

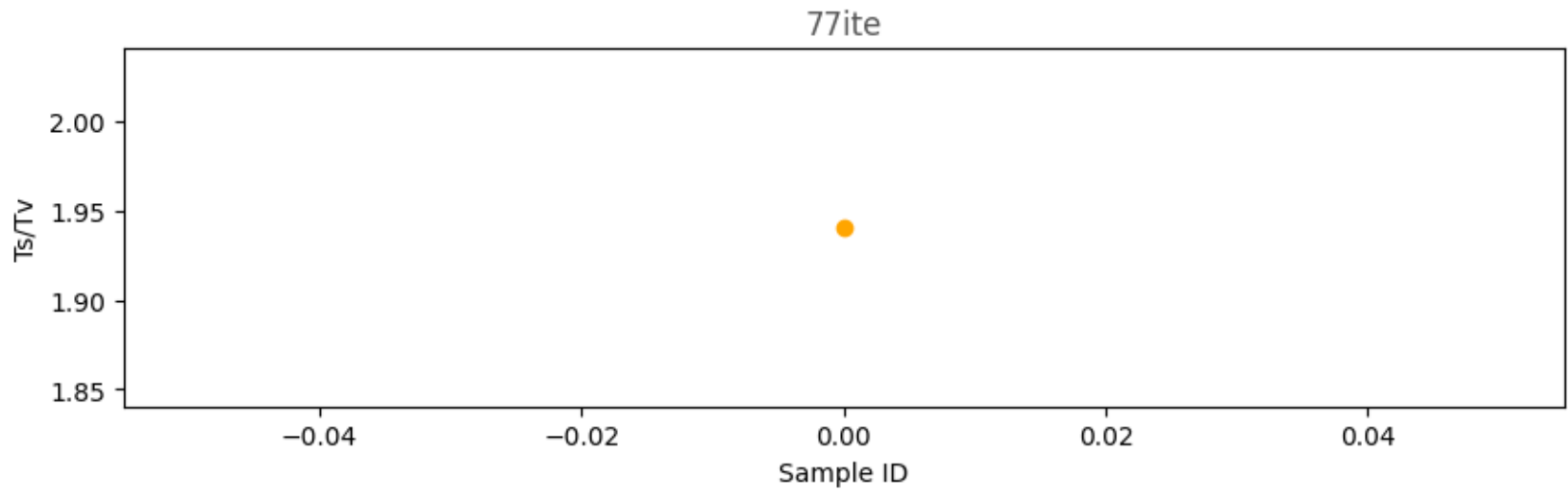
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
77ite	4,041,977	1.94	1.94	925,161	–	0	0
* frameshift ratio: out/(out+in)							

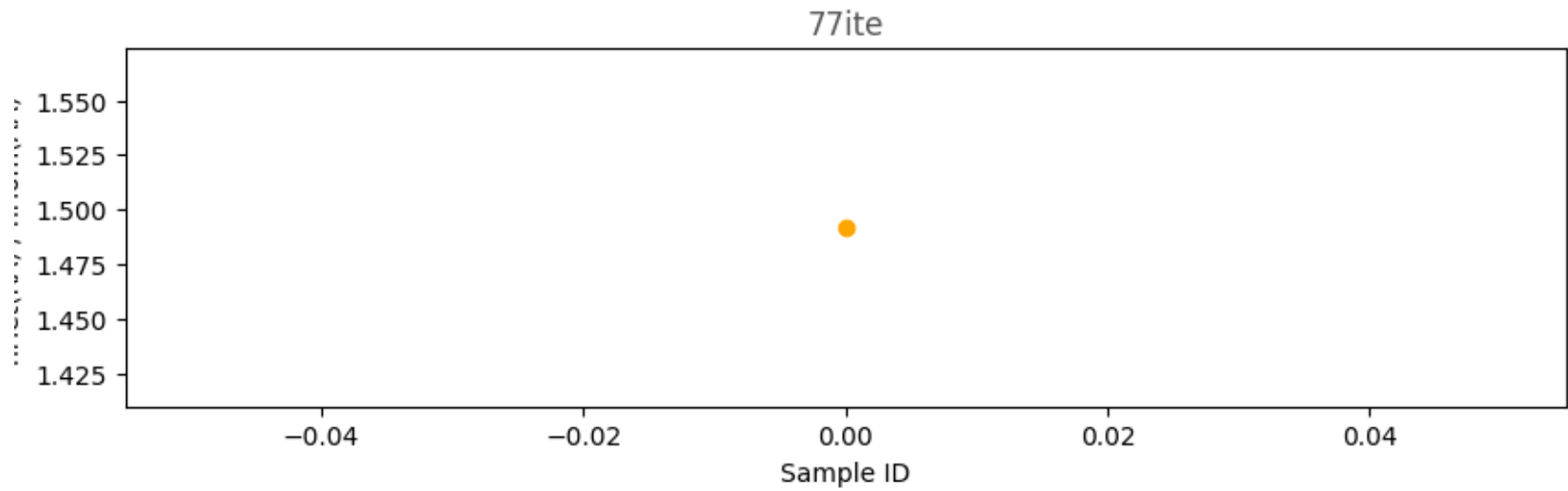
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
77ite	59.9%	1.91	63.4%	85,089	1,781

- 77ite .. /ngc/projects2/gm/data/archive/2022/variants/snv/77itecenm-103831406221-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00534-220202_A00559_BH75J2DSX3-EXT_LAB
KA_NGCWGS-NGCWGS03748_22RKG001630x01_snv_germline_raw.haplotype_caller.vcf.gz

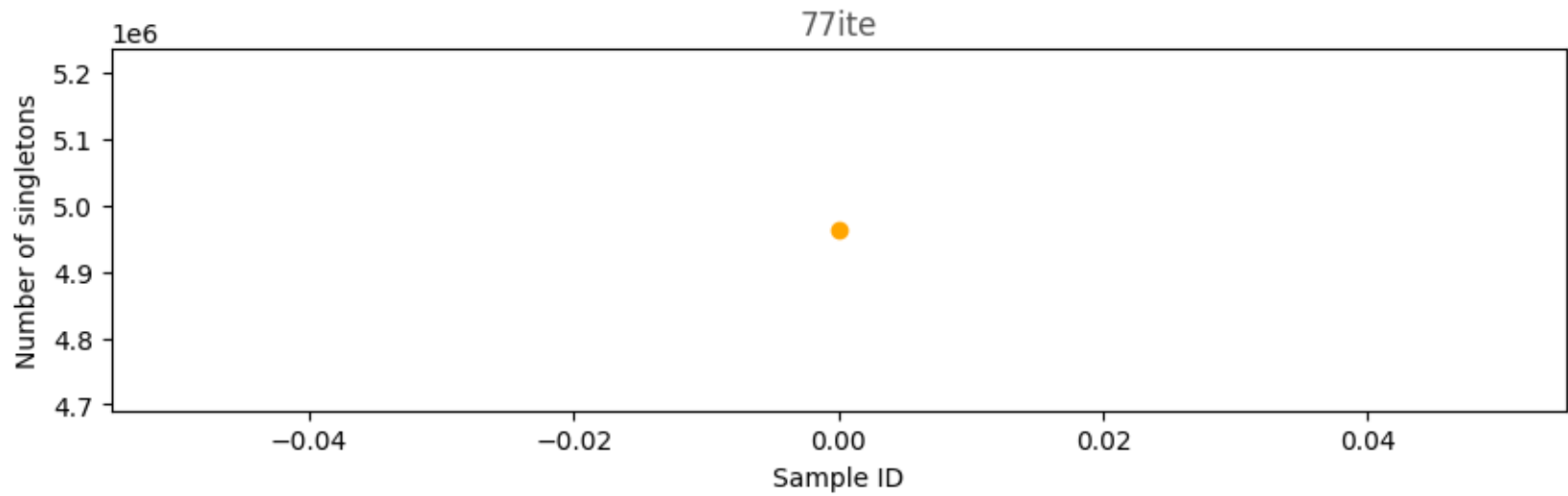
Ts/Tv by sample



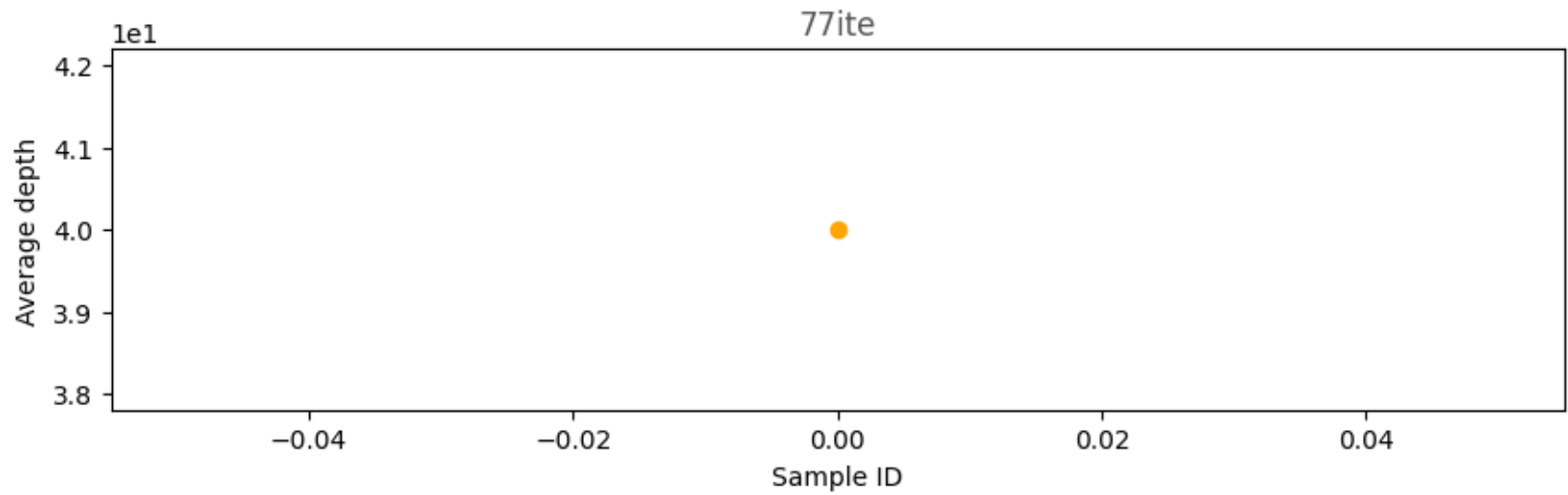
Hets vs non-ref Homs by sample



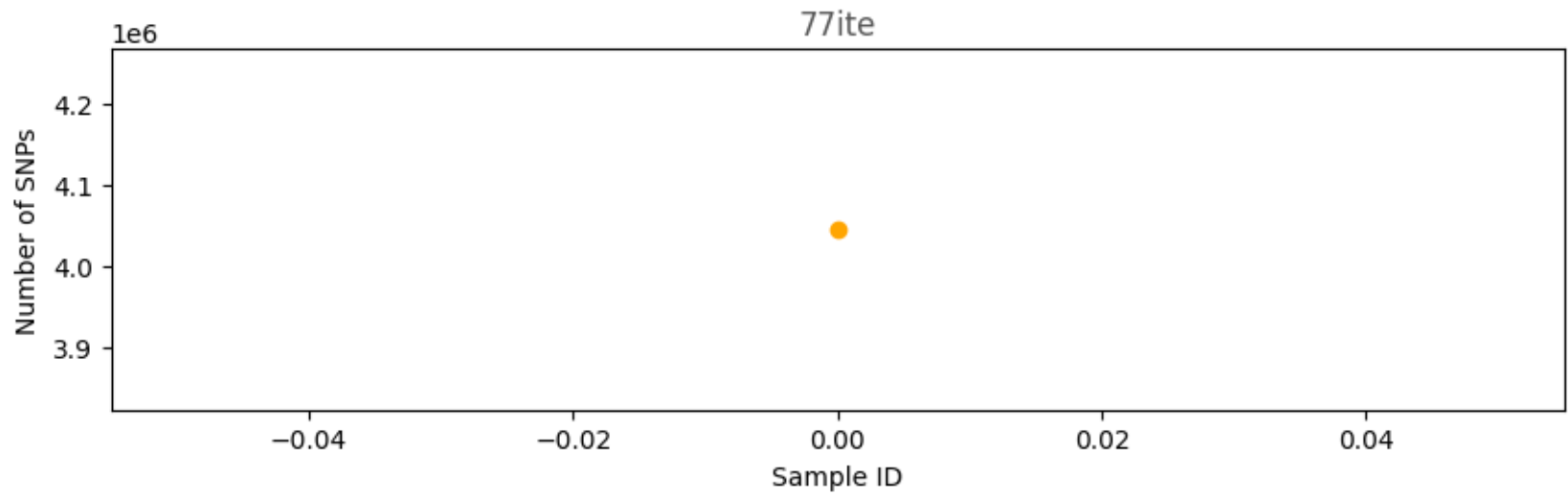
Singletons by sample (hets and homs)



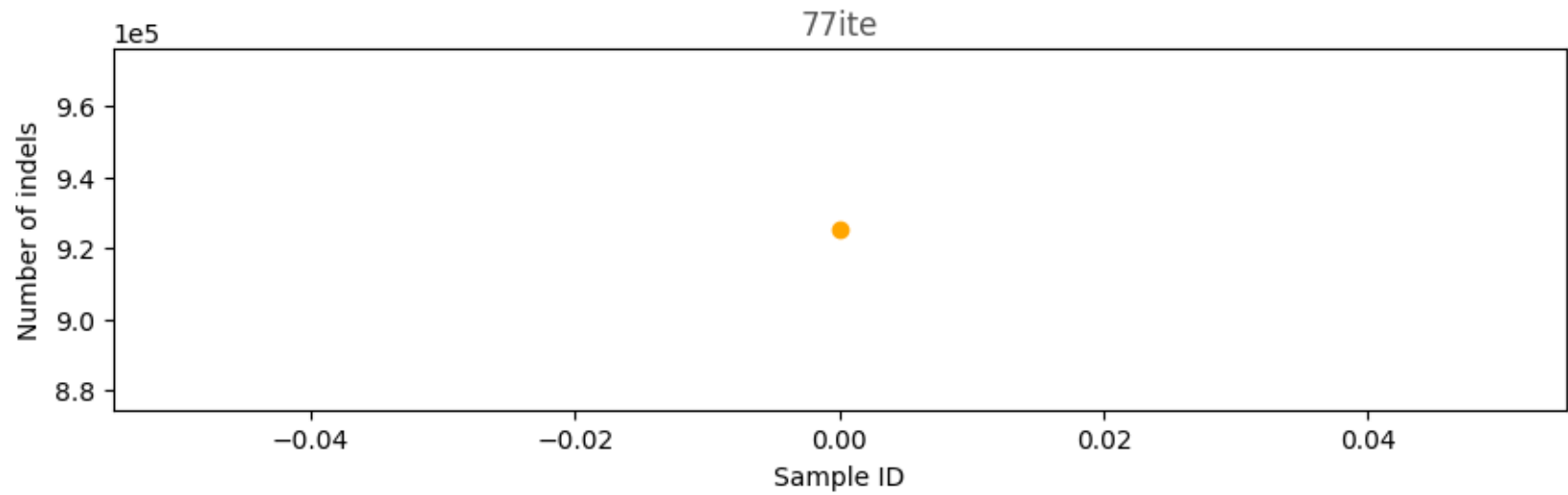
Average depth by sample



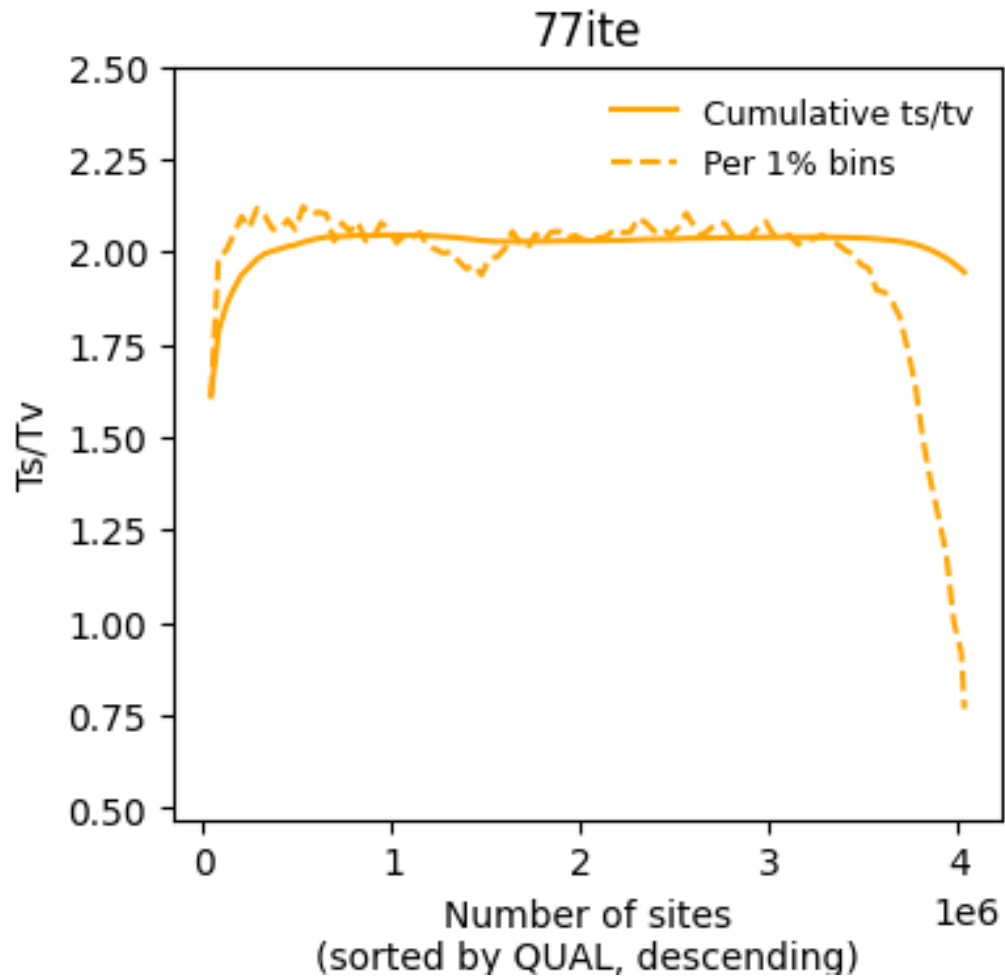
Number of SNPs by sample



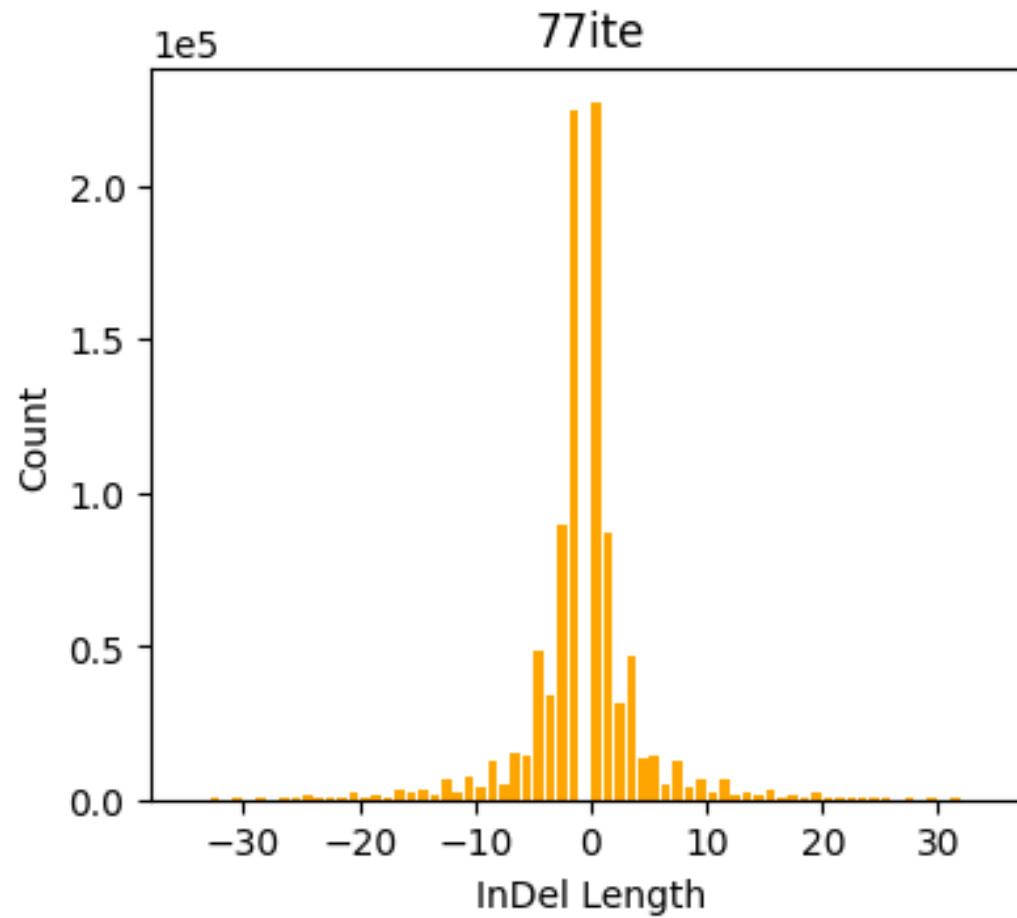
Number of indels by sample



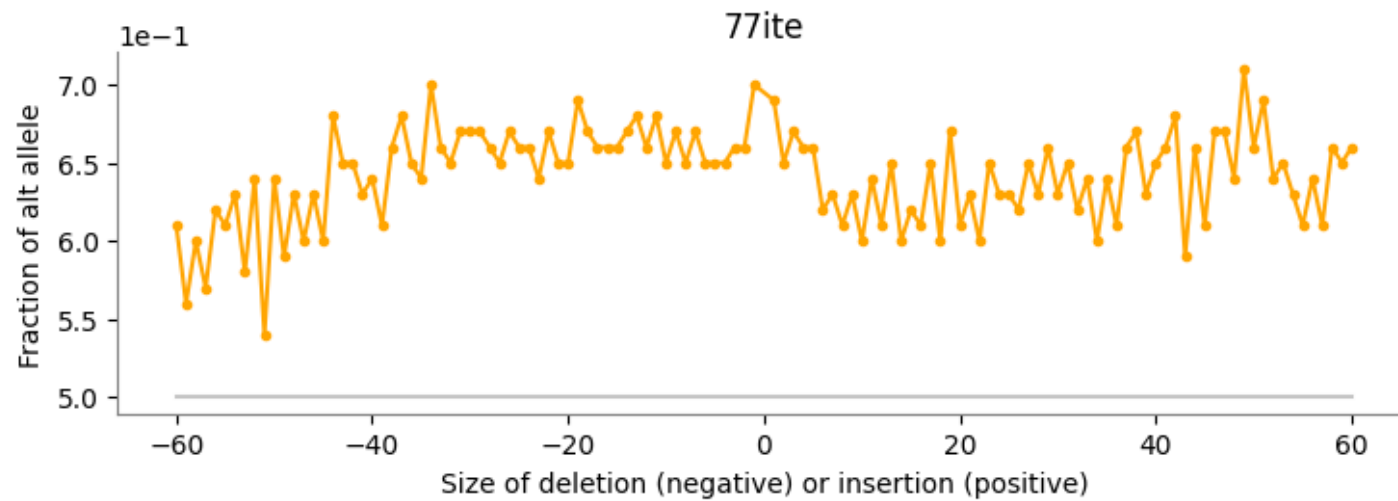
Ts/Tv stratified by QUAL



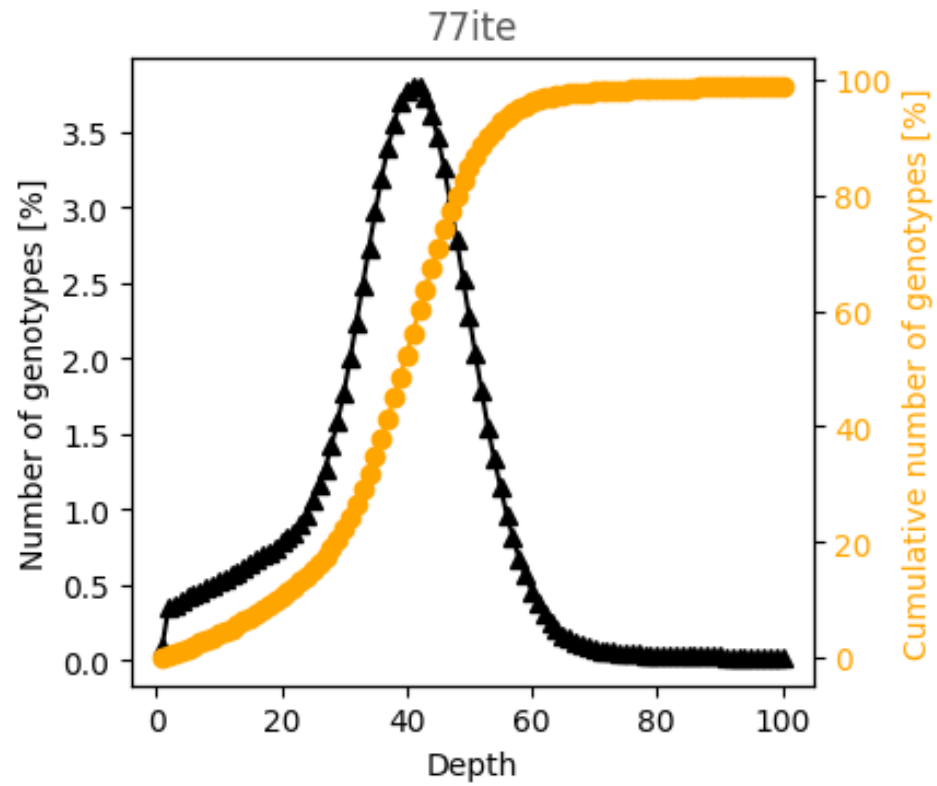
Indel distribution



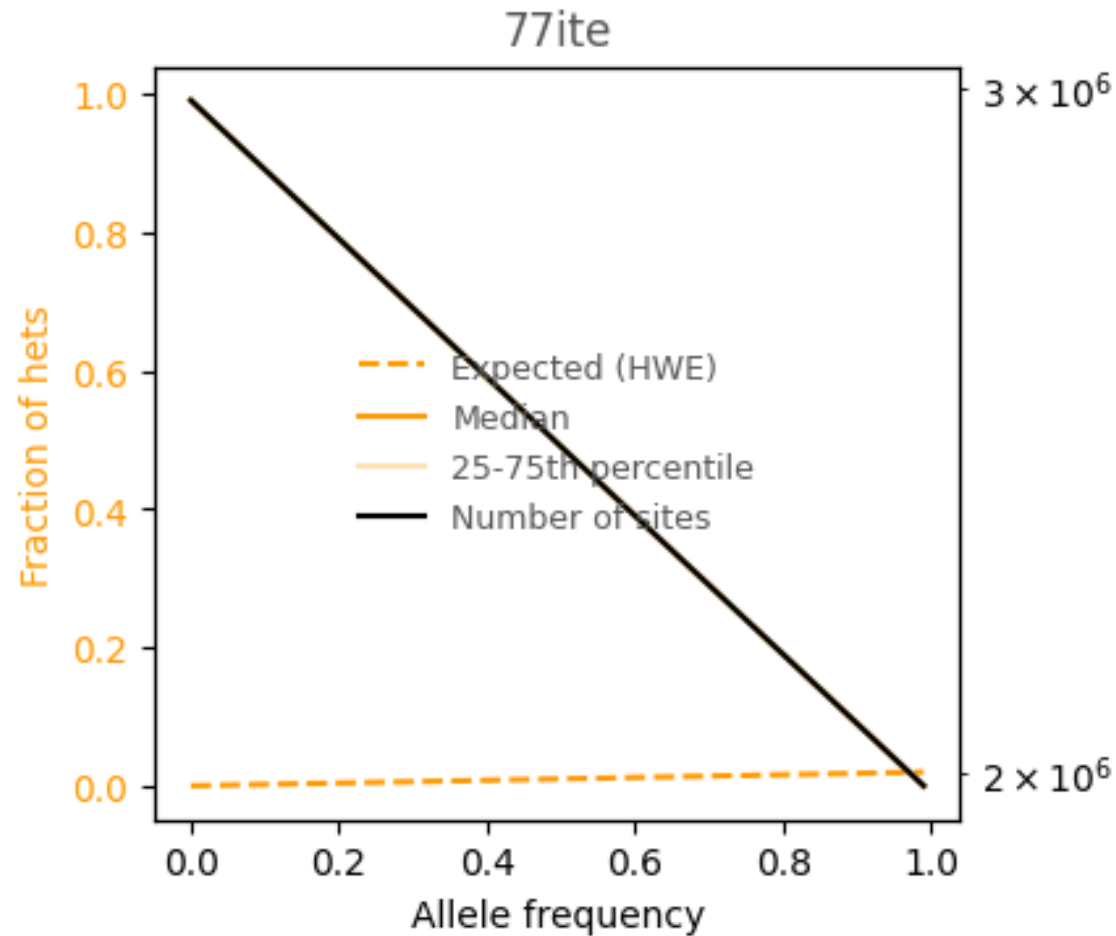
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

