

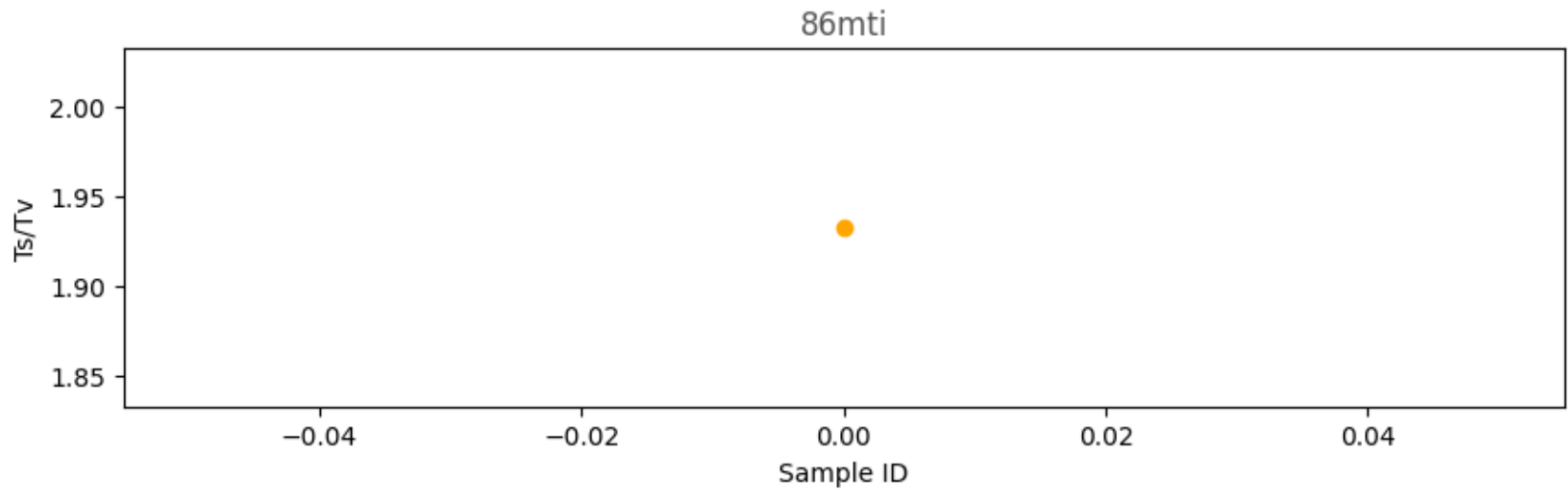
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
86mti	4,045,640	1.93	1.94	943,414	–	0	0
* frameshift ratio: out/(out+in)							

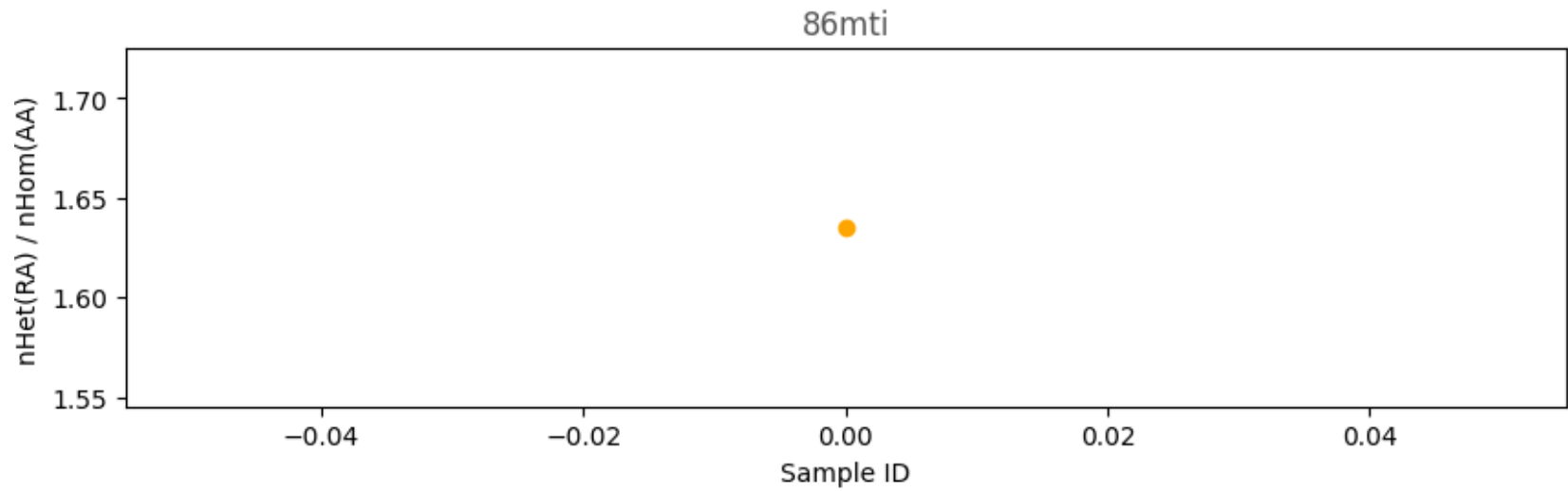
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
86mti	62.1%	1.90	66.5%	97,185	2,038

- 86mti .. /ngc/projects2/gm/data/archive/2022/variants/snv/86mtititm-103819278758-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211222_A01466_AH2CJCDSX3-RHGM_LABKA_WGSA
 KUT-WGSAKUT03554_21RKG031286x01_snv_germline_raw.haplotype_caller.vcf.gz

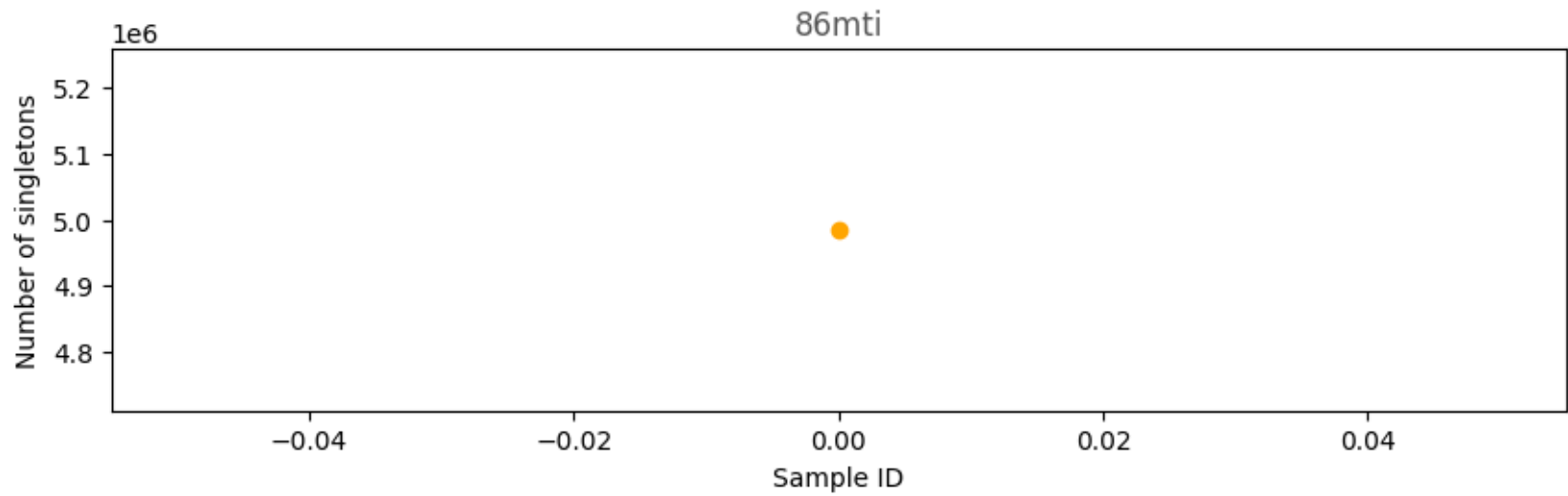
Ts/Tv by sample



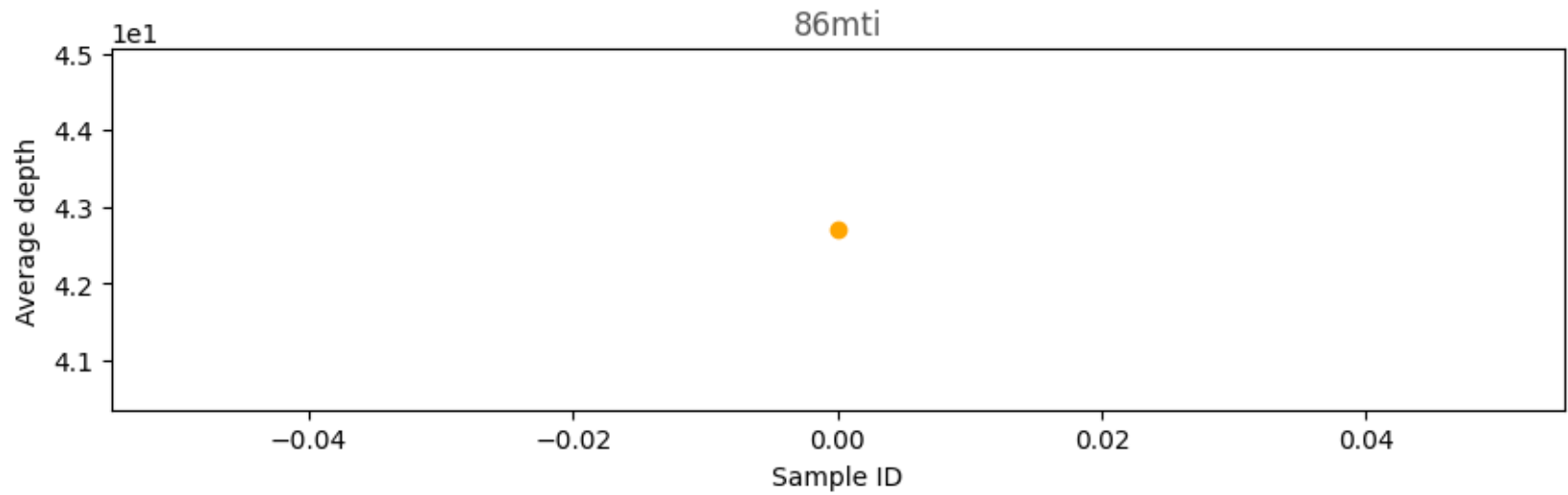
Hets vs non-ref Homs by sample



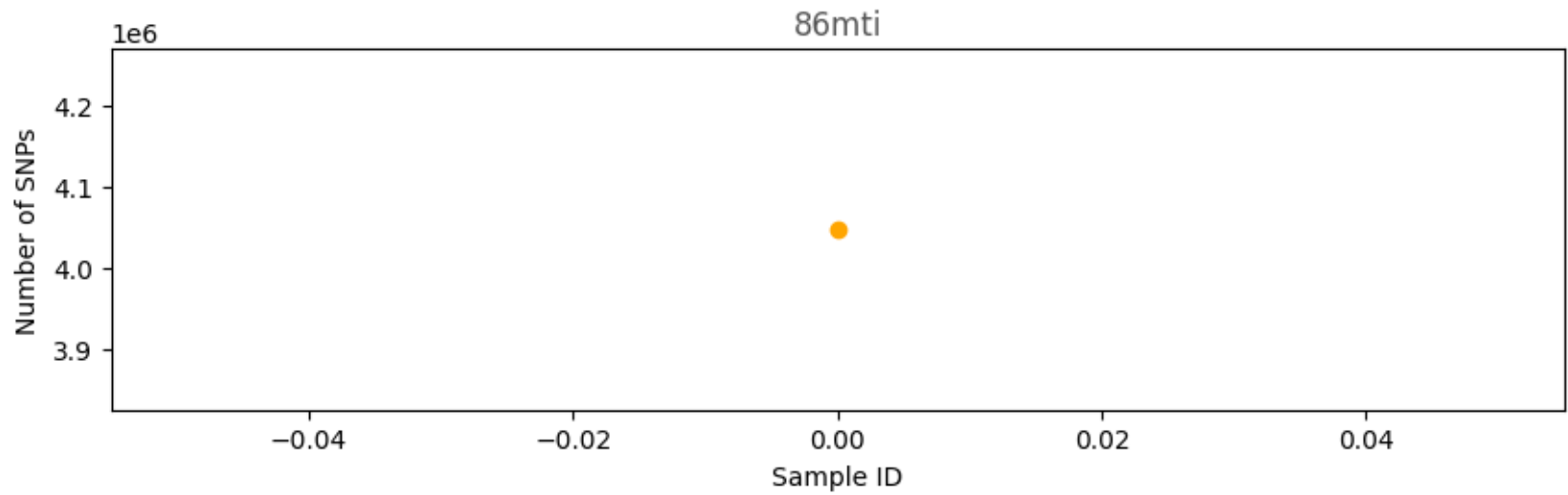
Singletons by sample (hets and homs)



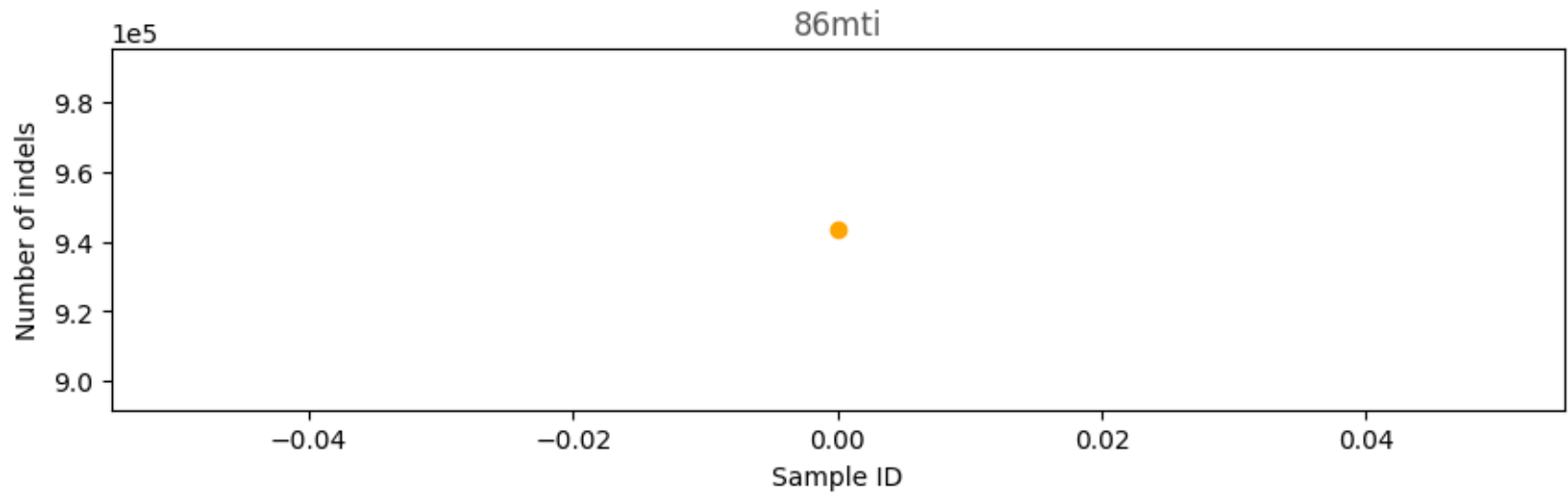
Average depth by sample



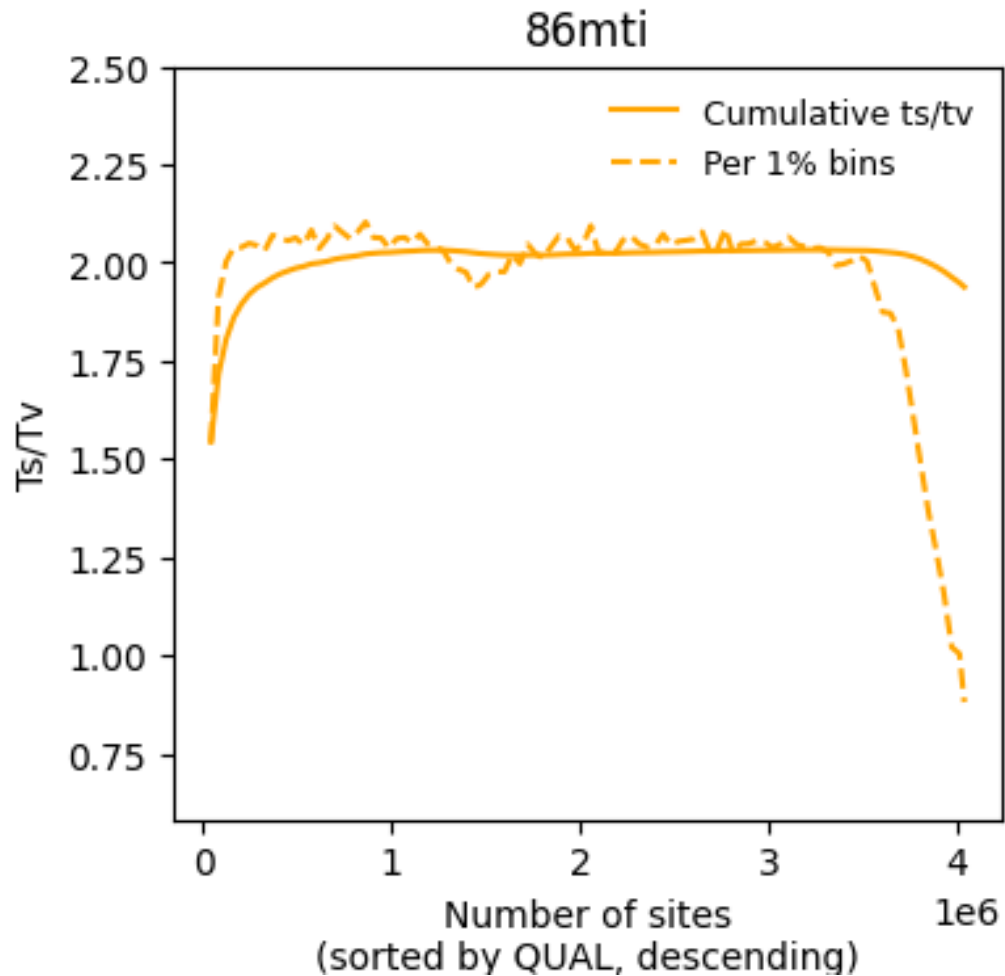
Number of SNPs by sample



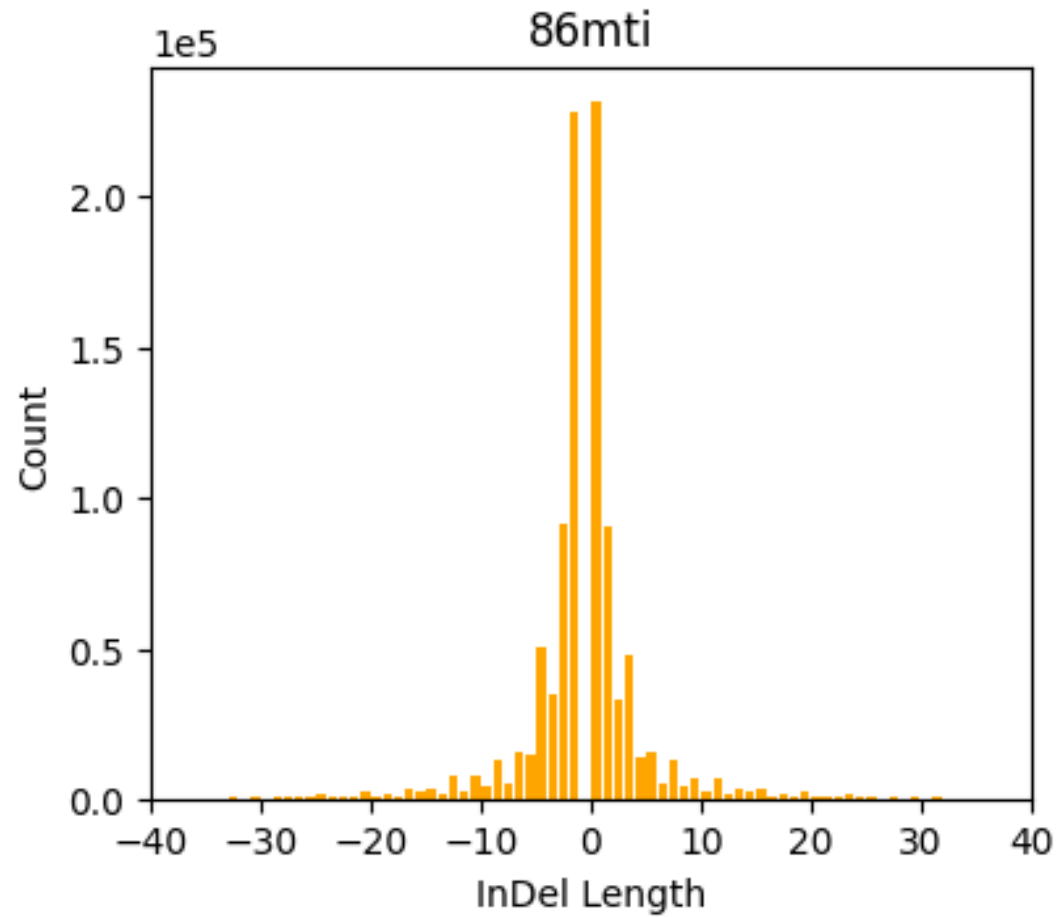
Number of indels by sample



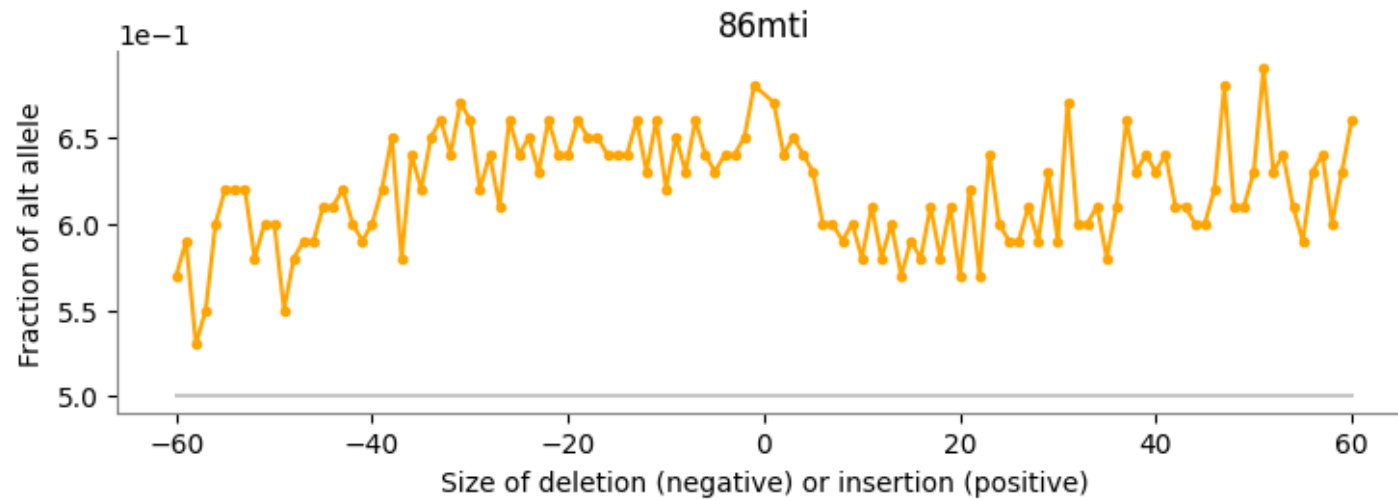
Ts/Tv stratified by QUAL



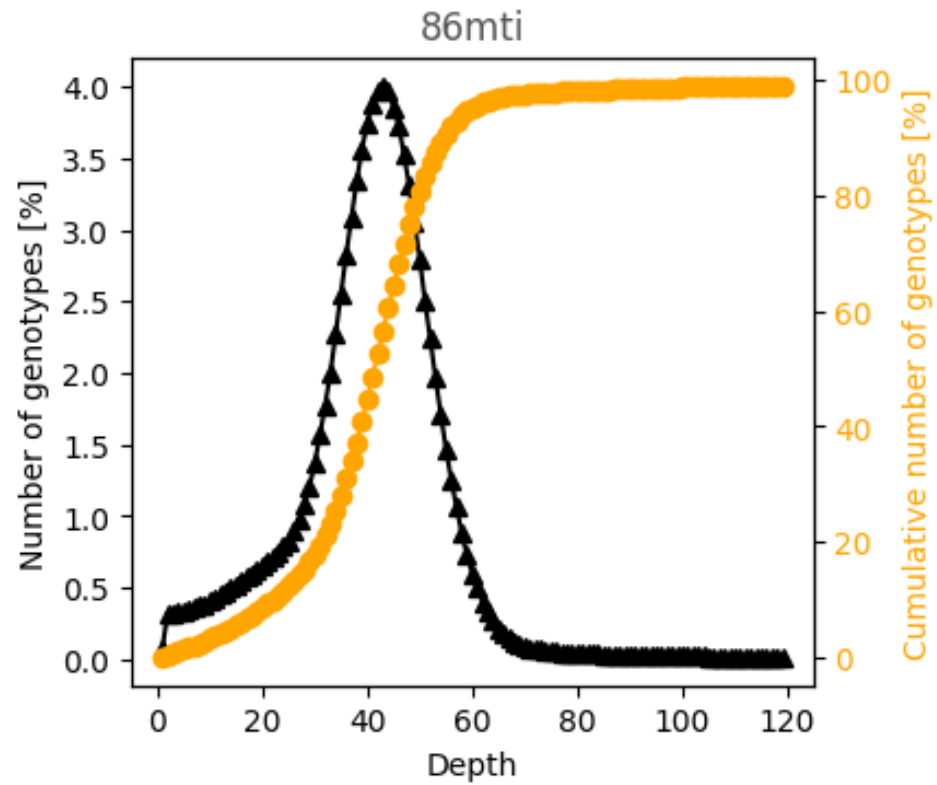
Indel distribution



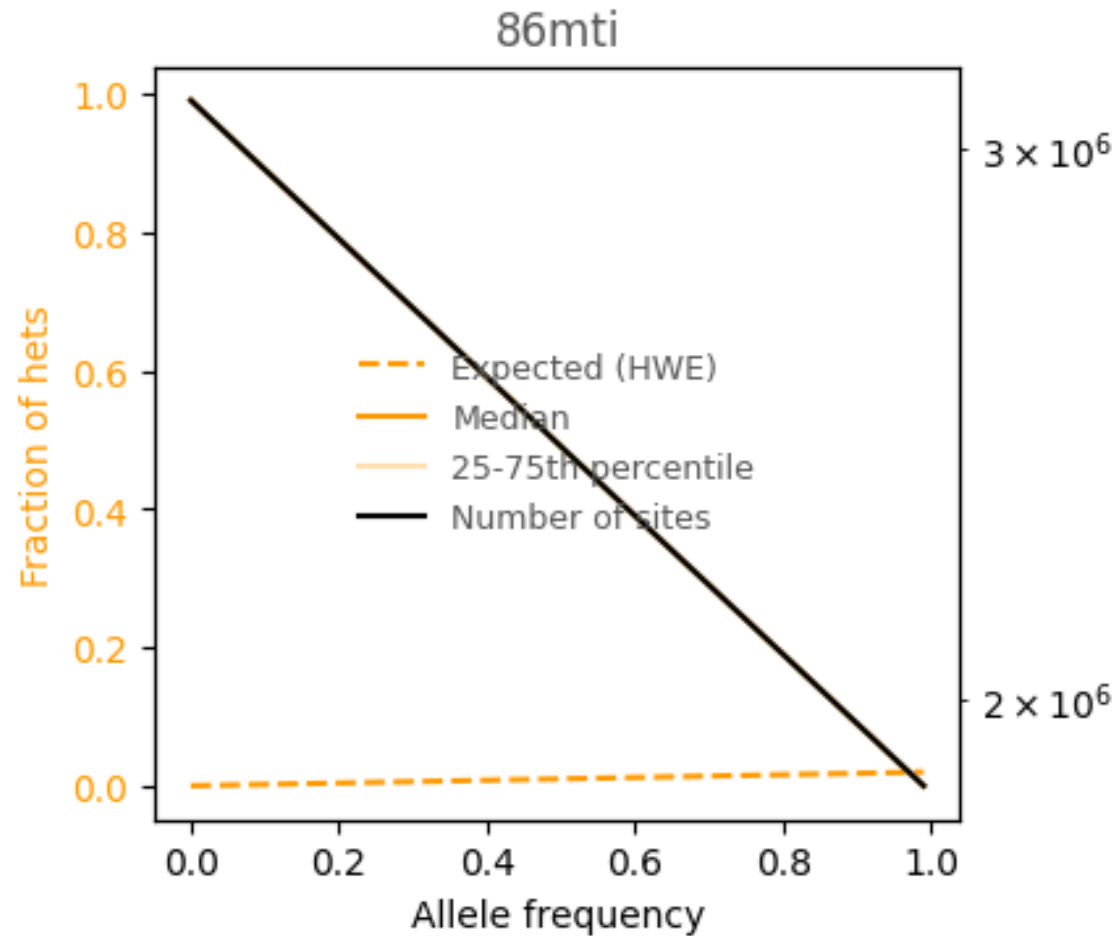
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

