

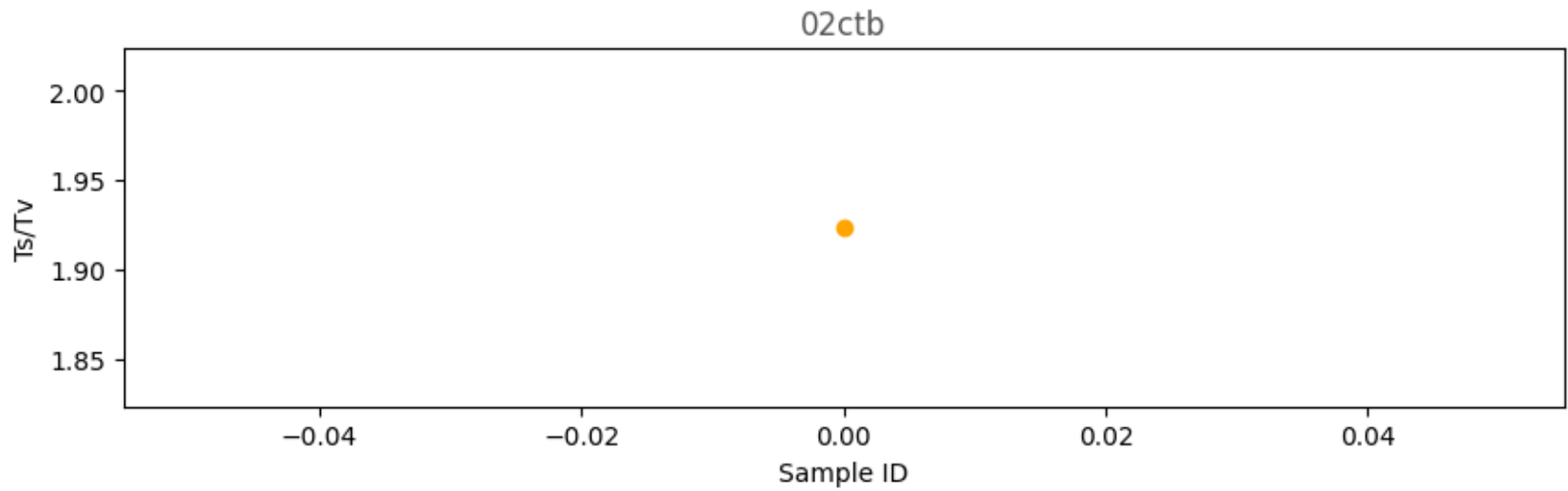
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
02ctb	4,127,878	1.92	1.93	961,158	–	0	0
* frameshift ratio: out/(out+in)							

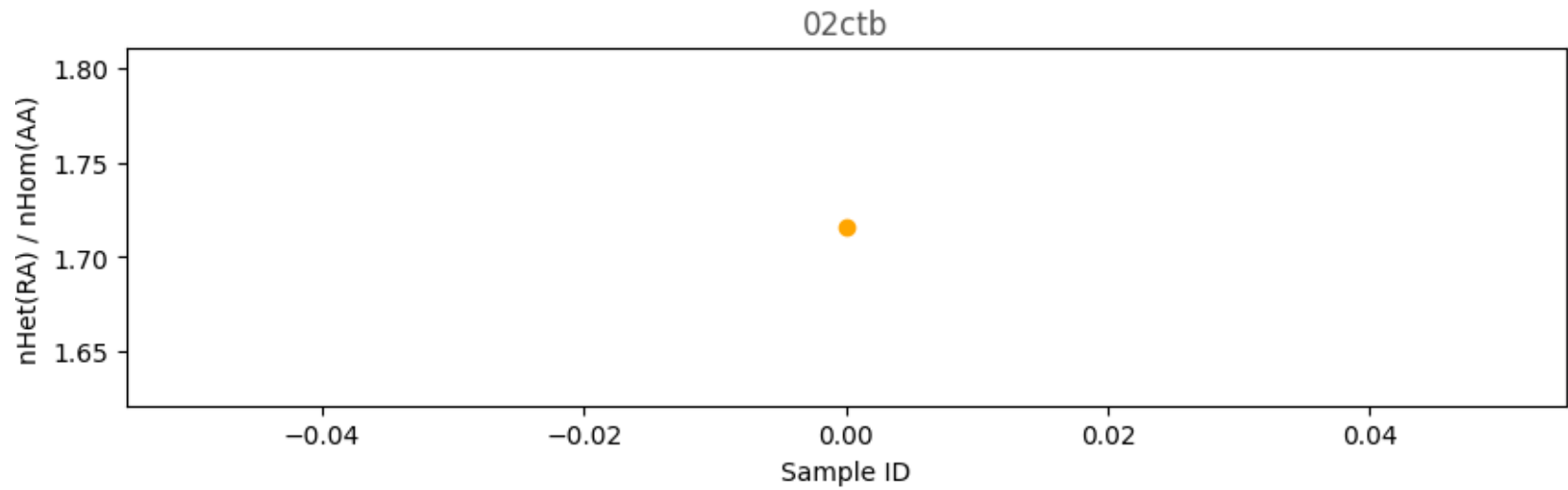
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
02ctb	63.2%	1.89	68.5%	103,812	2,191

- 02ctb .. /ngc/projects2/gm/data/archive/2022/variants/snv/02ctbedsf-103846429516-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220316_A01411_BHCF77DSX3-RHGM_LABKA_WGS-
 WGS03999_snv_germline_raw.haplotype_caller.vcf.gz

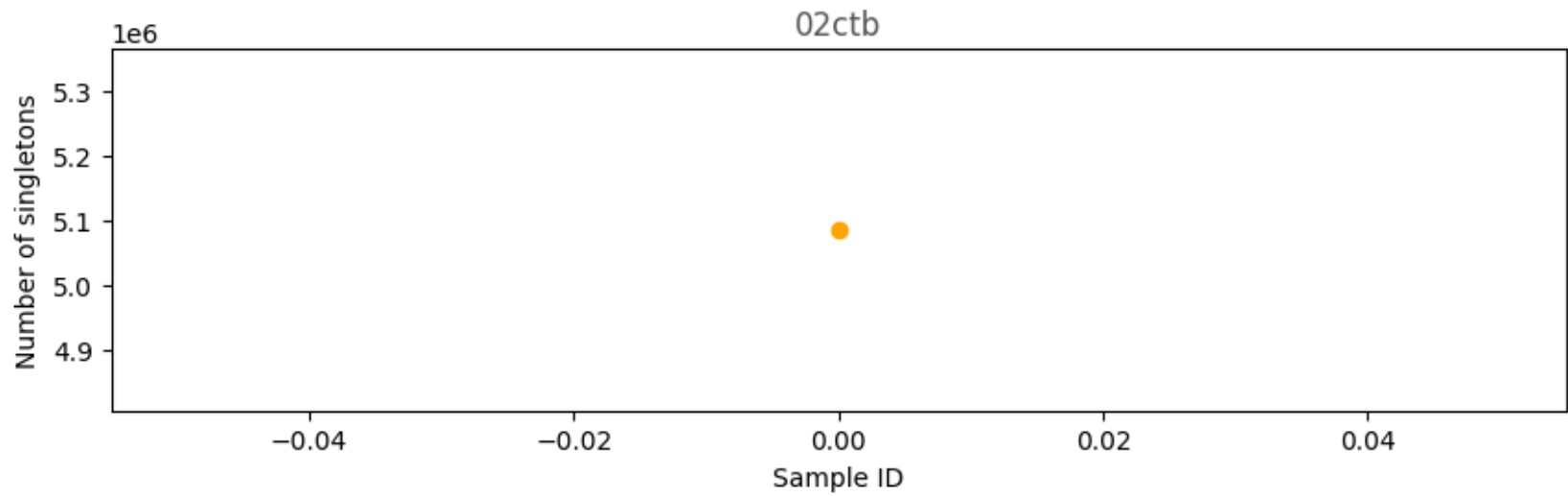
Ts/Tv by sample



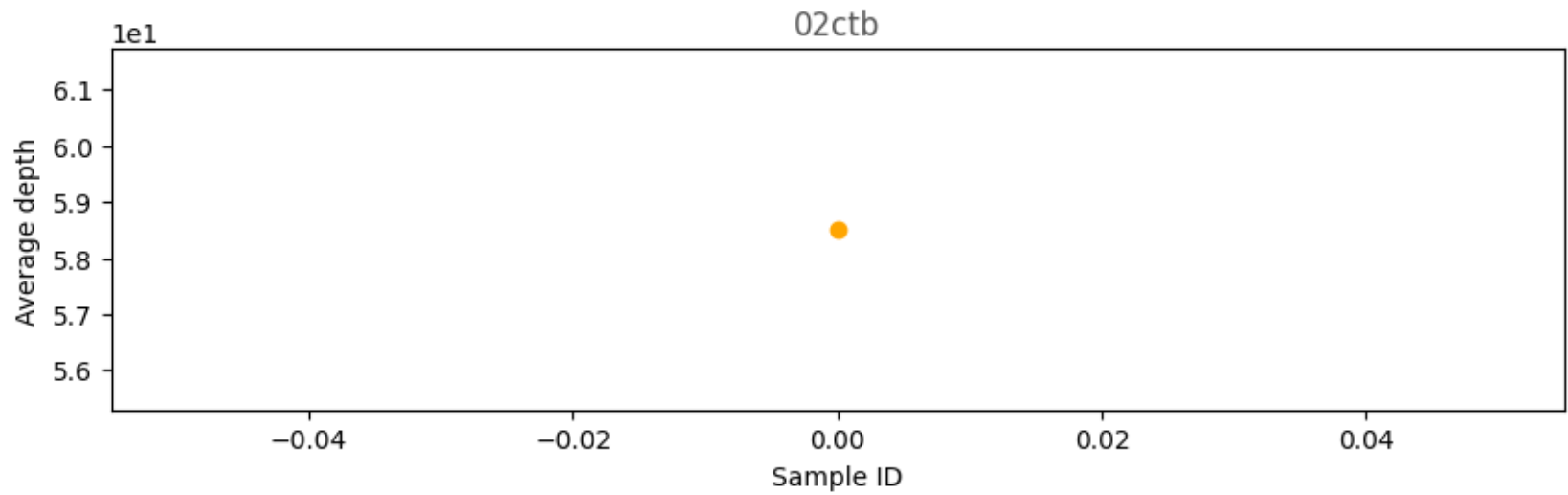
Hets vs non-ref Homs by sample



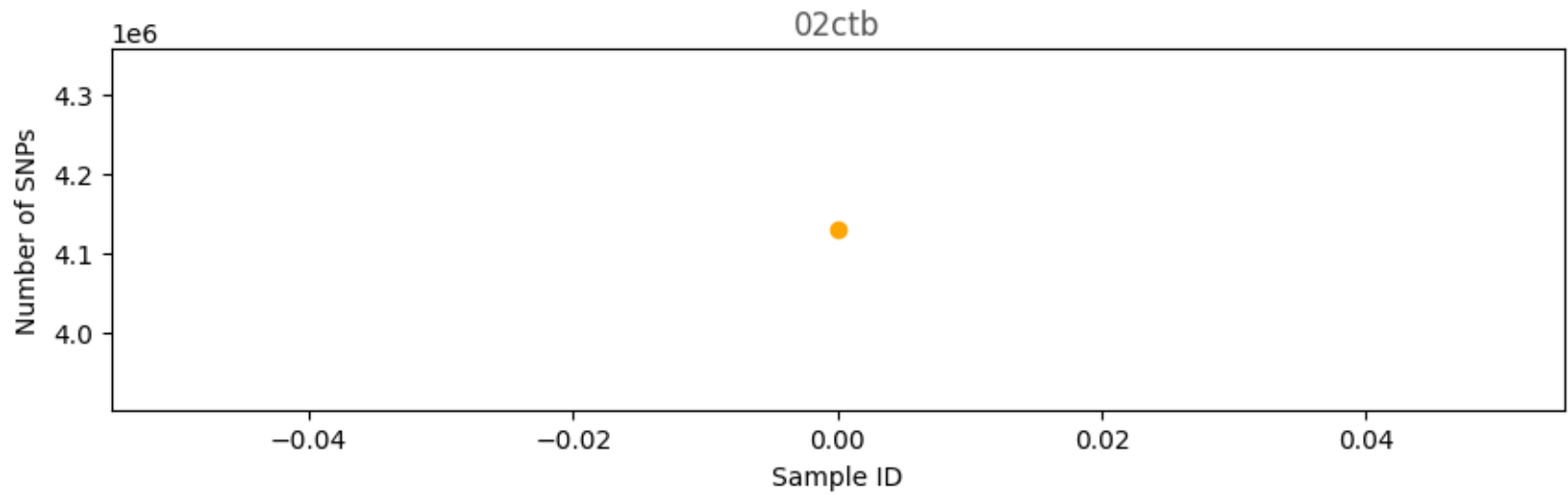
Singletons by sample (hets and homs)



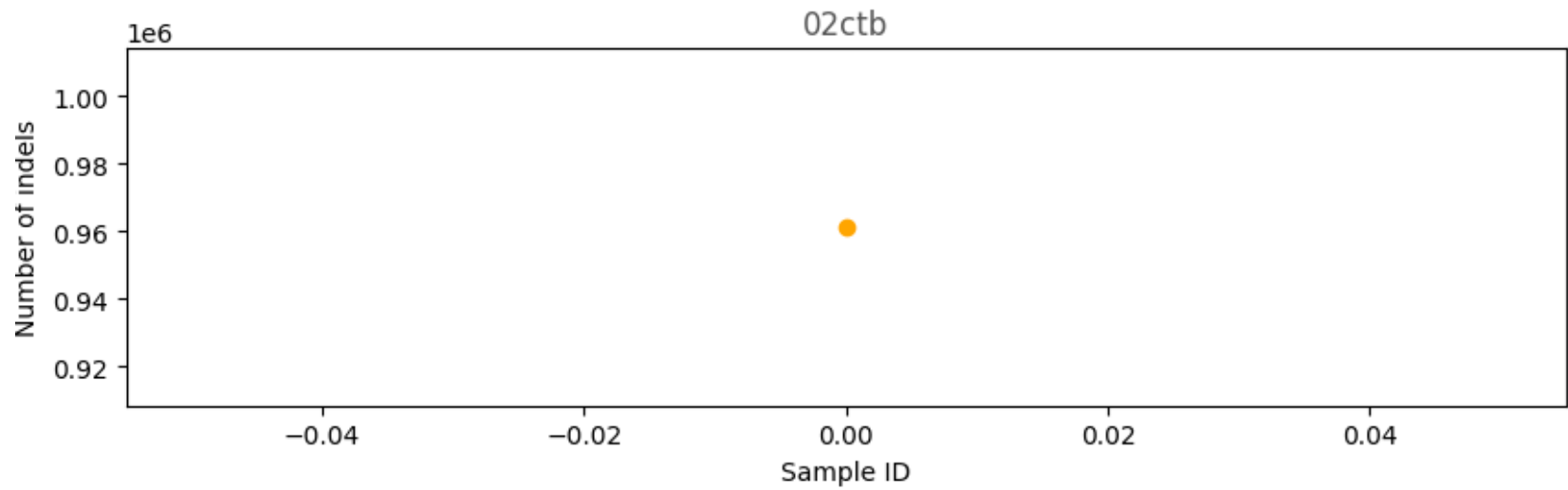
Average depth by sample



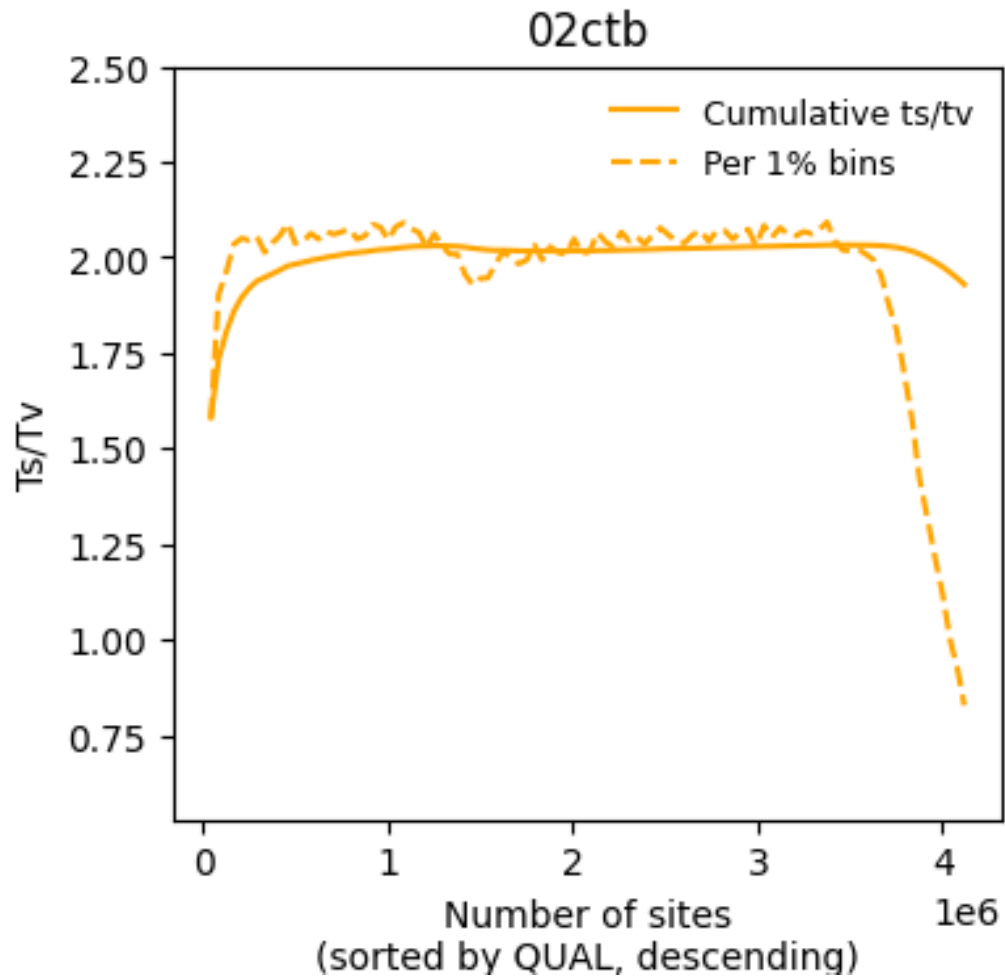
Number of SNPs by sample



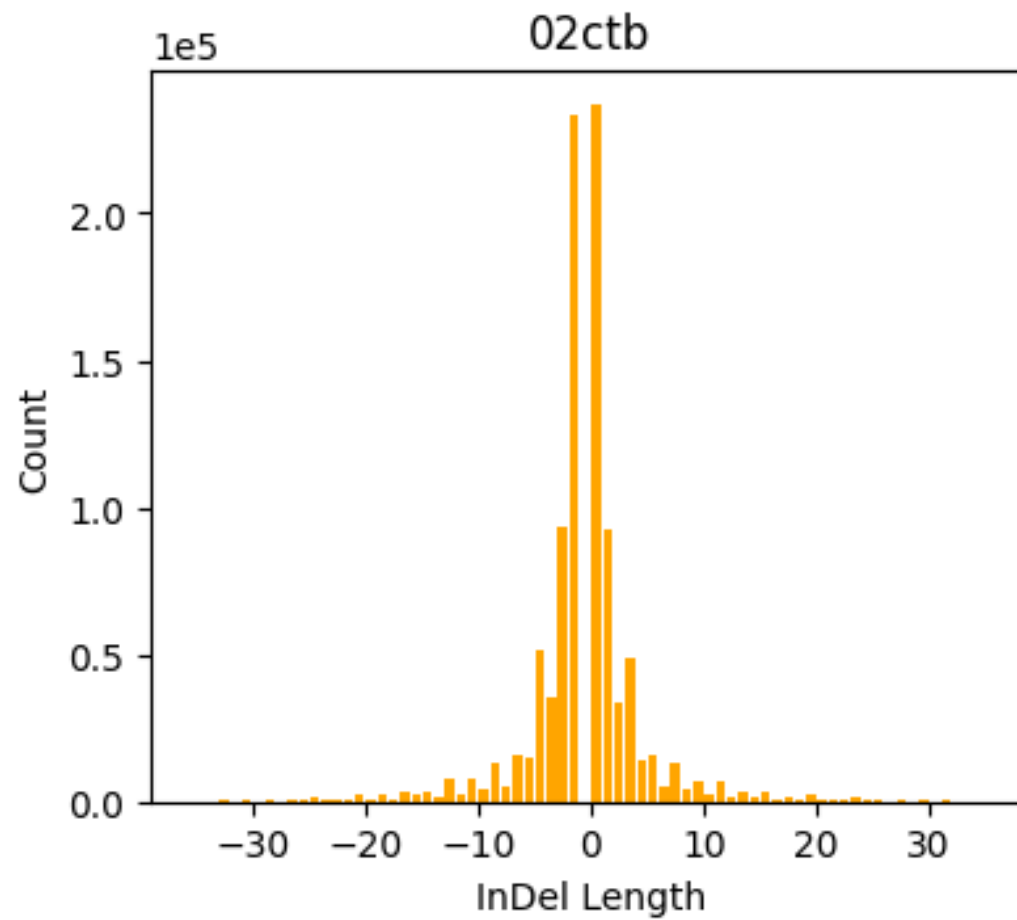
Number of indels by sample



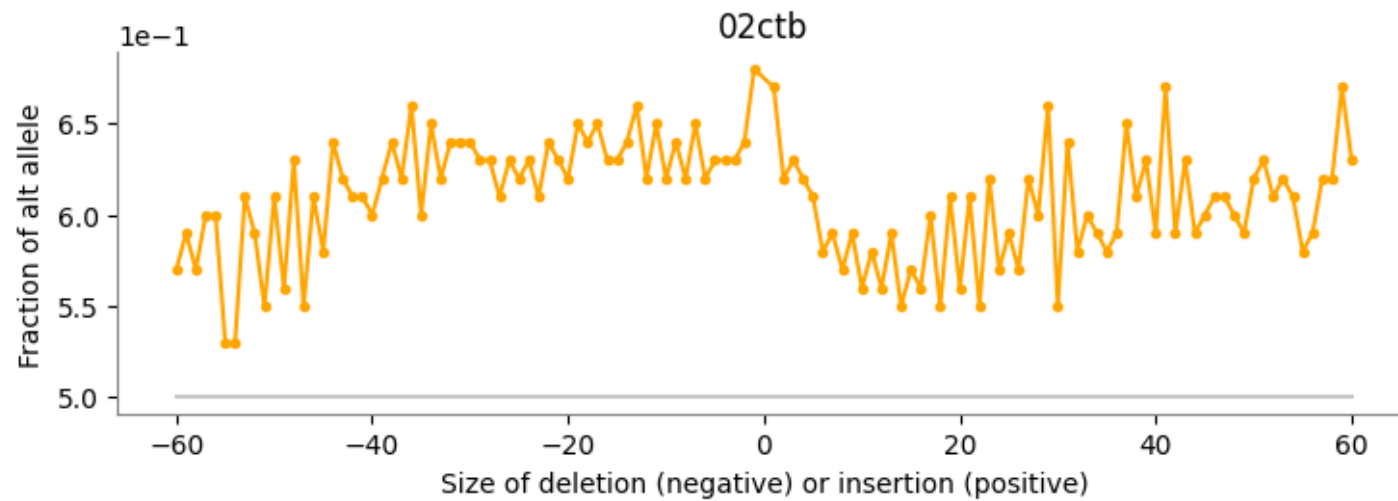
Ts/Tv stratified by QUAL



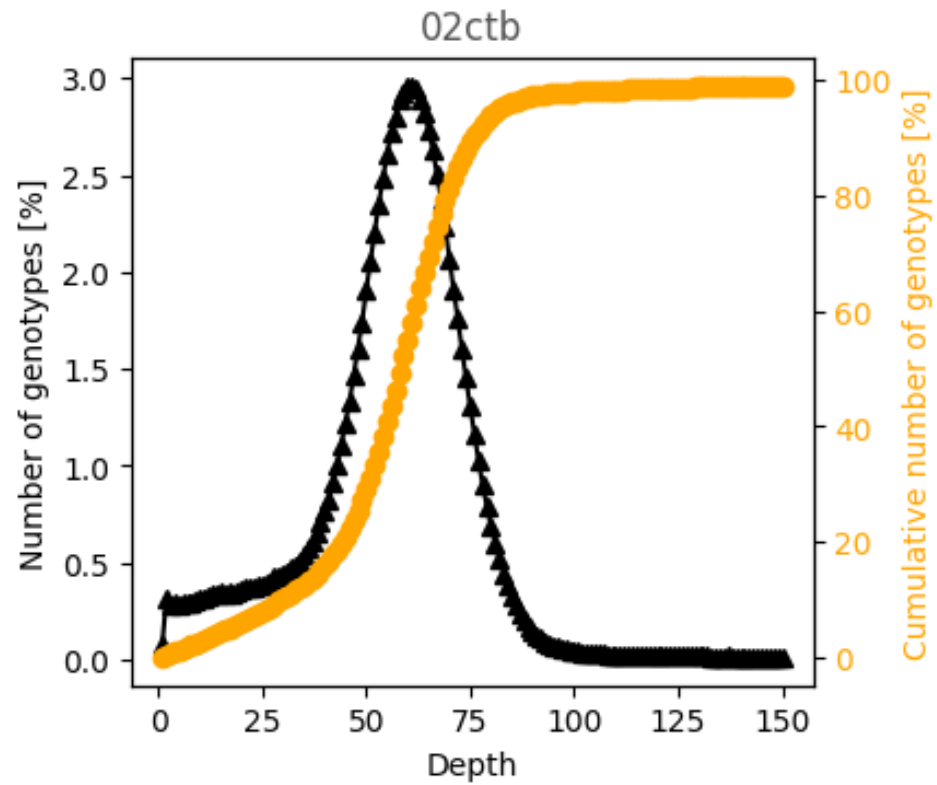
Indel distribution



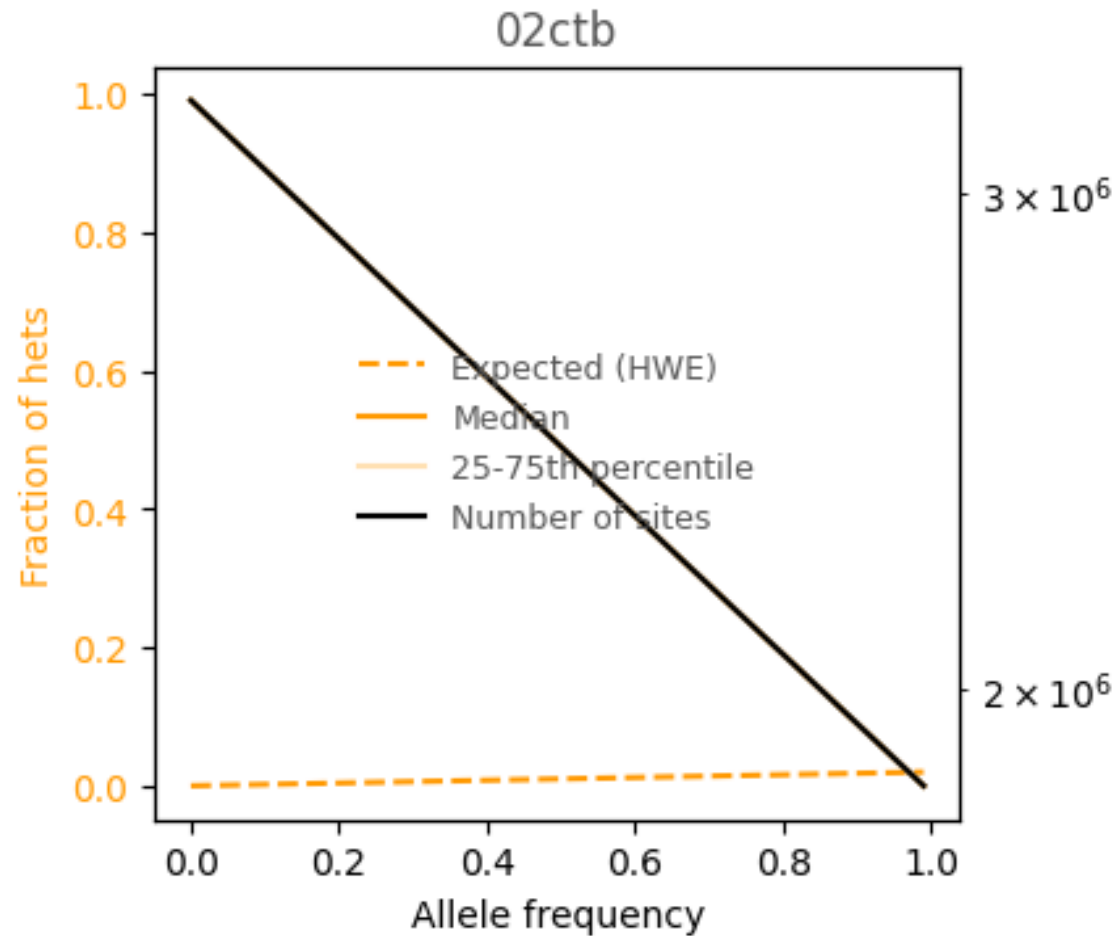
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

