Summary Numbers

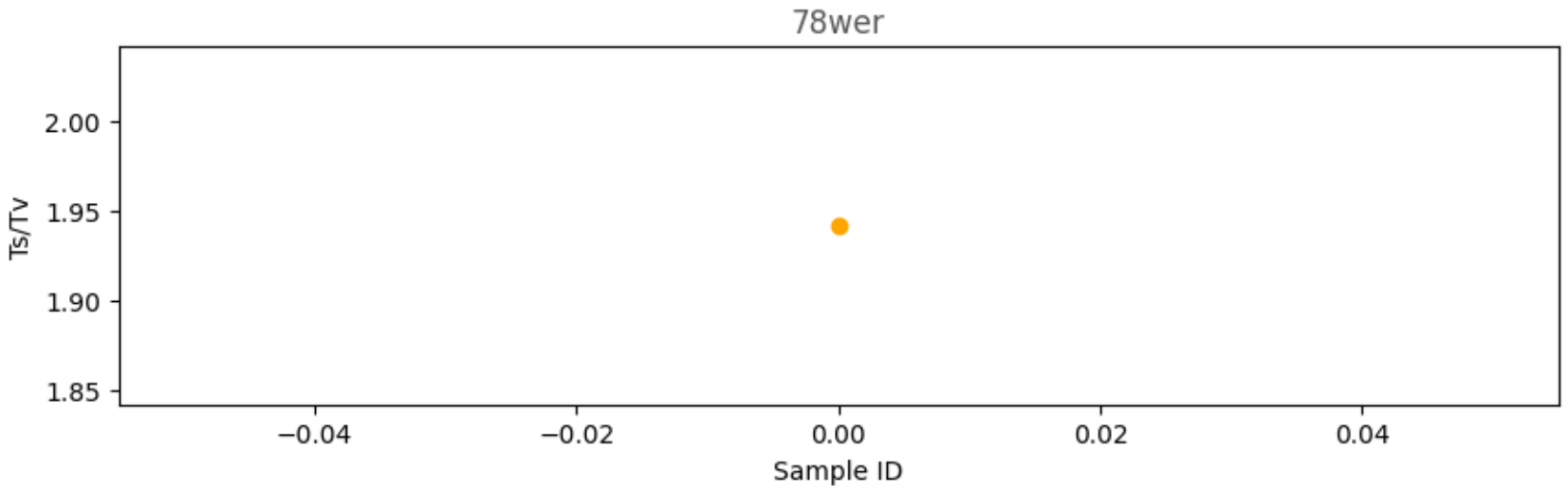
	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
78wer	4,031,806	1.94	1.95	937,989	–	0	0

* frameshift ratio: out/(out+in)

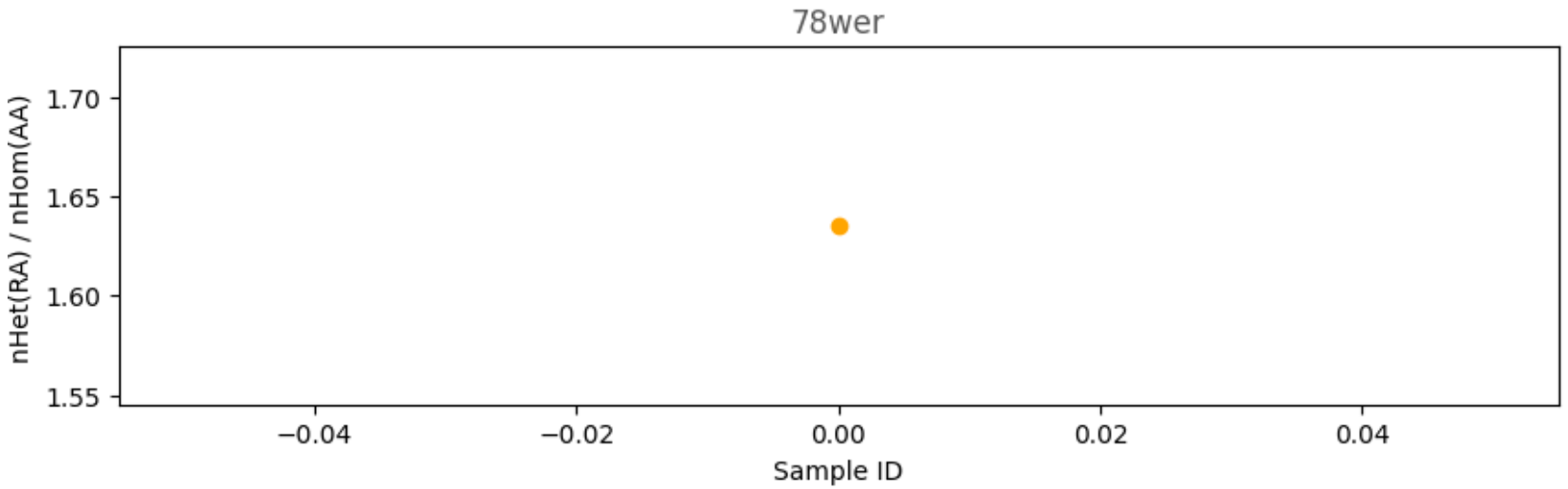
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
78wer	62.1%	1.92	66.4%	94,450	1,901

- 78wer .. `/ngc/projects2/gm/data/archive/2022/variants/snv/78werdtem-103825441290-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00506-220119_A01176_AH2CTLDSX3-EXT_LABKA_NGCWGS-NGCWGS03679_21RKG030103x01_snv_germline_raw.haplotype_caller.vcf.gz`

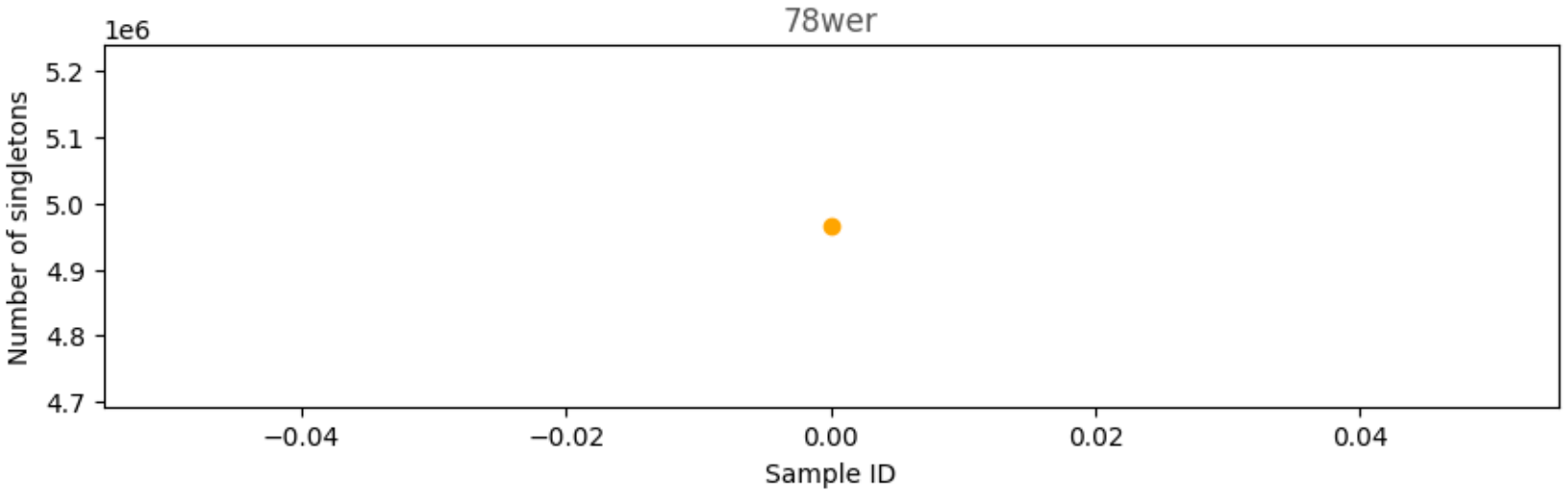
Ts/Tv by sample



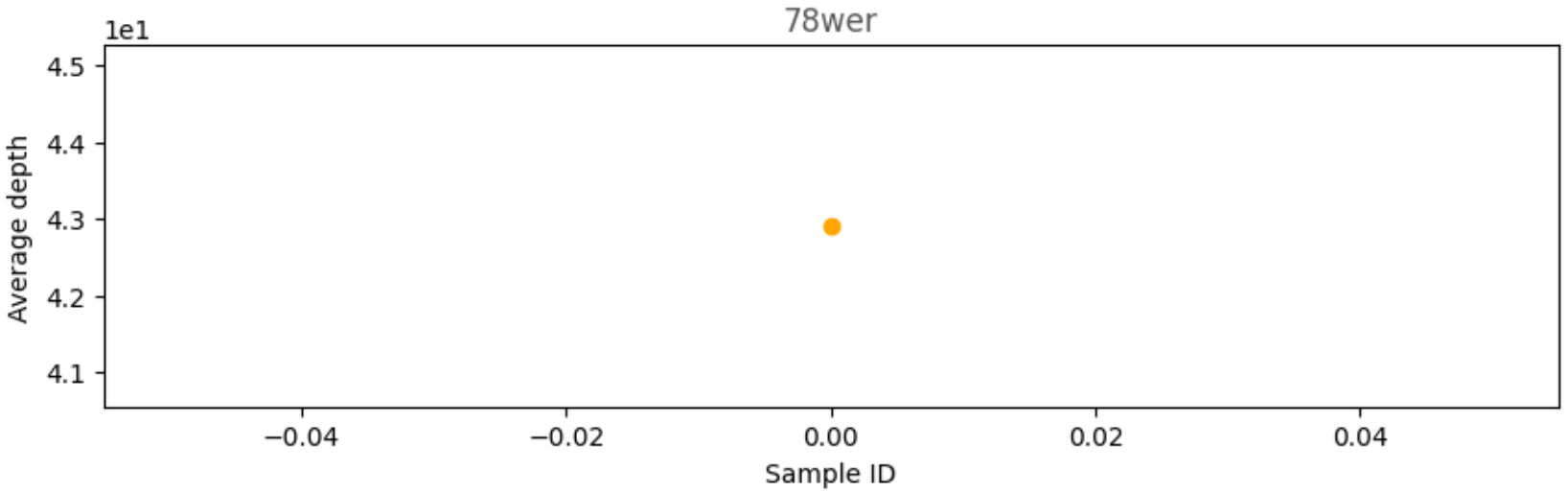
Hets vs non-ref Homs by sample



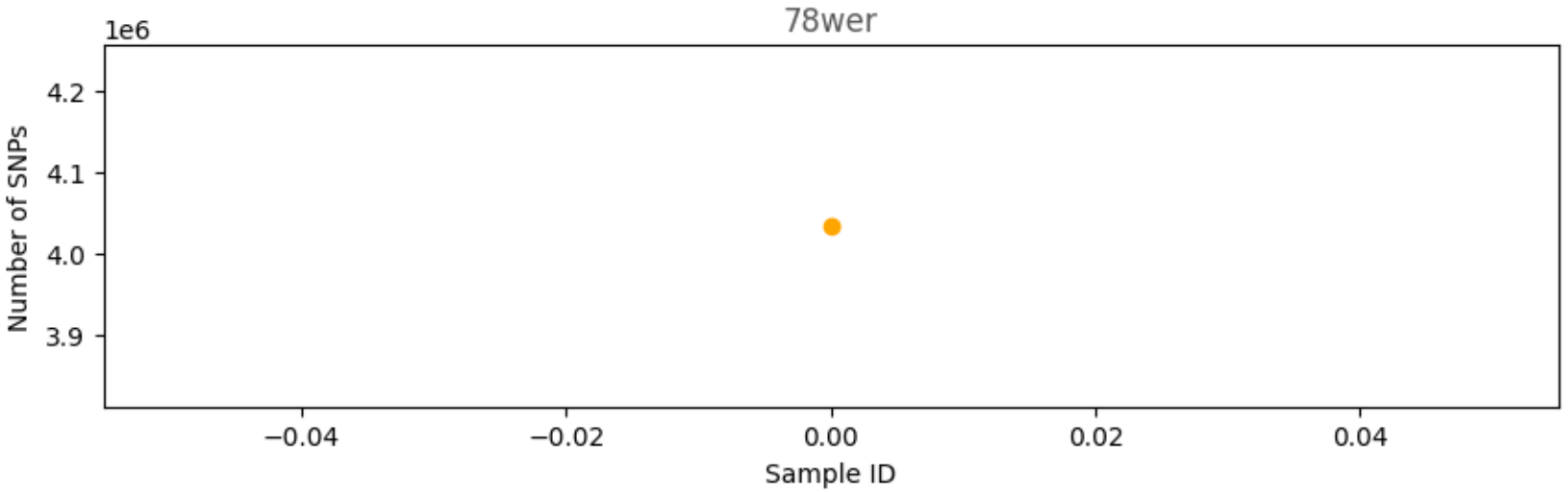
Singletons by sample (hets and homs)



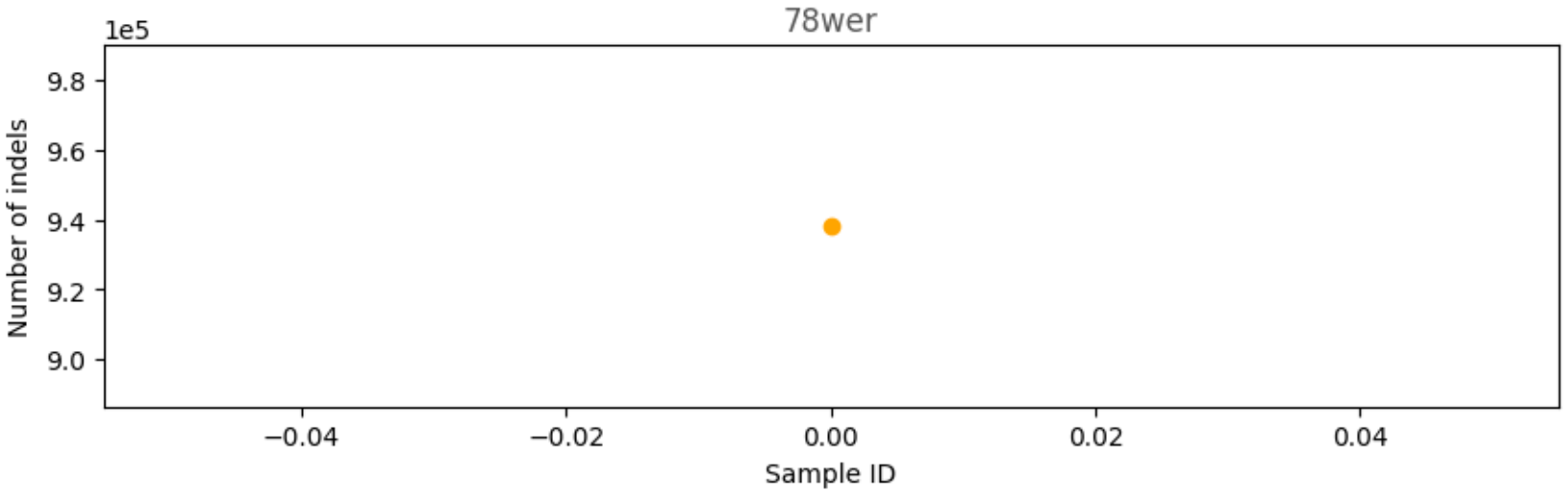
Average depth by sample



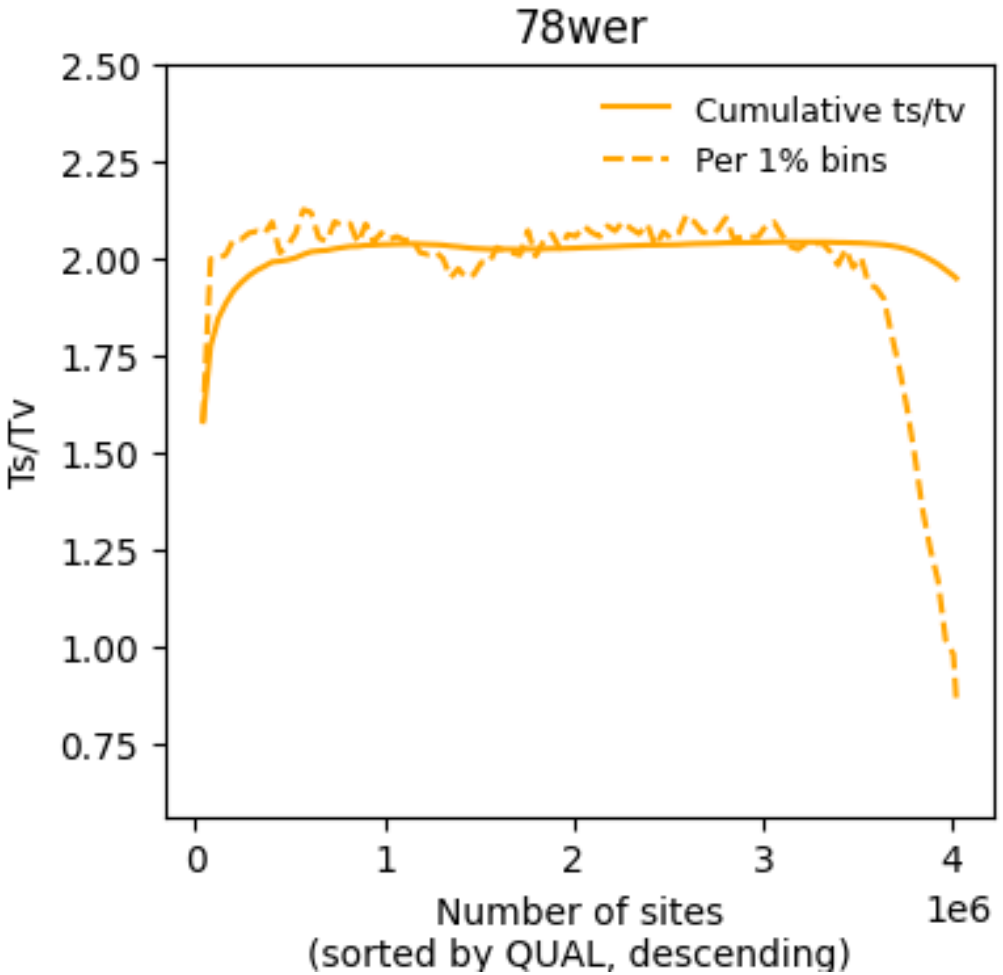
Number of SNPs by sample



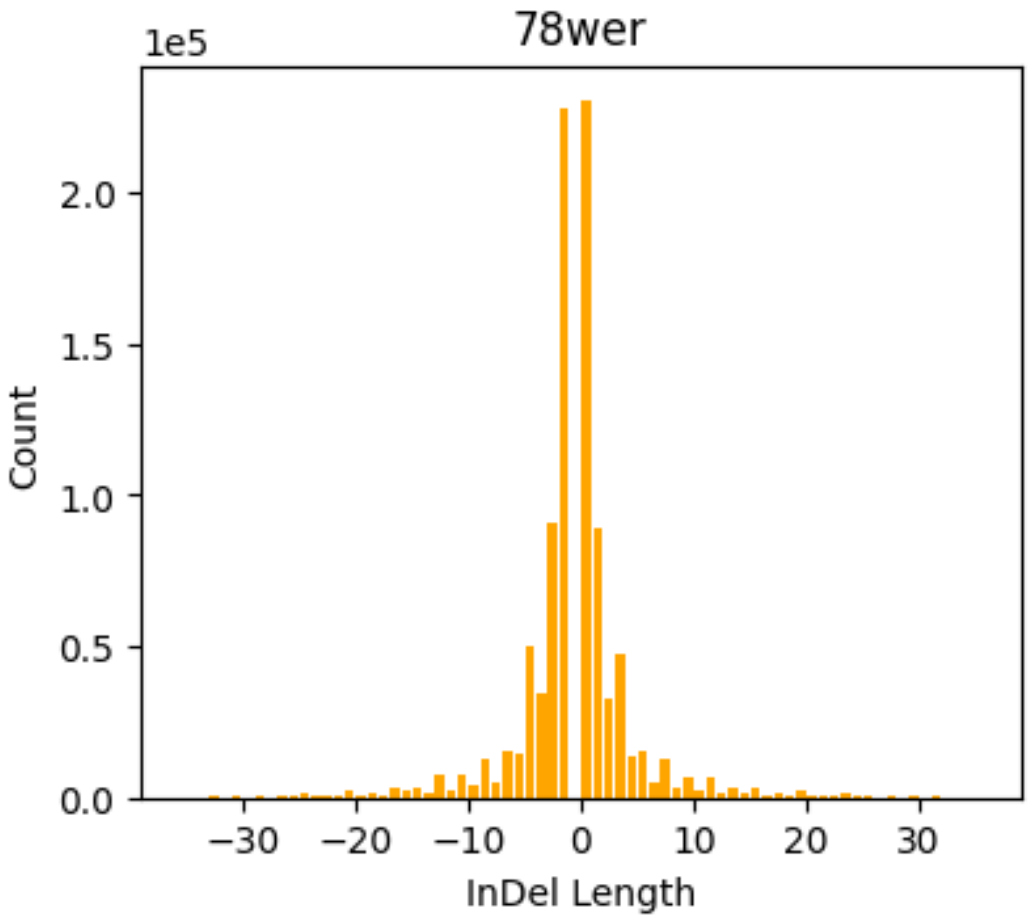
Number of indels by sample



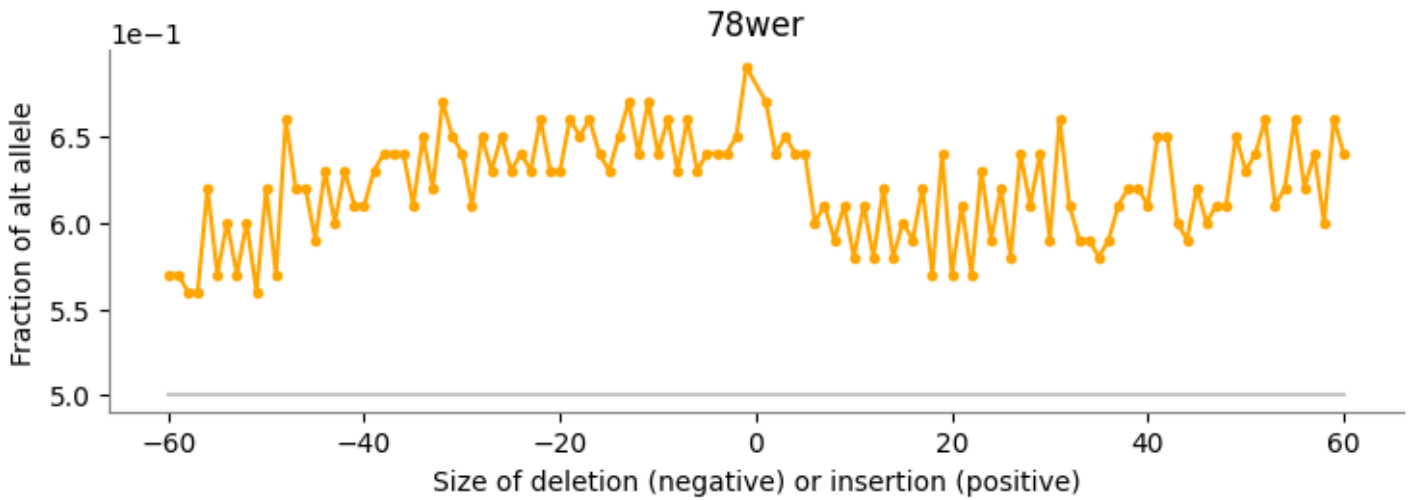
Ts/Tv stratified by QUAL



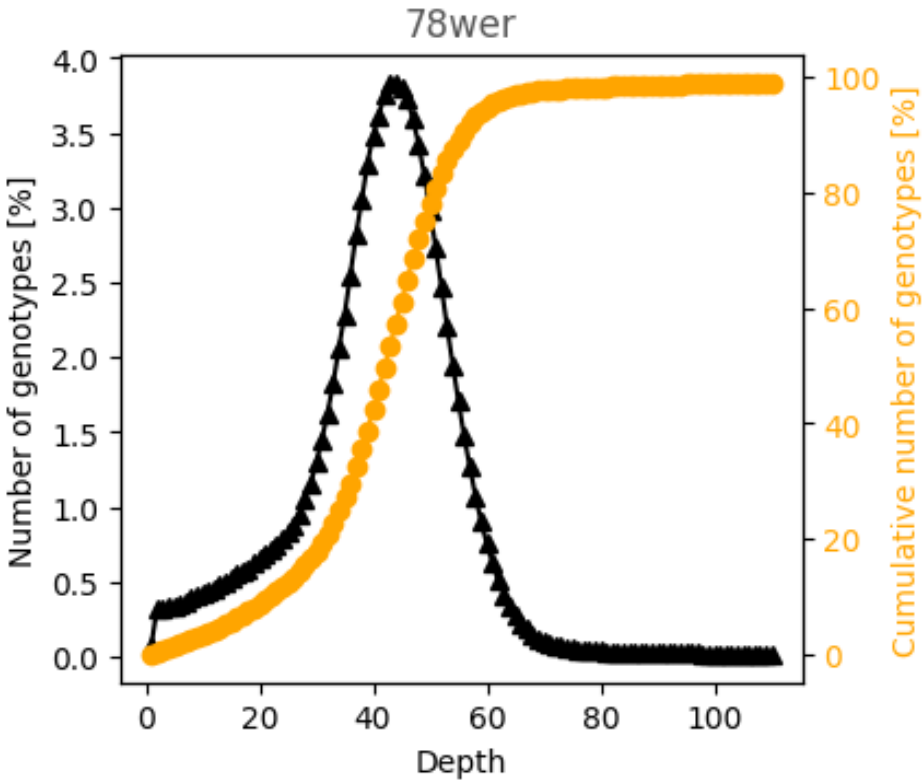
Indel distribution



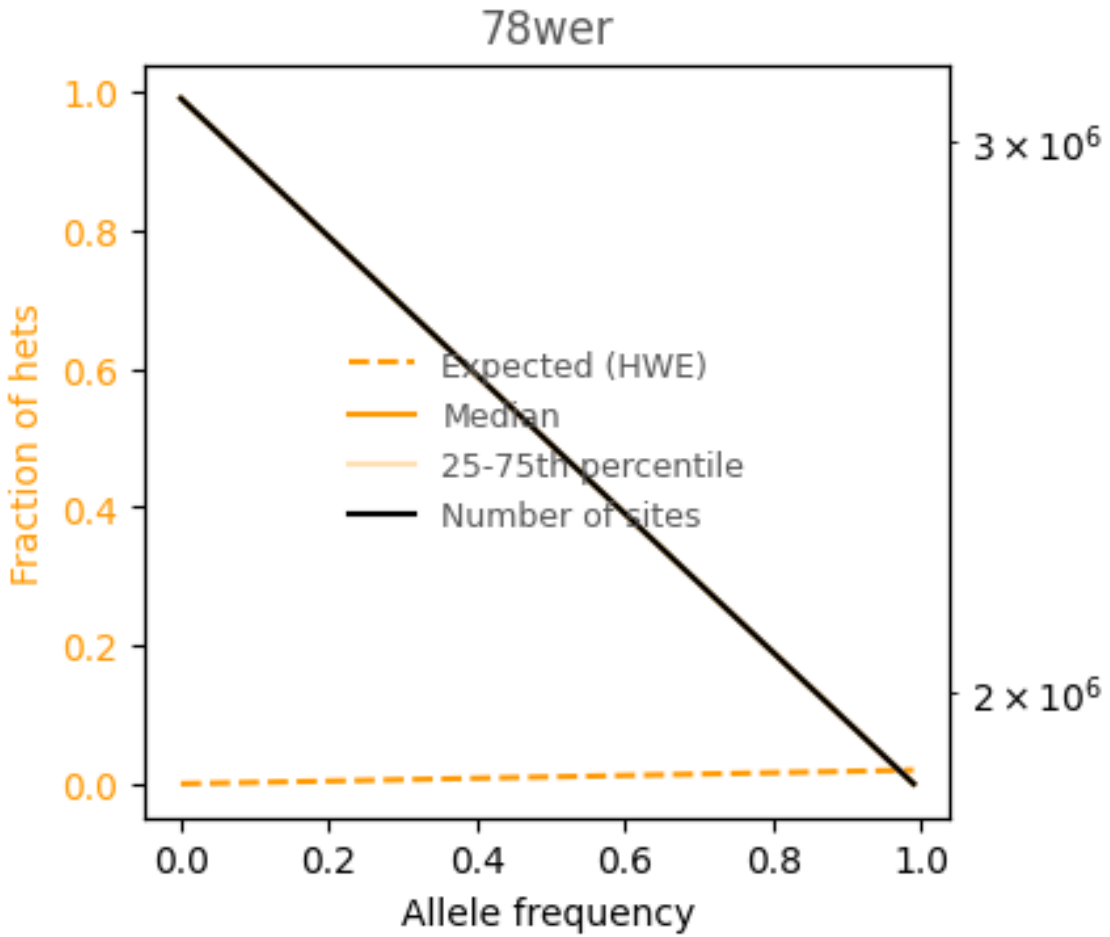
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types