

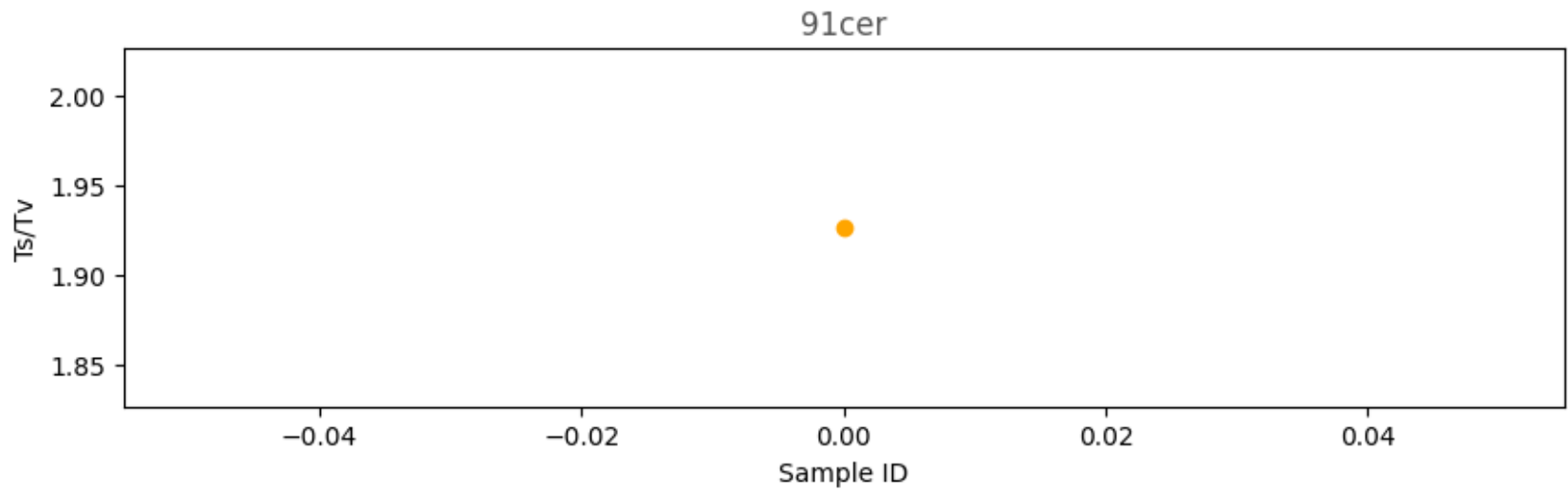
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
91cer	4,100,896	1.93	1.93	955,396	–	0	0
* frameshift ratio: out/(out+in)							

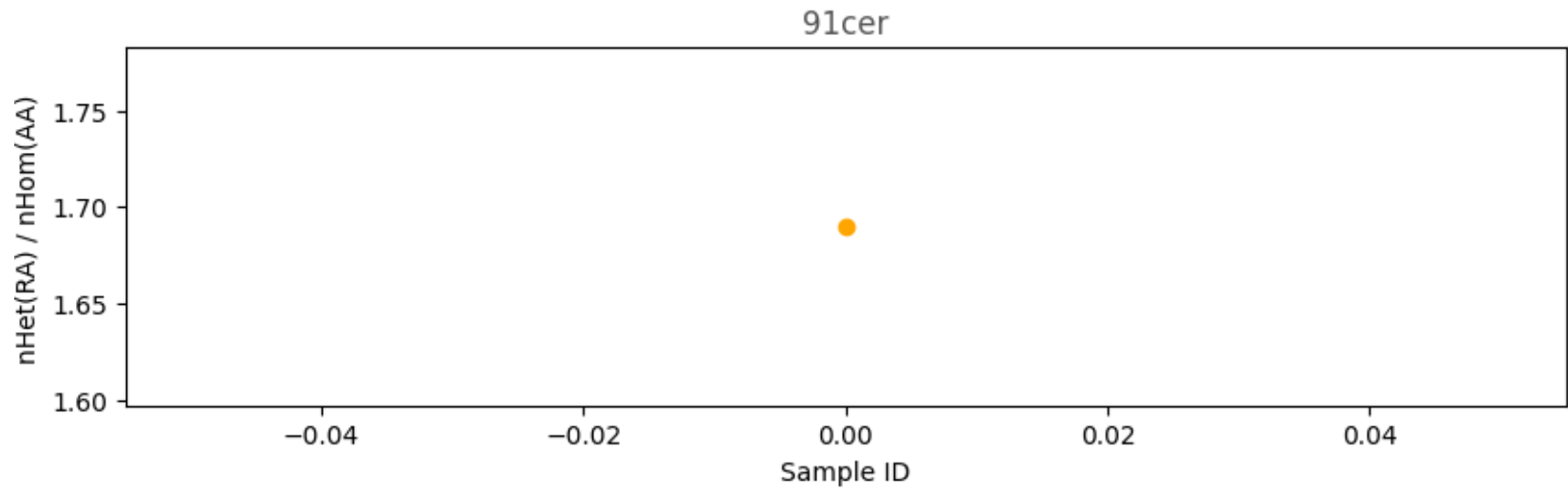
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
91cer	62.8%	1.90	68.0%	100,627	2,114

- 91cer .. /ngc/projects2/gm/data/archive/2022/variants/snv/91cerkizf-103860132452-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220422_A01176_BH2CKMDSX3-RHGM_LABKA_WGSA_KUT-WGSAKUT04164_22RKG009560x01_snv_germline_raw.haplotype_caller.vcf.gz

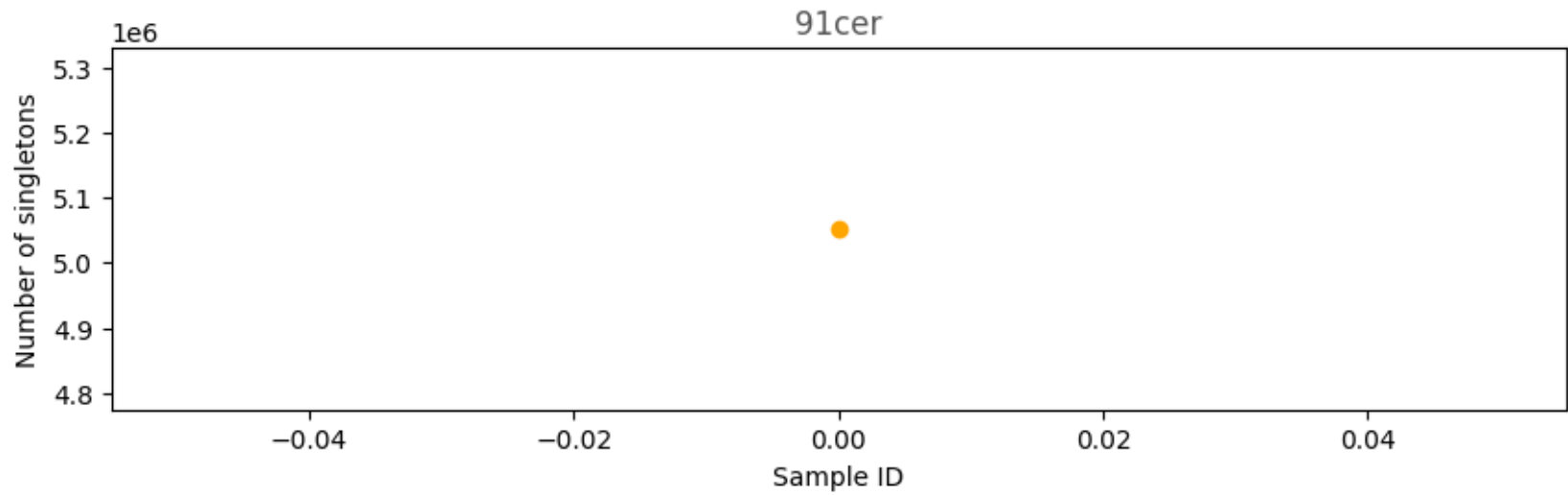
Ts/Tv by sample



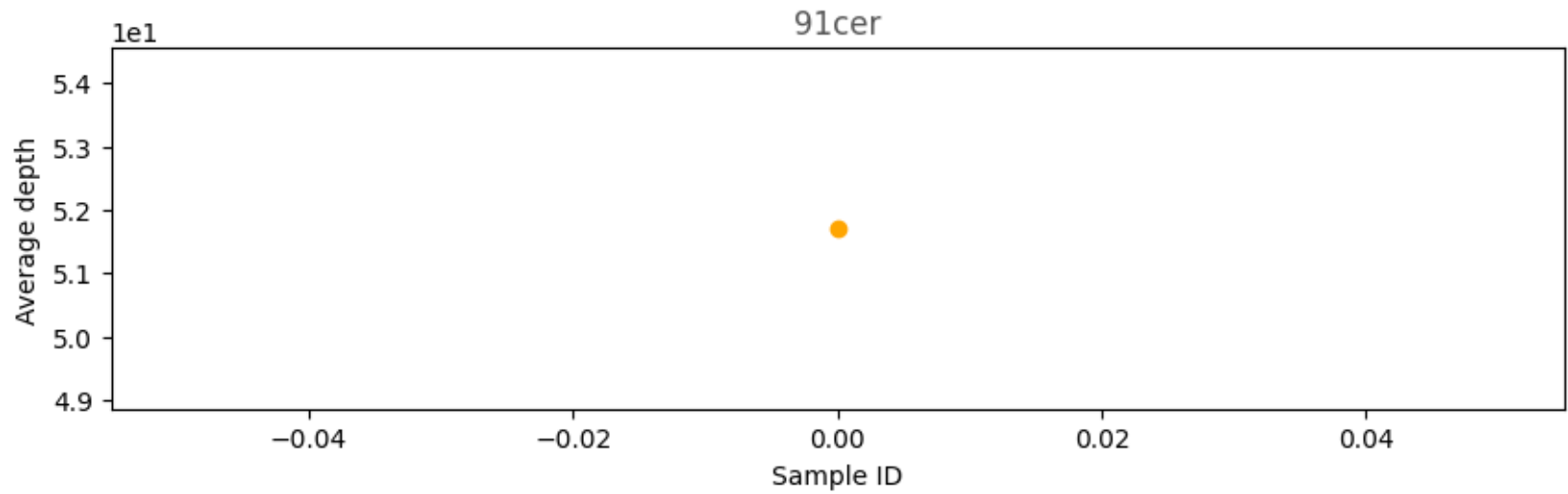
Hets vs non-ref Homs by sample



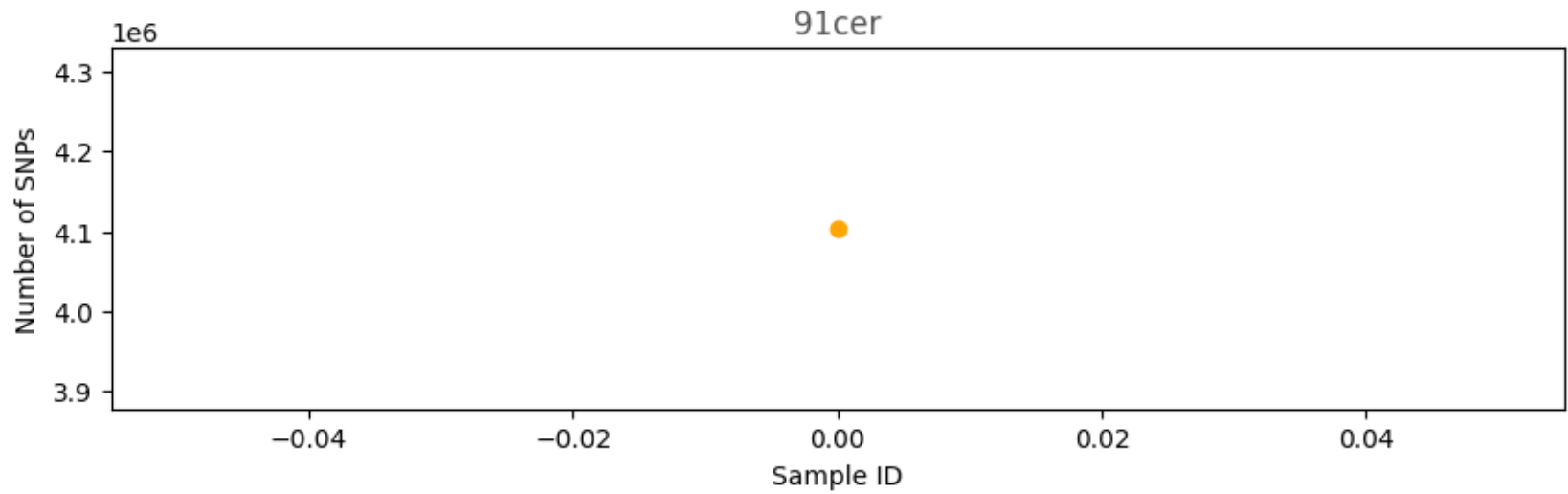
Singletons by sample (hets and homs)



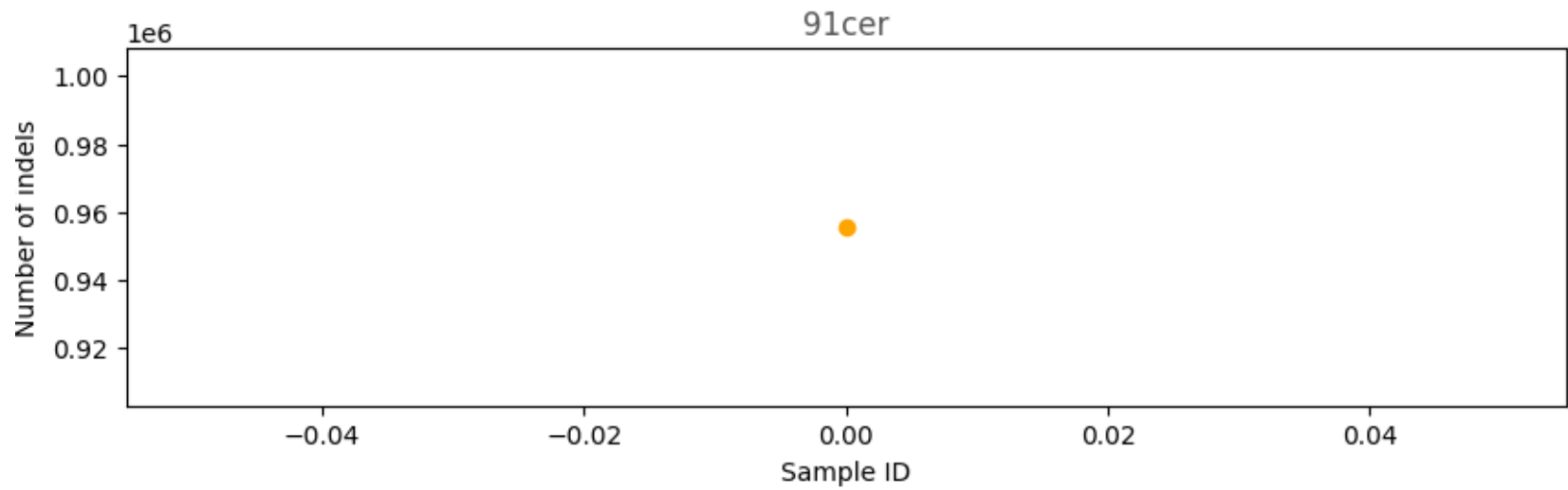
Average depth by sample



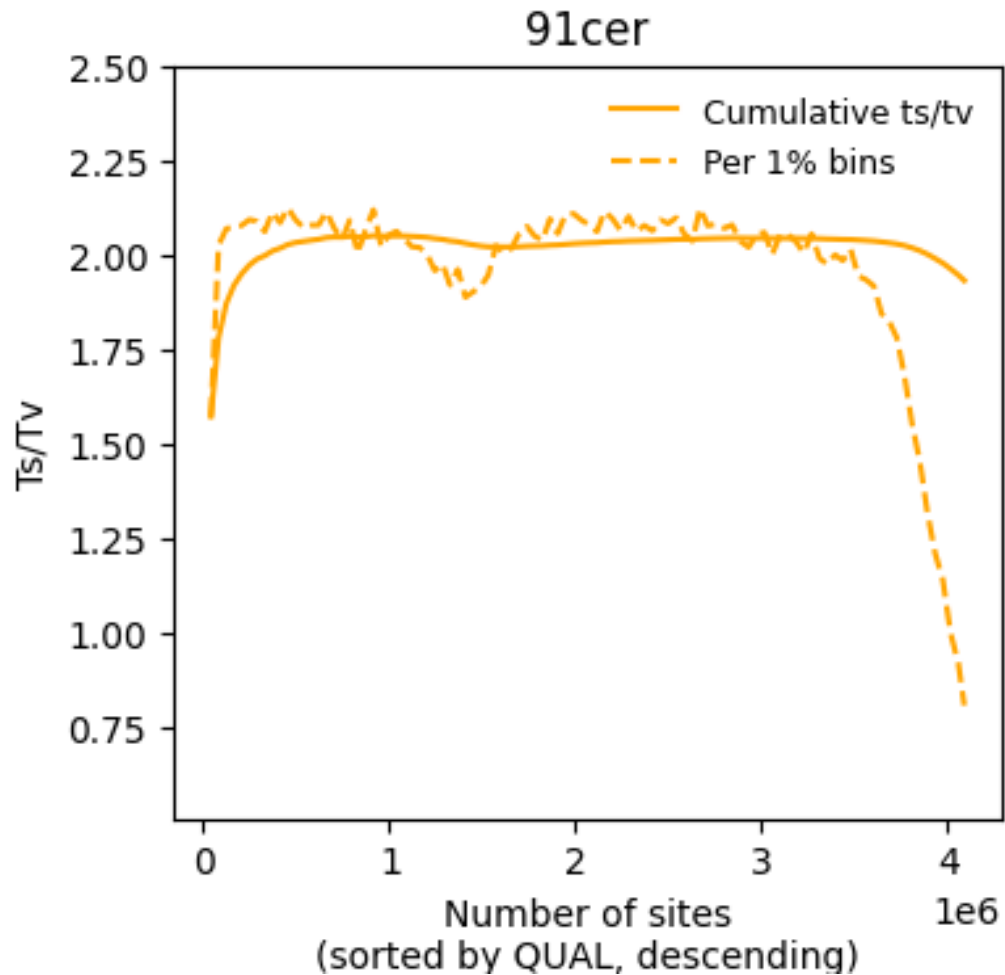
Number of SNPs by sample



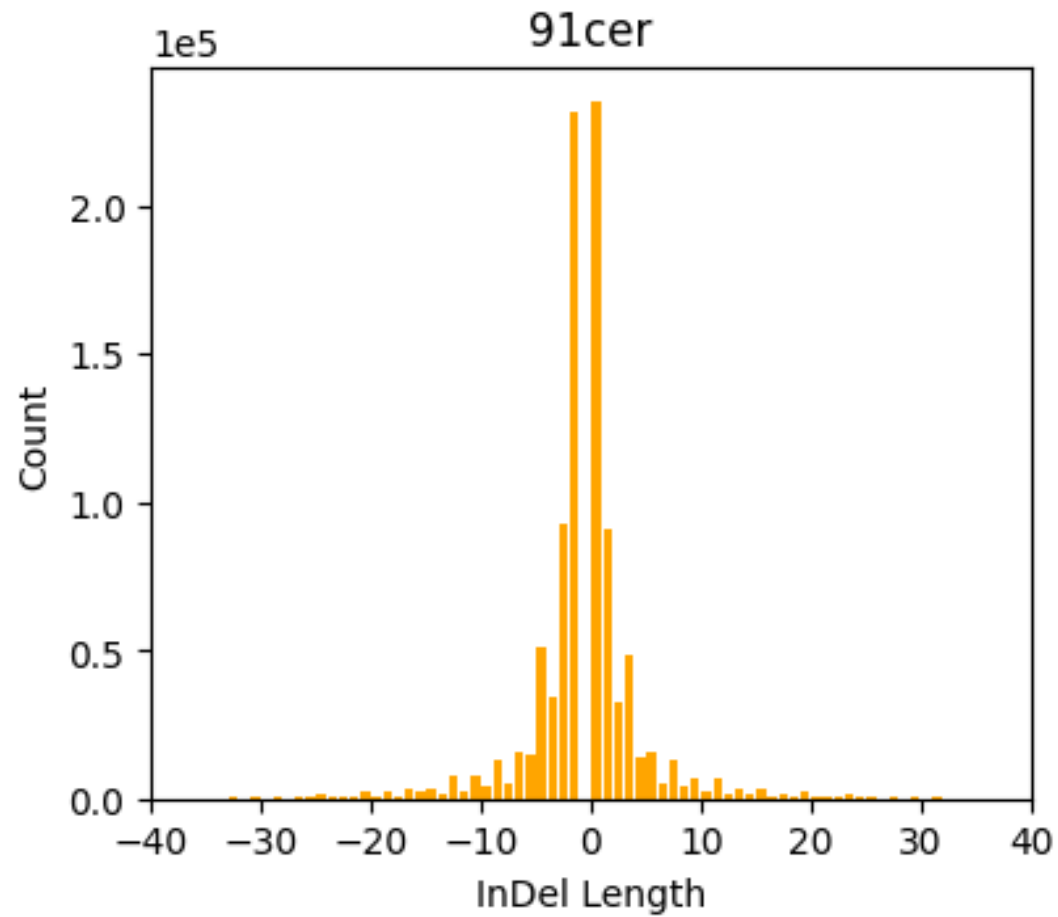
Number of indels by sample



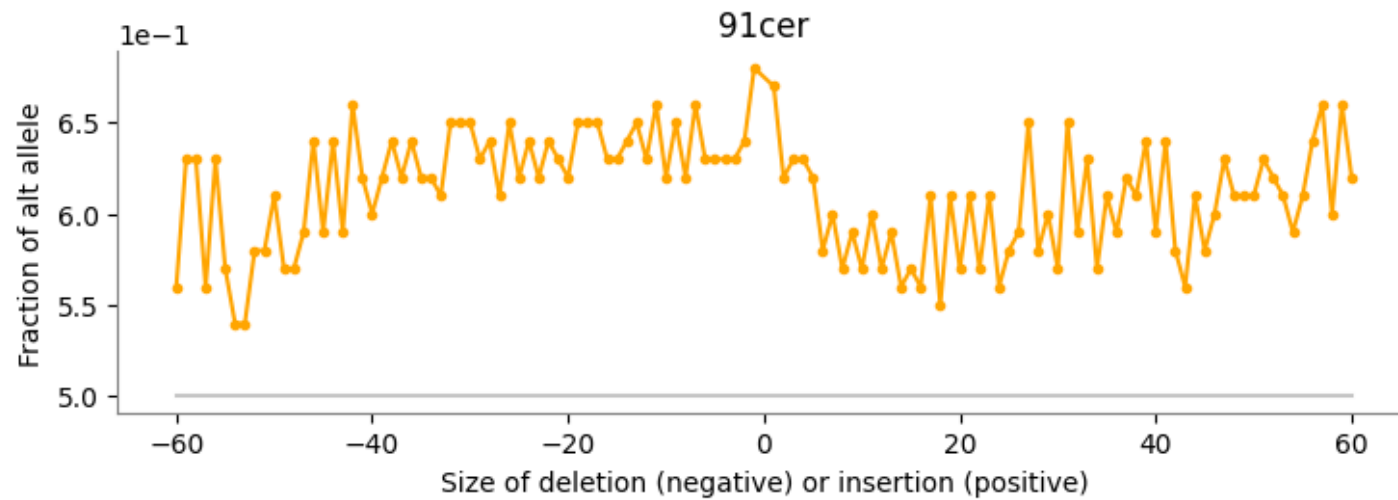
Ts/Tv stratified by QUAL



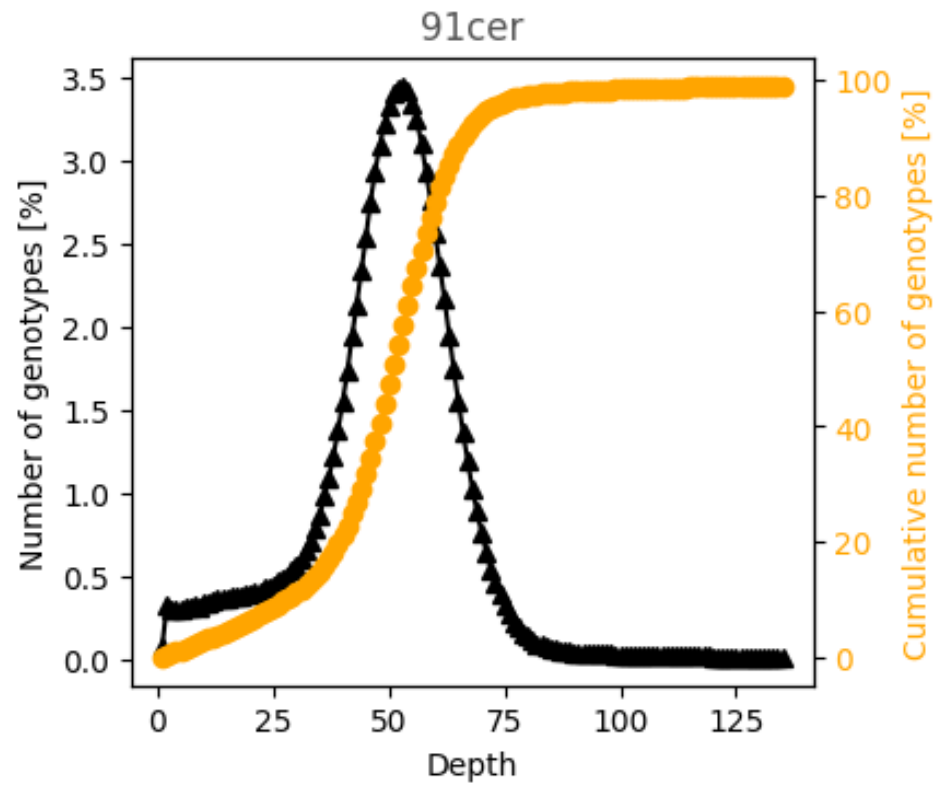
Indel distribution



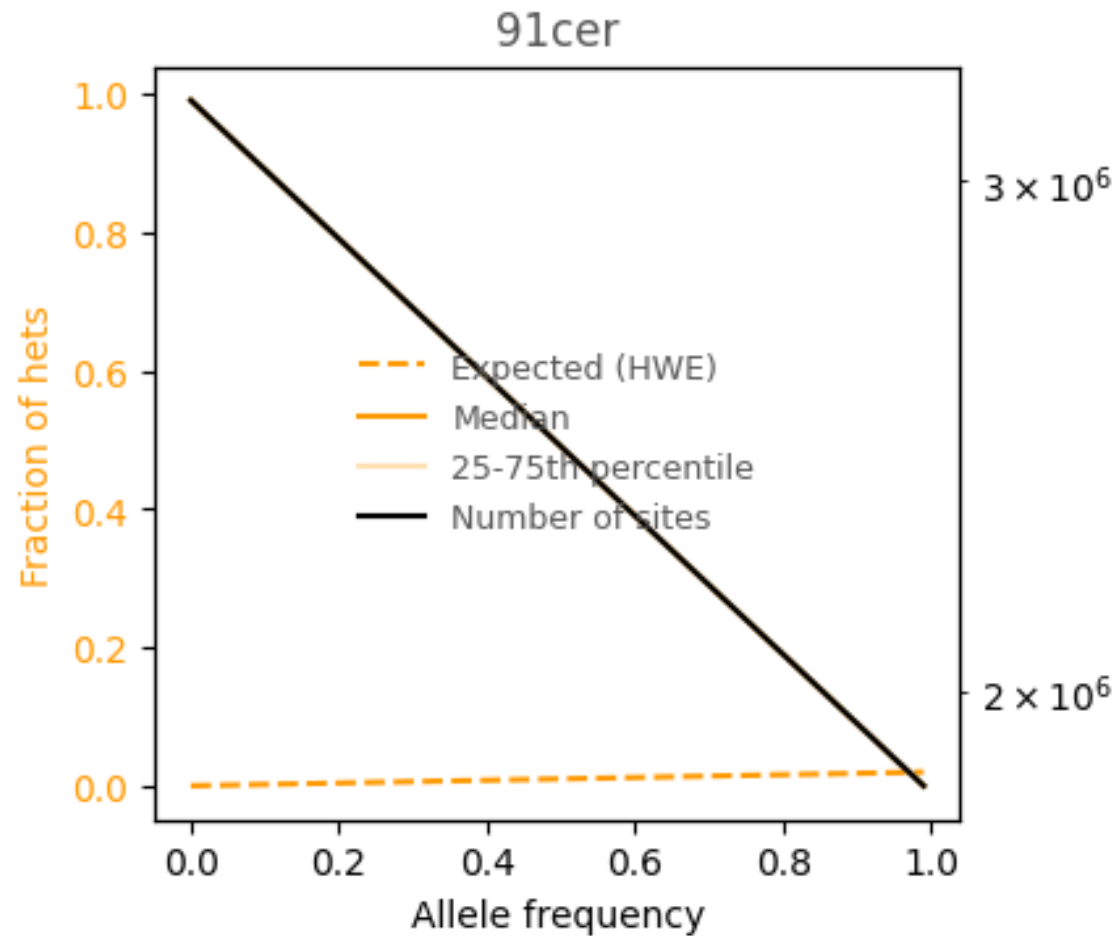
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

