

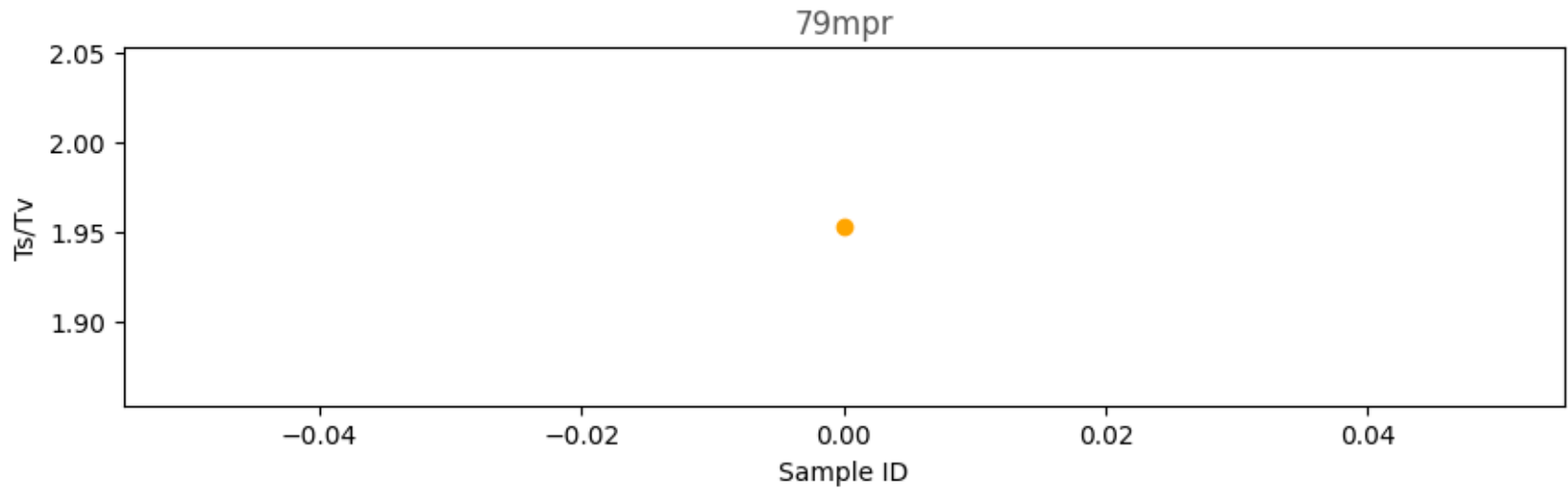
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
79mpr	4,979,553	1.95	1.96	1,142,757	–	0	0
* frameshift ratio: out/(out+in)							

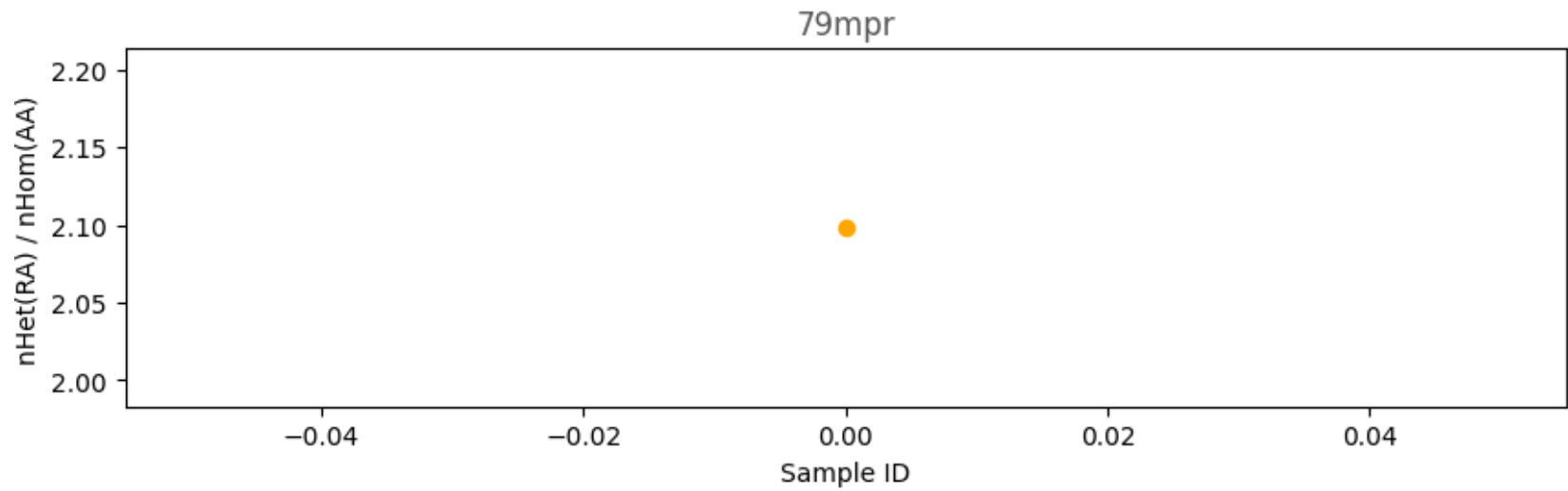
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
79mpr	67.7%	1.93	74.0%	149,956	2,985

- 79mpr .. /ngc/projects2/gm/data/archive/2022/variants/snv/79mpresif-110297641003-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02018-221021_A01411_AH3MJYDSX5-EXT_LAB_KA_NGCWGS-NGCWGS05891_17RKG013139_snv_germline_raw.haplotype_caller.vcf.gz

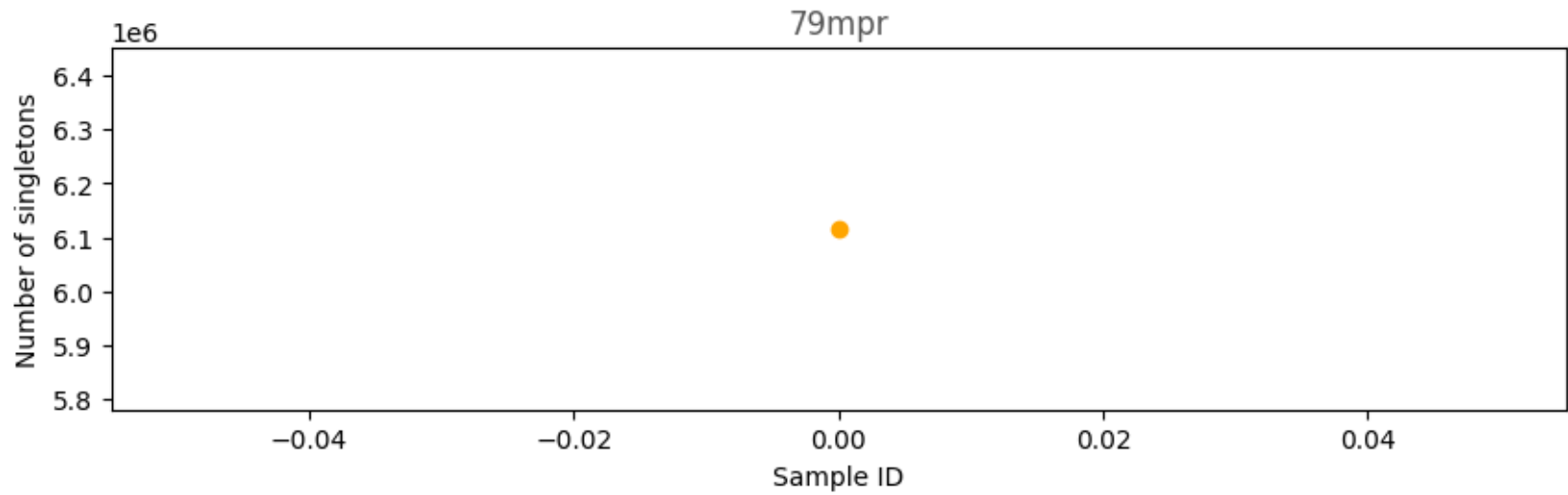
Ts/Tv by sample



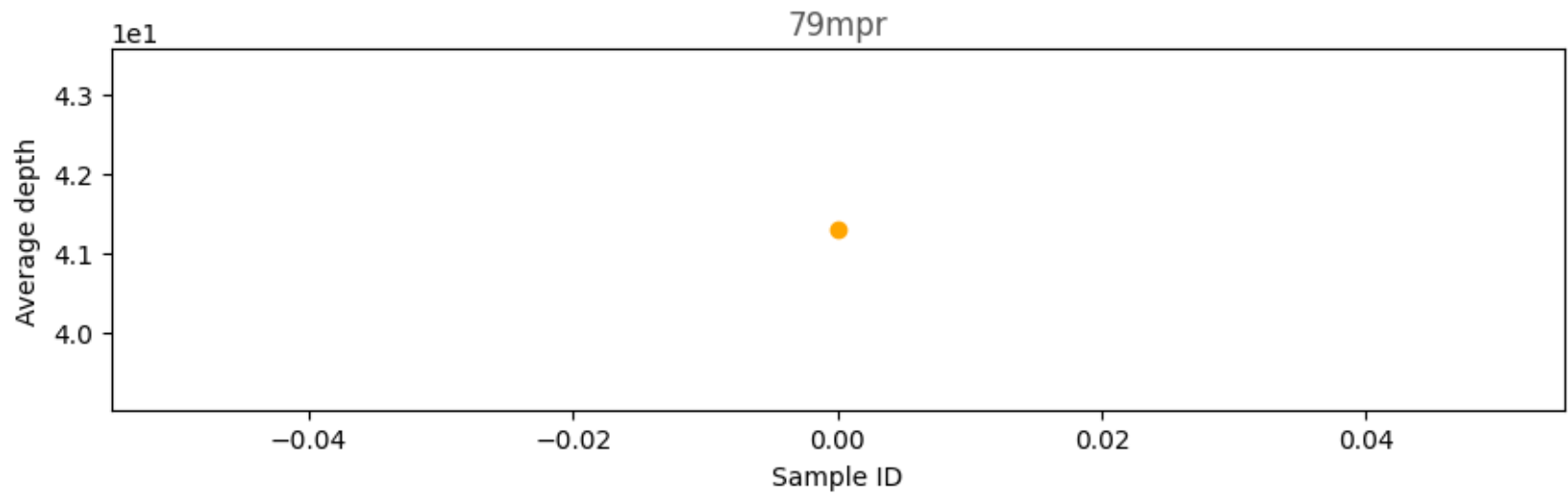
Hets vs non-ref Homs by sample



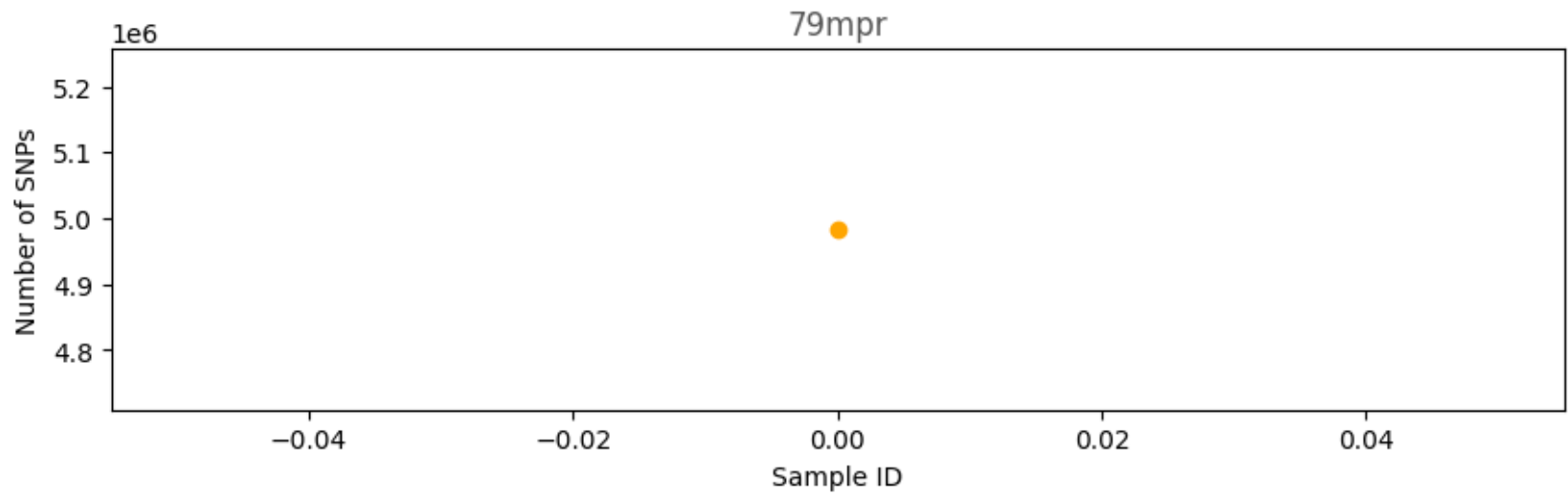
Singletons by sample (hets and homs)



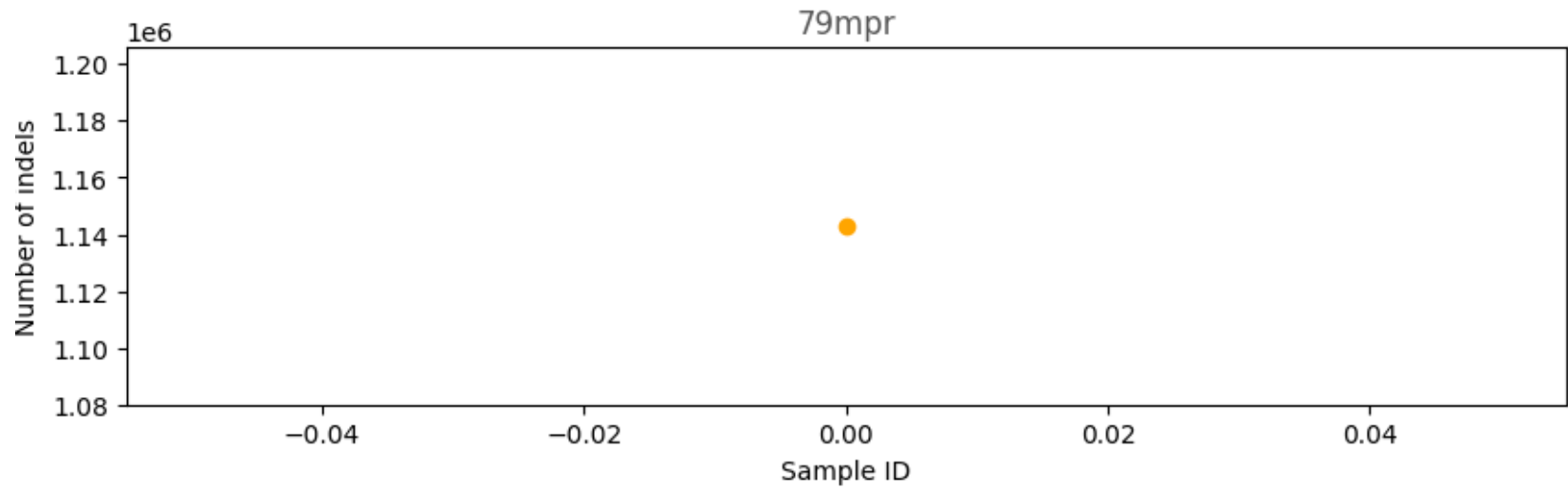
Average depth by sample



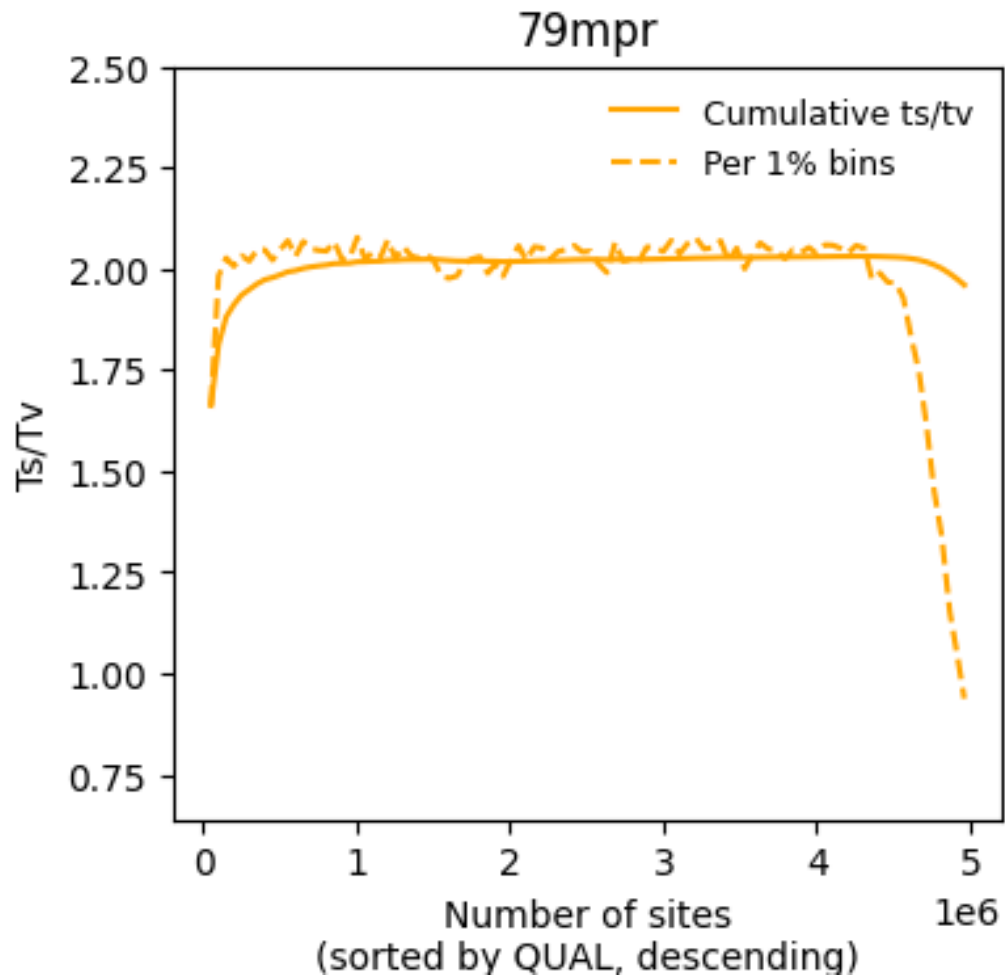
Number of SNPs by sample



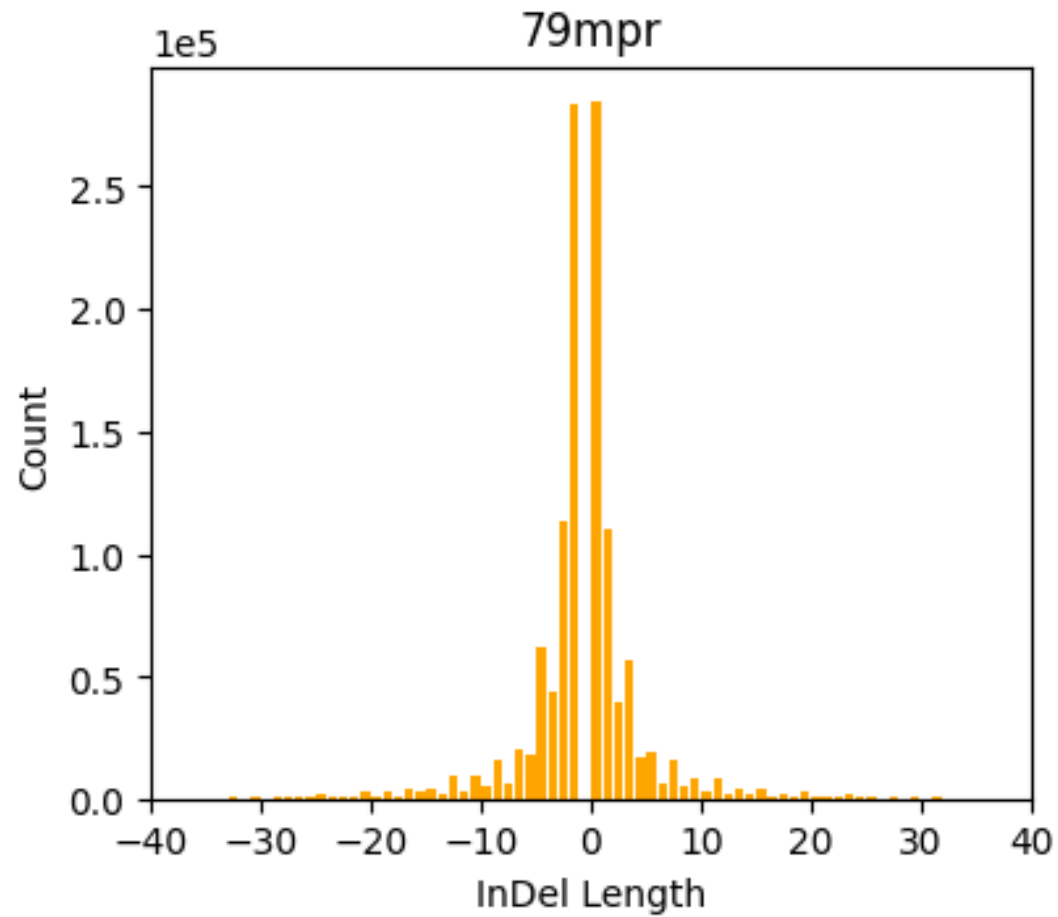
Number of indels by sample



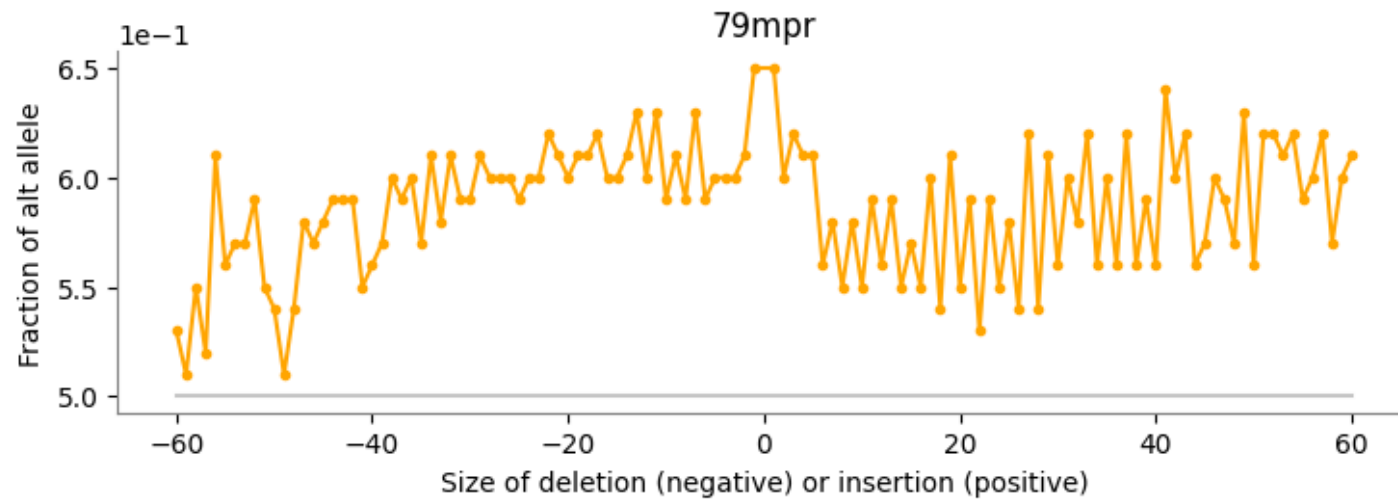
Ts/Tv stratified by QUAL



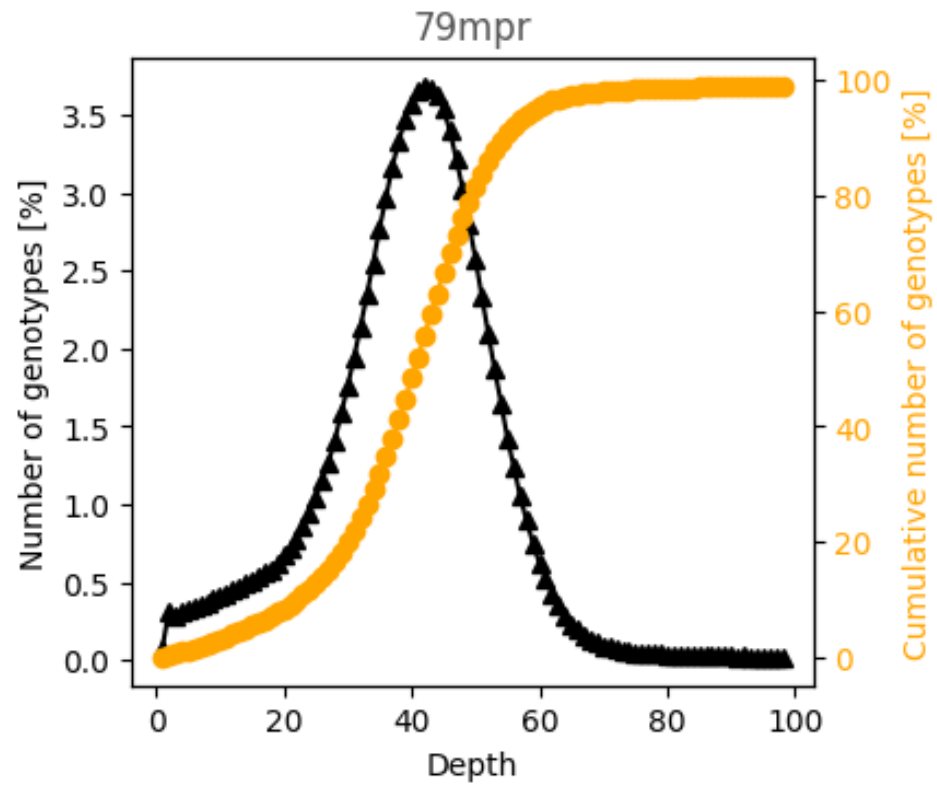
Indel distribution



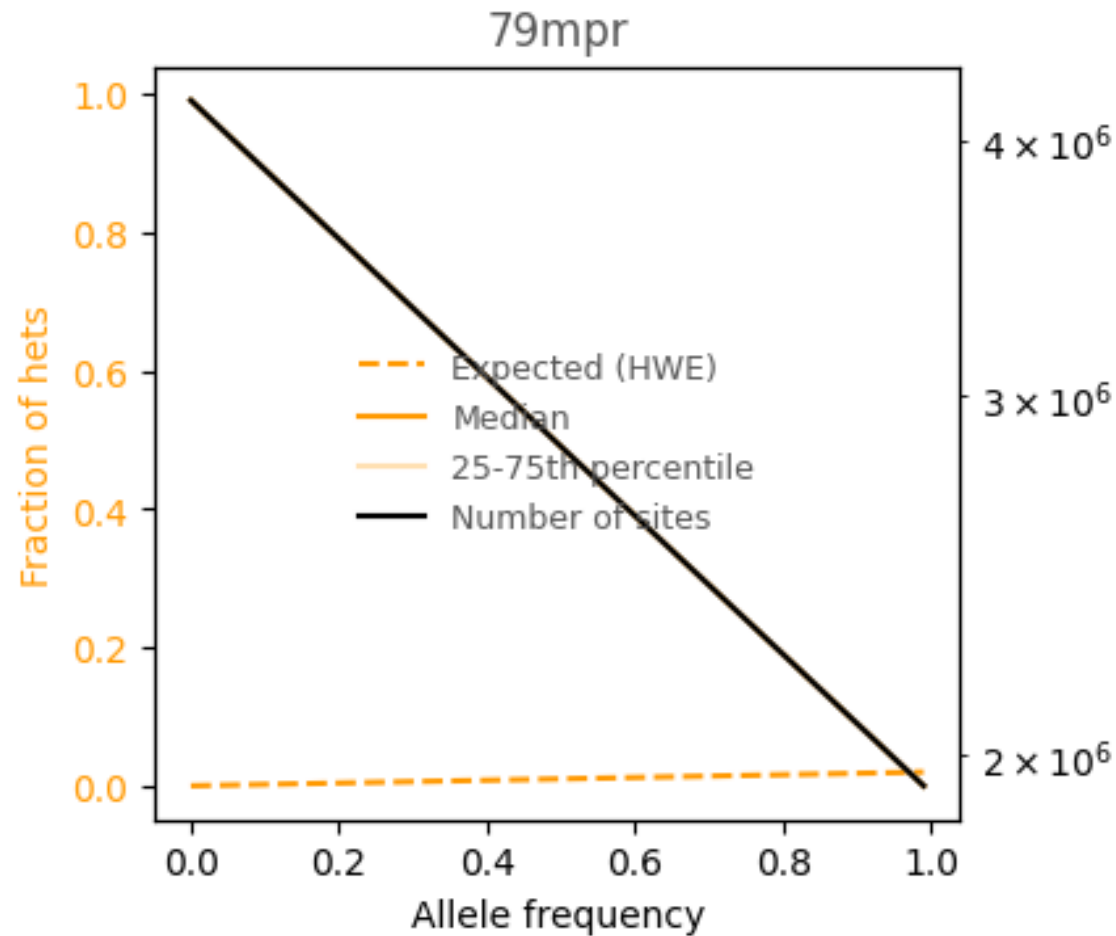
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

