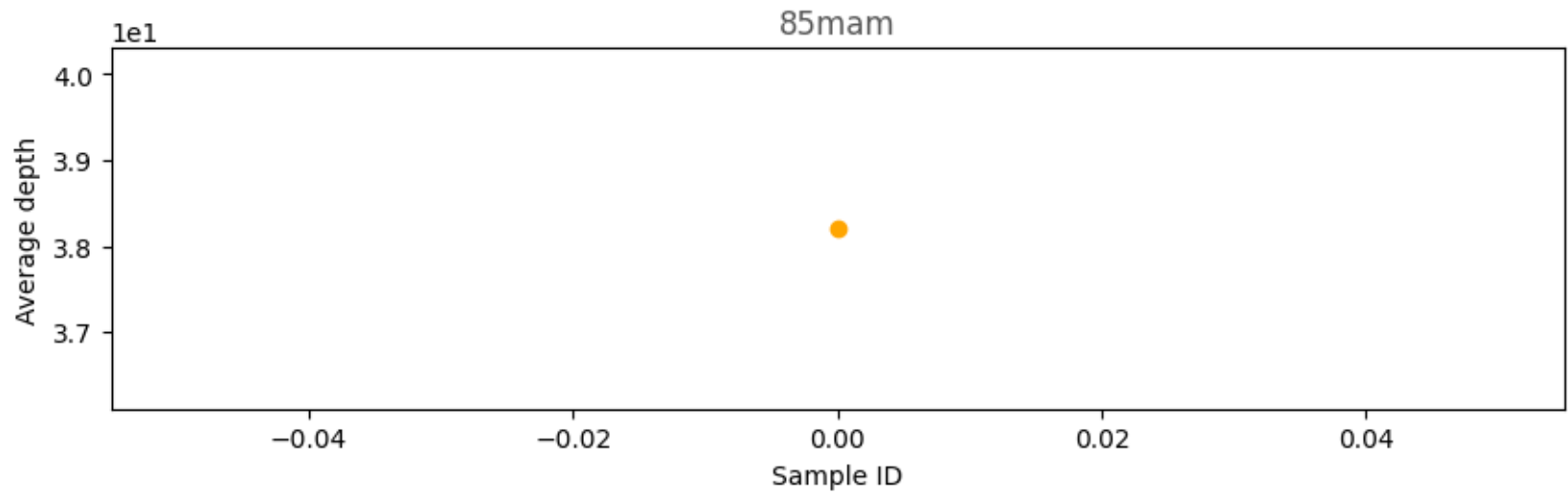
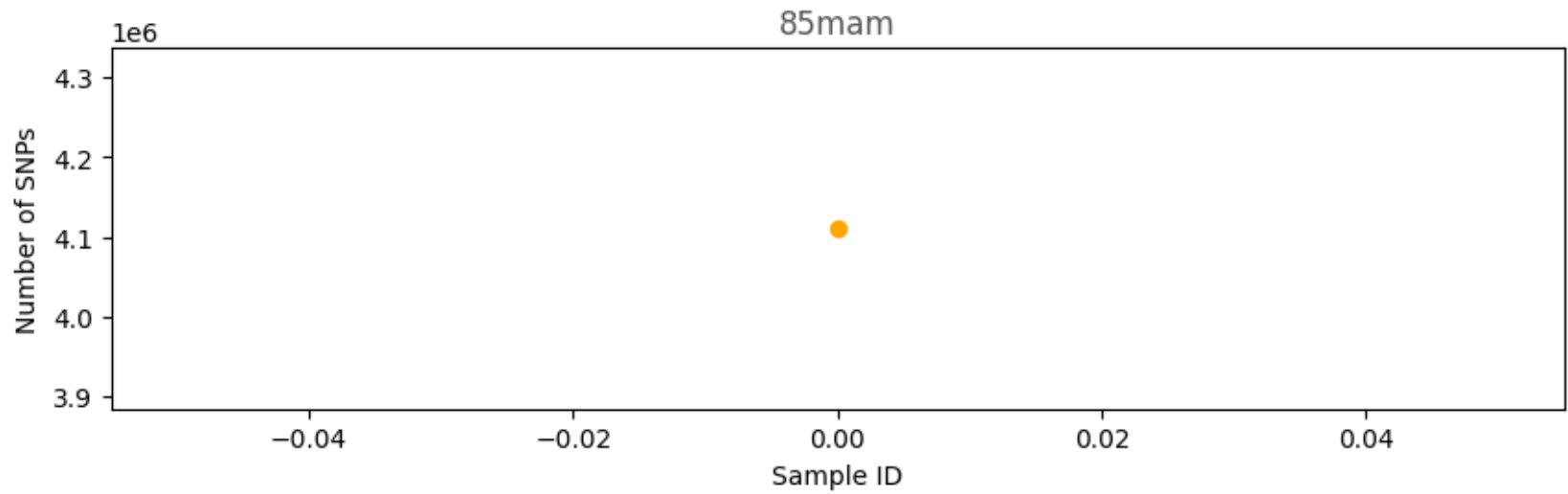# Ts/Tv by sample
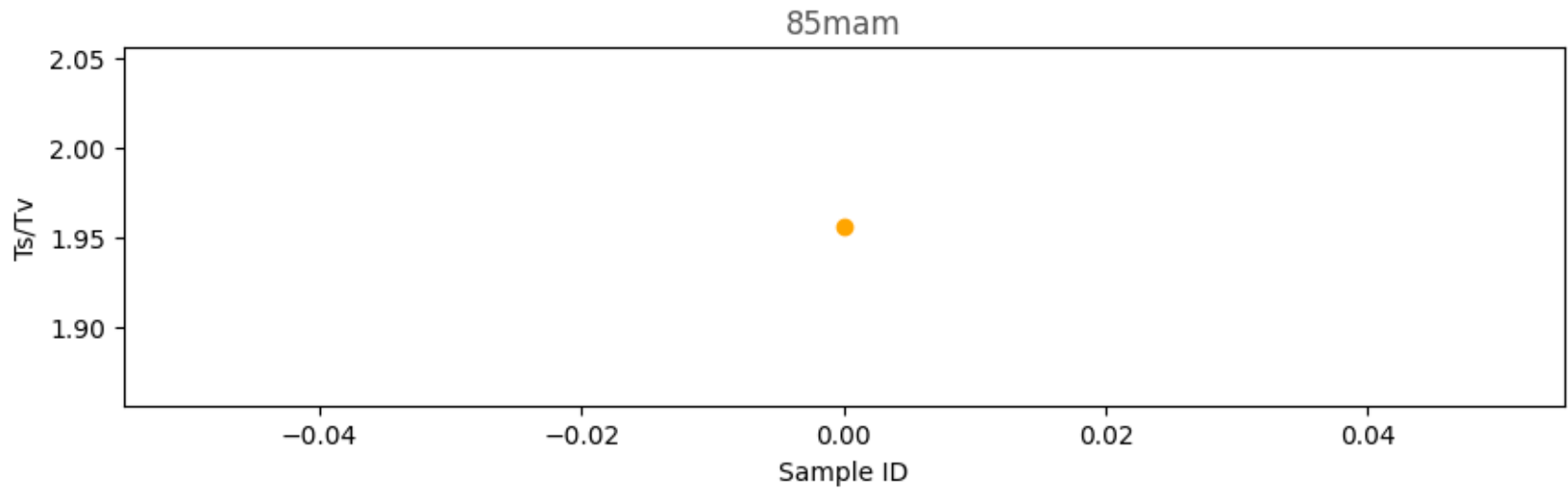
Hets vs non-ref Homs by sample
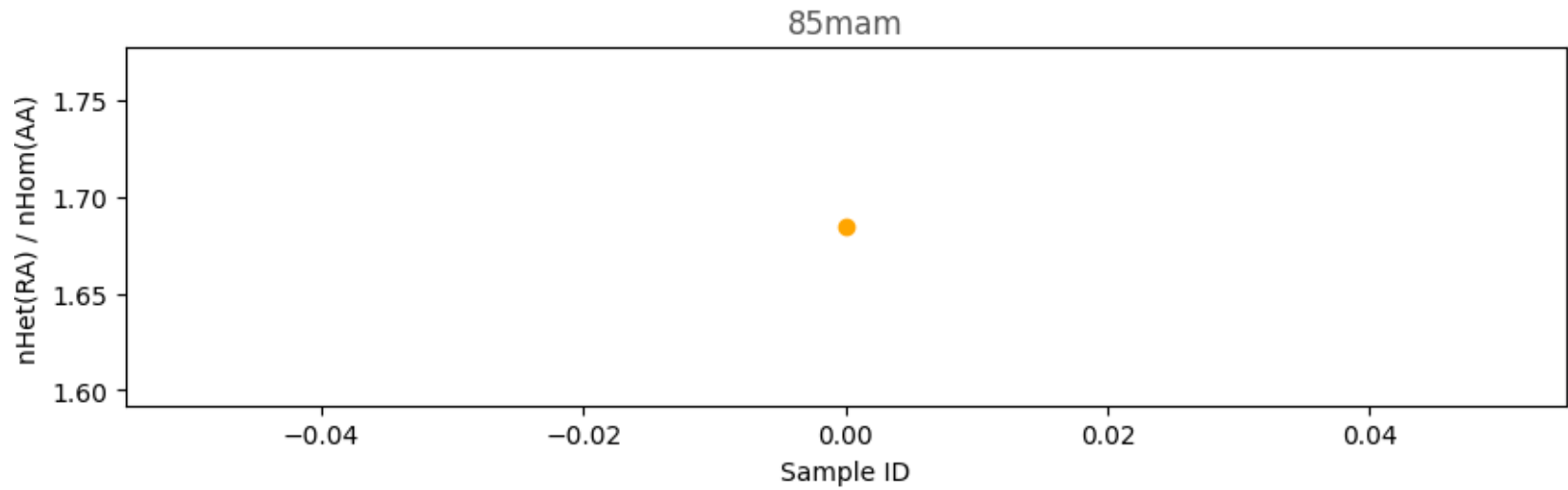
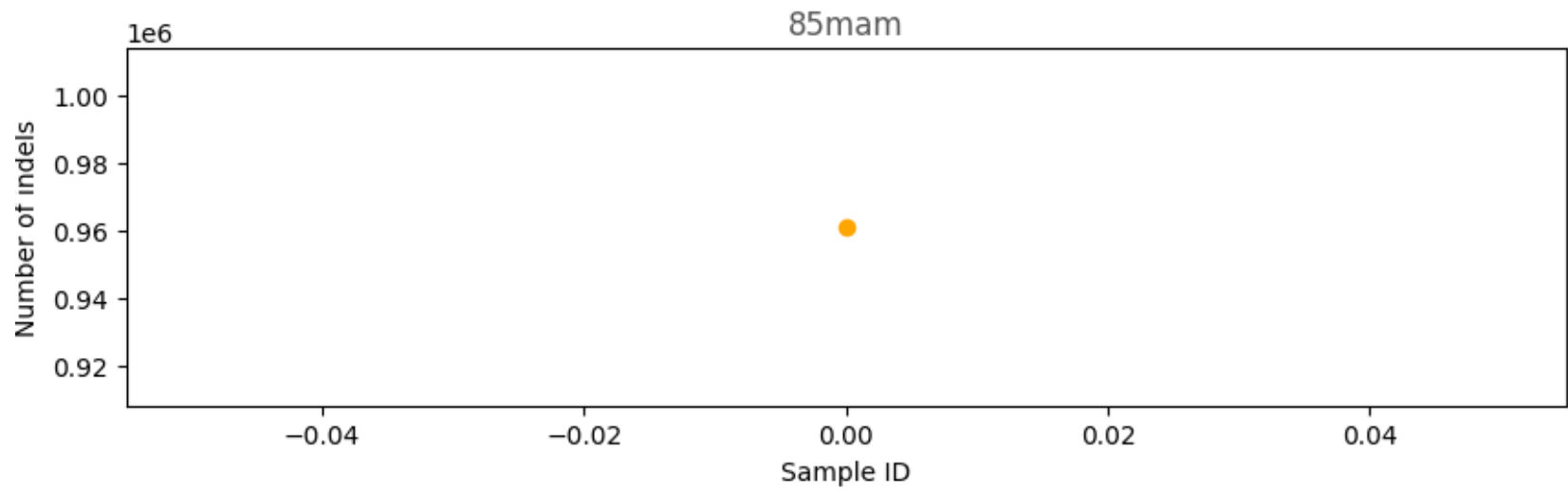# Singletons by sample (hets and homs)
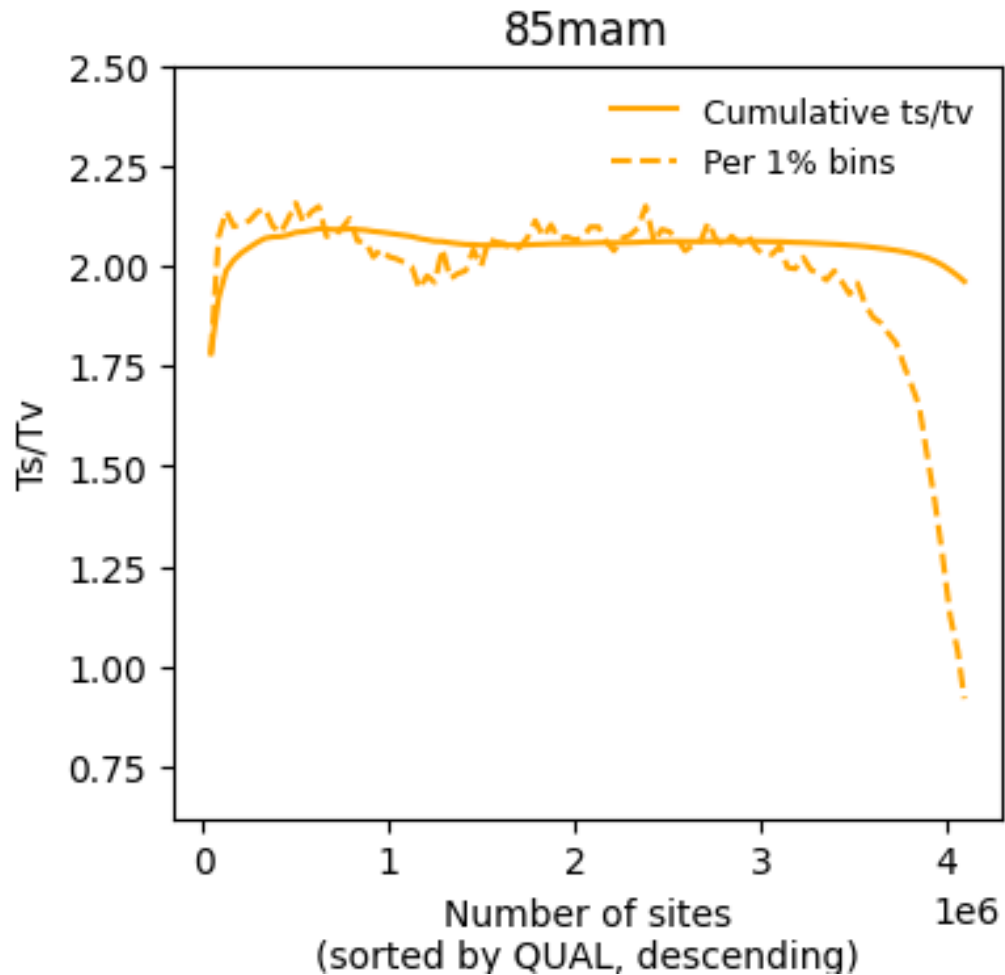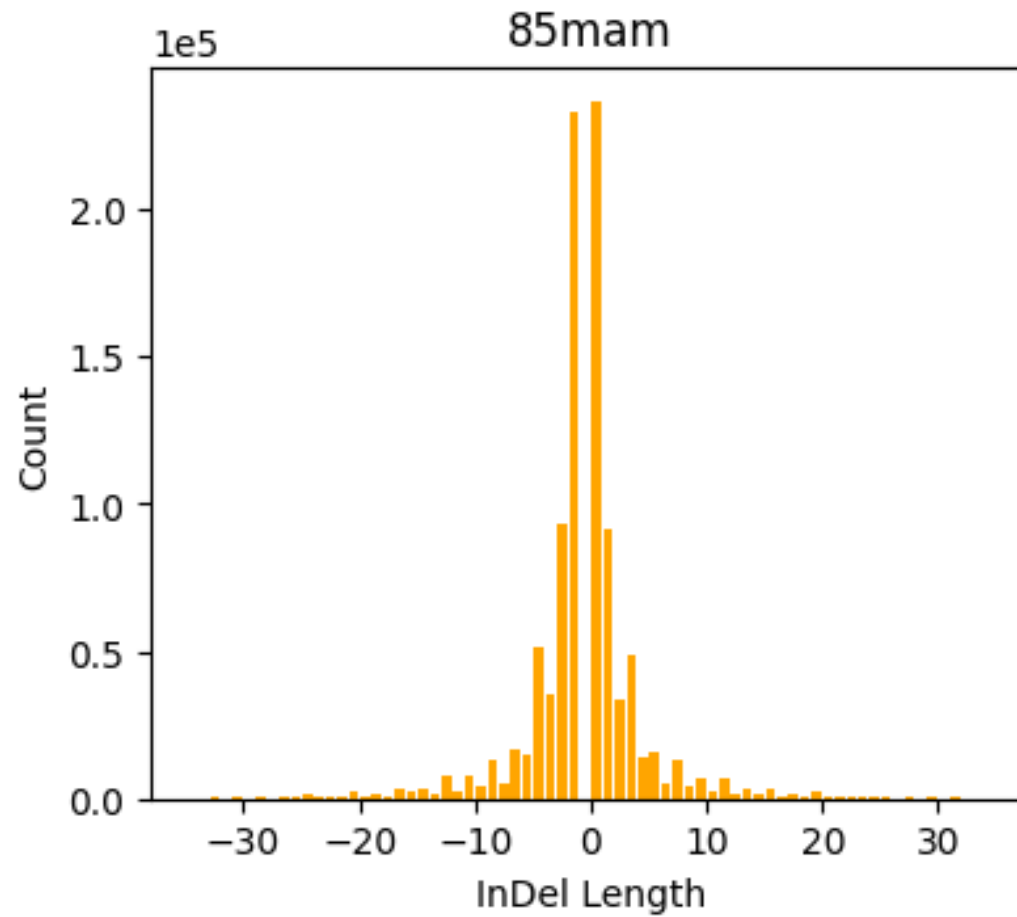
Average depth by sample
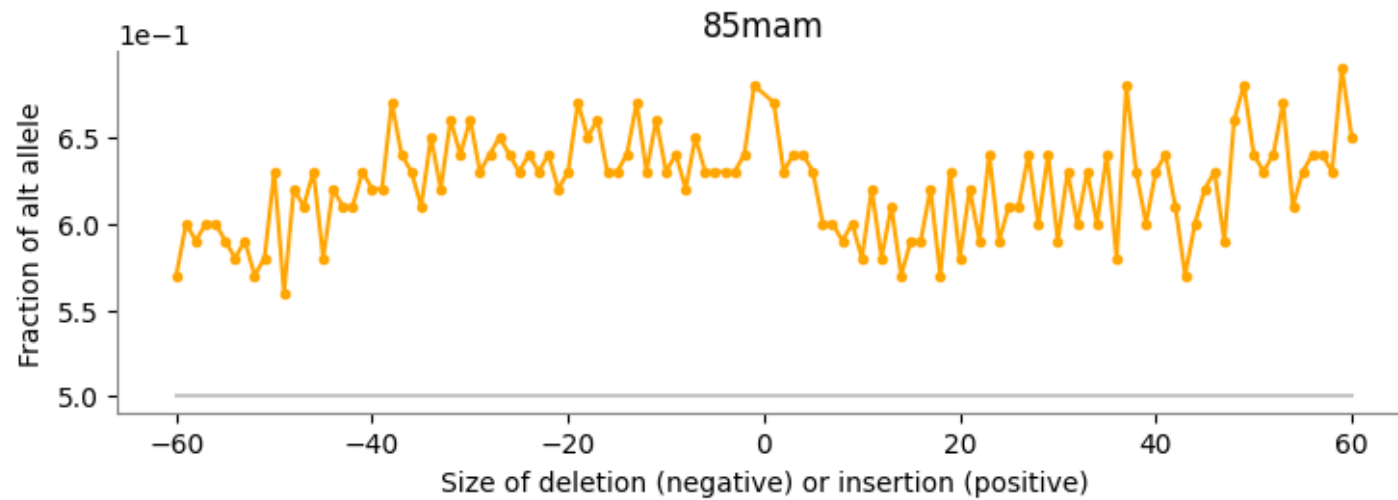
Number of SNPs by sample

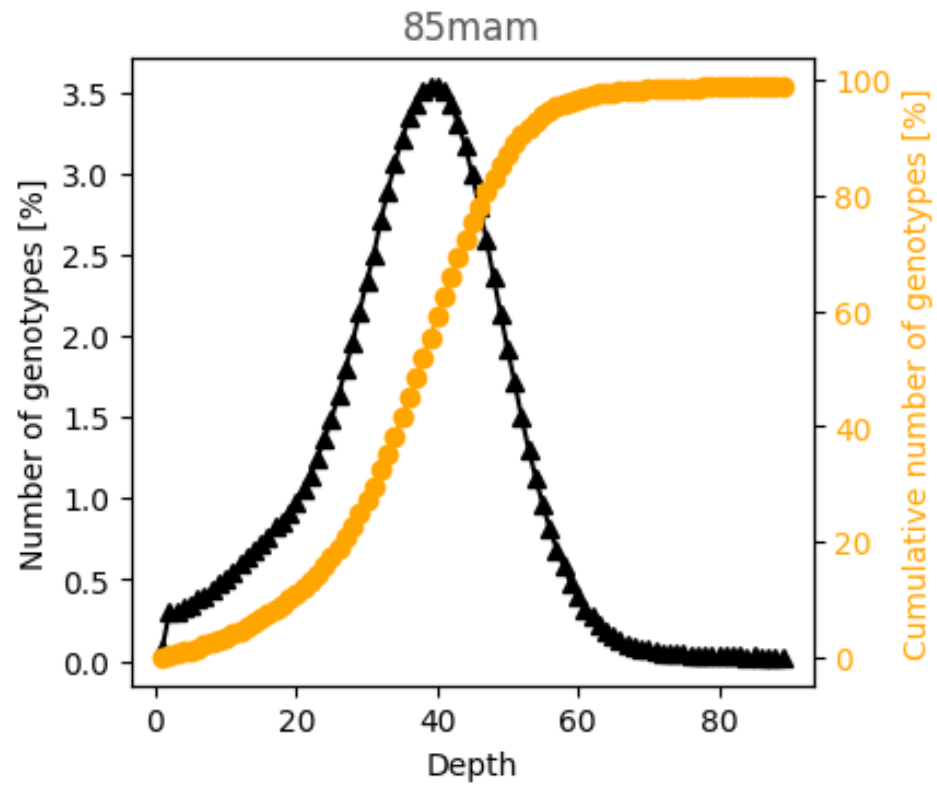# Number of indels by sample
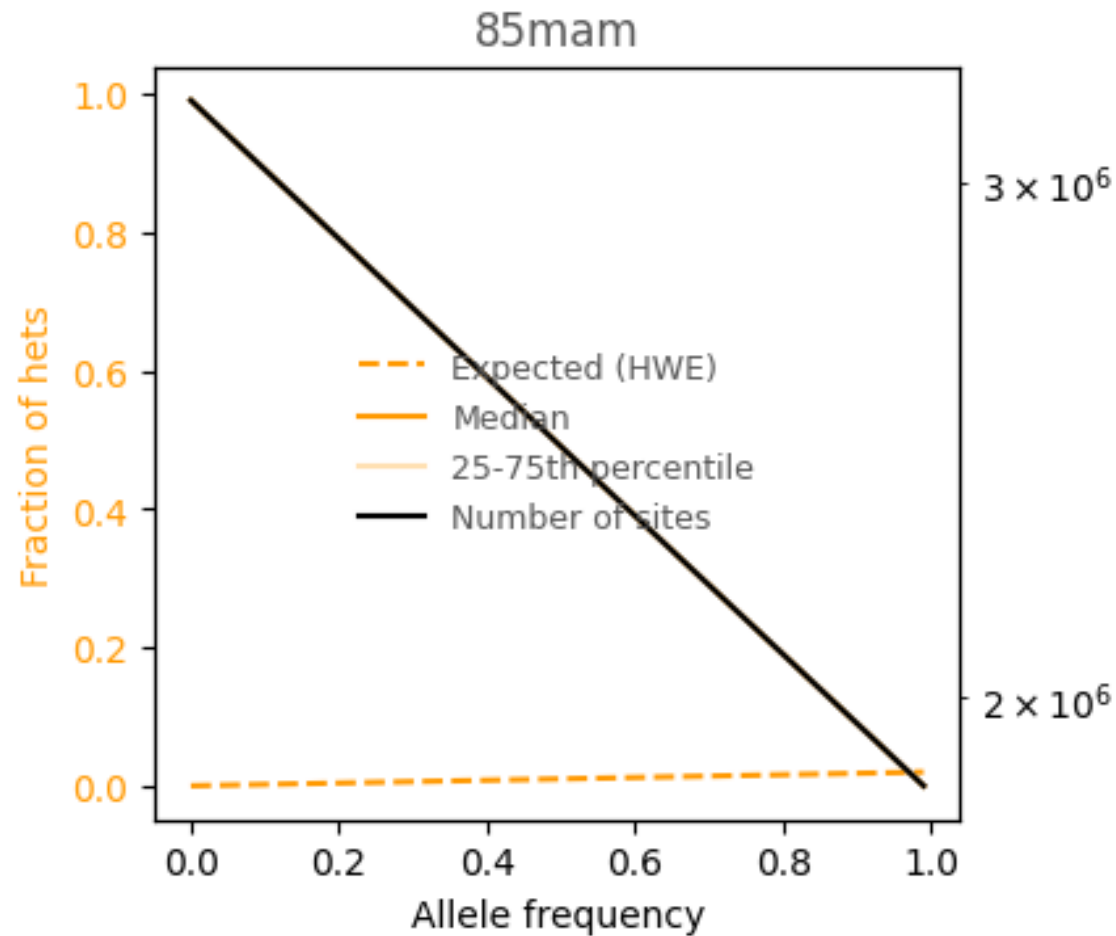
Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

85mam

Substitution types