

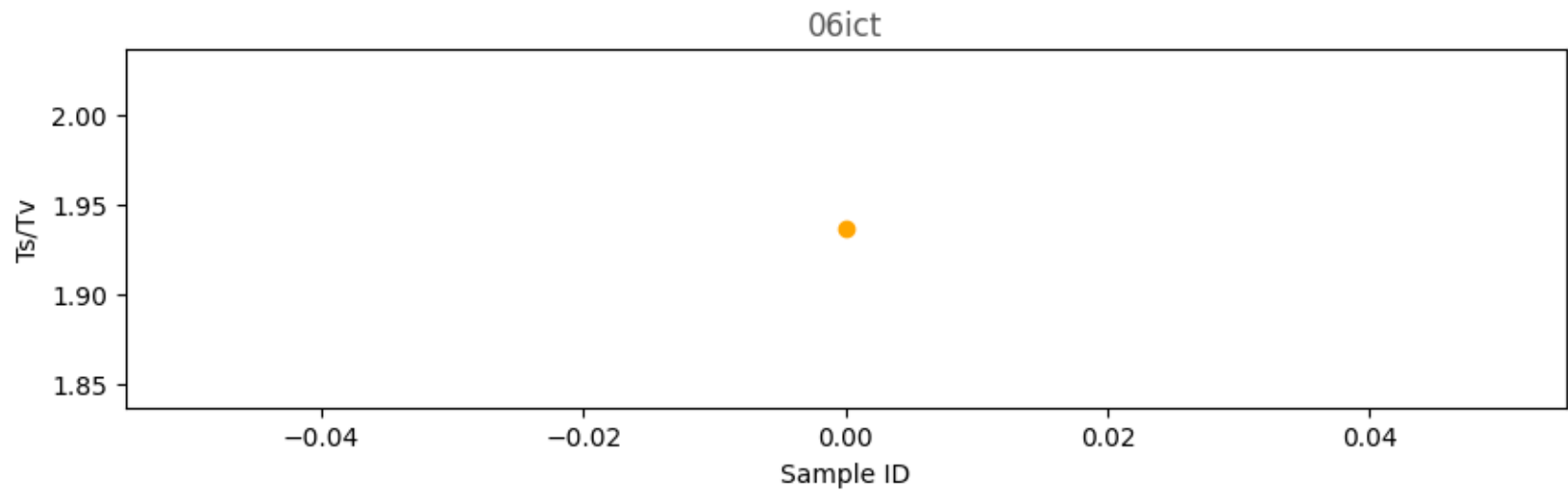
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
06ict	4,040,020	1.94	1.94	927,741	–	0	0
* frameshift ratio: out/(out+in)							

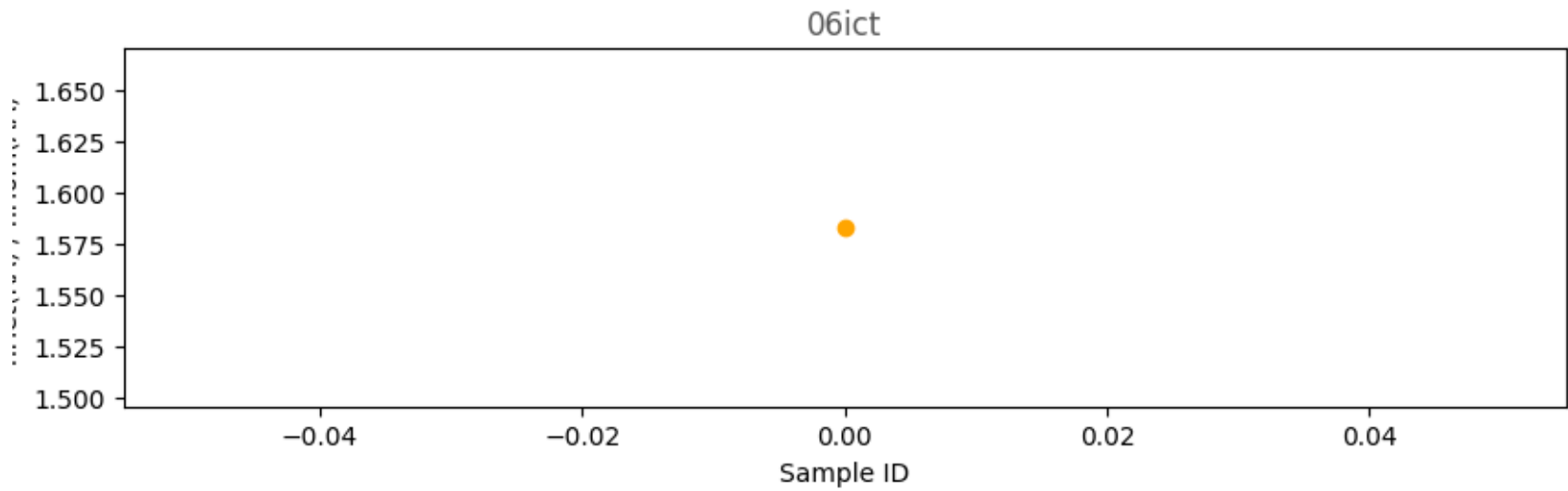
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
06ict	61.3%	1.90	65.3%	90,459	1,947

- 06ict .. /ngc/projects2/gm/data/archive/2022/variants/snv/06ictweem-103872139830-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00815-220525_A00559_BHJ5JJDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS04321_snv_germline_raw.haplotype_caller.vcf.gz

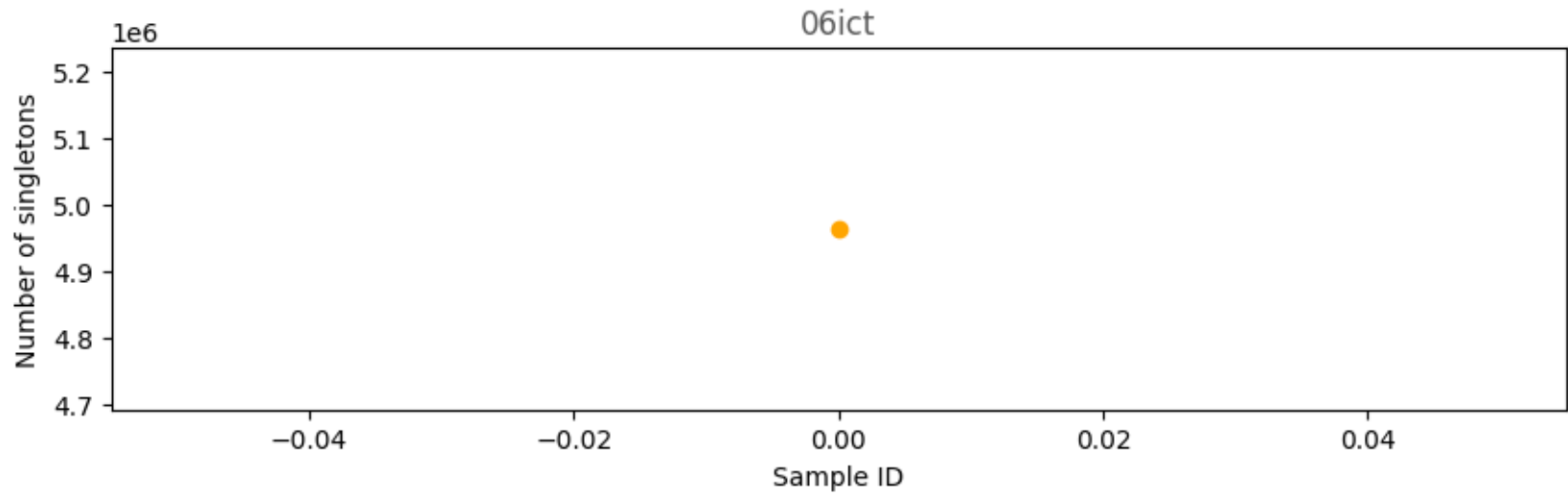
Ts/Tv by sample



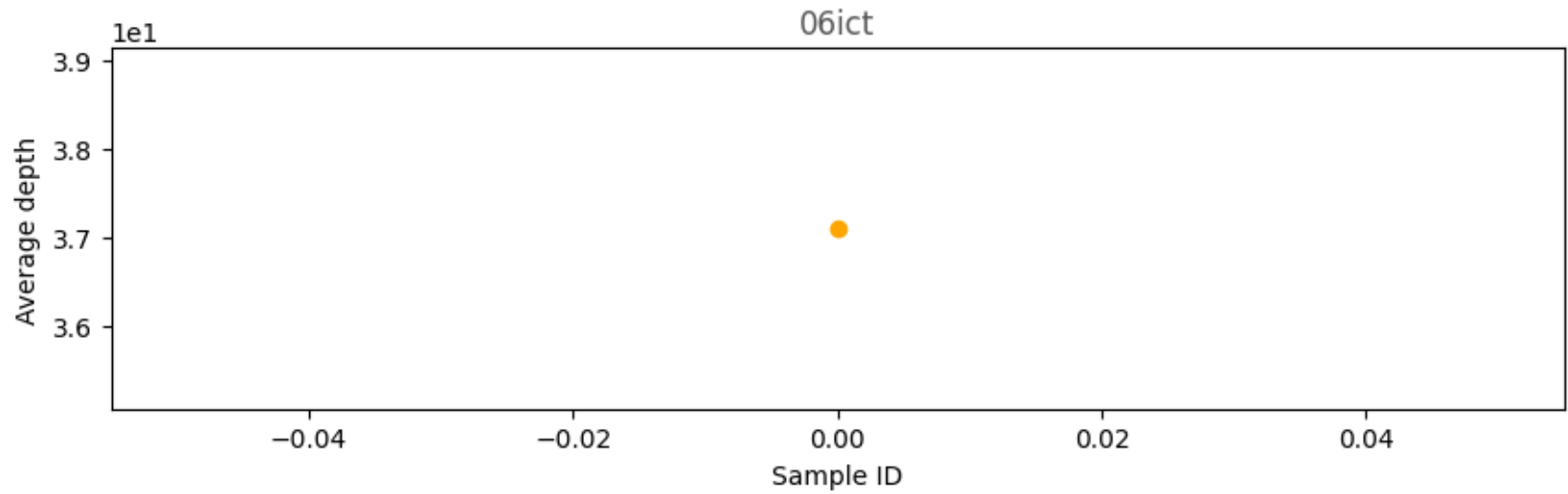
Hets vs non-ref Homs by sample



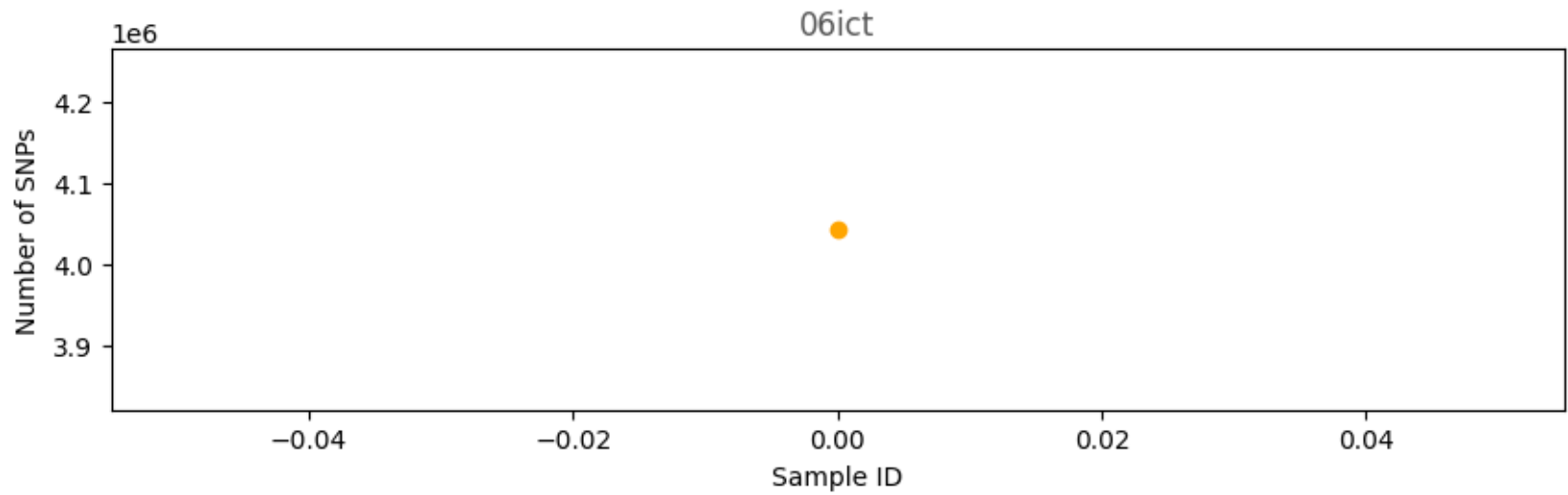
Singletons by sample (hets and homs)



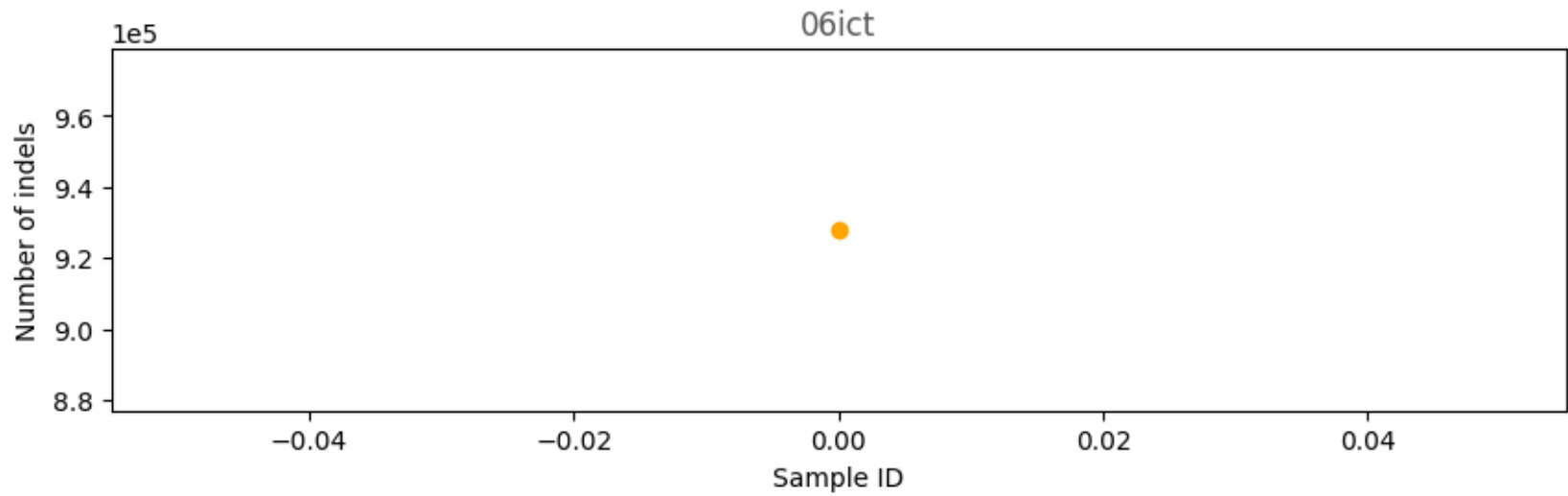
Average depth by sample



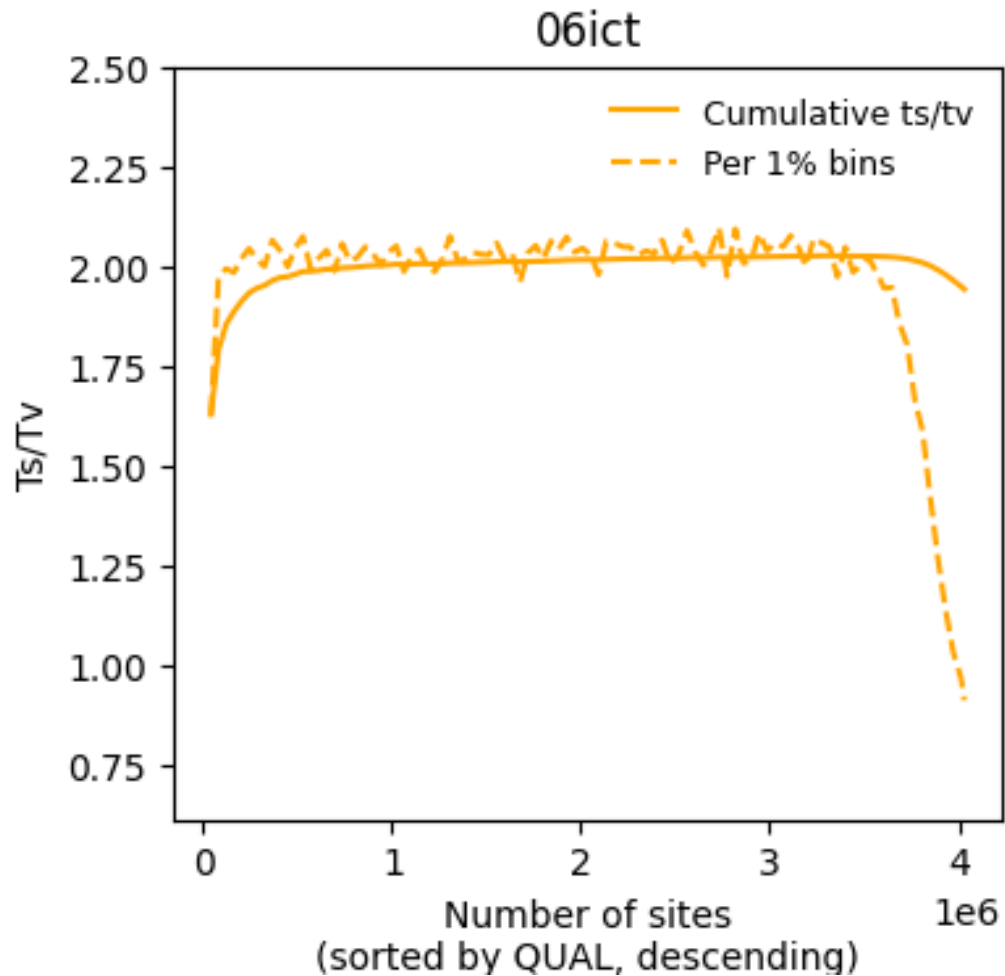
Number of SNPs by sample



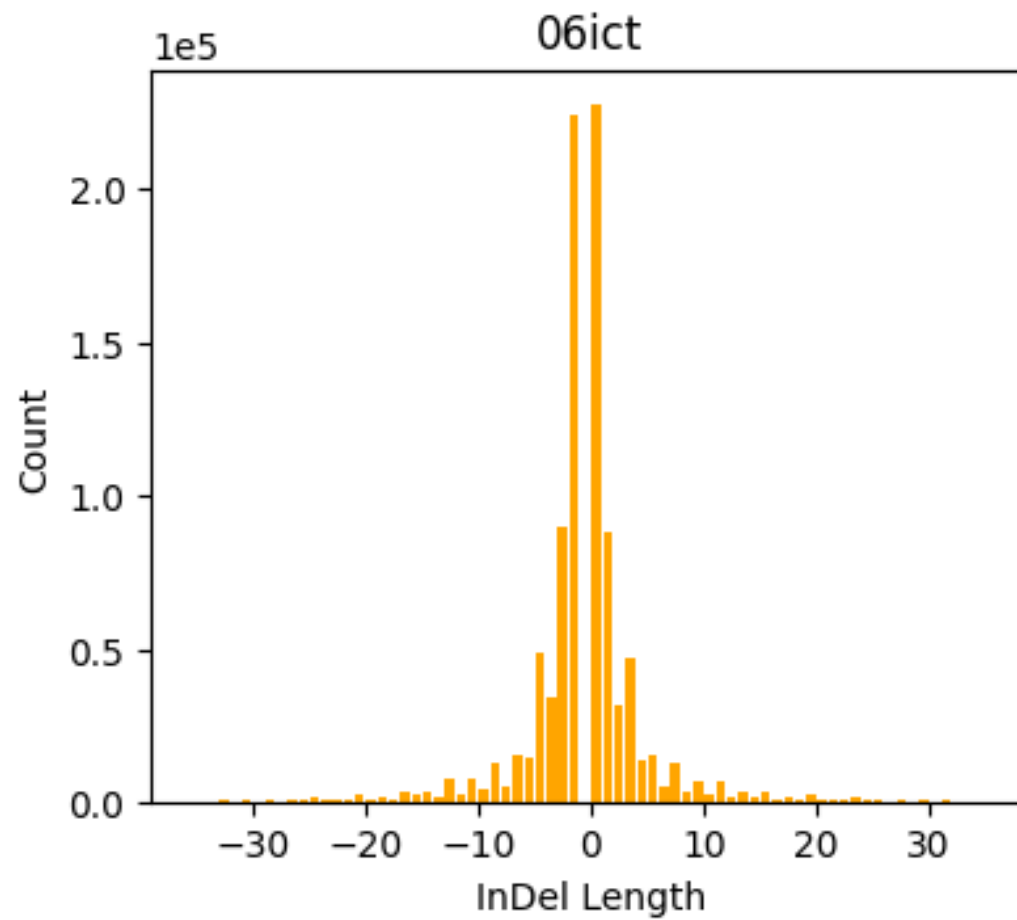
Number of indels by sample



Ts/Tv stratified by QUAL



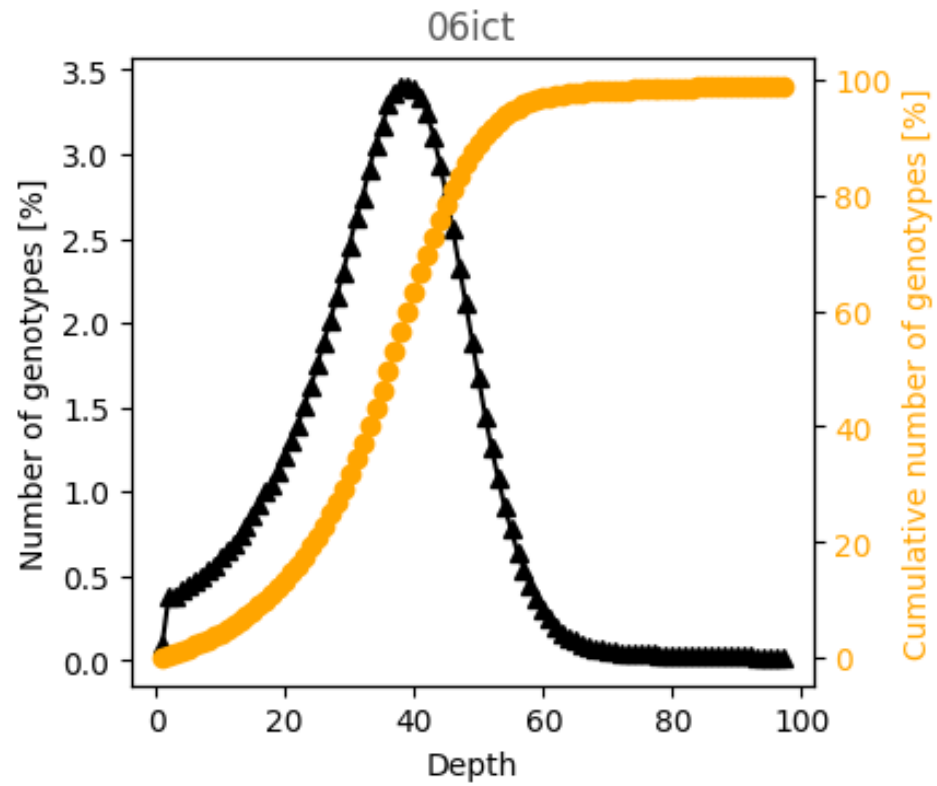
Indel distribution



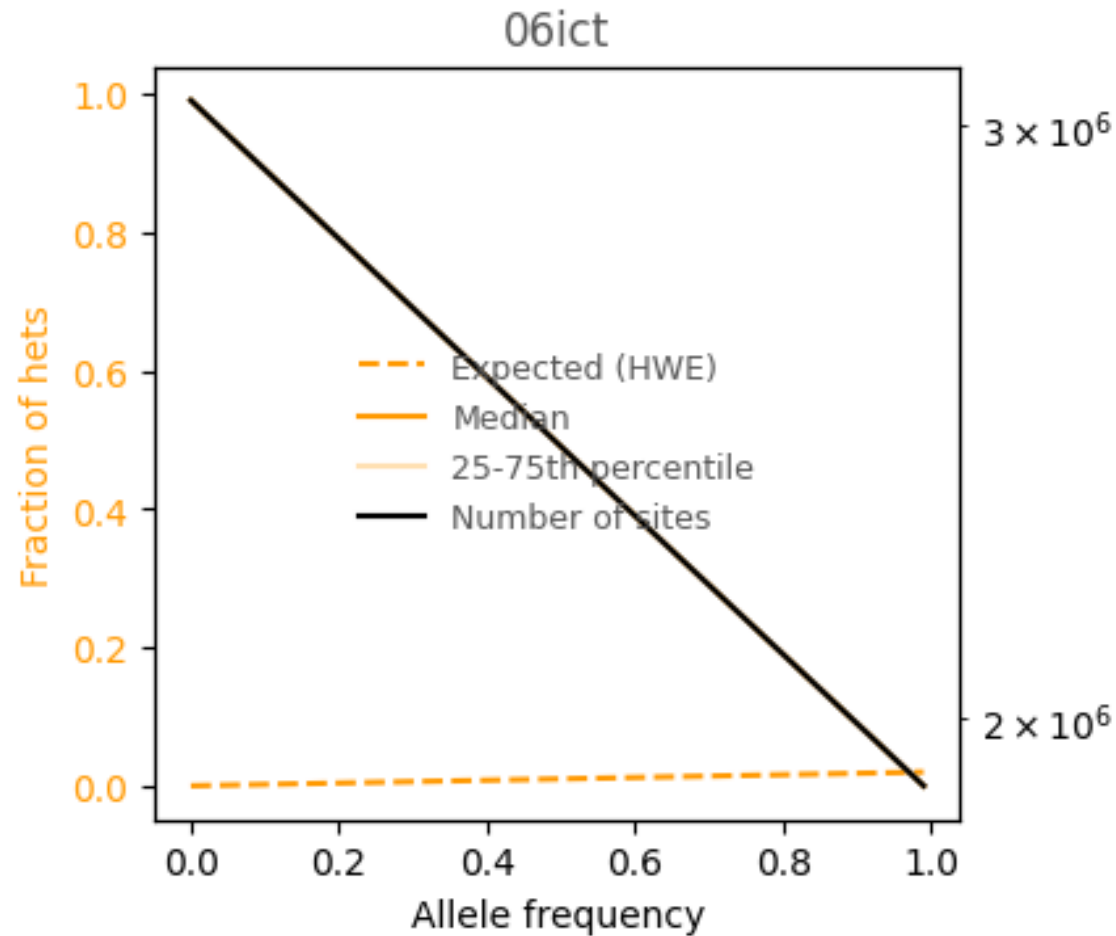
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

