

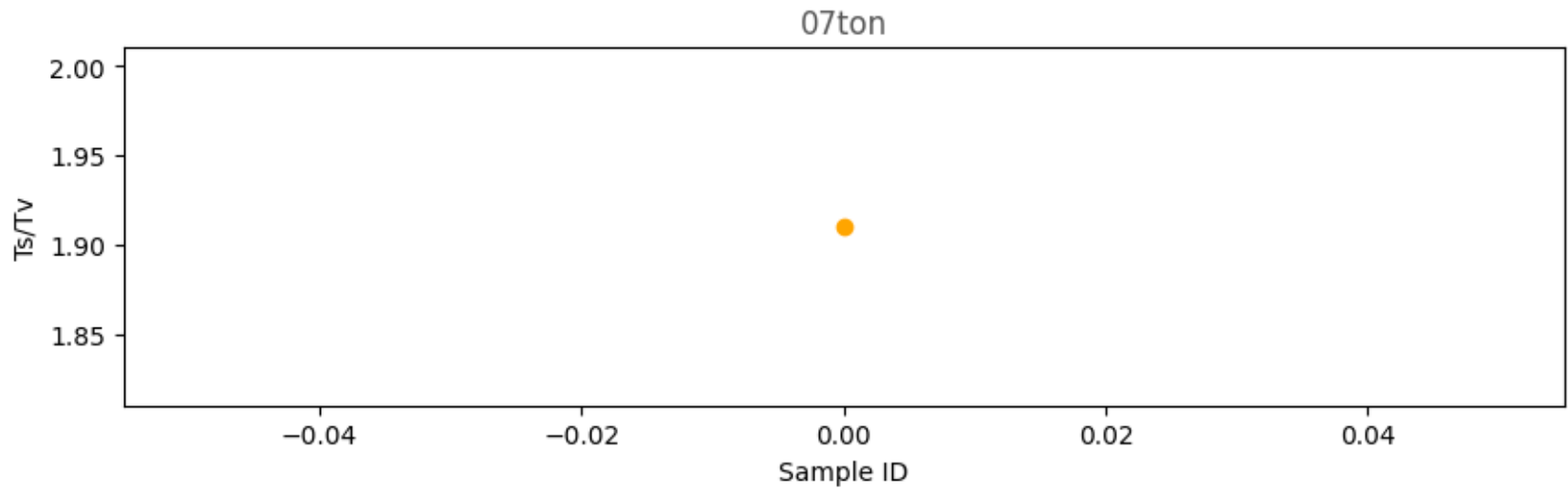
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
07ton	4,107,281	1.91	1.92	960,867	–	0	0
* frameshift ratio: out/(out+in)							

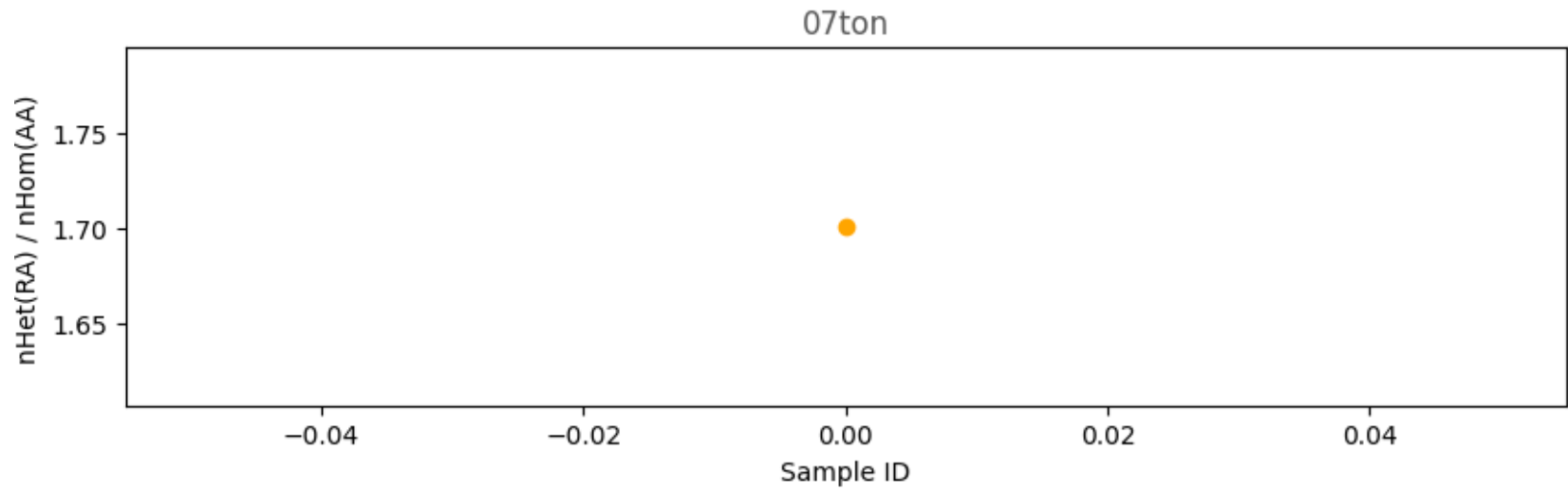
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
07ton	63.0%	1.87	68.1%	106,520	2,126

- 07ton .. /ngc/projects2/gm/data/archive/2022/variants/snv/07toncofm-103850424272-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220324_A01176_AHCFCNDSX3-RHGM_LABKA_WGS-
 WGS04032_snv_germline_raw.haplotype_caller.vcf.gz

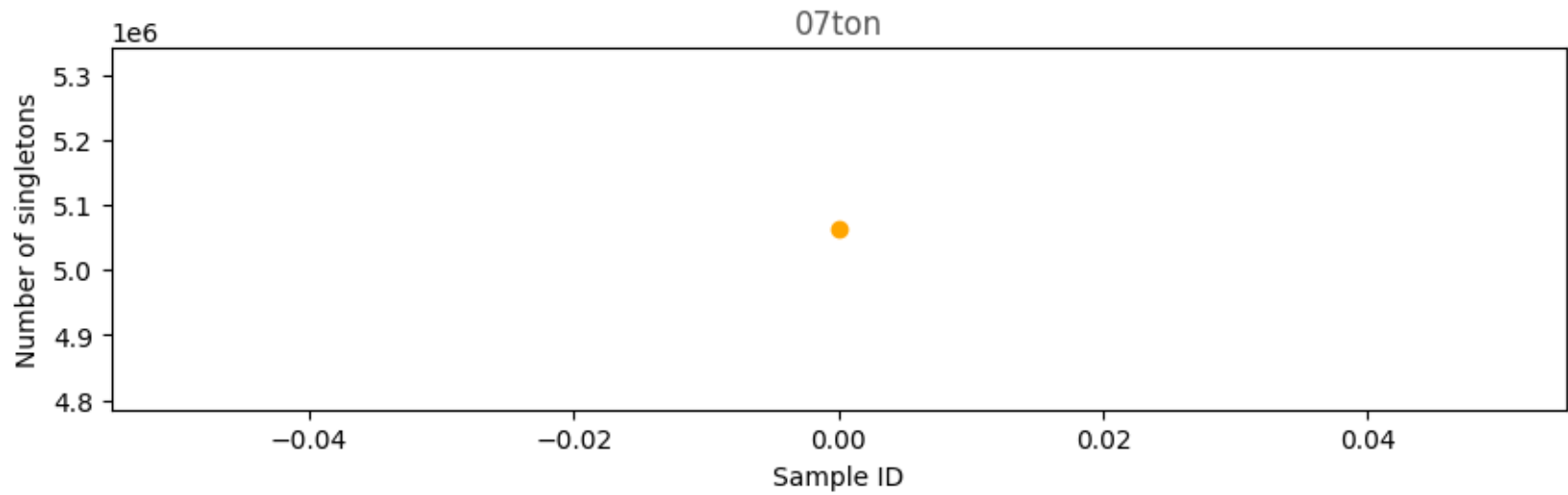
Ts/Tv by sample



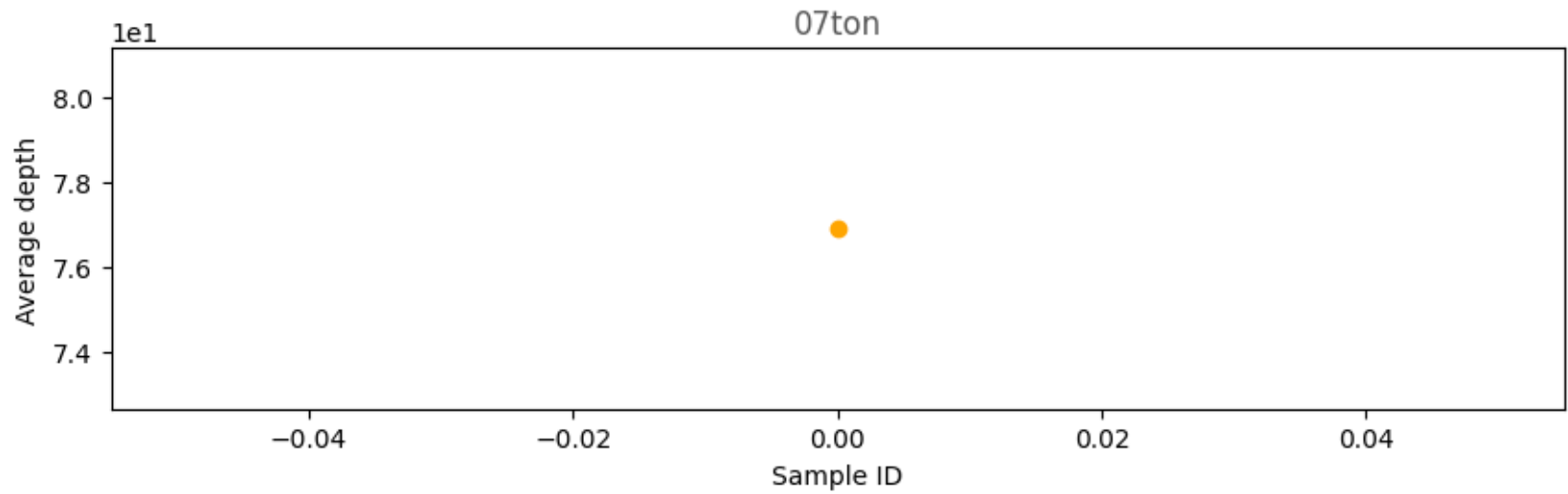
Hets vs non-ref Homs by sample



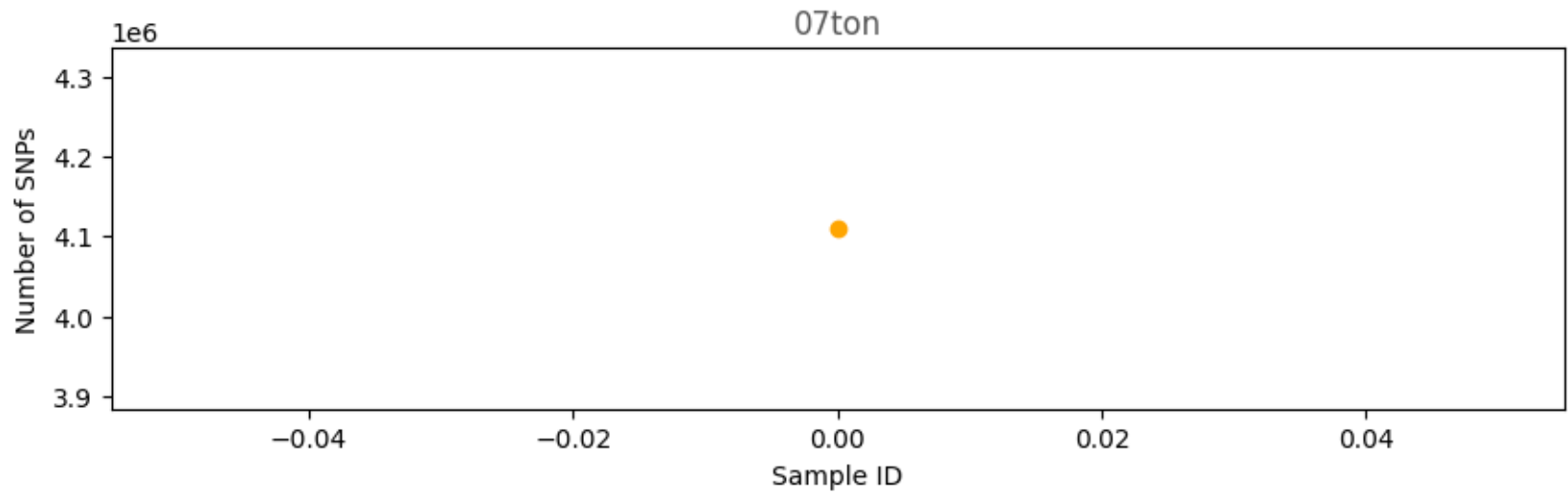
Singletons by sample (hets and homs)



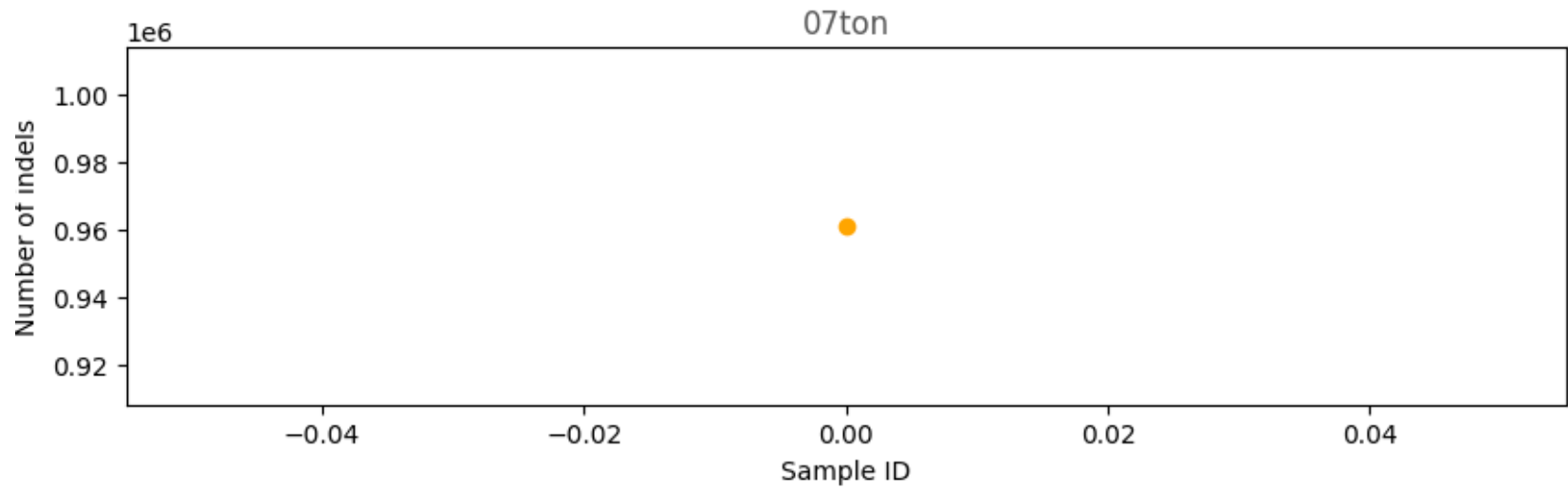
Average depth by sample



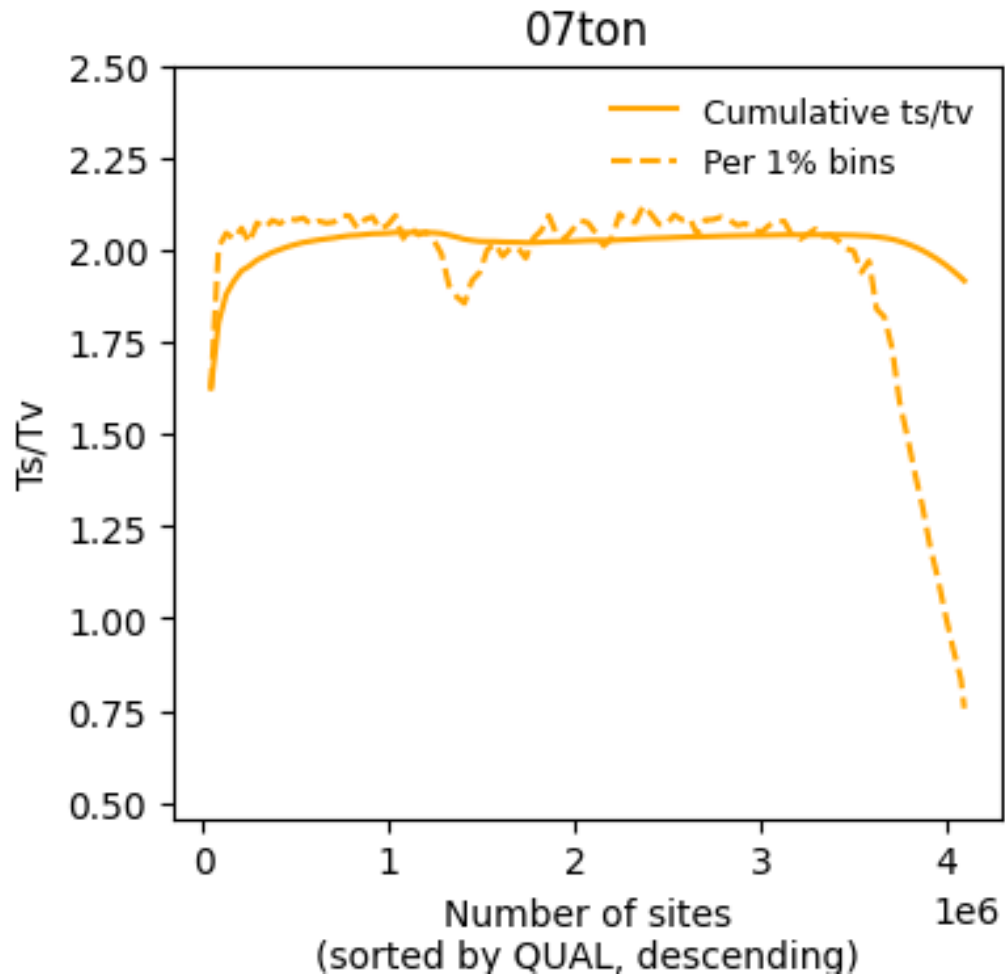
Number of SNPs by sample



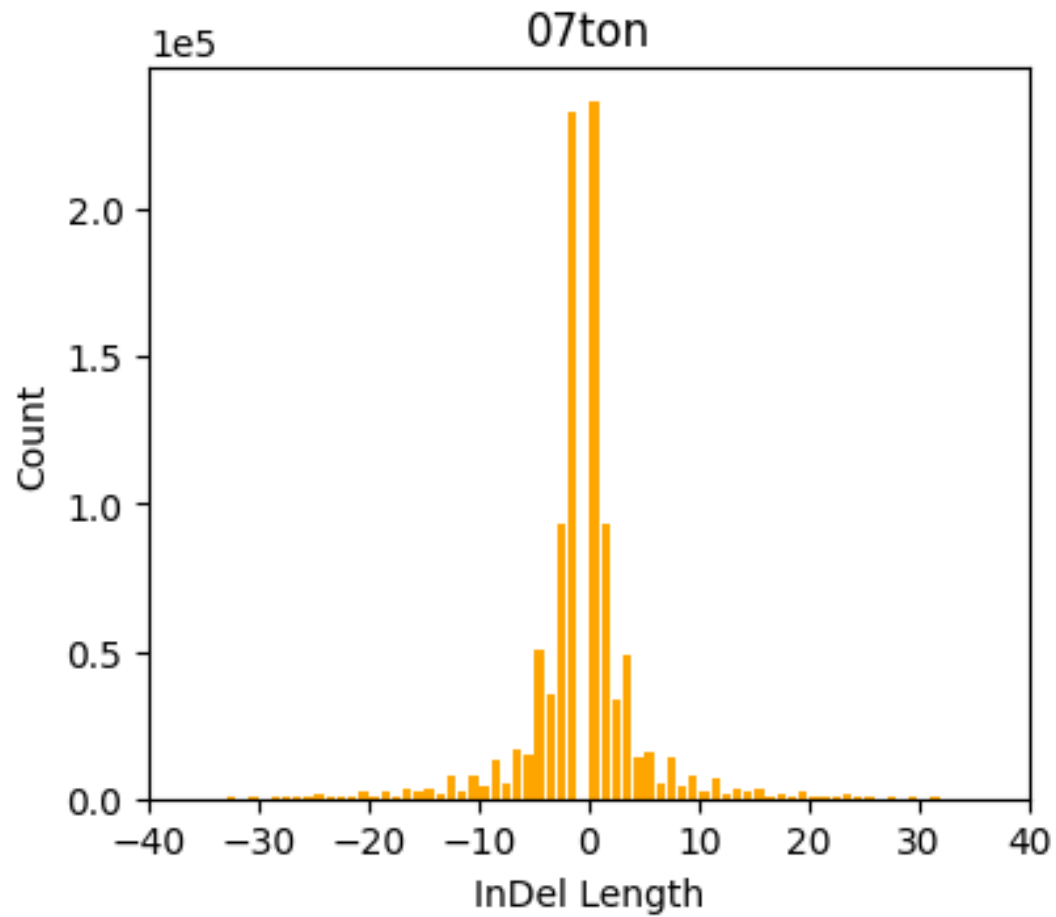
Number of indels by sample



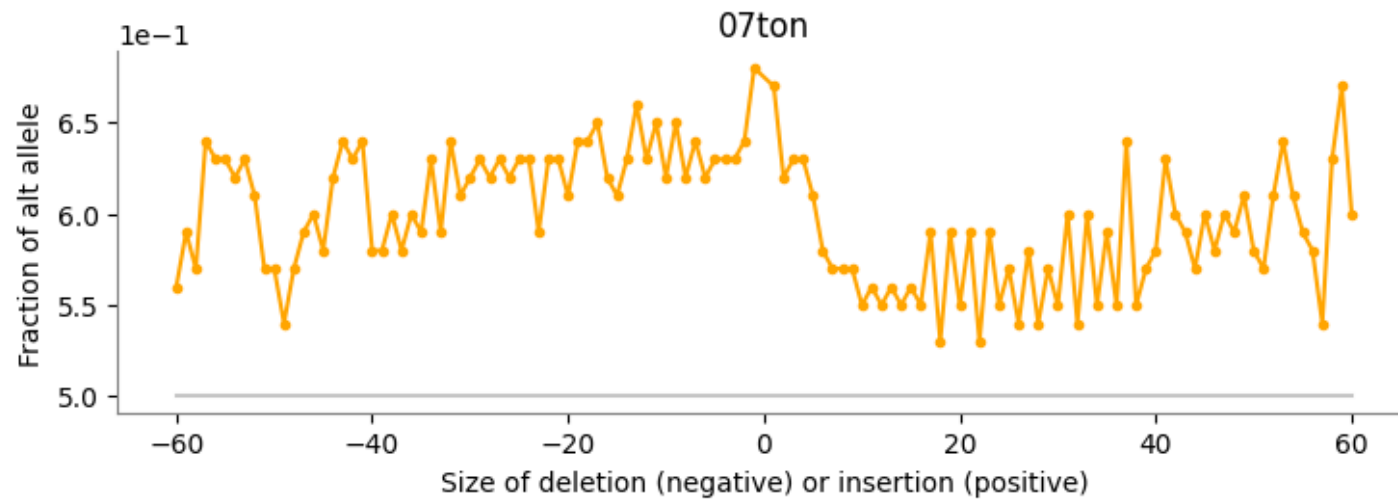
Ts/Tv stratified by QUAL



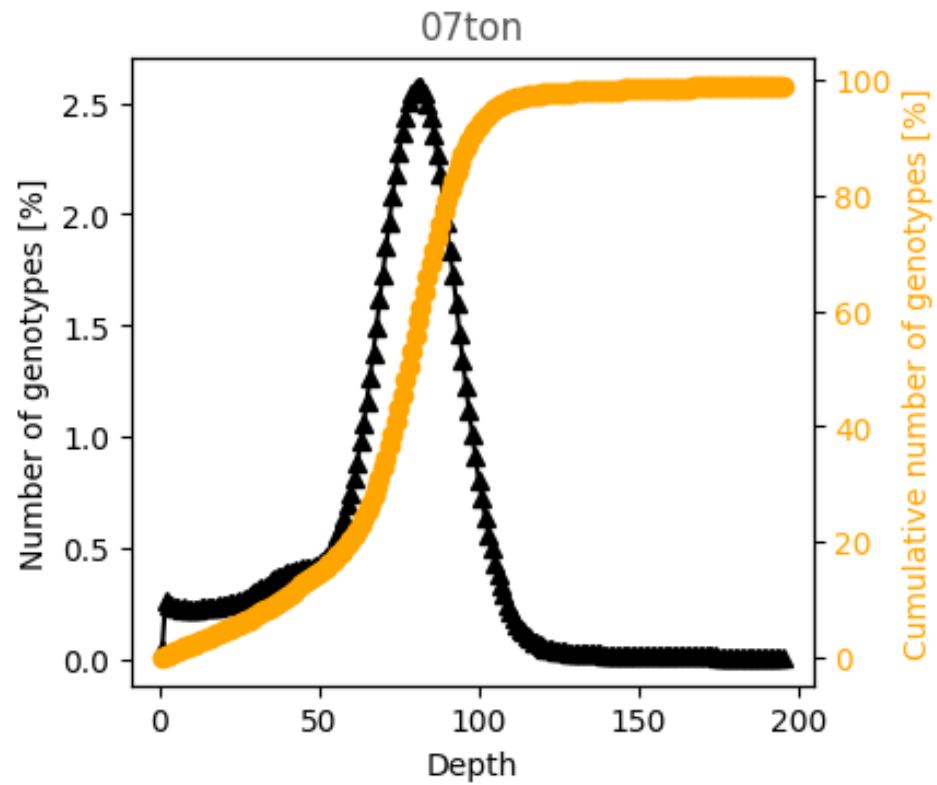
Indel distribution



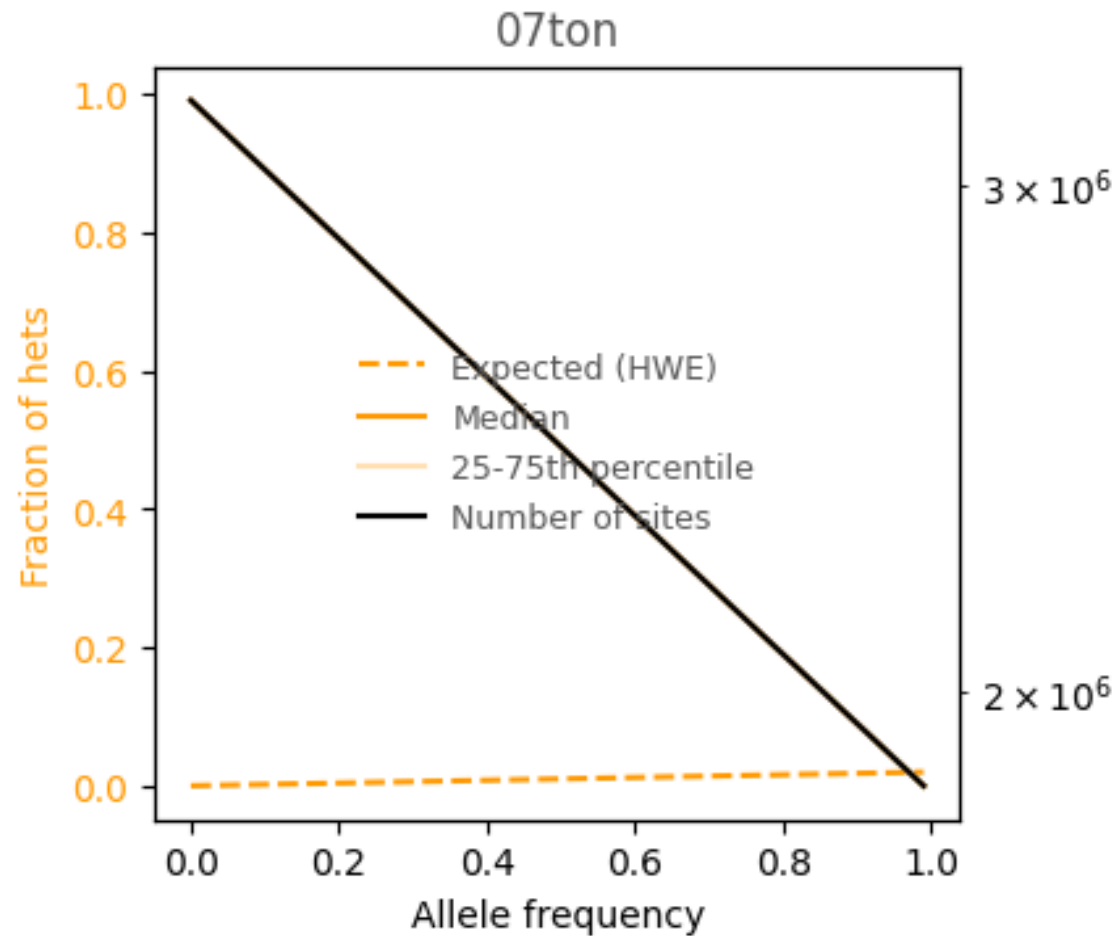
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

