Summary Numbers

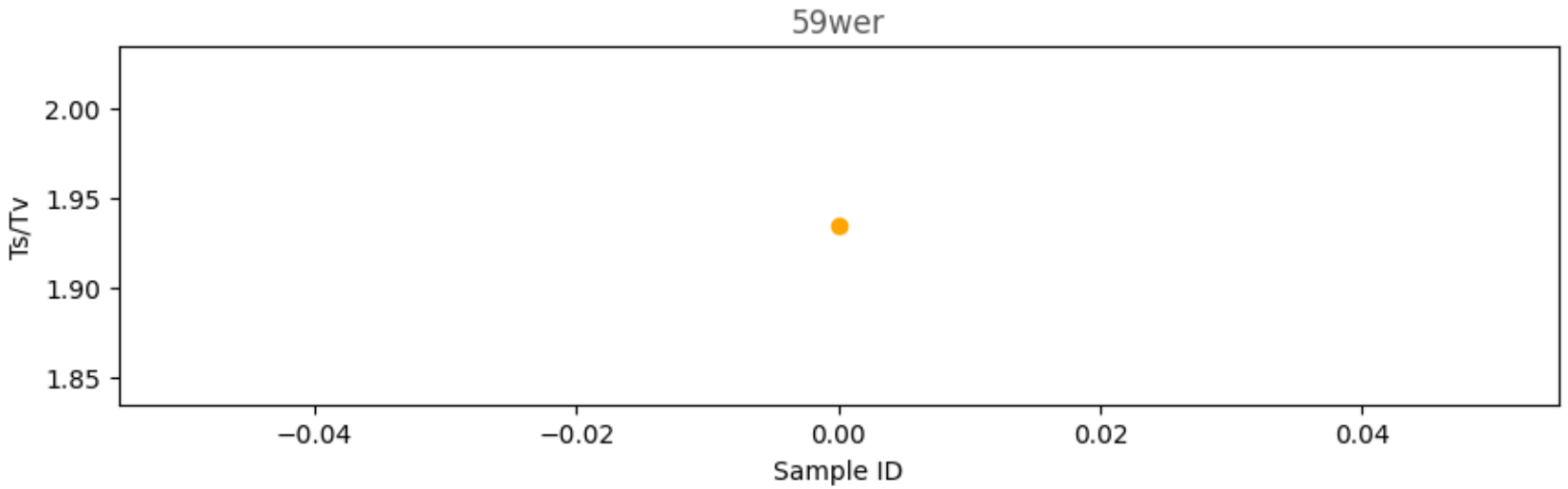
	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
59wer	4,073,330	1.93	1.94	939,829	–	0	0

* frameshift ratio: out/(out+in)

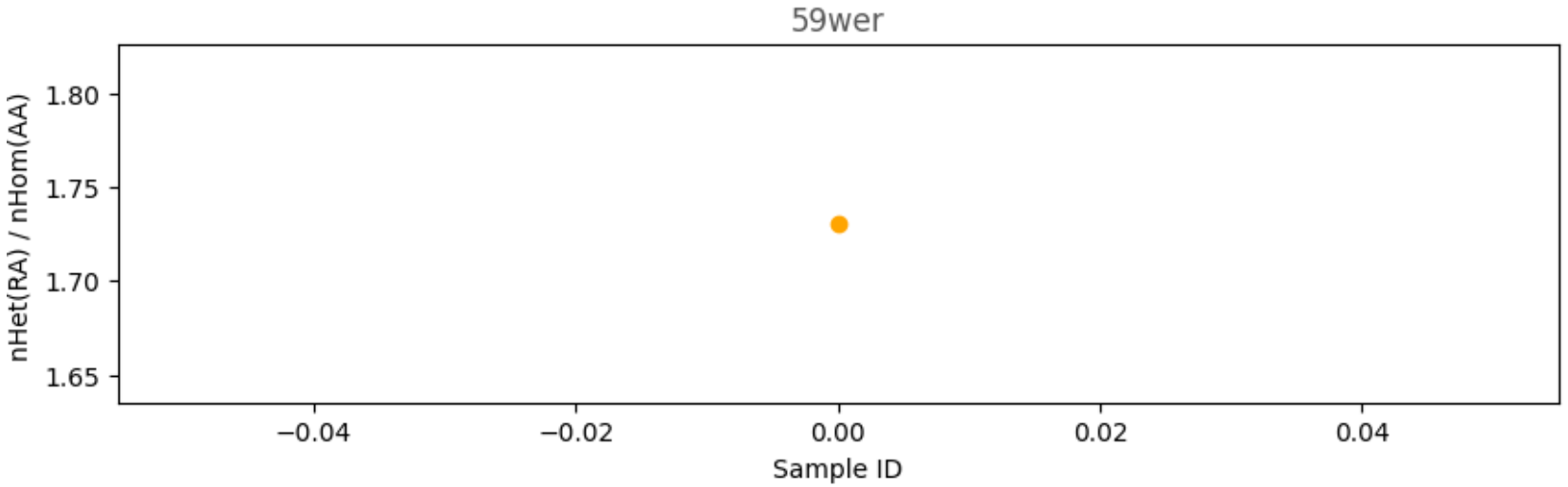
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
59wer	63.4%	1.91	67.8%	93,861	2,047

- 59wer .. `/ngc/projects2/gm/data/archive/2022/variants/snv/59weribaf-103833689964-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220202_A01176_AH52HTDMXY-RHGM_LABKA_WGS-WGS03783_snv_germline_raw.haplotype_caller.vcf.gz`

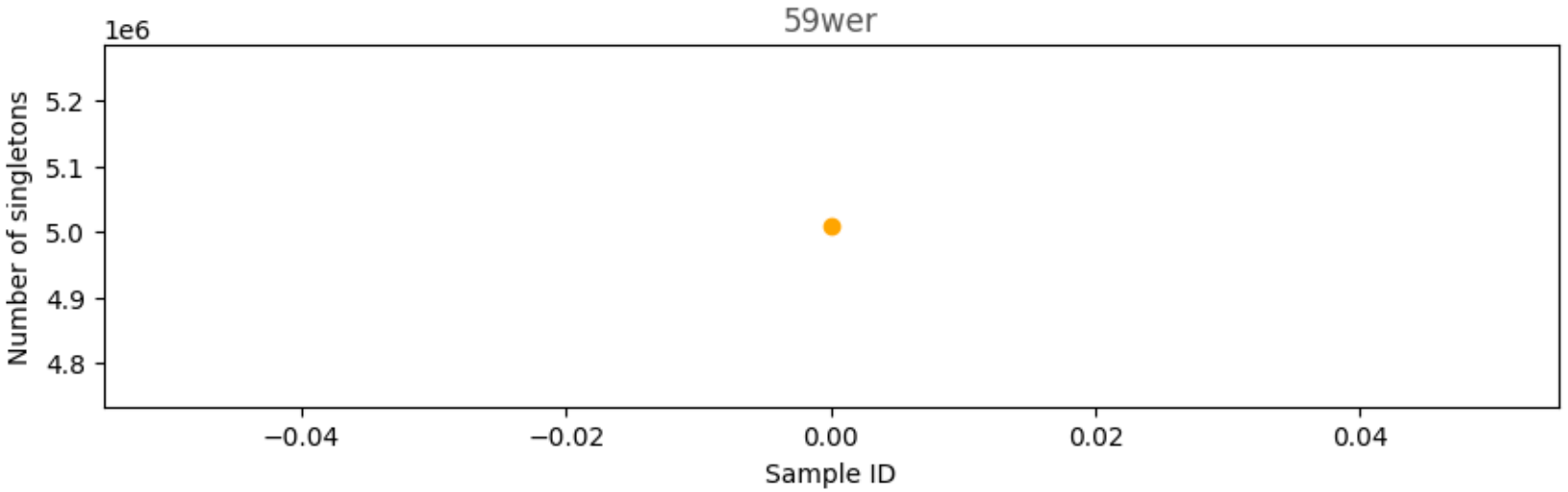
Ts/Tv by sample



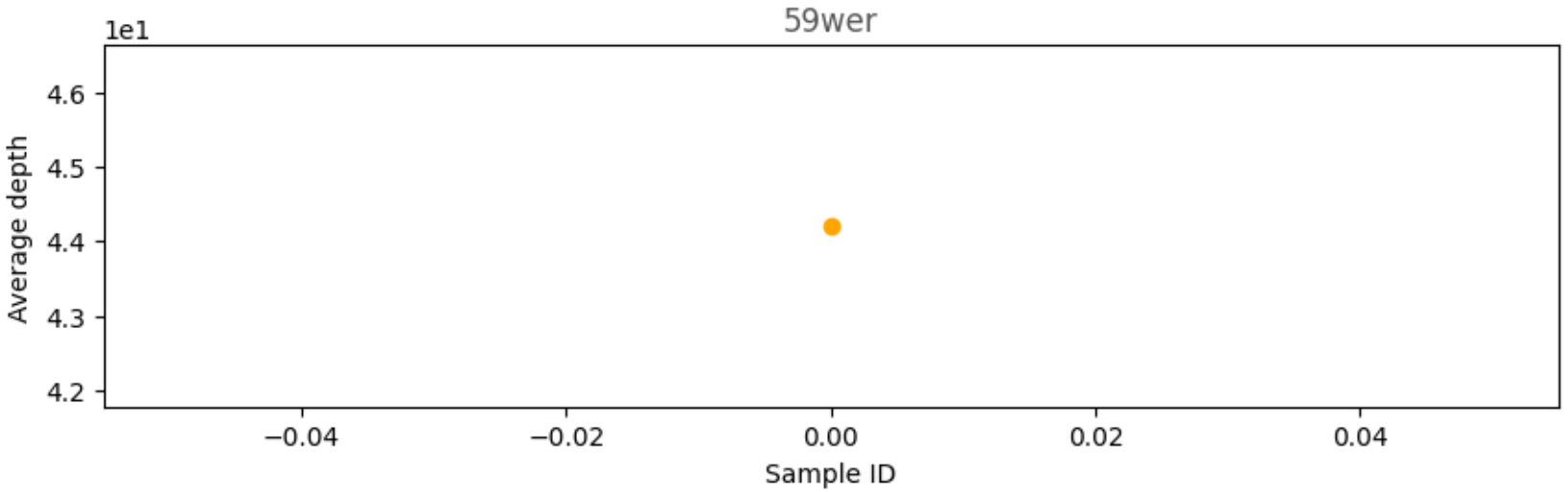
Hets vs non-ref Homs by sample



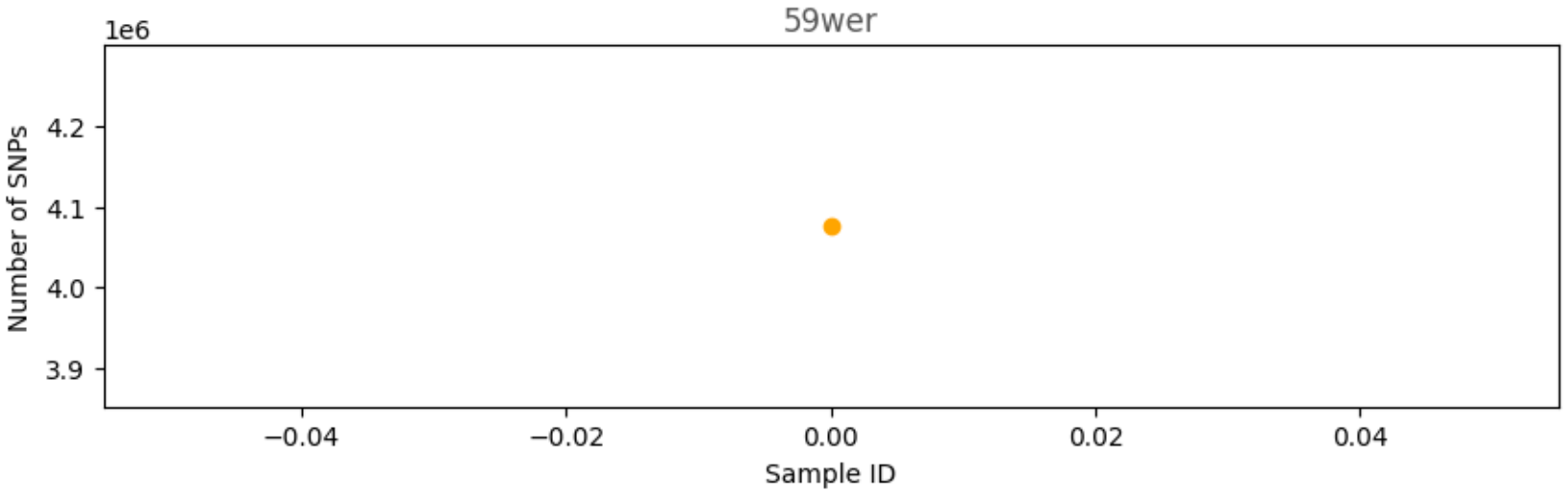
Singletons by sample (hets and homs)



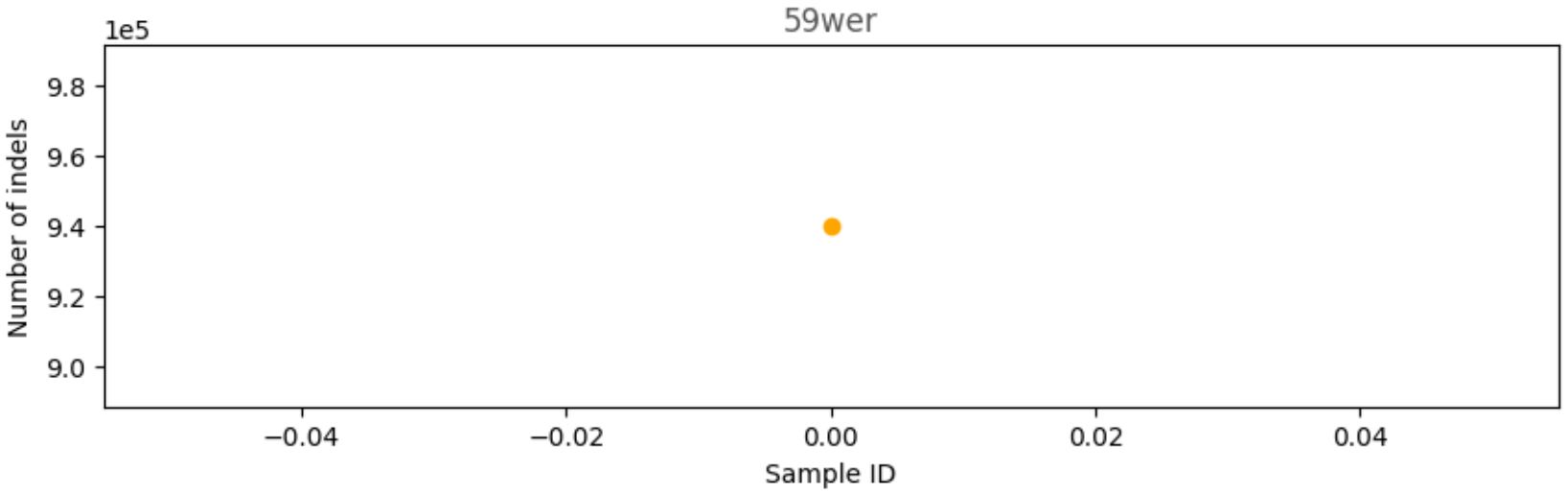
Average depth by sample



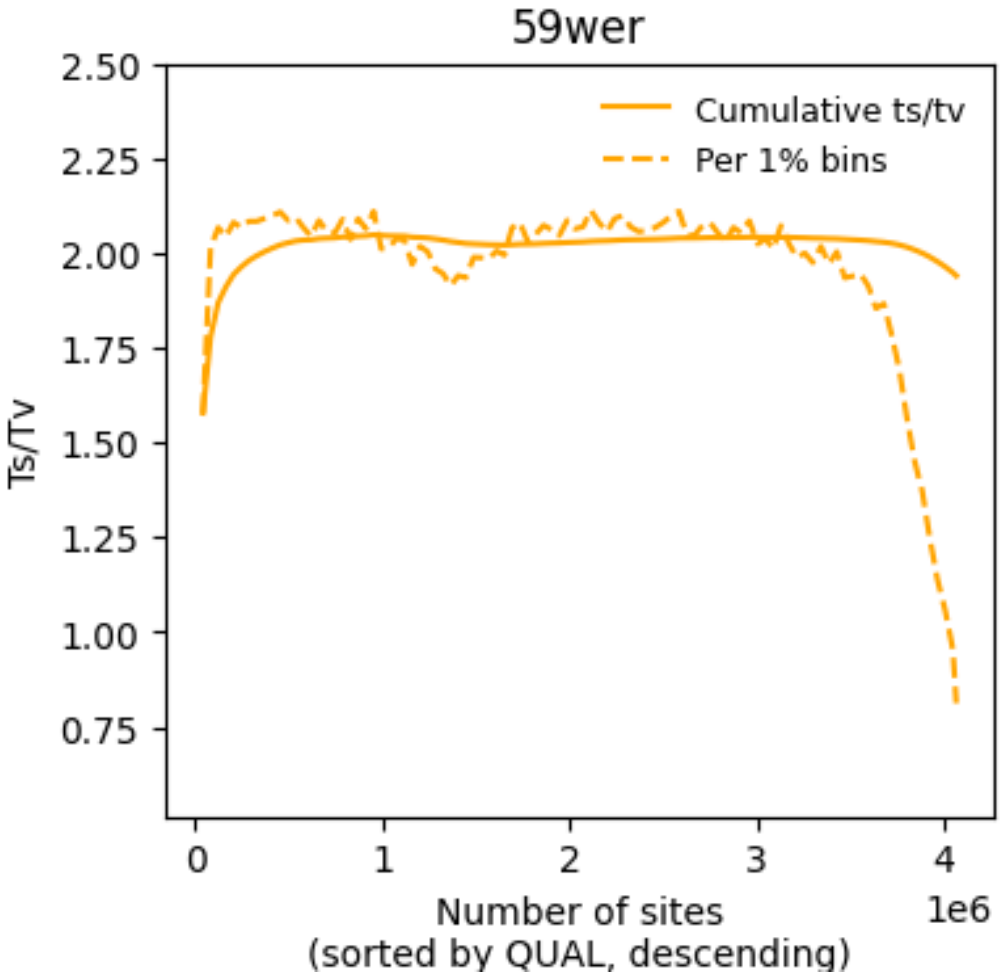
Number of SNPs by sample



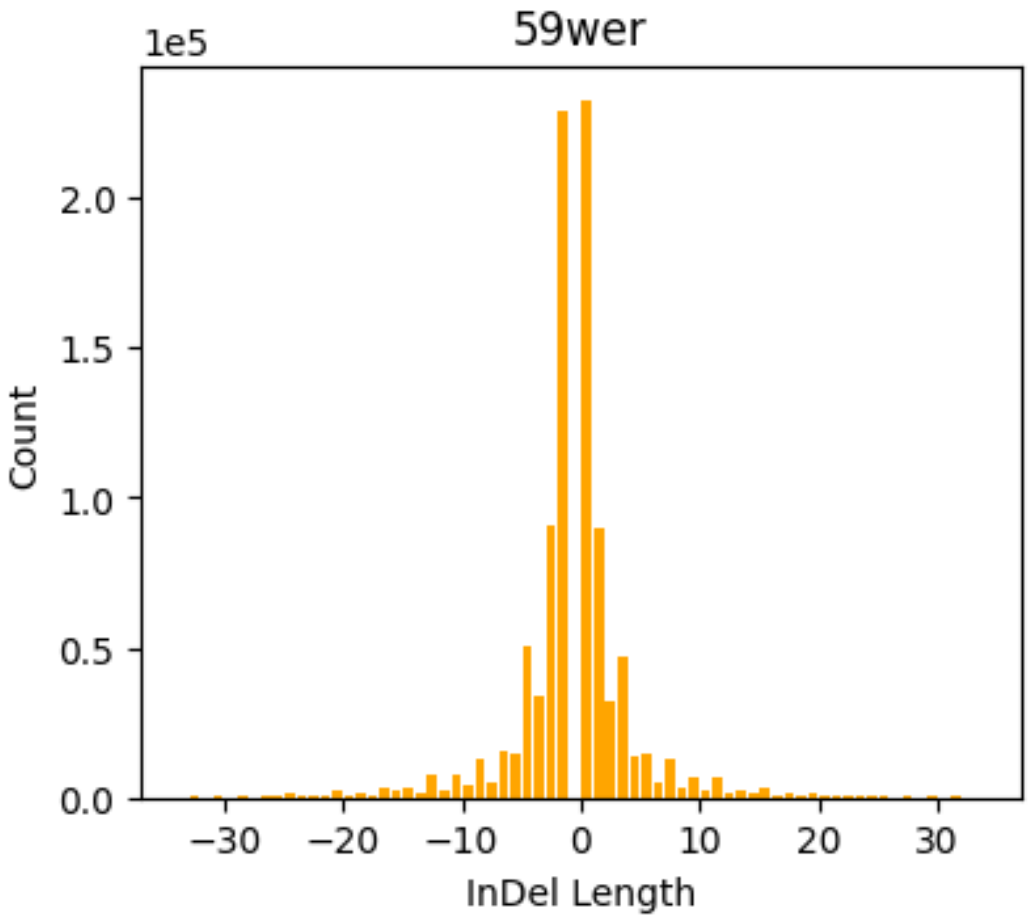
Number of indels by sample



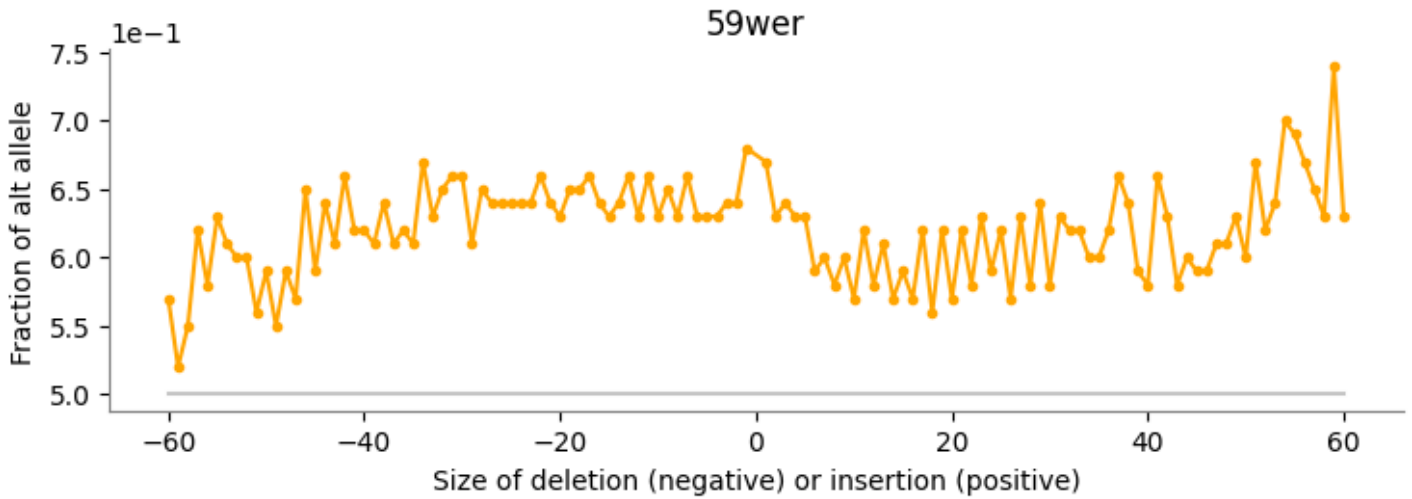
Ts/Tv stratified by QUAL



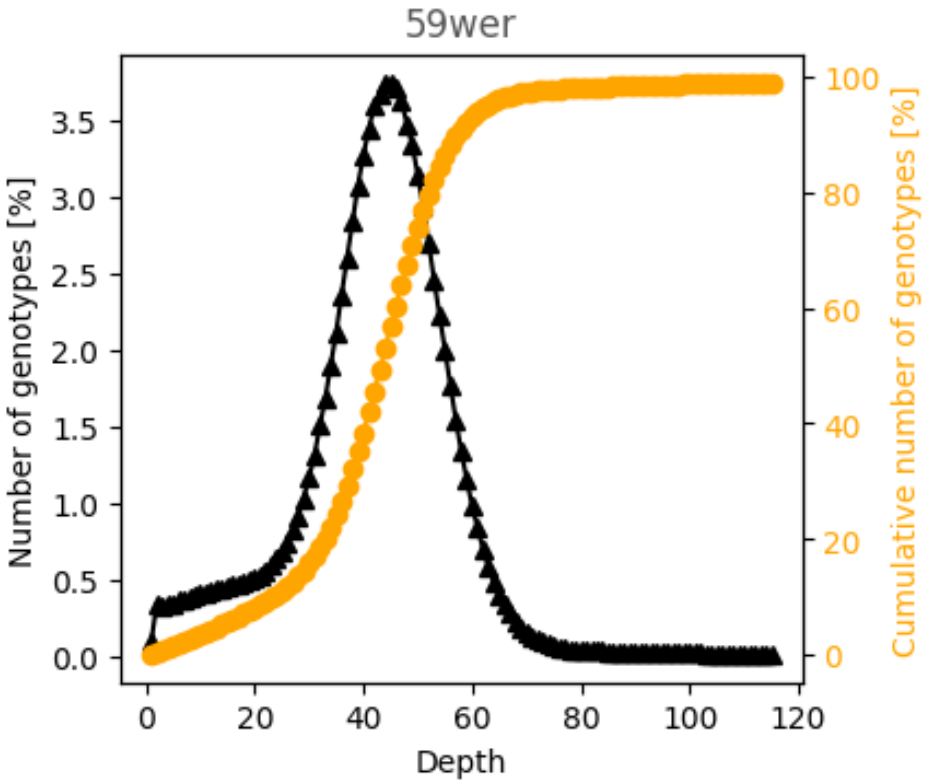
Indel distribution



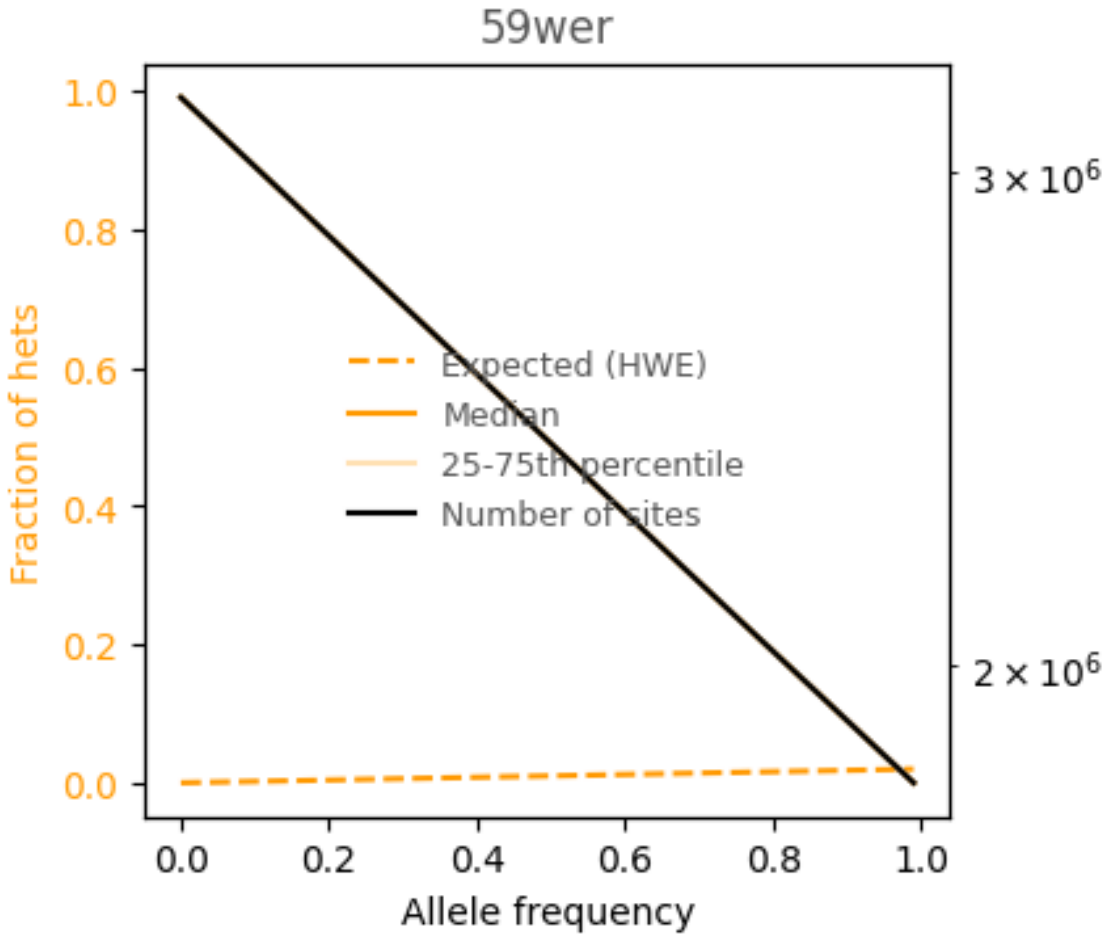
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types