

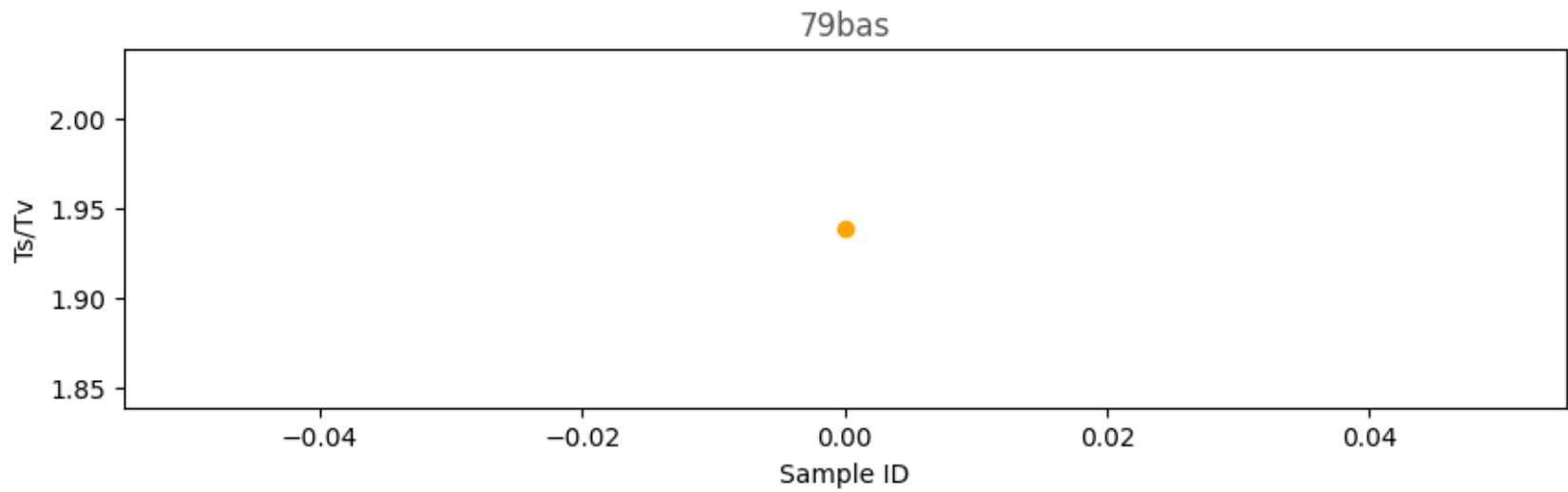
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
79bas	3,997,193	1.94	1.94	912,651	–	0	0
* frameshift ratio: out/(out+in)							

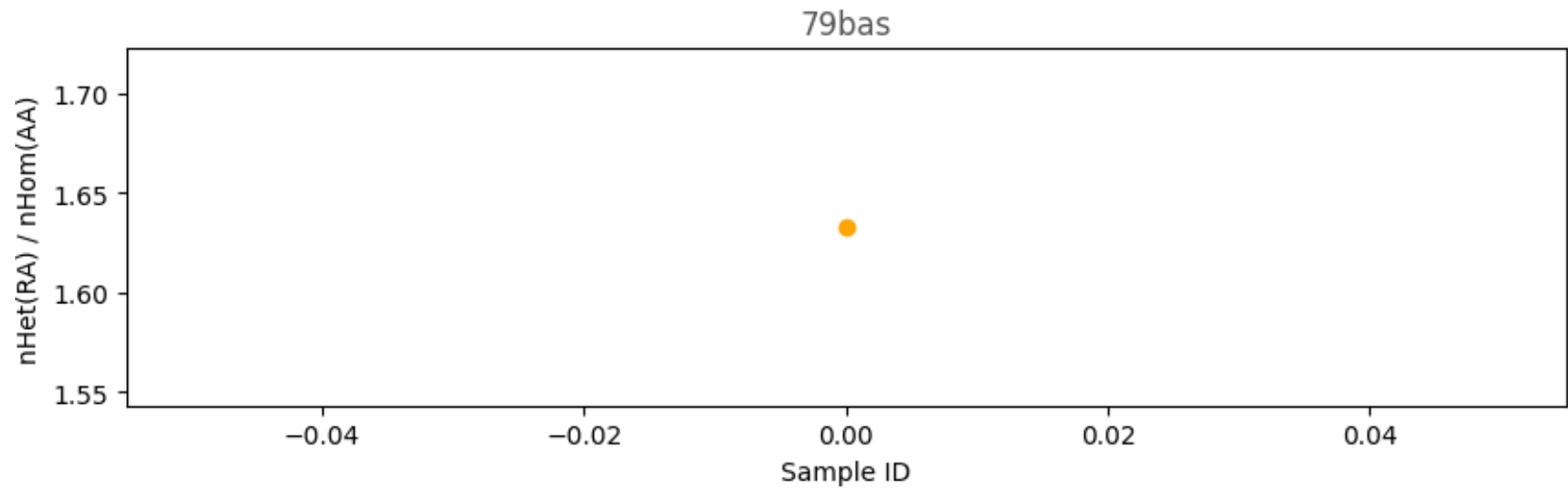
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
79bas	62.0%	1.91	65.4%	85,371	1,728

- 79bas .. /ngc/projects2/gm/data/archive/2022/variants/snv/79baswanm-103831404474-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00550-220202_A00559_BH75J2DSX3-EXT_LAB
KA_NGCWGS-NGCWGS03764_21RKG031906x01_snv_germline_raw.haplotype_caller.vcf.gz

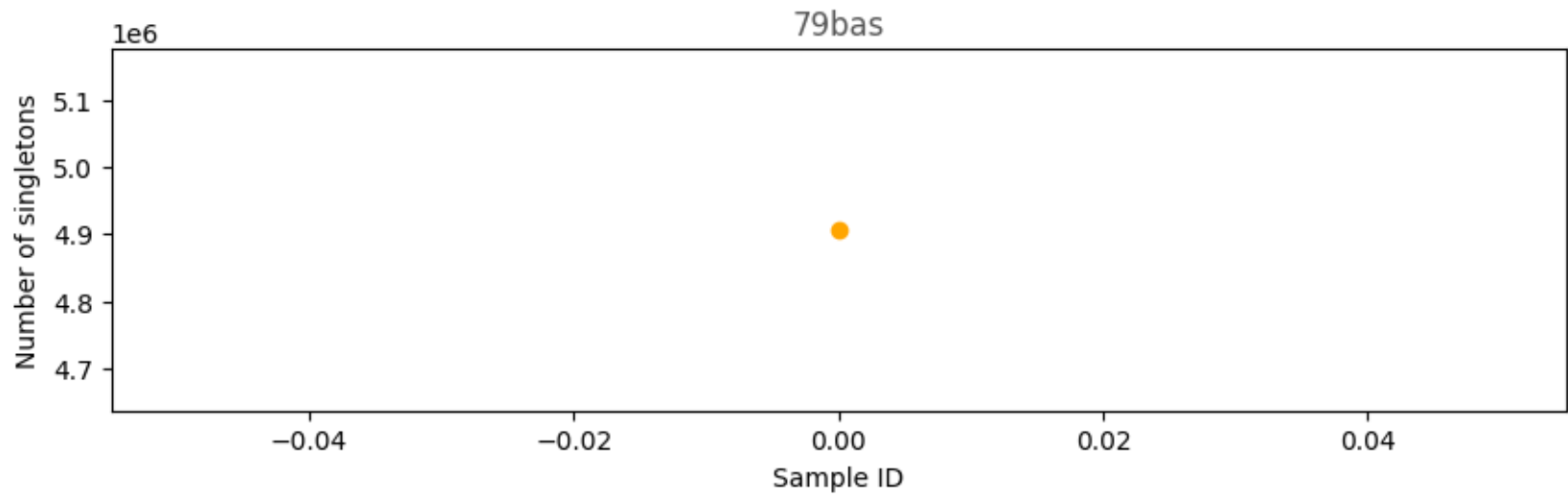
Ts/Tv by sample



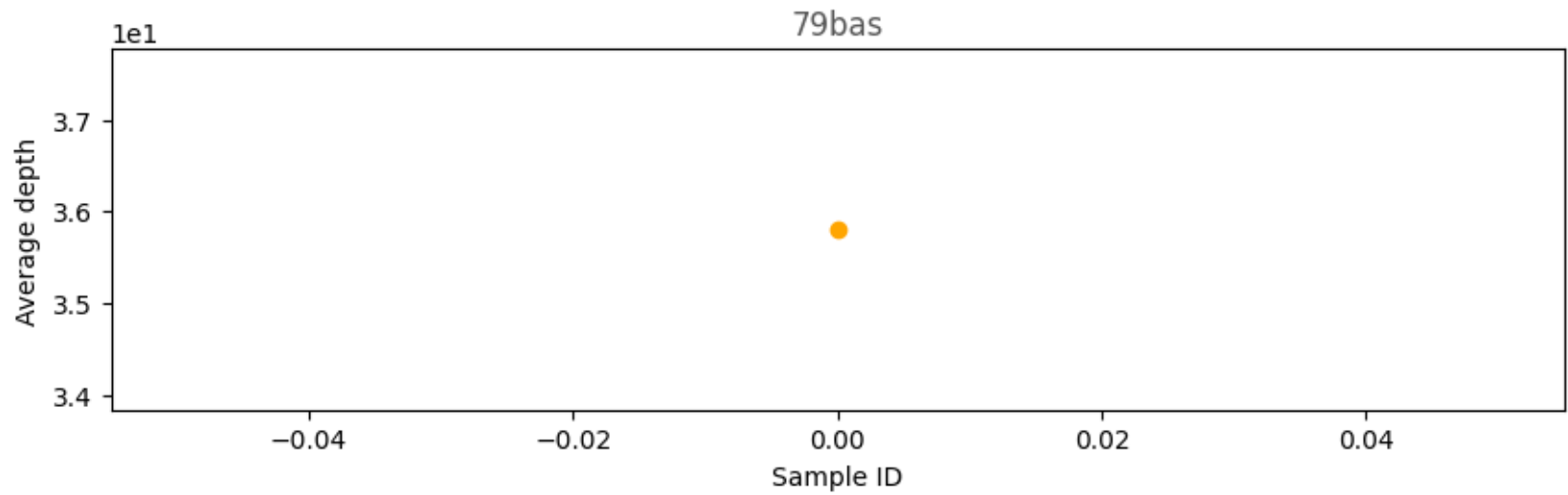
Hets vs non-ref Homs by sample



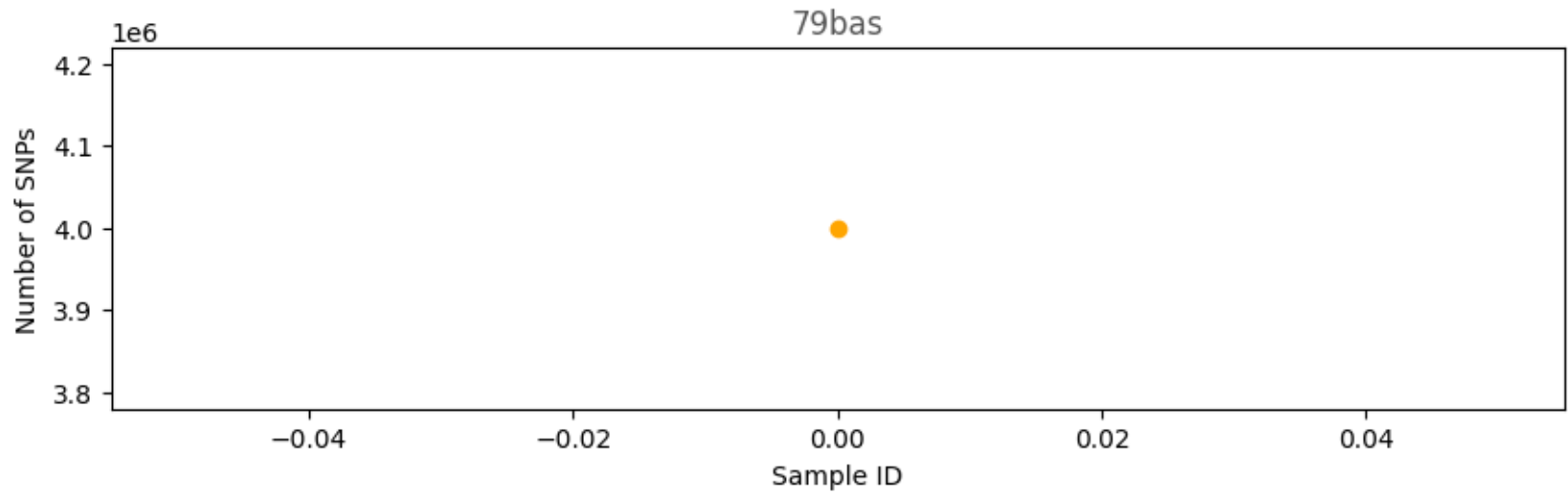
Singletons by sample (hets and homs)



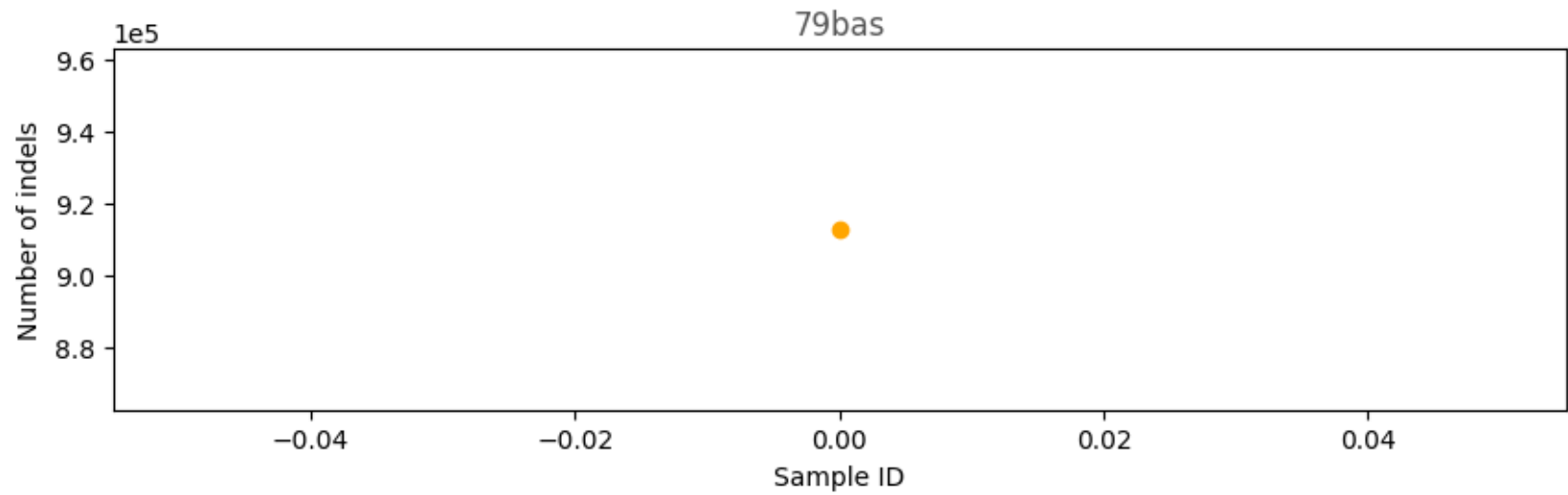
Average depth by sample



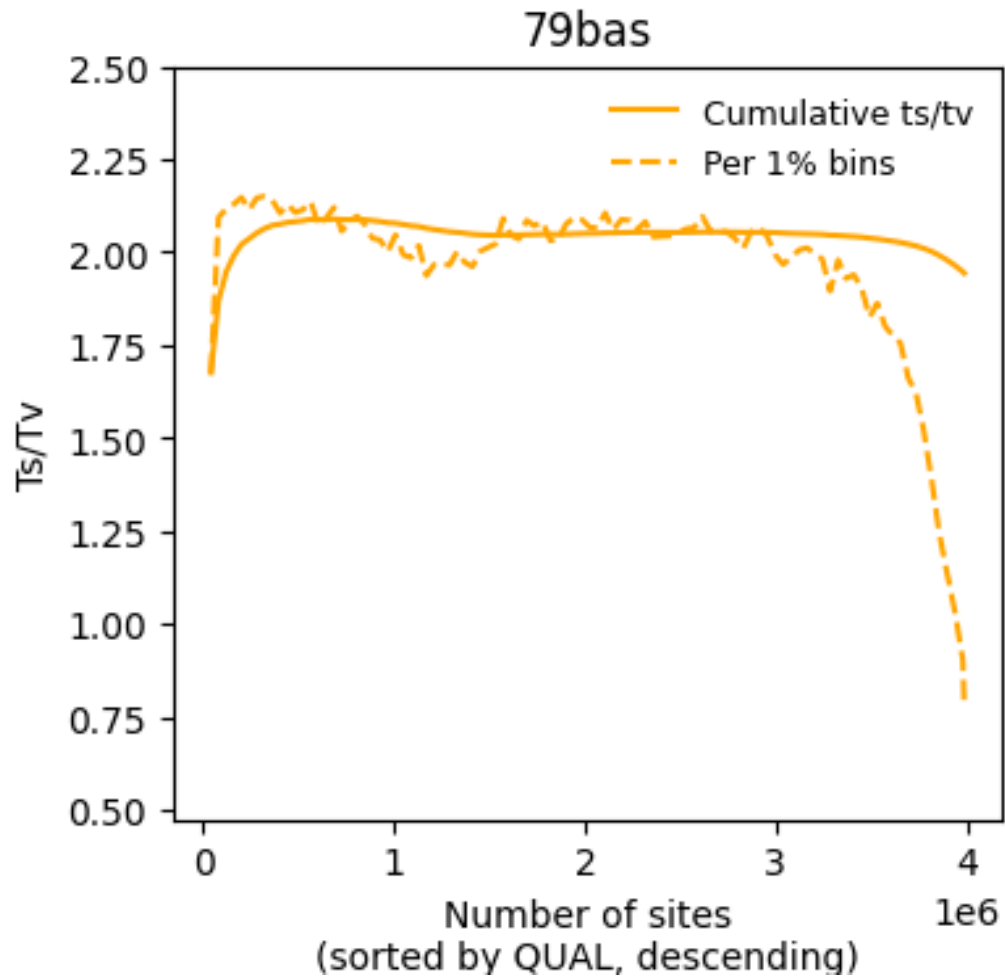
Number of SNPs by sample



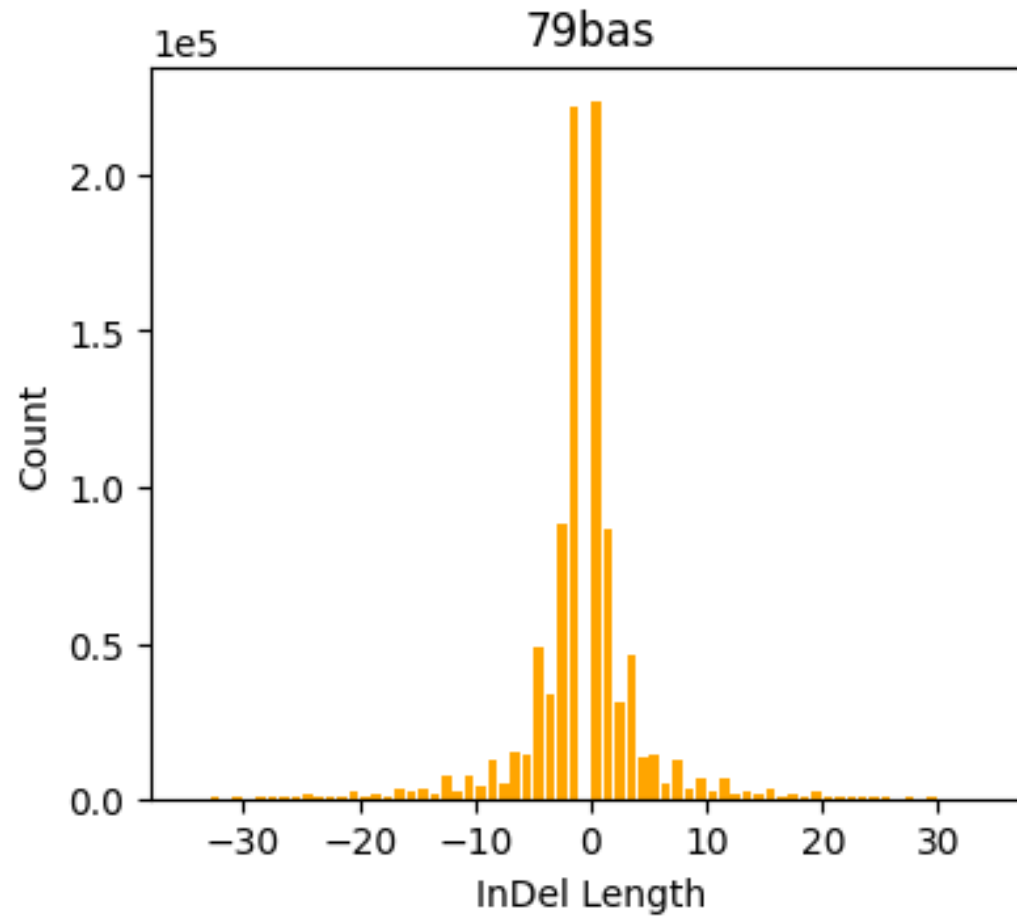
Number of indels by sample



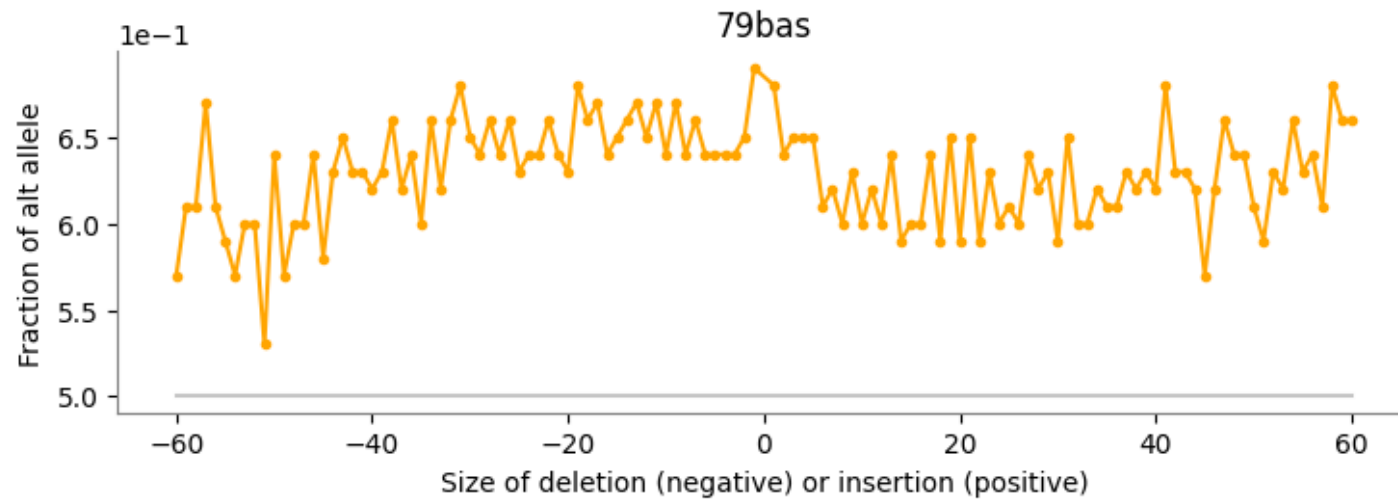
Ts/Tv stratified by QUAL



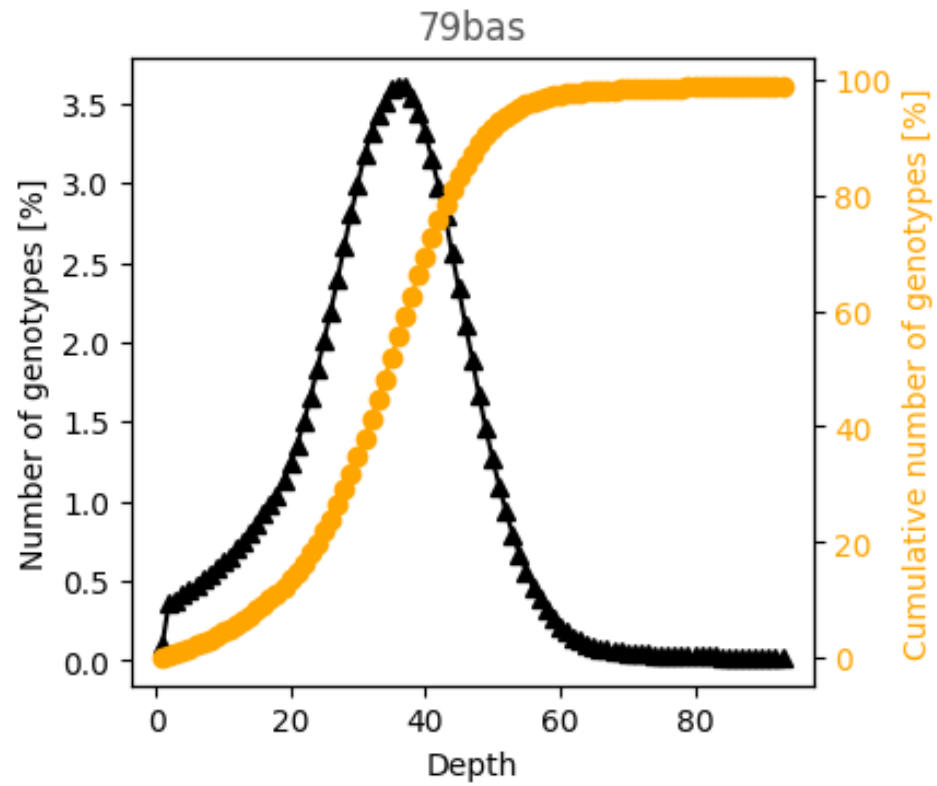
Indel distribution



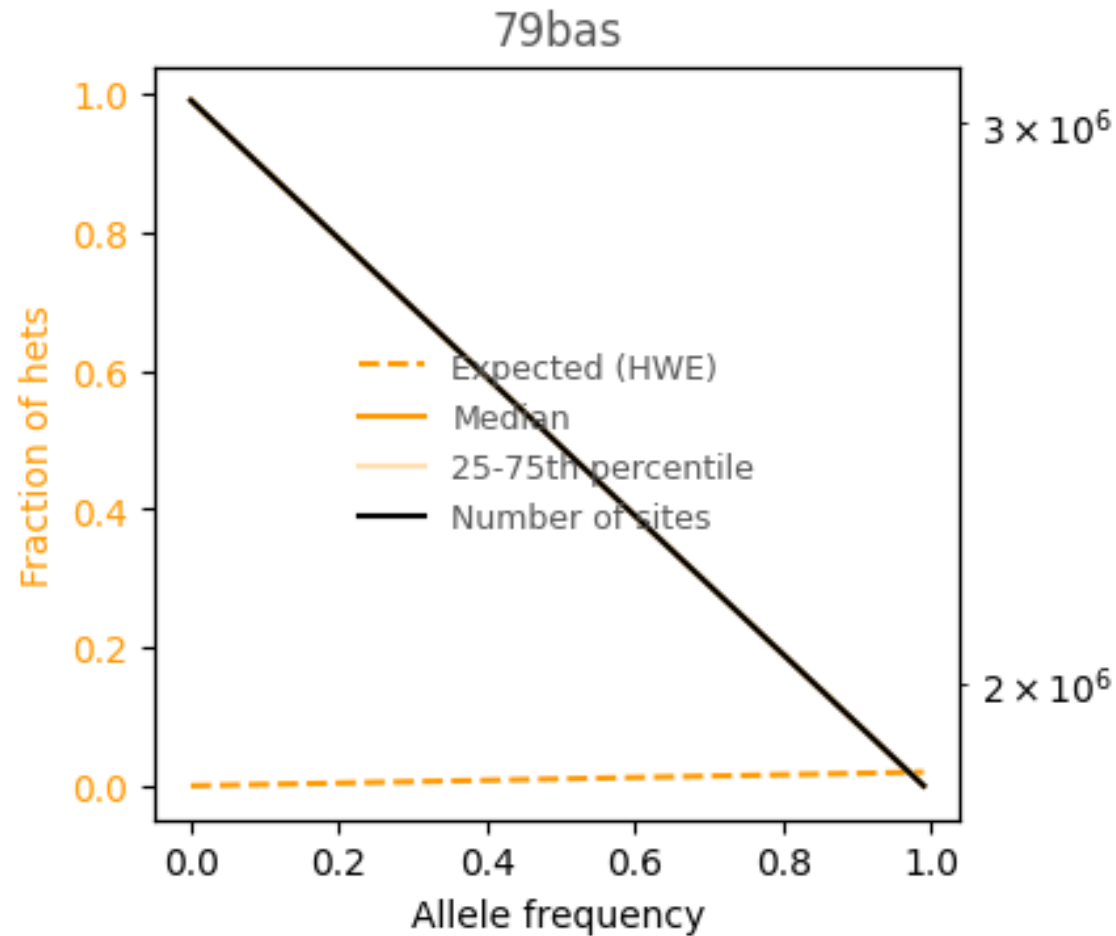
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

