

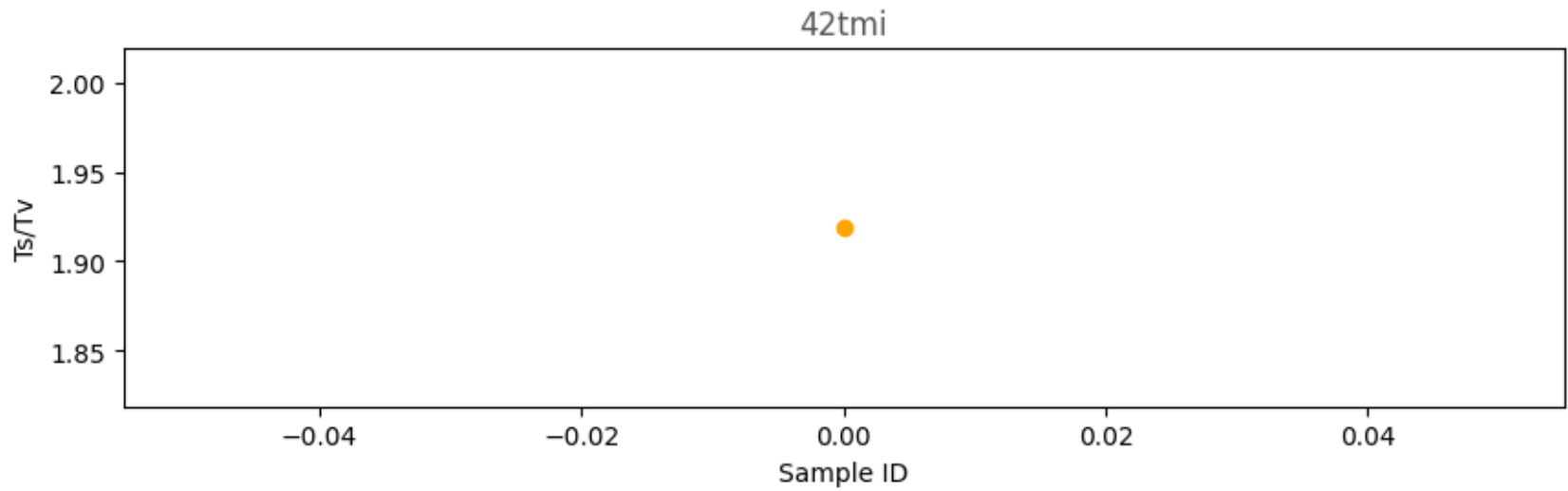
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
42tmi	4,101,541	1.92	1.92	959,754	–	0	0
* frameshift ratio: out/(out+in)							

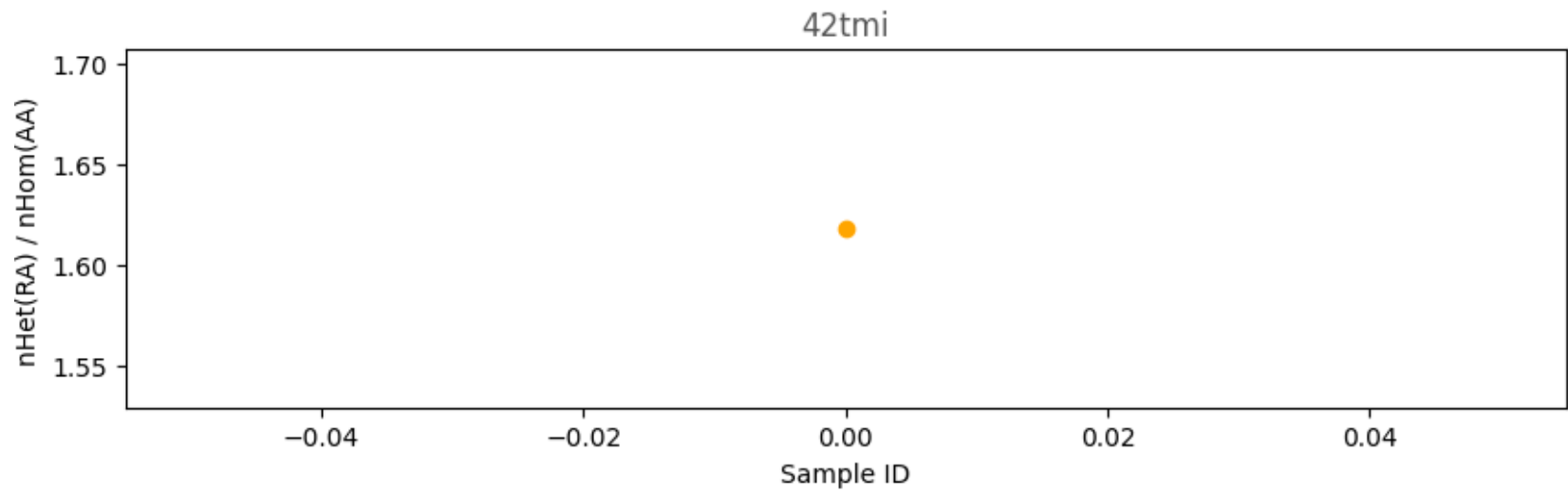
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
42tmi	61.8%	1.88	67.0%	103,559	2,103

- 42tmi .. /ngc/projects2/gm/data/archive/2022/variants/snv/42tmitthm-110292254907-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01961-221012_A00559_AH3FGGDSX5-EXT_LAB
 KA_NGCWGS-NGCWGS05779_snv_germline_raw.haplotype_caller.vcf.gz

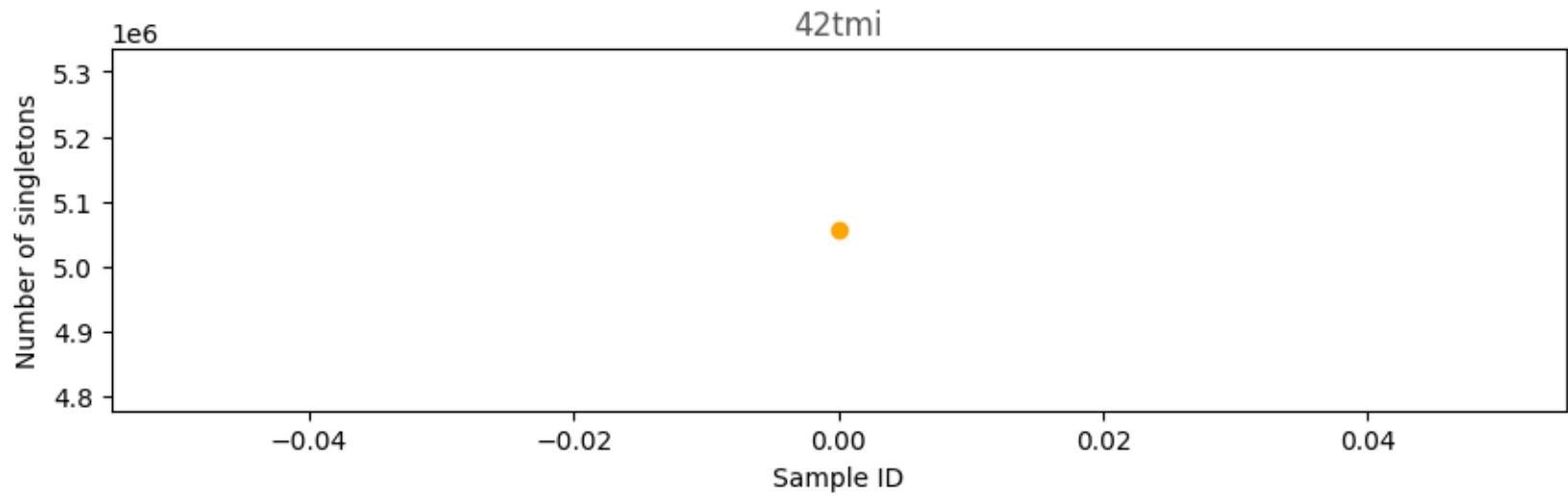
Ts/Tv by sample



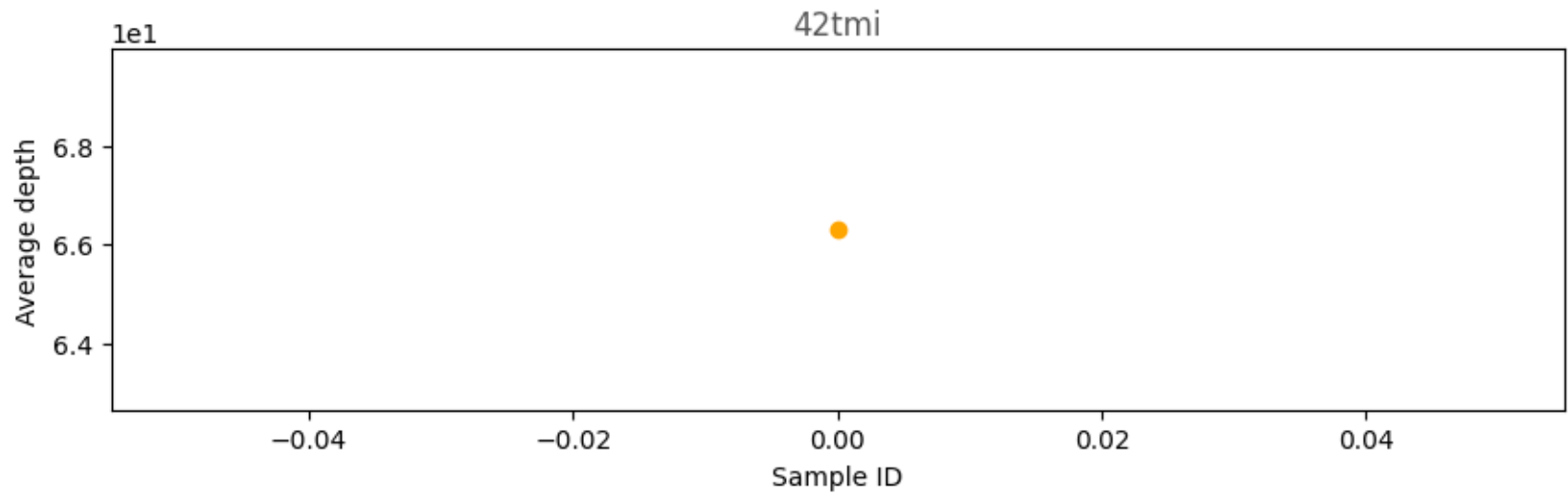
Hets vs non-ref Homs by sample



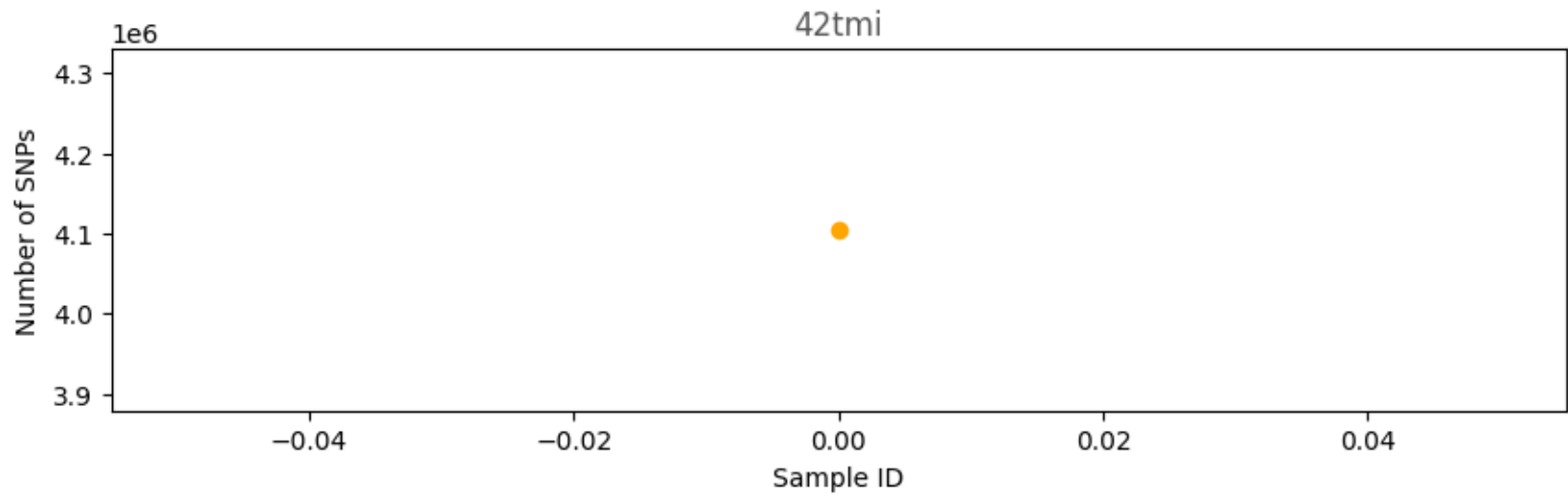
Singletons by sample (hets and homs)



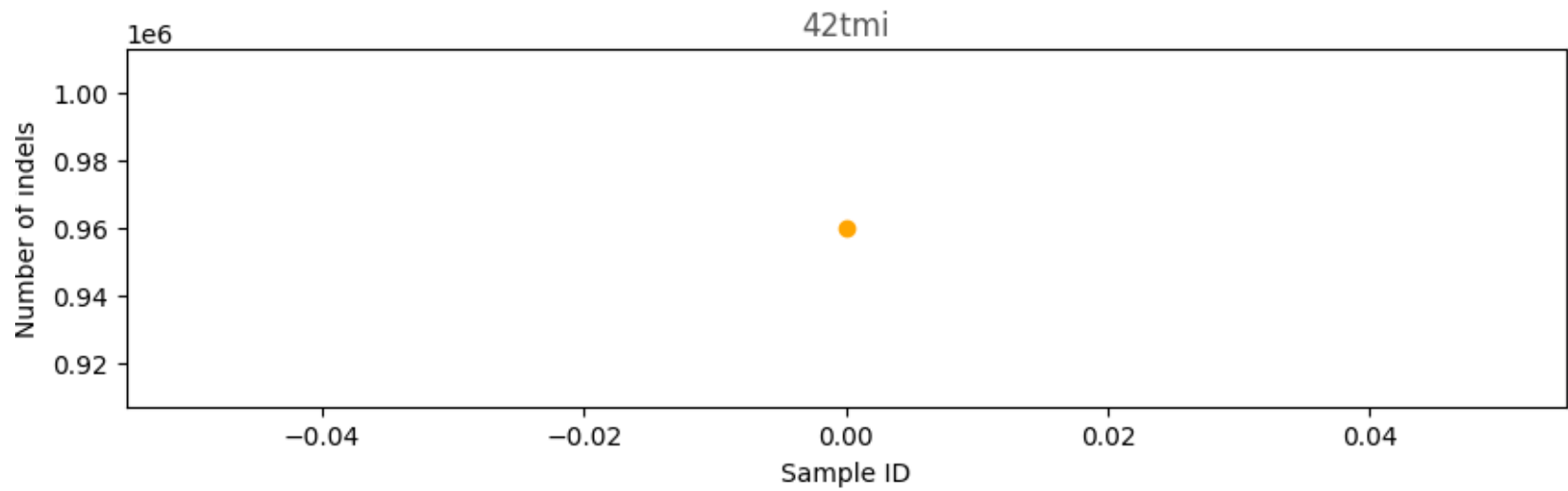
Average depth by sample



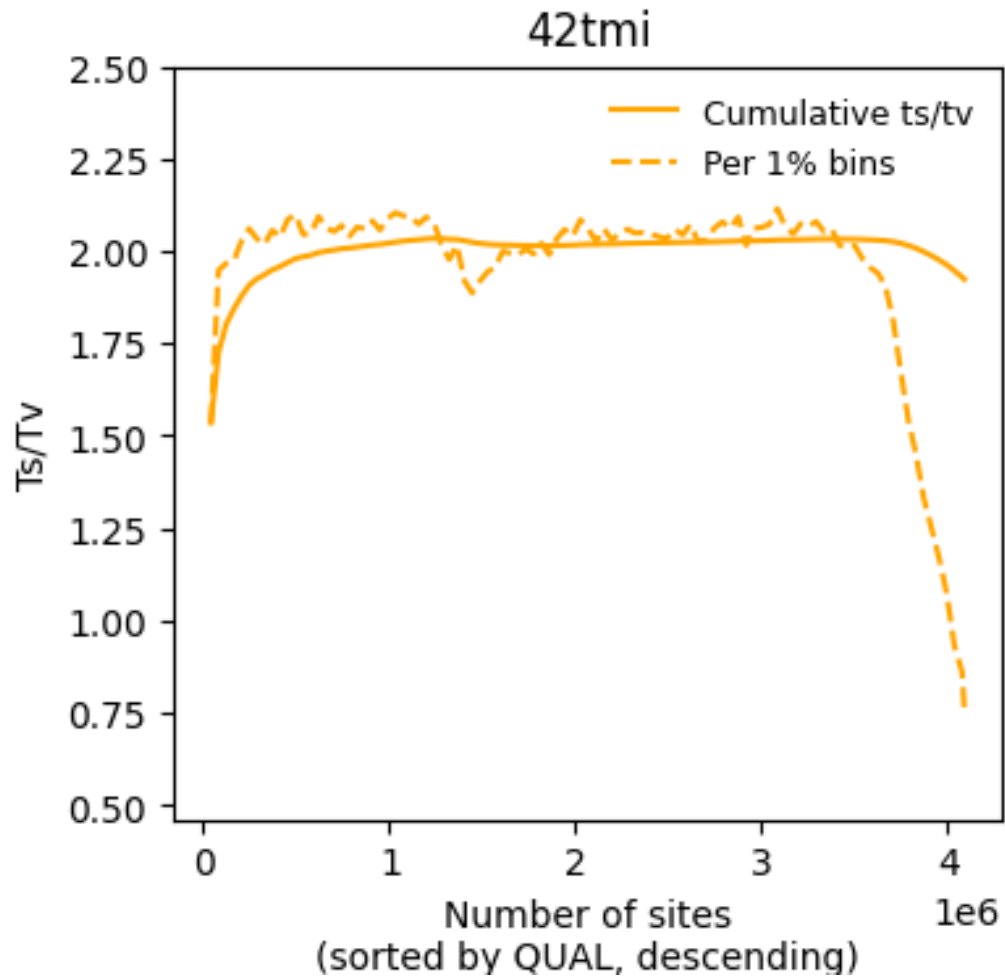
Number of SNPs by sample



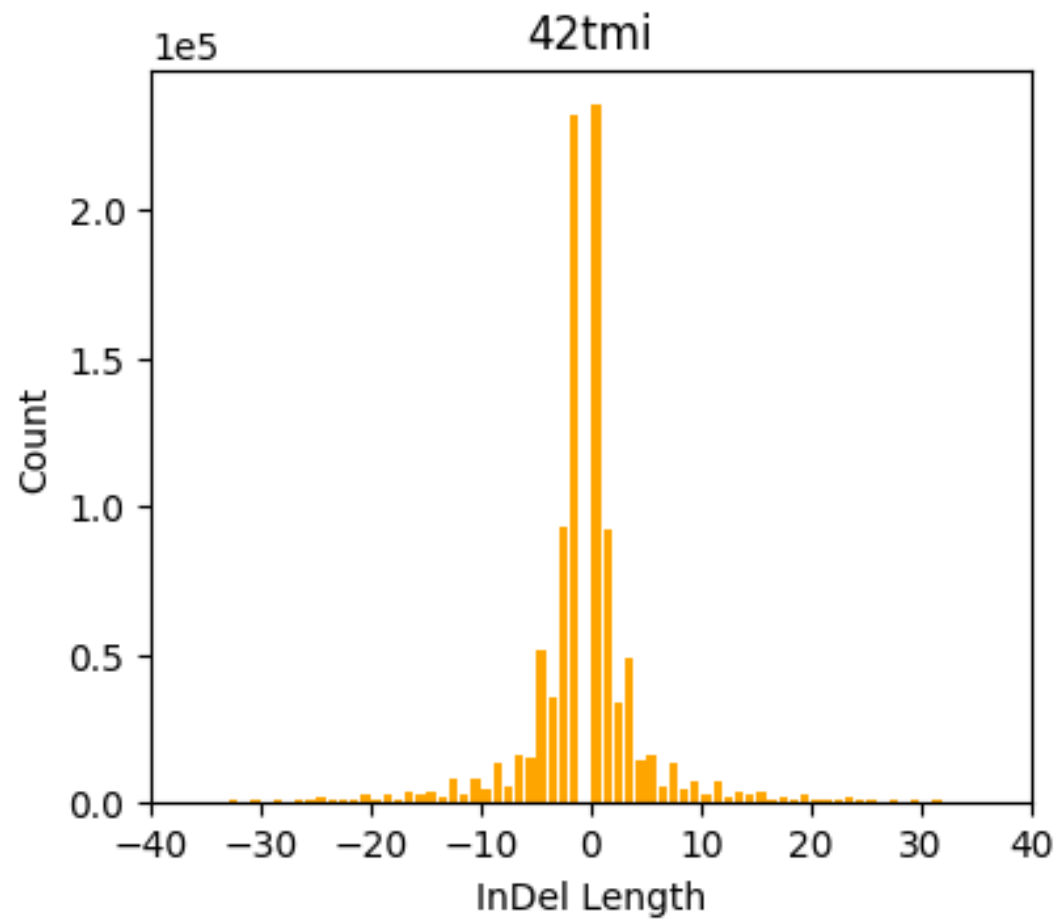
Number of indels by sample



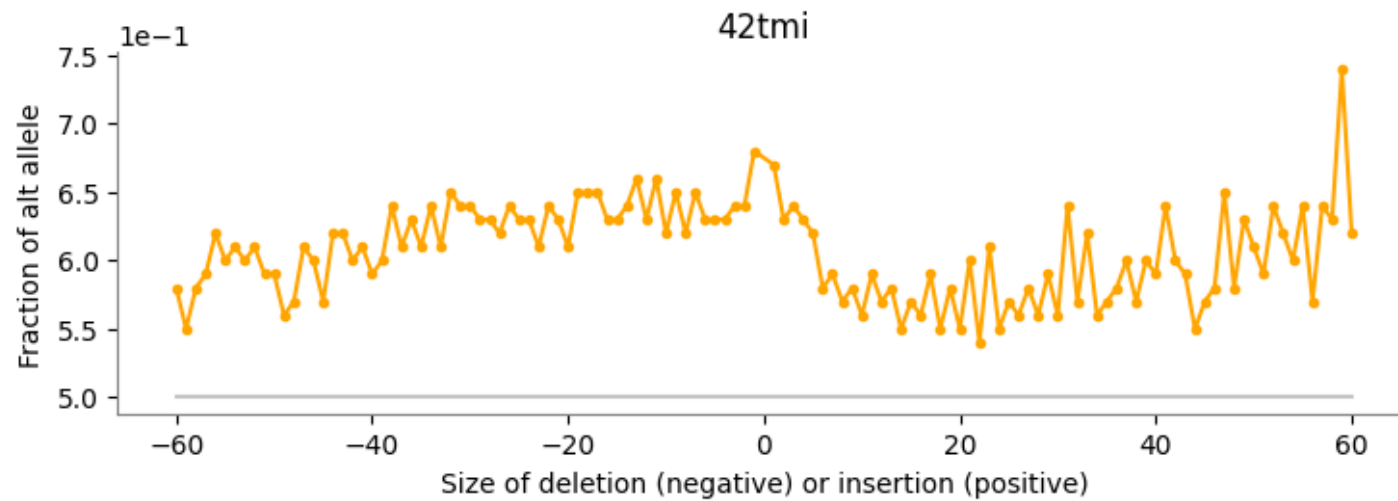
Ts/Tv stratified by QUAL



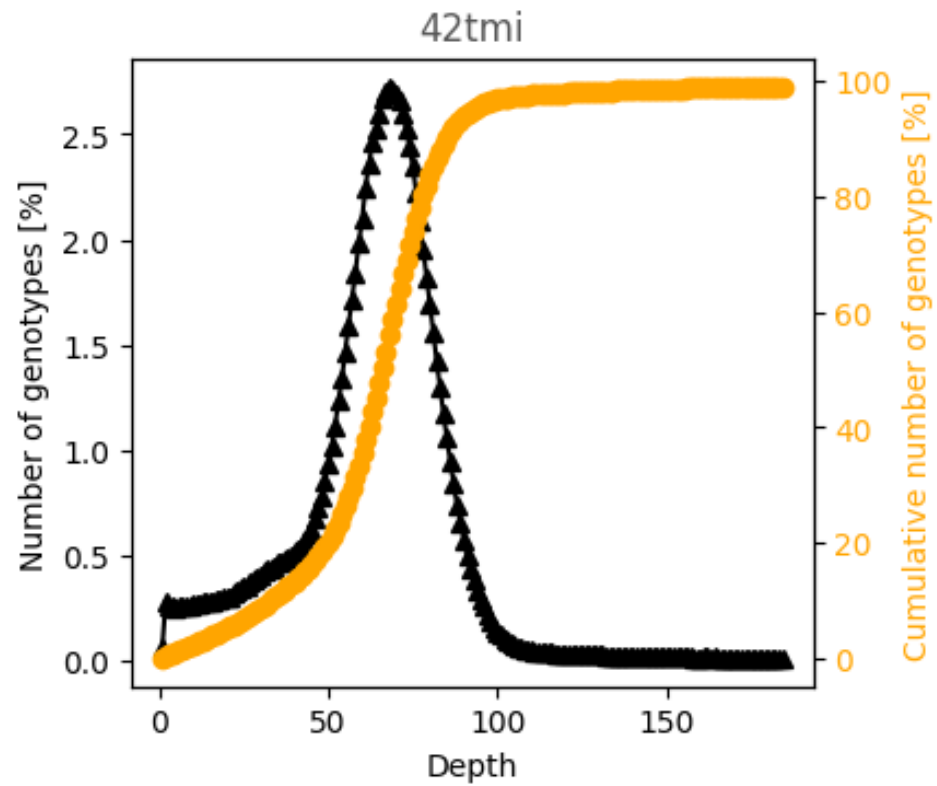
Indel distribution



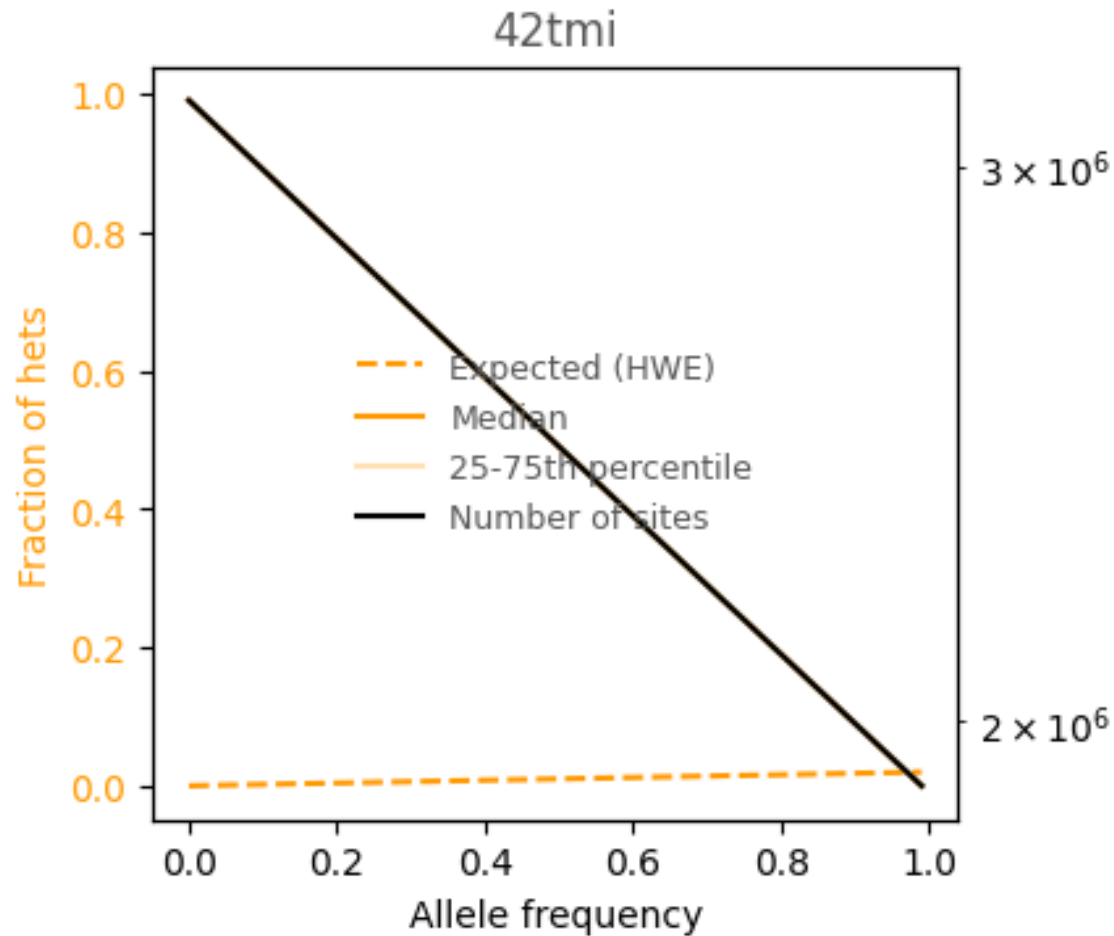
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

