

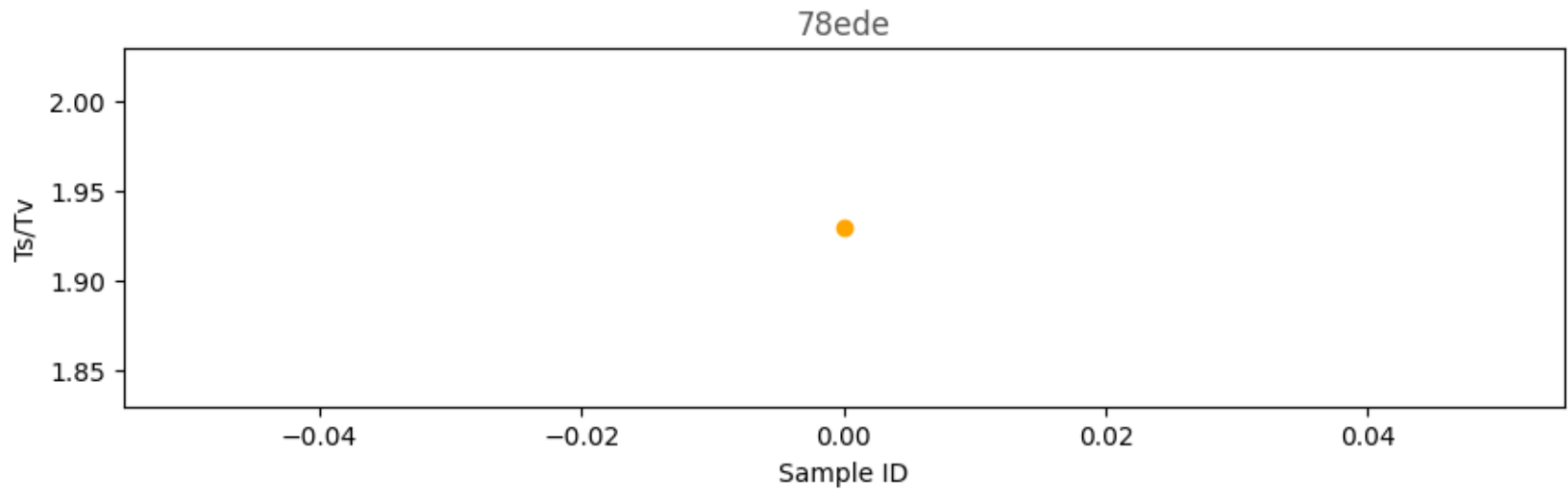
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
78ede	4,074,694	1.93	1.93	949,175	–	0	0
* frameshift ratio: out/(out+in)							

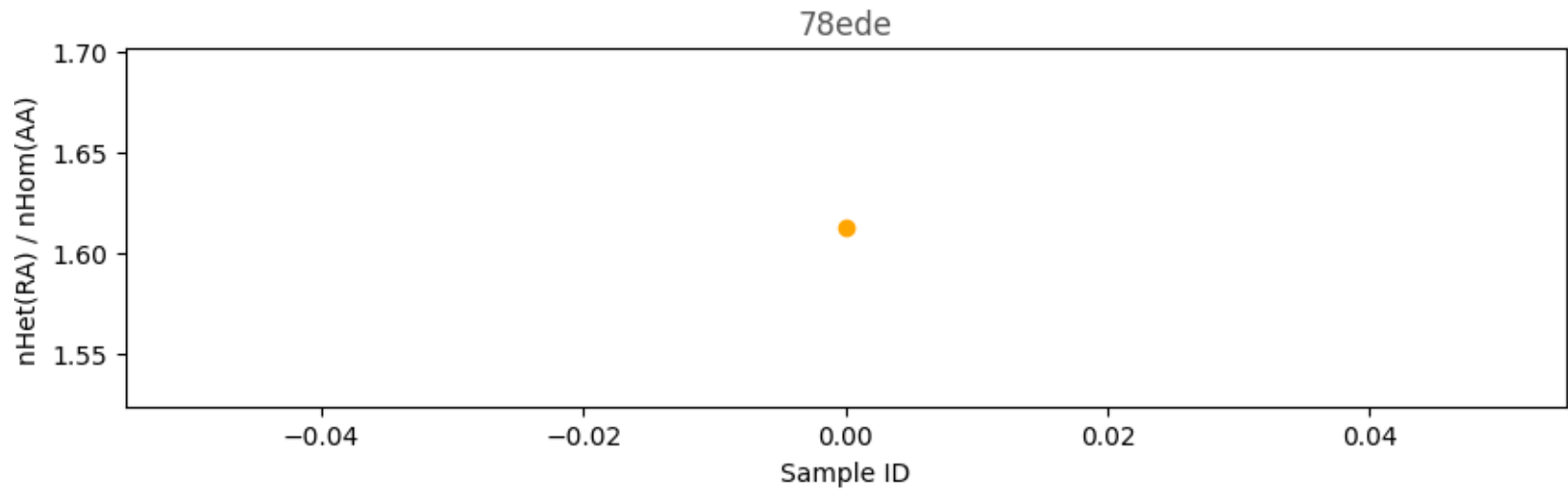
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
78ede	61.7%	1.90	66.4%	98,503	2,049

- 78ede .. /ngc/projects2/gm/data/archive/2022/variants/snv/78edeeotm-103911978635-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01668-220921_A01176_BHYLVHDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS05461_snv_germline_raw.haplotype_caller.vcf.gz

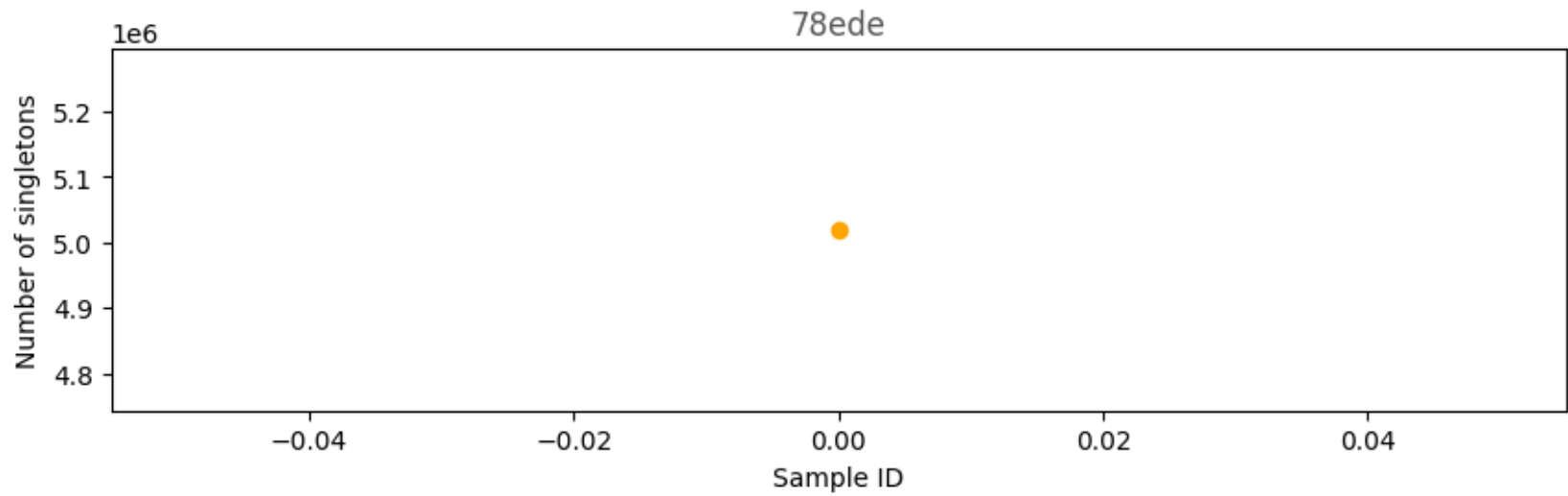
Ts/Tv by sample



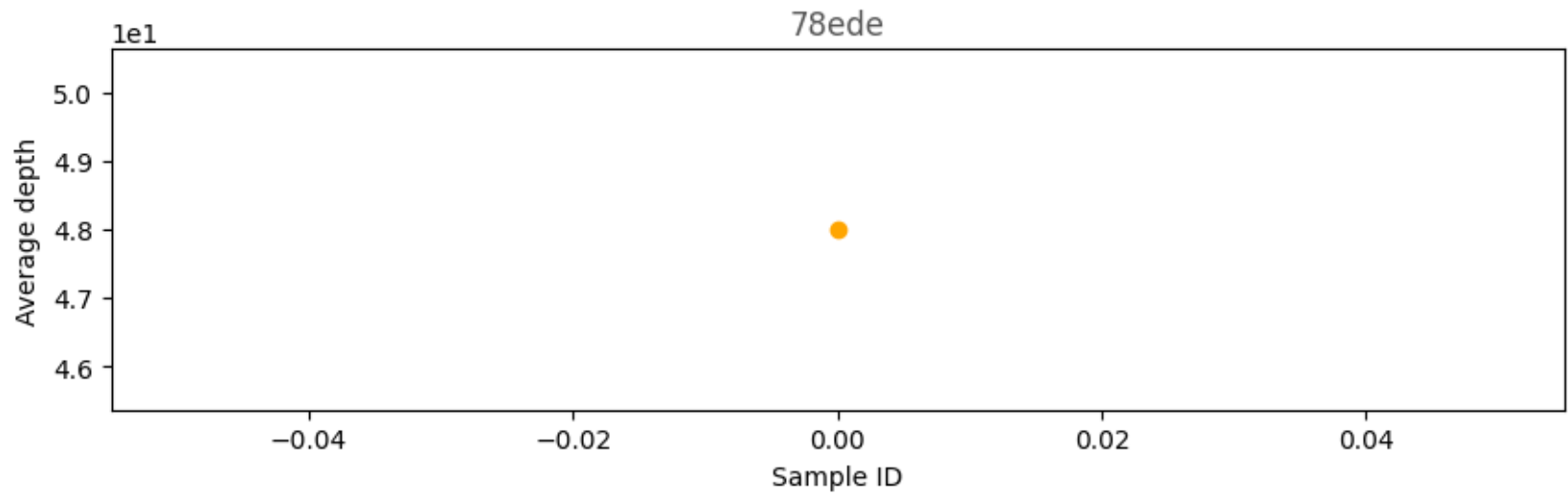
Hets vs non-ref Homs by sample



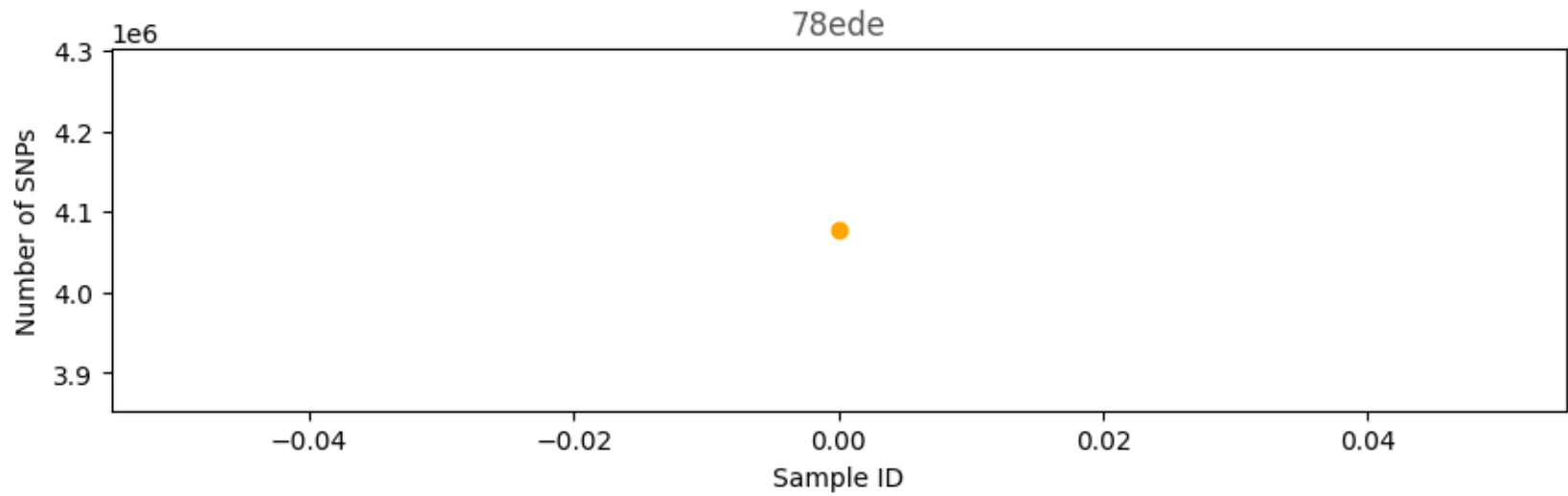
Singletons by sample (hets and homs)



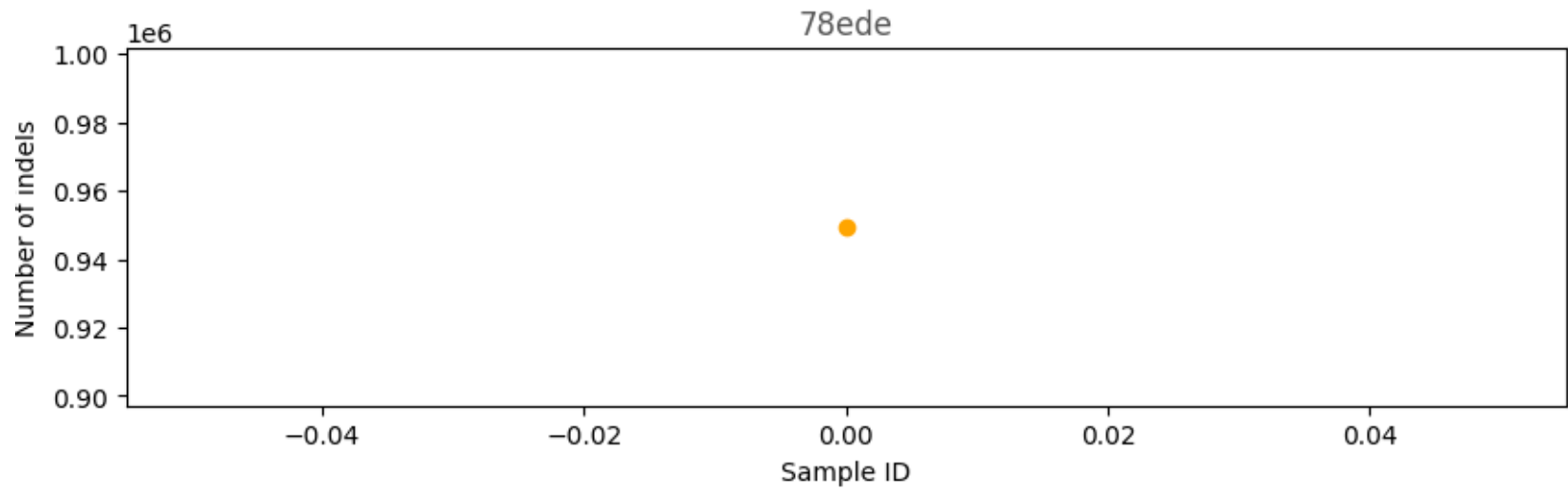
Average depth by sample



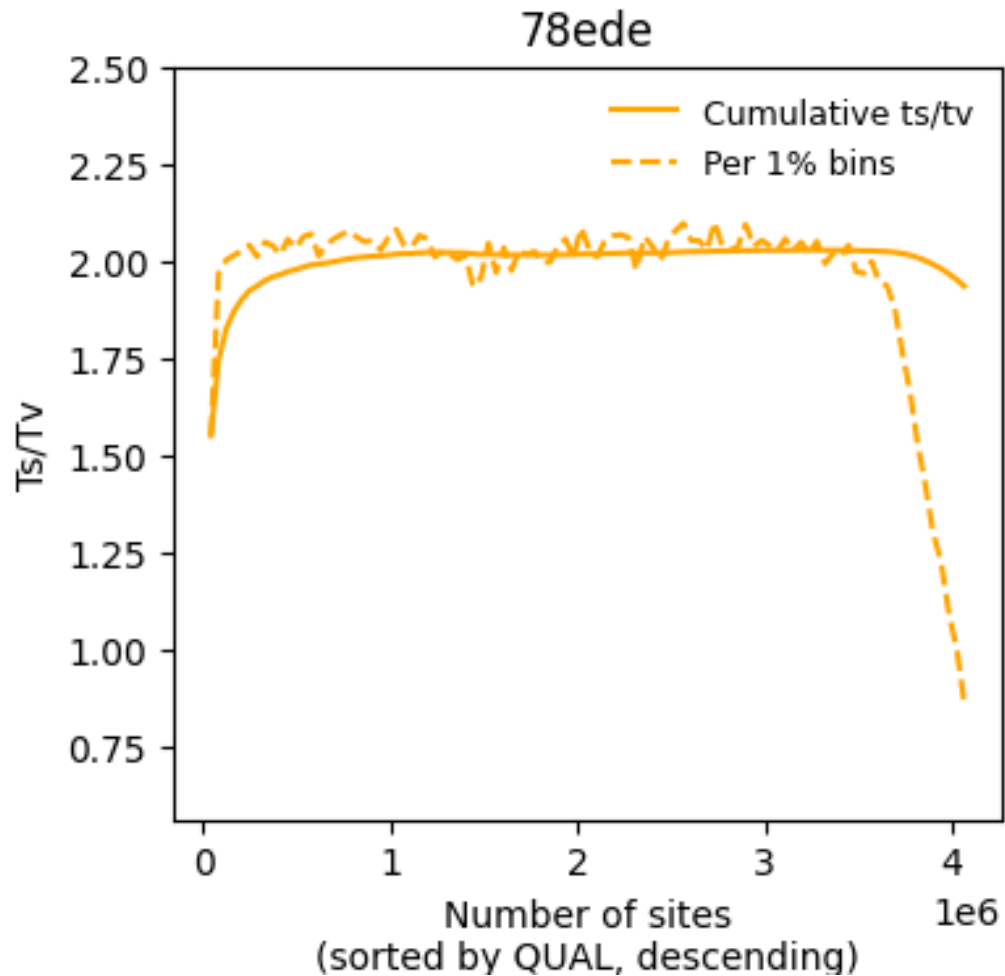
Number of SNPs by sample



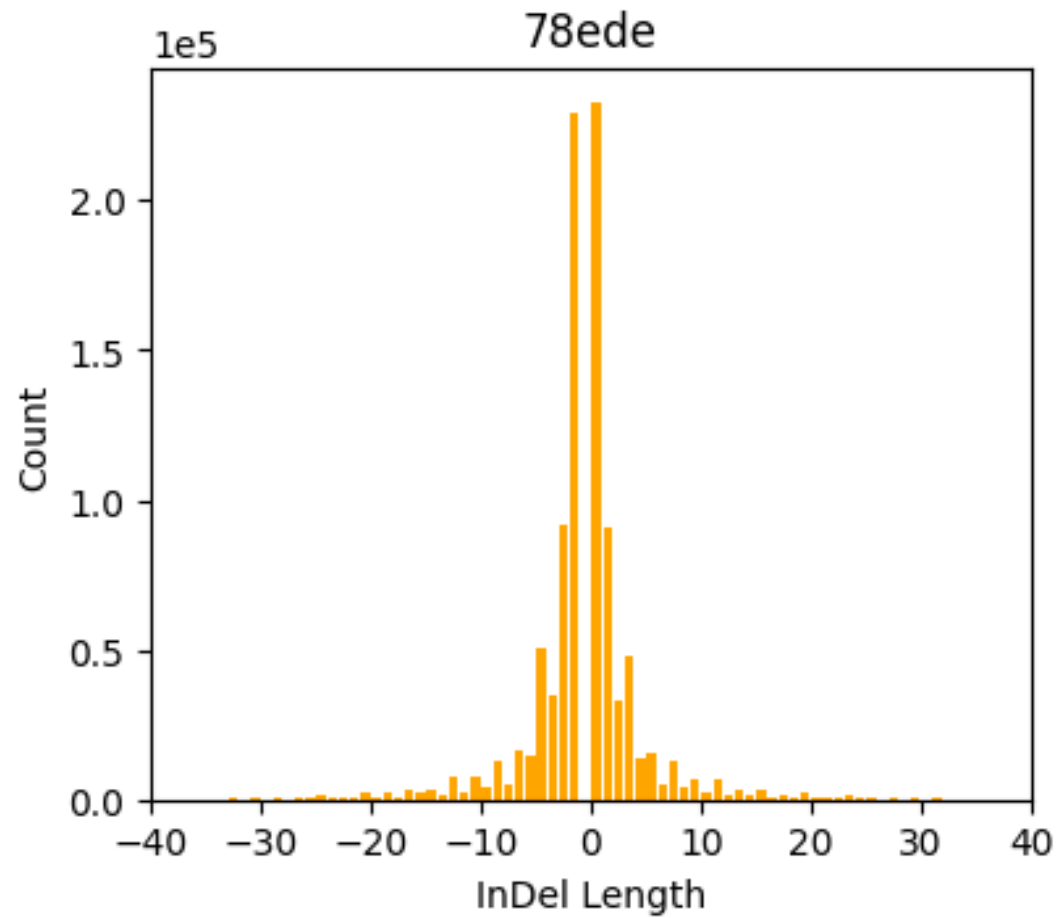
Number of indels by sample



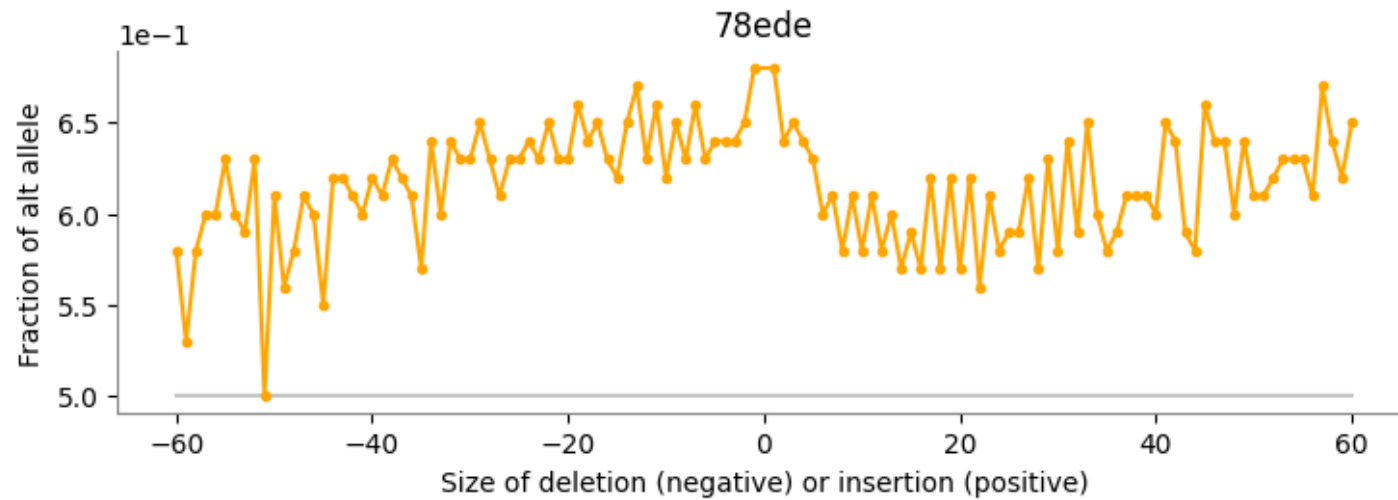
Ts/Tv stratified by QUAL



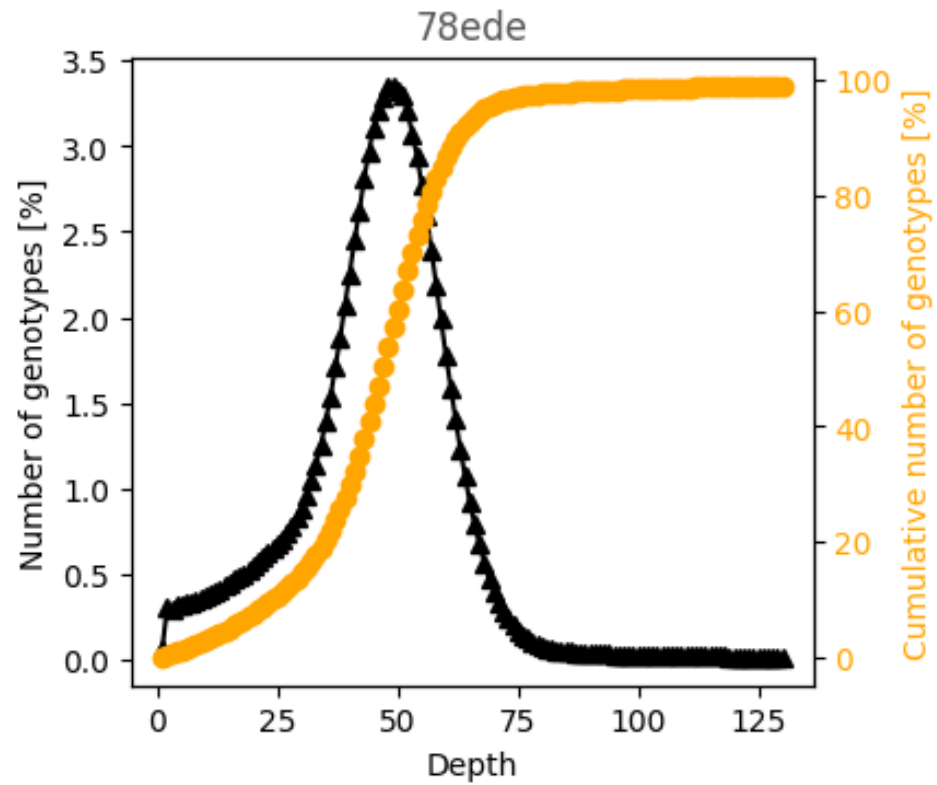
Indel distribution



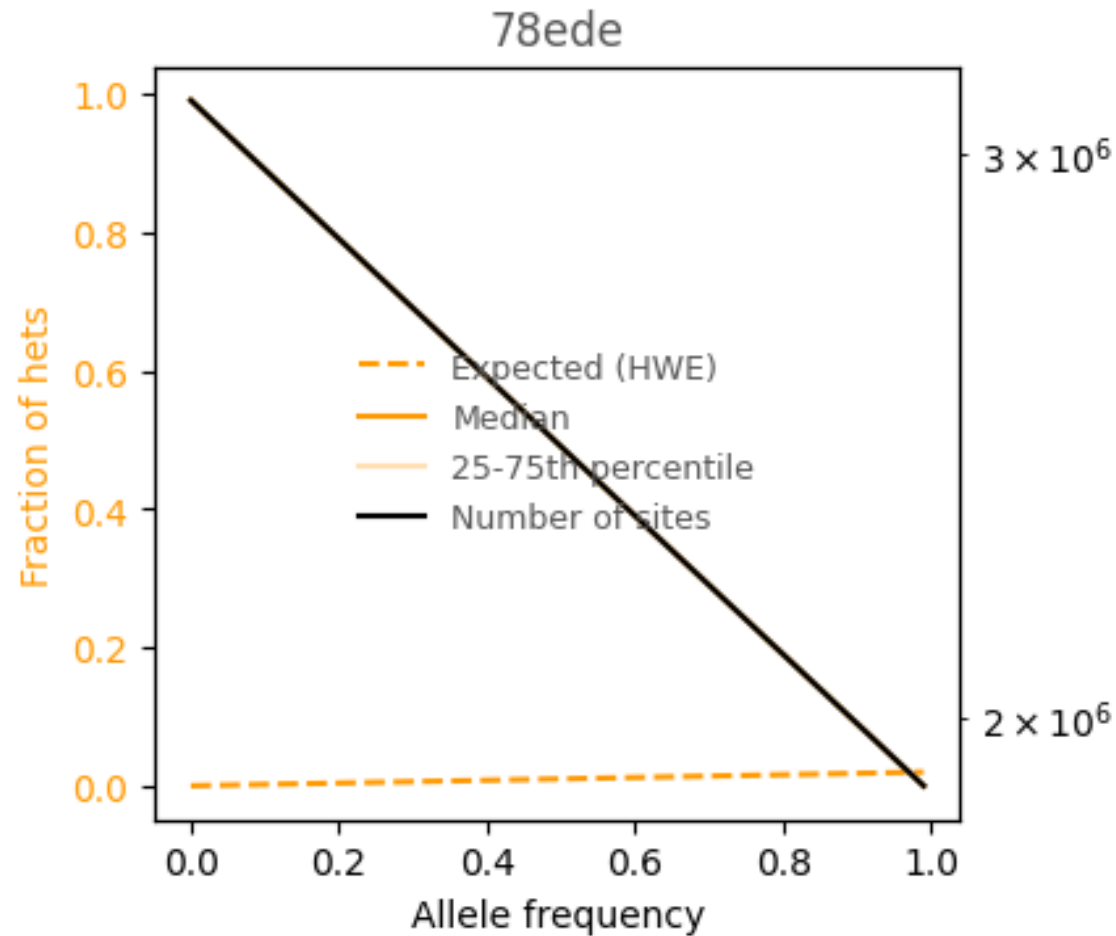
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

