

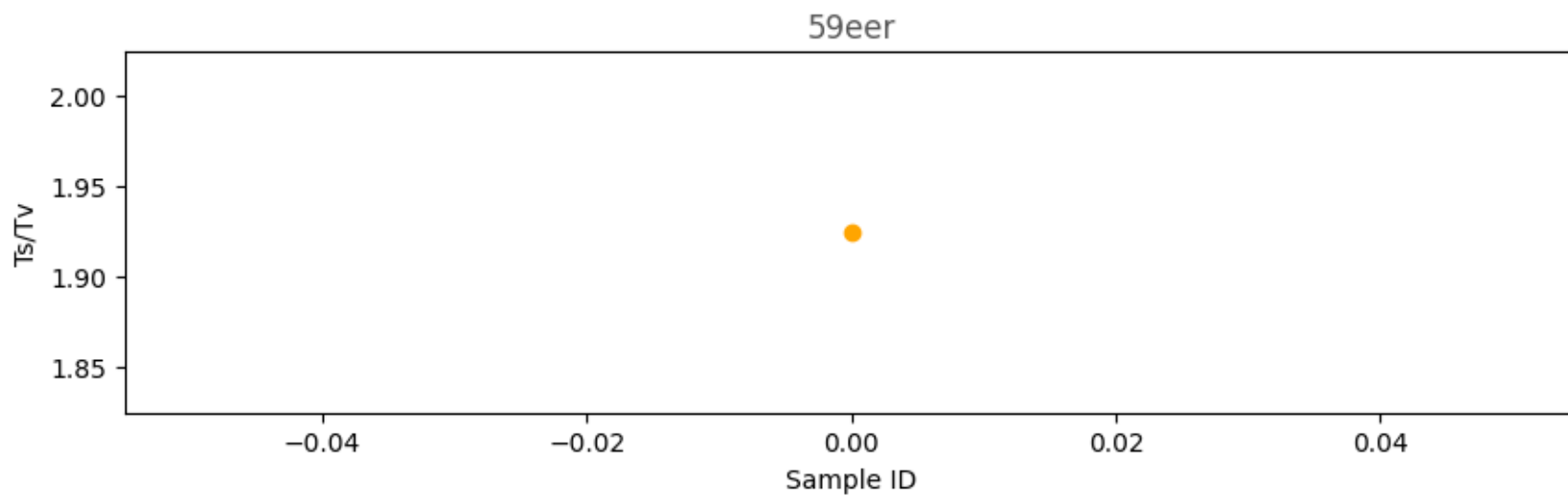
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
59eer	4,081,698	1.92	1.93	947,881	–	0	0
* frameshift ratio: out/(out+in)							

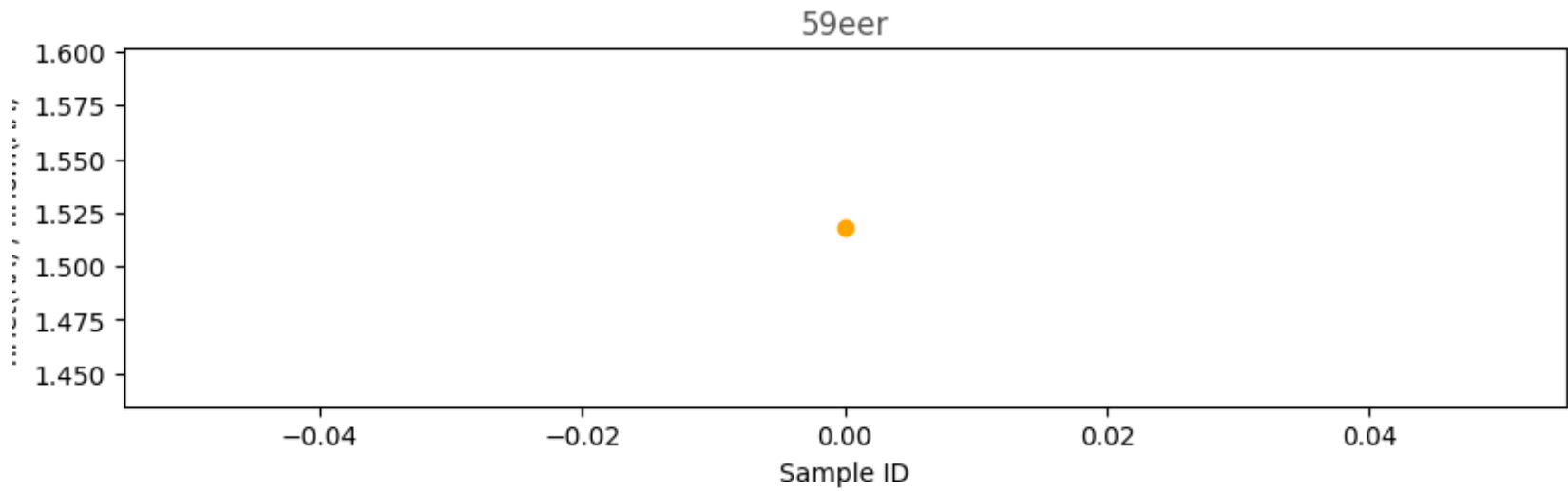
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
59eer	60.3%	1.89	65.1%	96,549	2,085

- 59eer .. /ngc/projects2/gm/data/archive/2022/variants/snv/59eerstof-110296767640-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02106-221026_A00559_AH3MCCDSX5-EXT_LAB
 KA_NGCWGS-NGCWGS05967_snv_germline_raw.haplotype_caller.vcf.gz

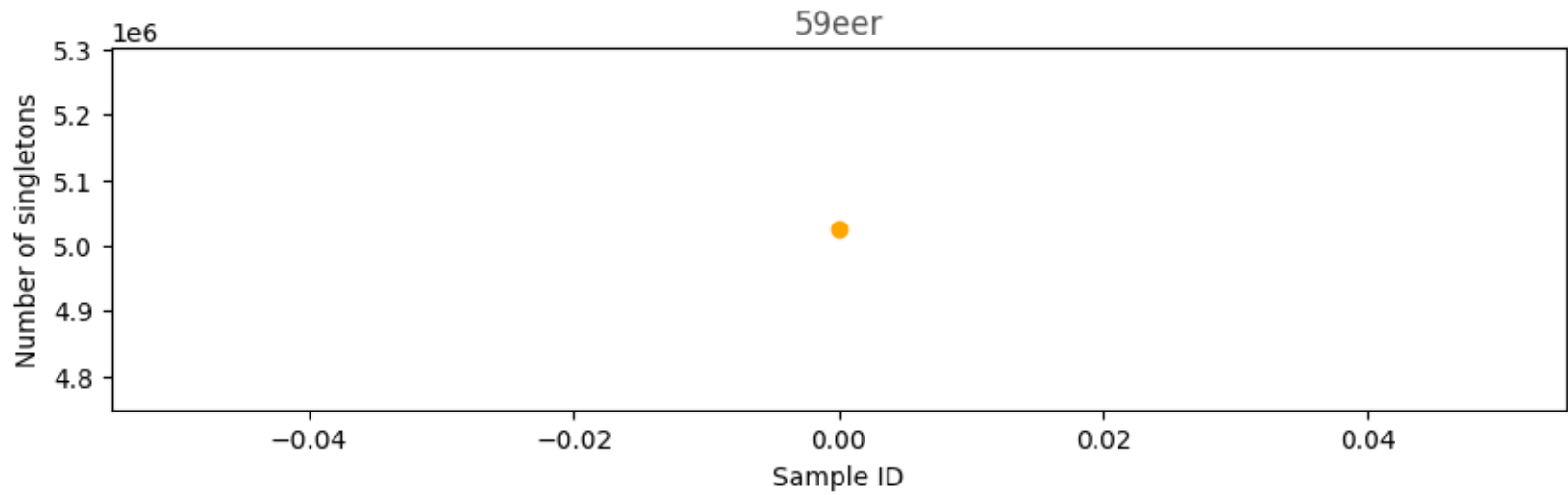
Ts/Tv by sample



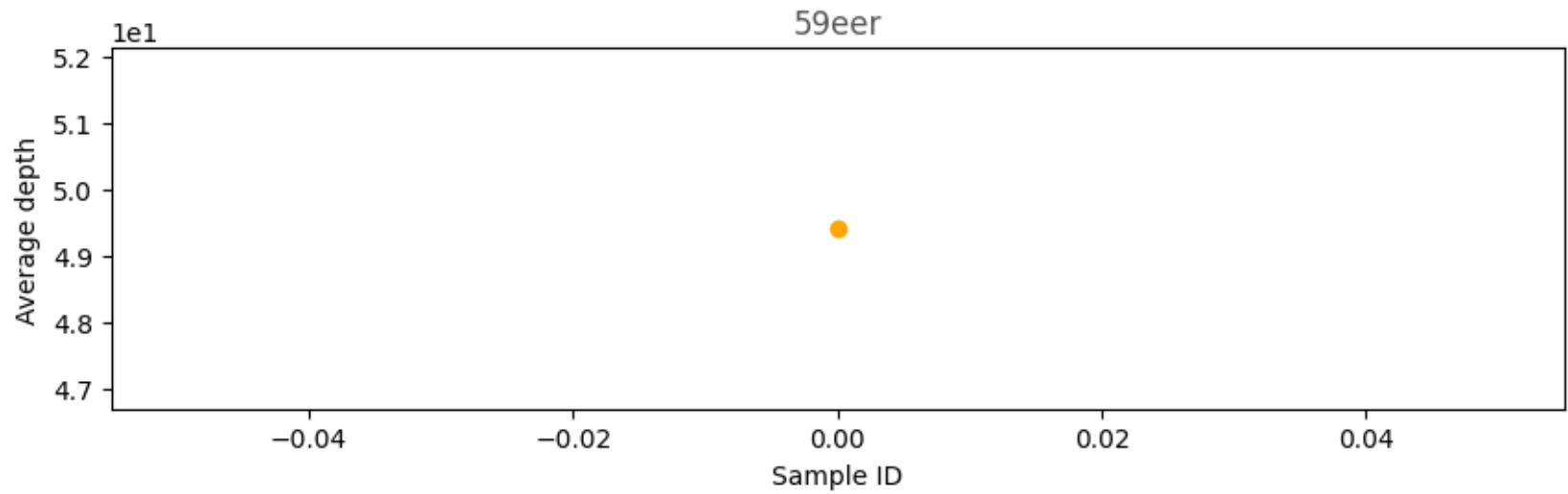
Hets vs non-ref Homs by sample



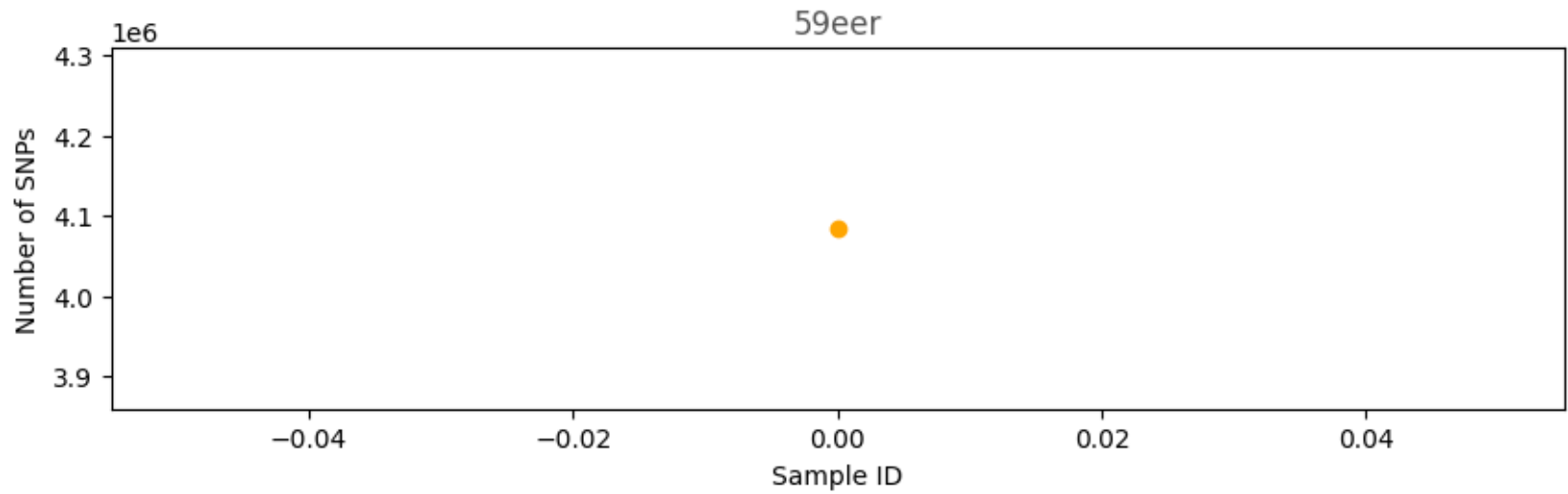
Singletons by sample (hets and homs)



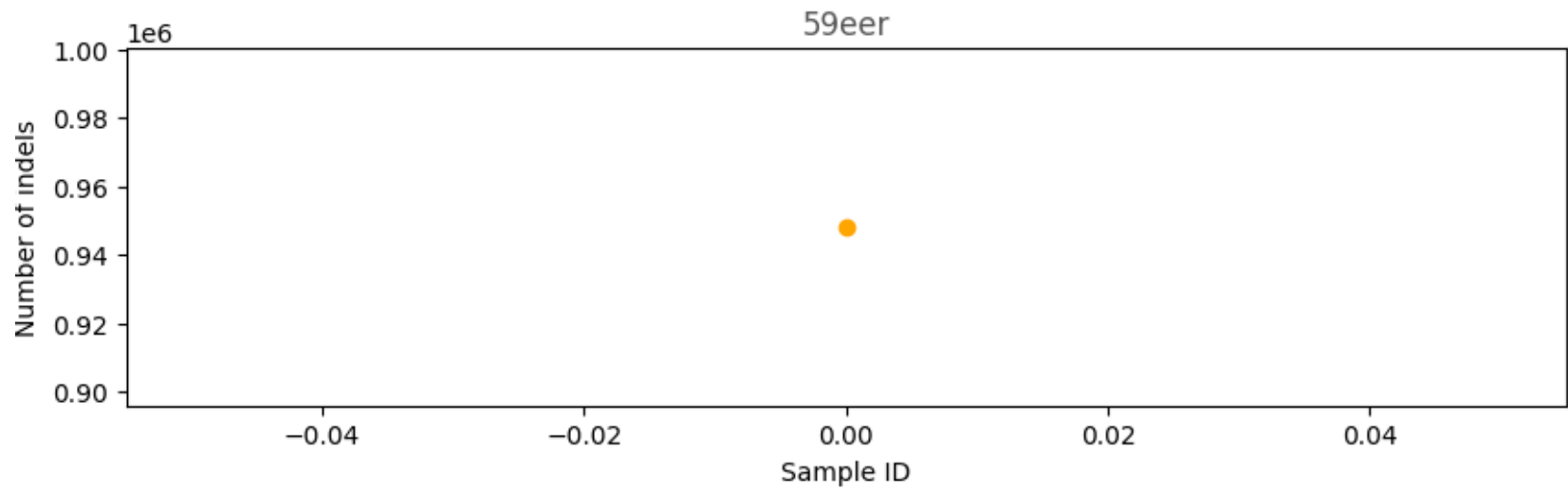
Average depth by sample



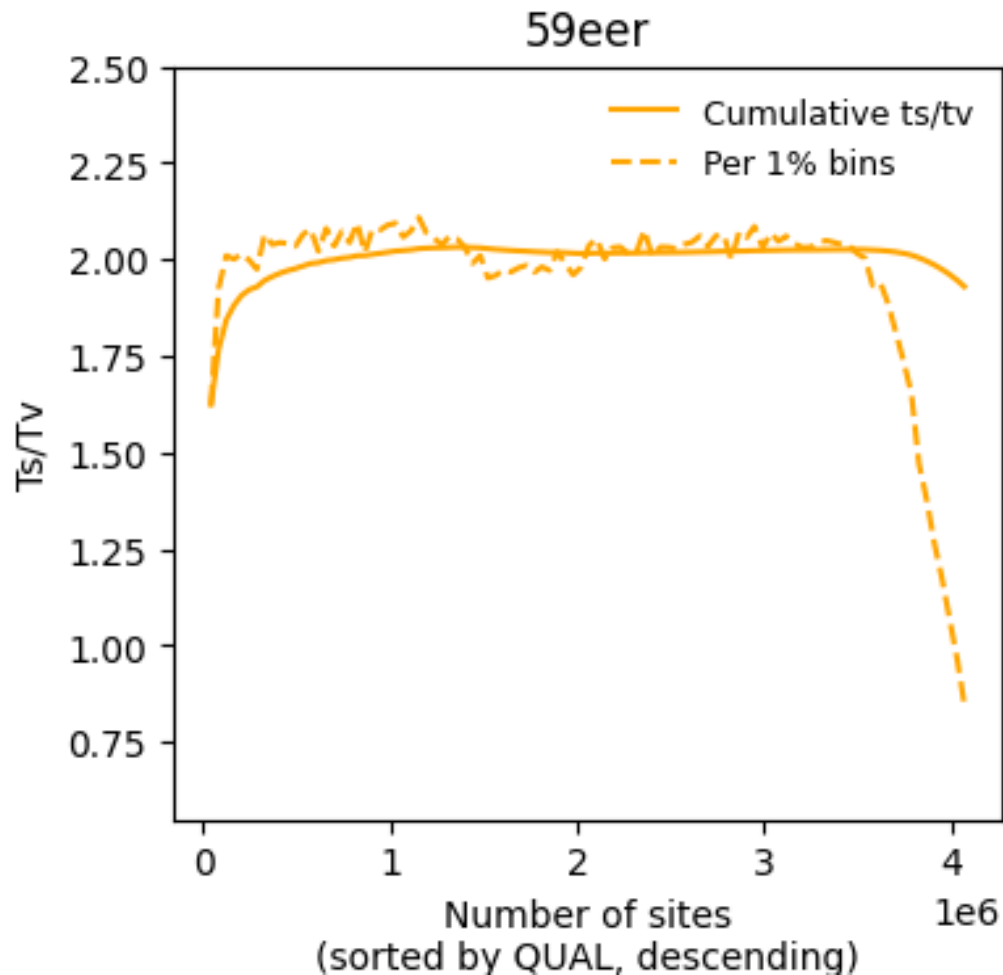
Number of SNPs by sample



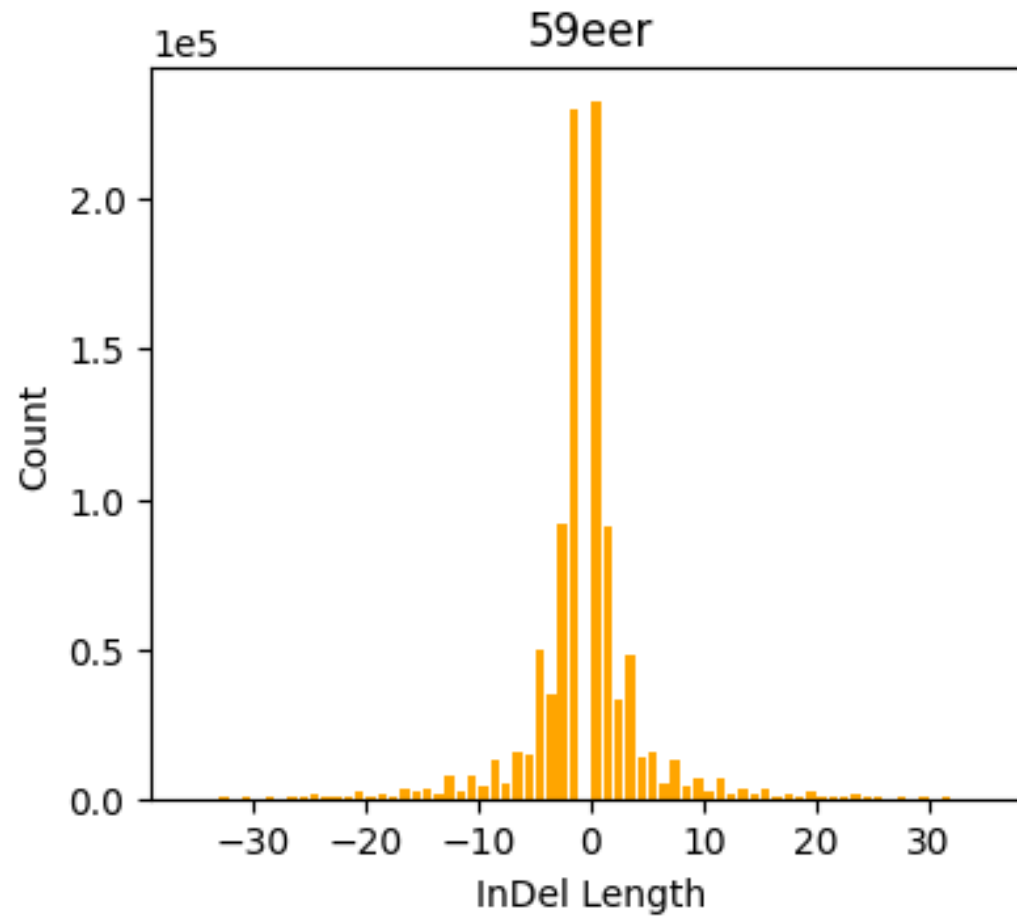
Number of indels by sample



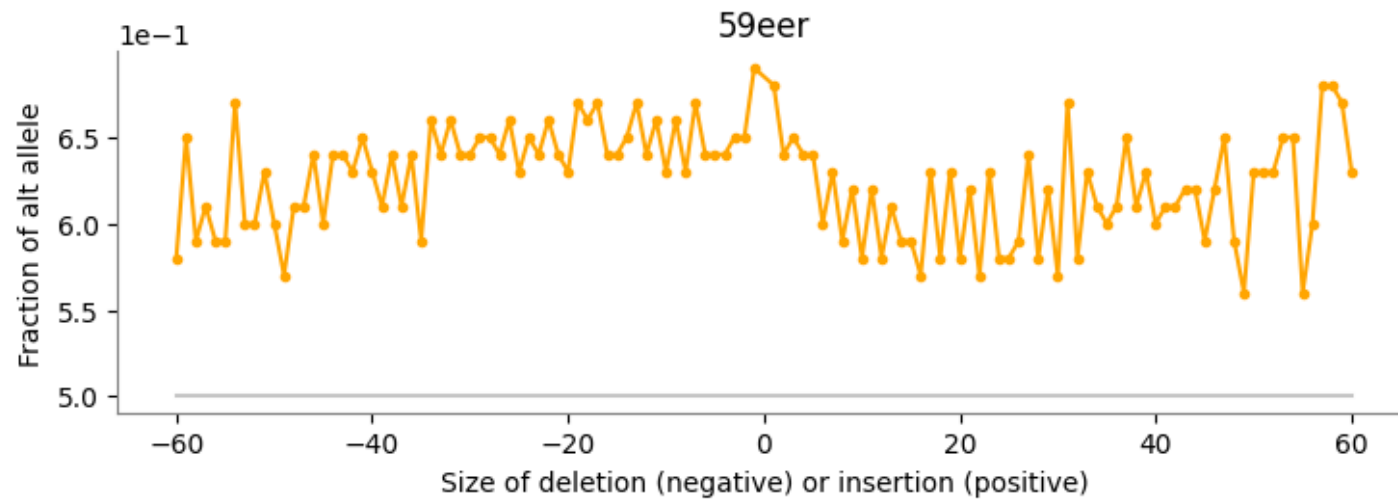
Ts/Tv stratified by QUAL



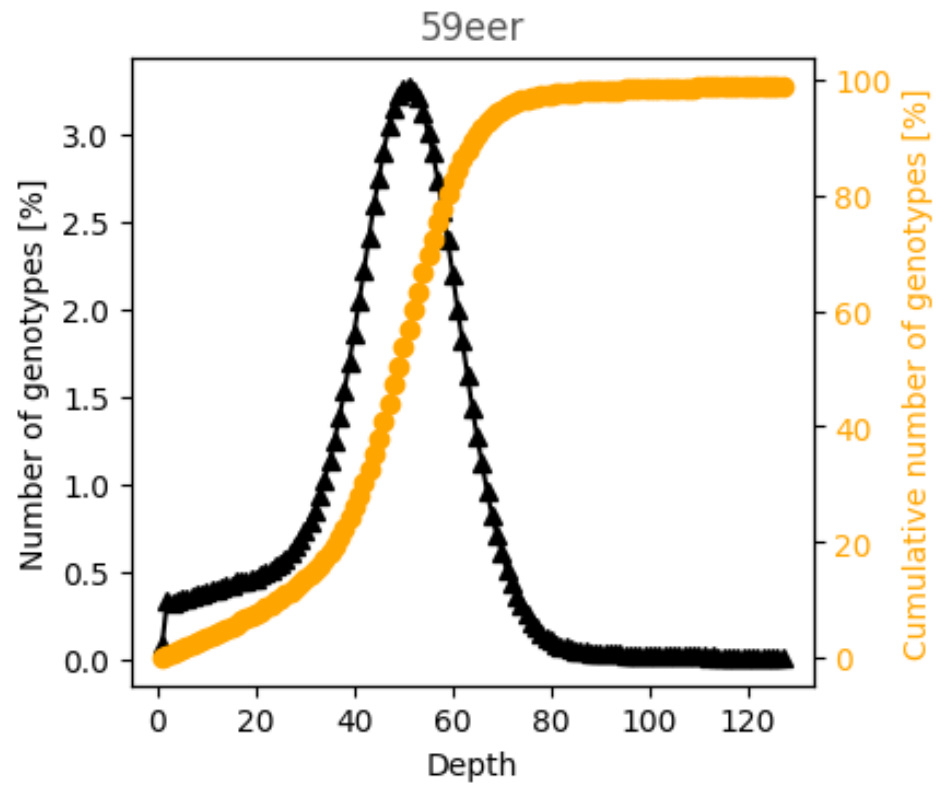
Indel distribution



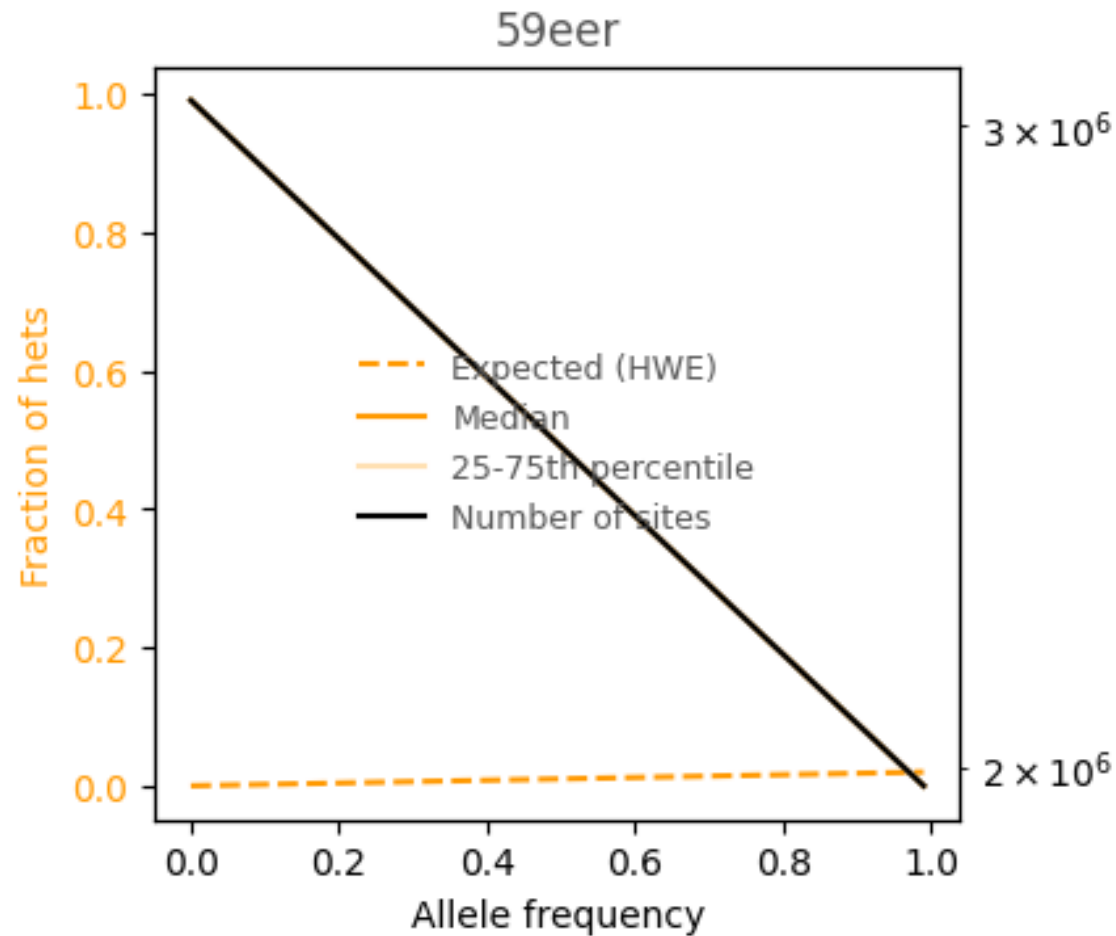
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

