

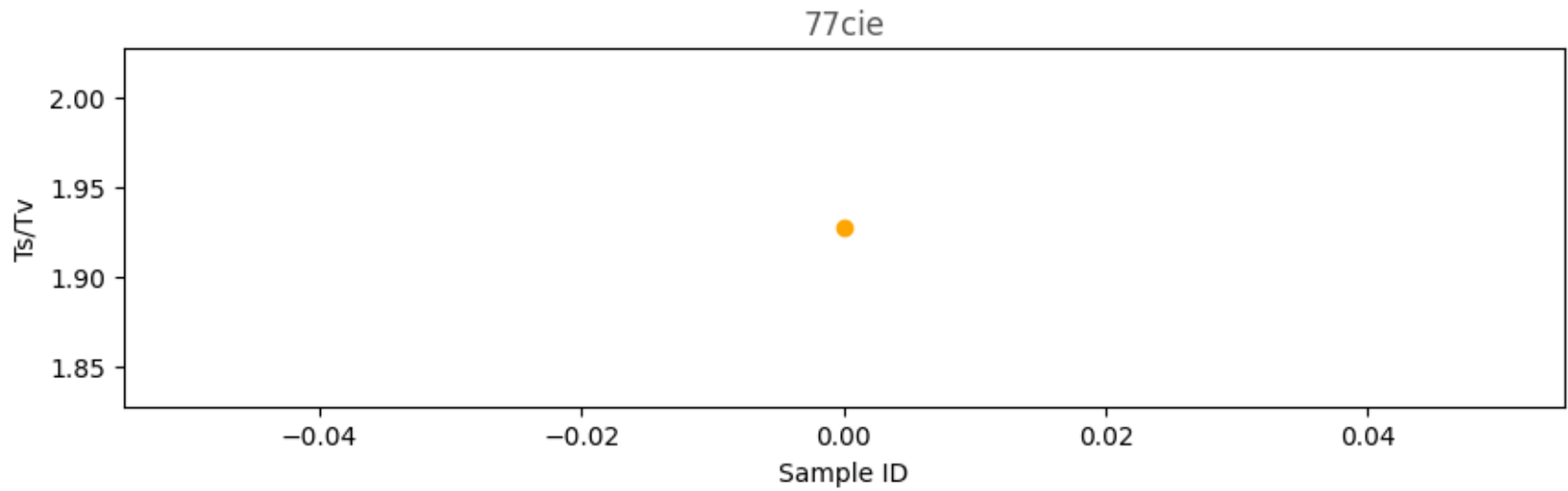
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
77cie	4,125,032	1.93	1.93	967,032	–	0	0
* frameshift ratio: out/(out+in)							

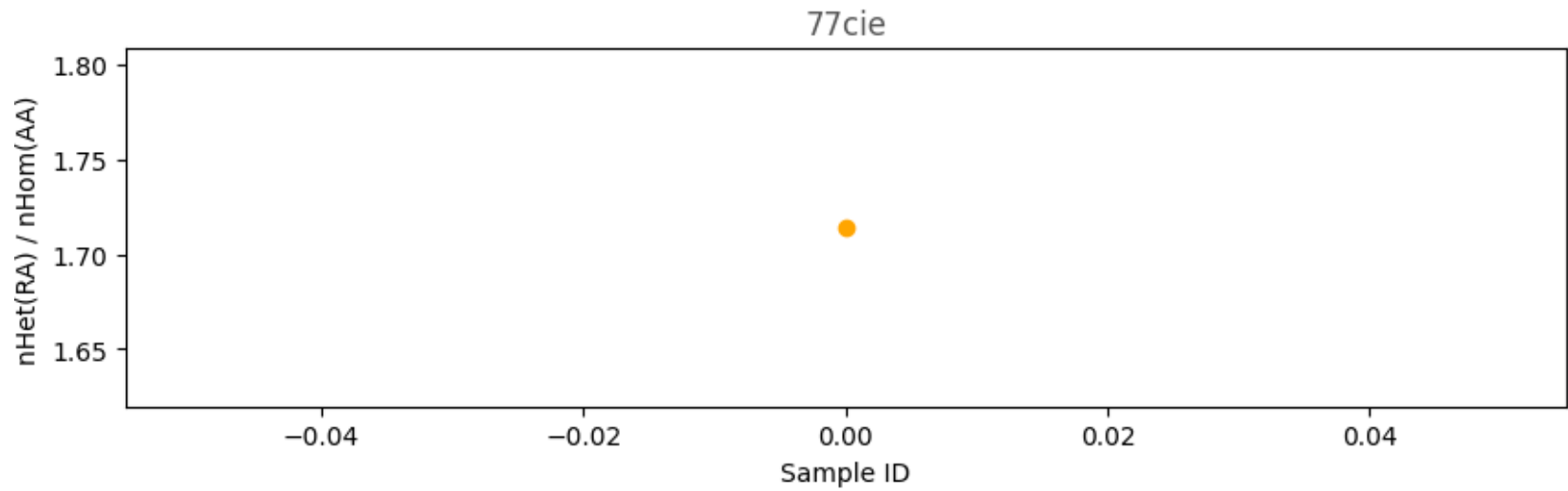
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
77cie	63.2%	1.90	68.8%	108,128	2,332

- 77cie .. /ngc/projects2/gm/data/archive/2022/variants/snv/77cieblef-110295668052-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01989-221012_A01176_AH2CWMSX5-EXT_LAB_KA_NGCWGS-NGCWGS05810_snv_germline_raw.haplotype_caller.vcf.gz

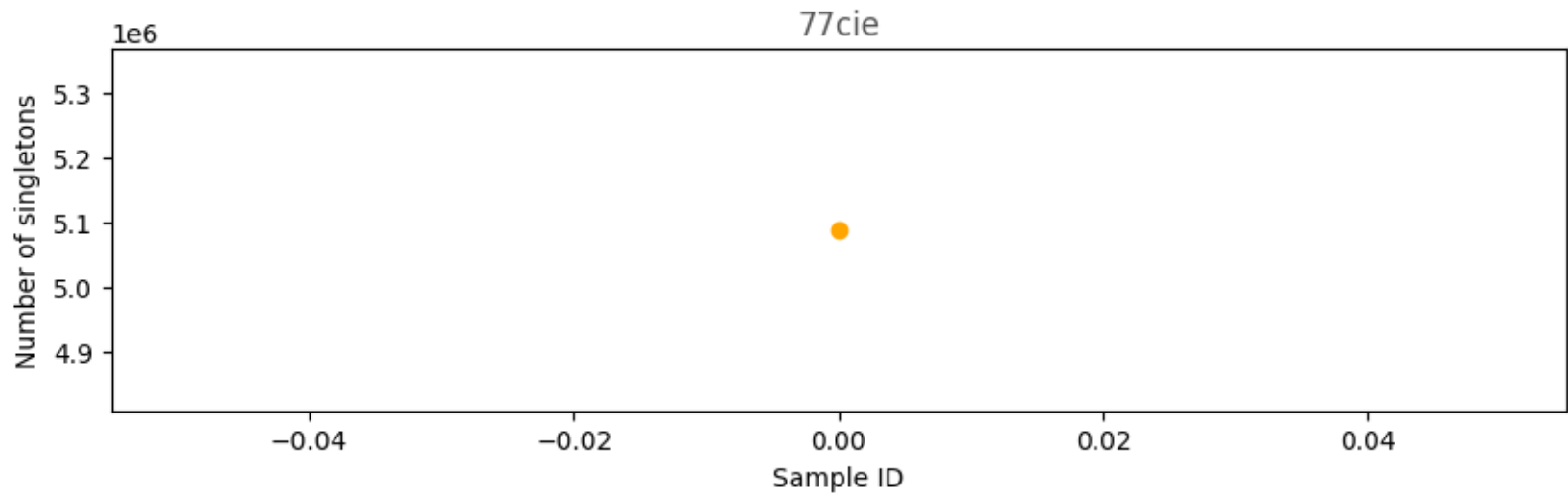
Ts/Tv by sample



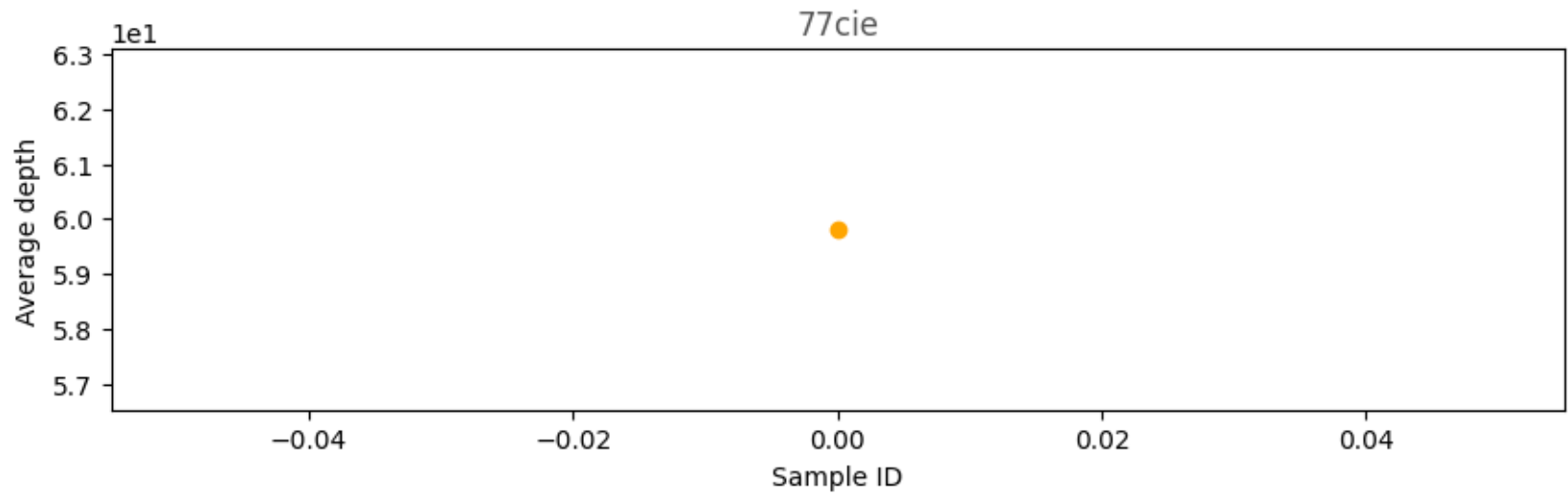
Hets vs non-ref Homs by sample



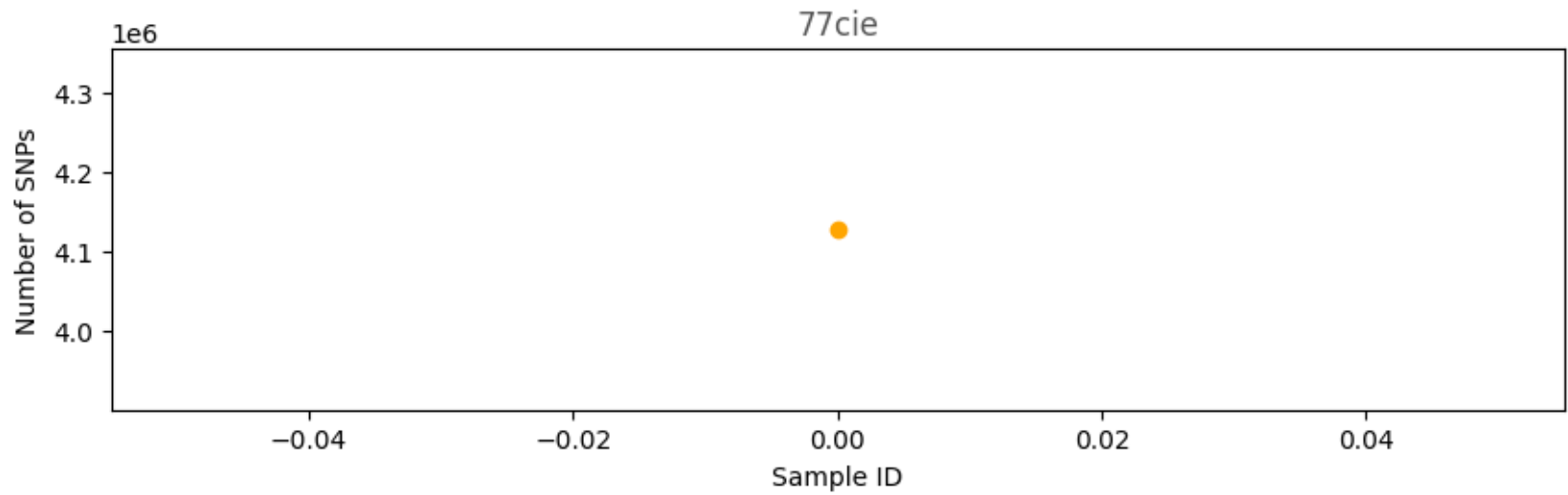
Singletons by sample (hets and homs)



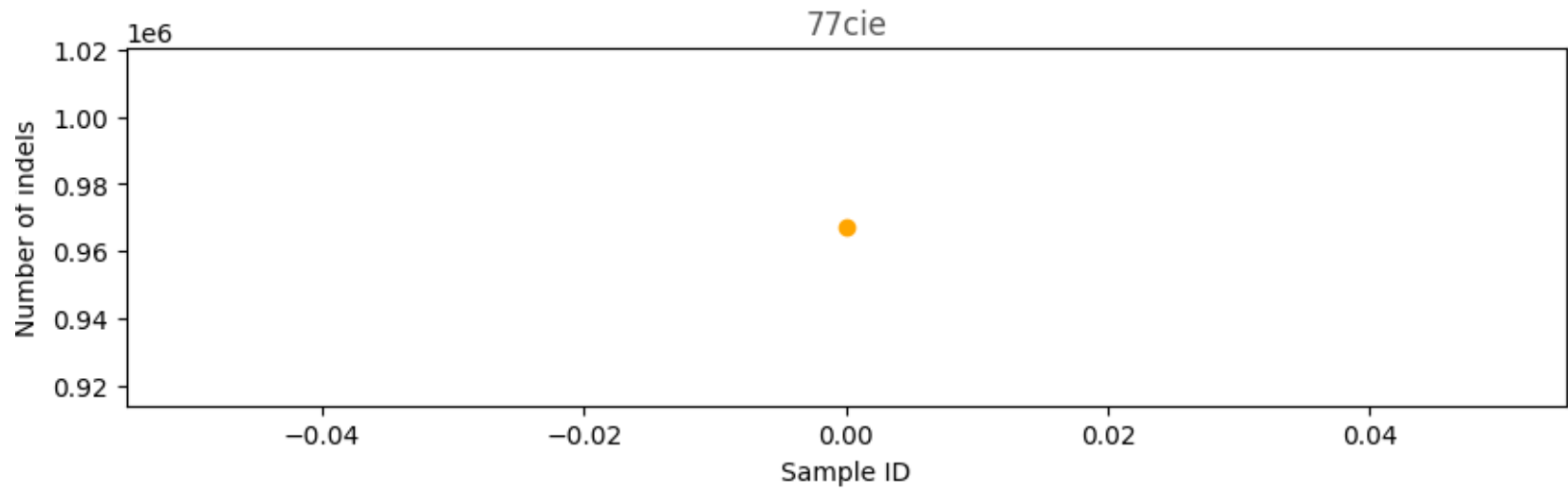
Average depth by sample



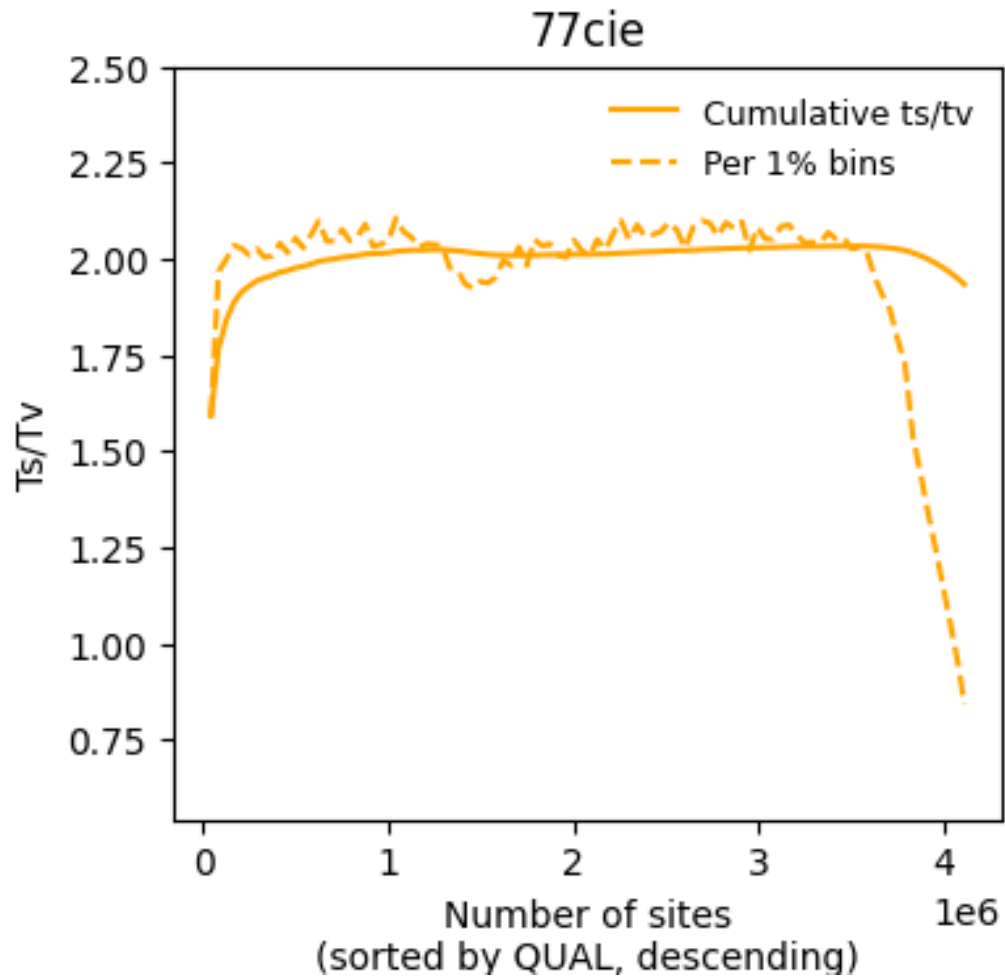
Number of SNPs by sample



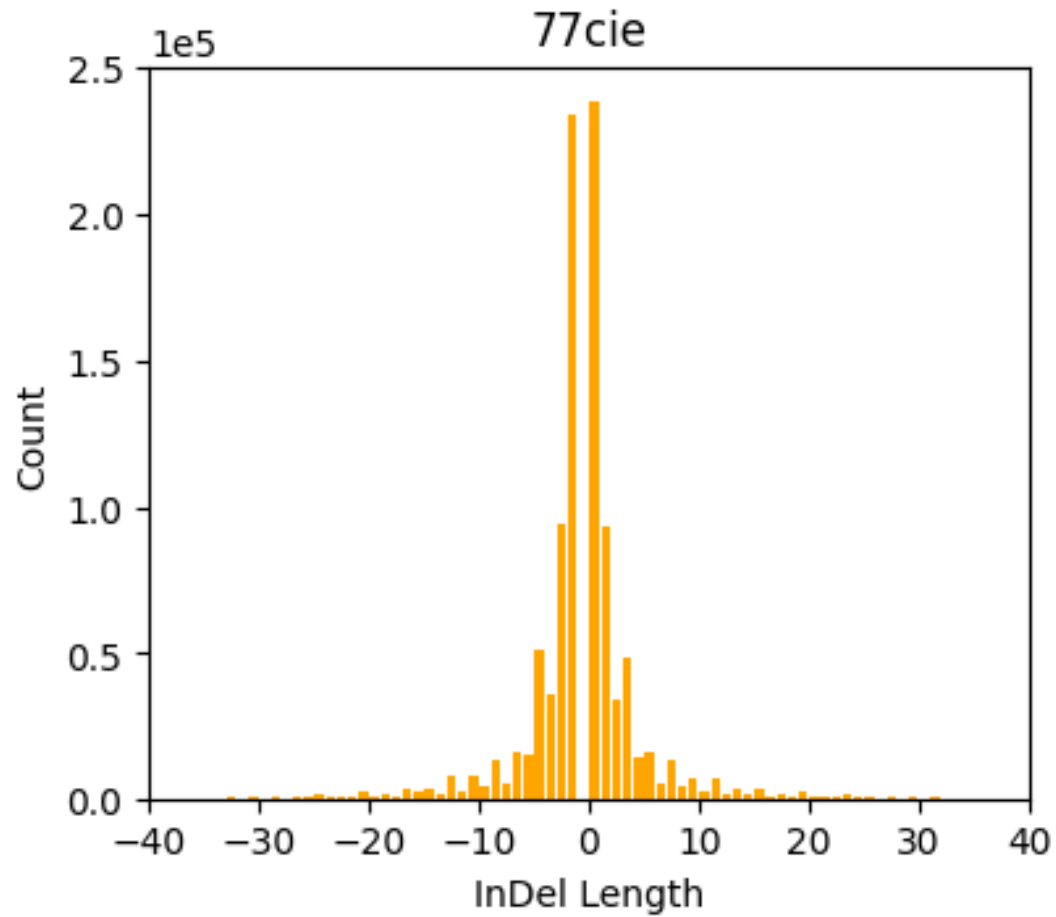
Number of indels by sample



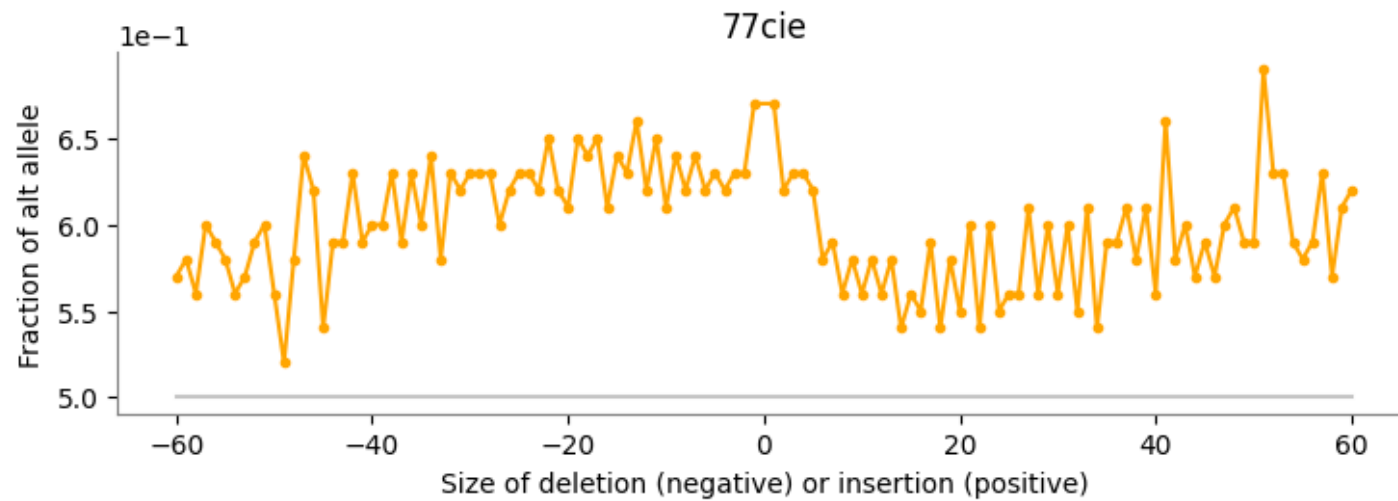
Ts/Tv stratified by QUAL



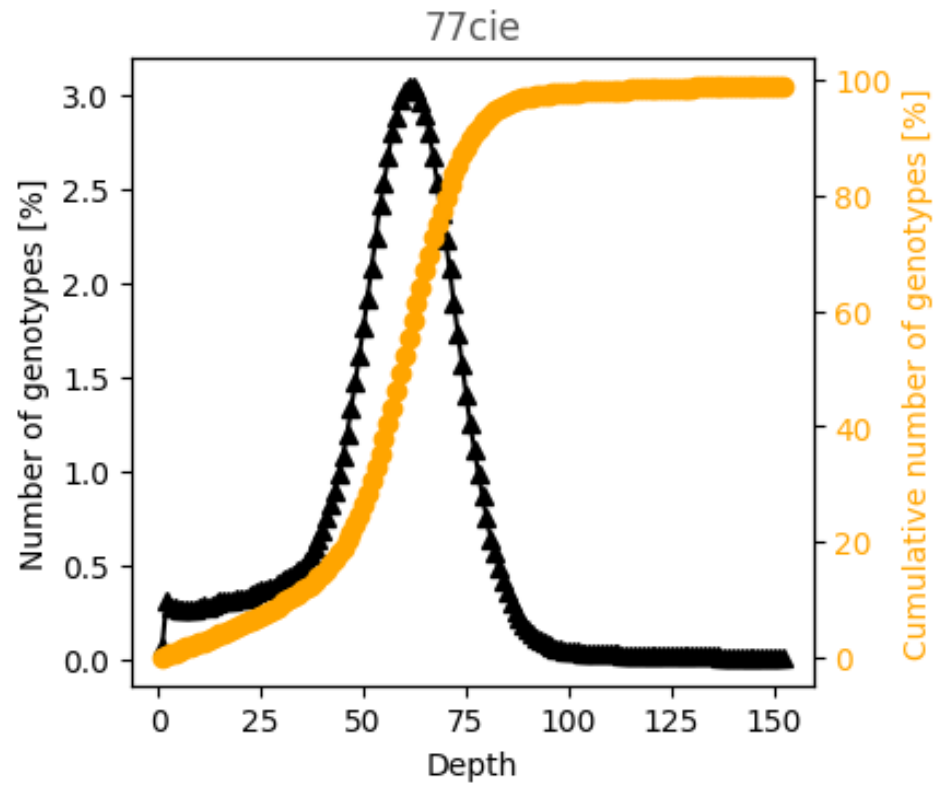
Indel distribution



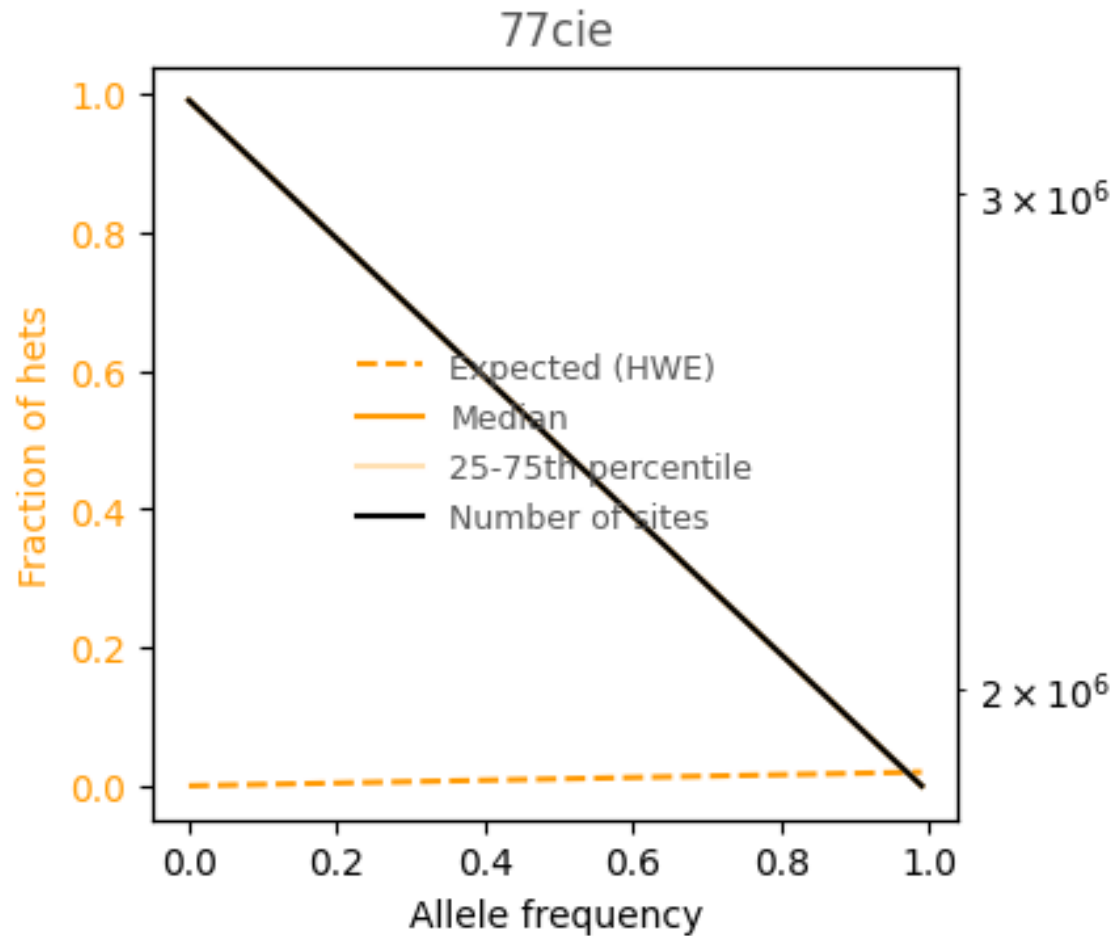
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

