

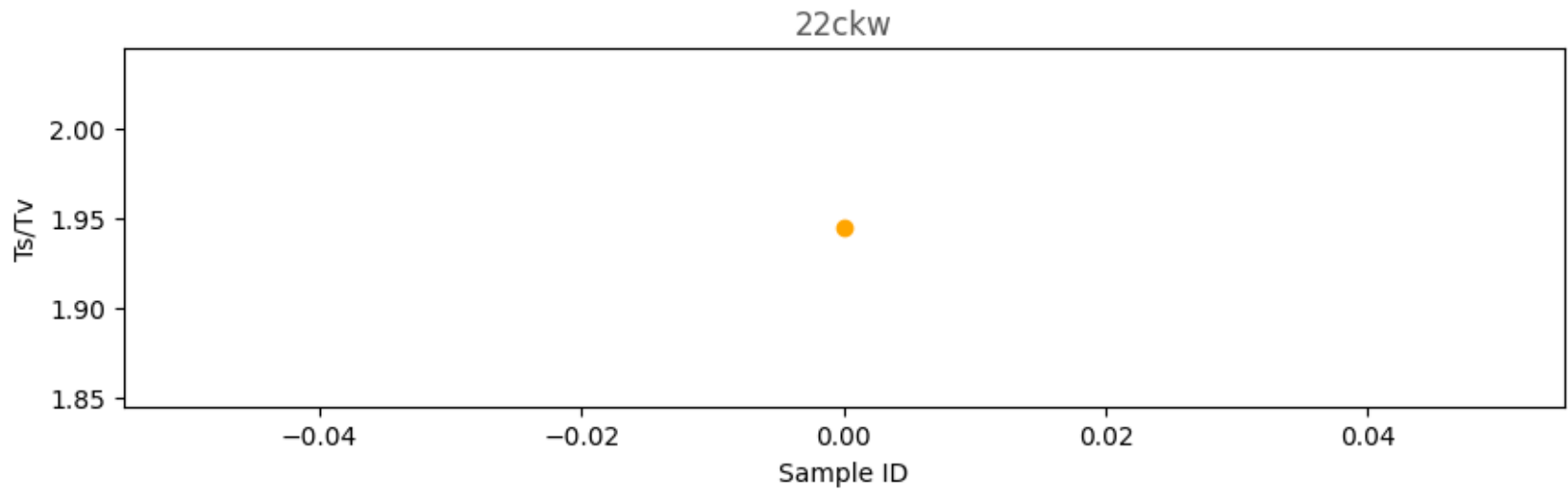
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
22ckw	4,015,450	1.94	1.95	922,872	–	0	0
* frameshift ratio: out/(out+in)							

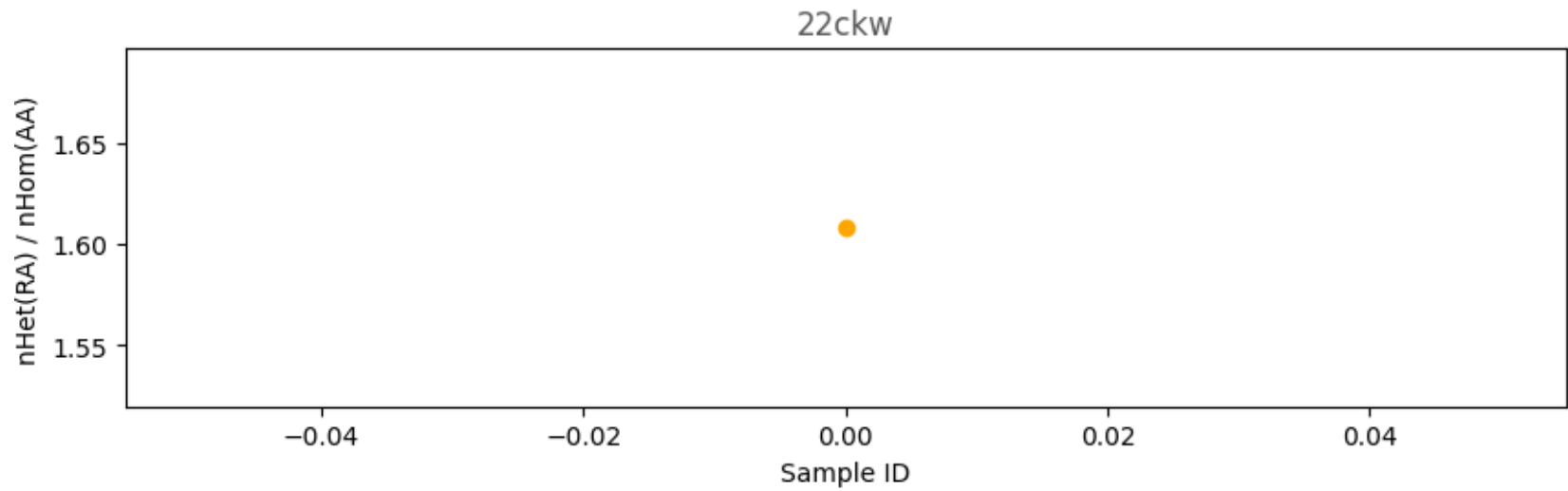
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
22ckw	61.7%	1.92	65.2%	86,491	1,856

- 22ckw .. /ngc/projects2/gm/data/archive/2022/variants/snv/22ckweprm-103884637538-Normal_B
load_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01459-220914_A01411_AHTNL5DSX3-EXT_LAB
KA_NGCWGS-NGCWGS05233_snv_germline_raw.haplotype_caller.vcf.gz

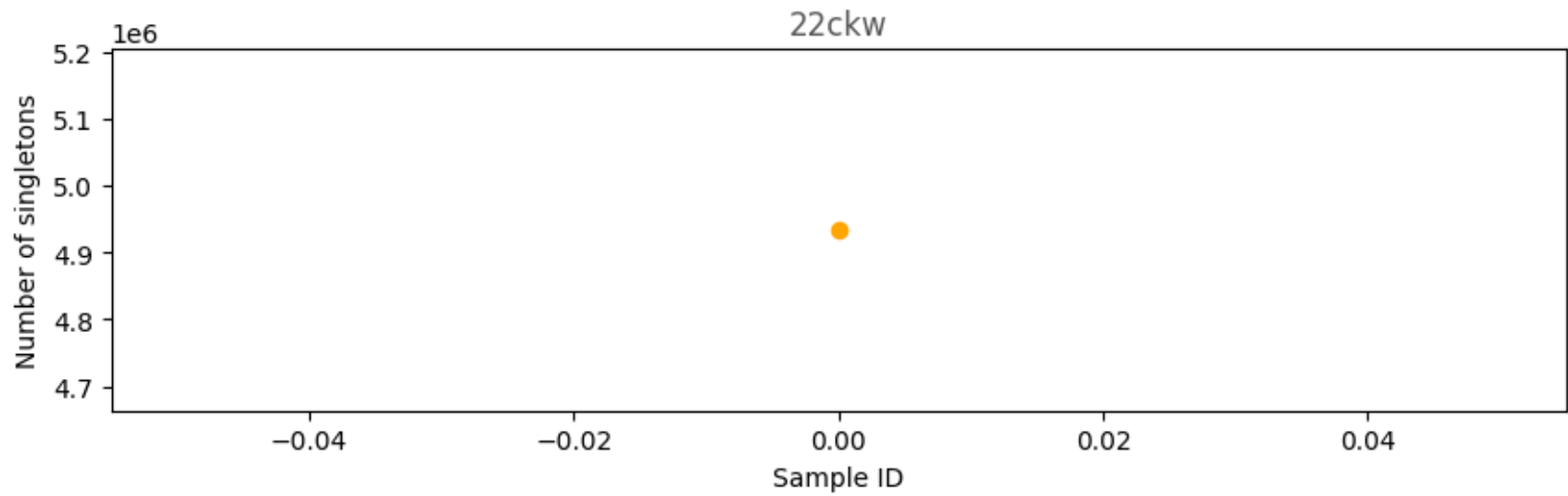
Ts/Tv by sample



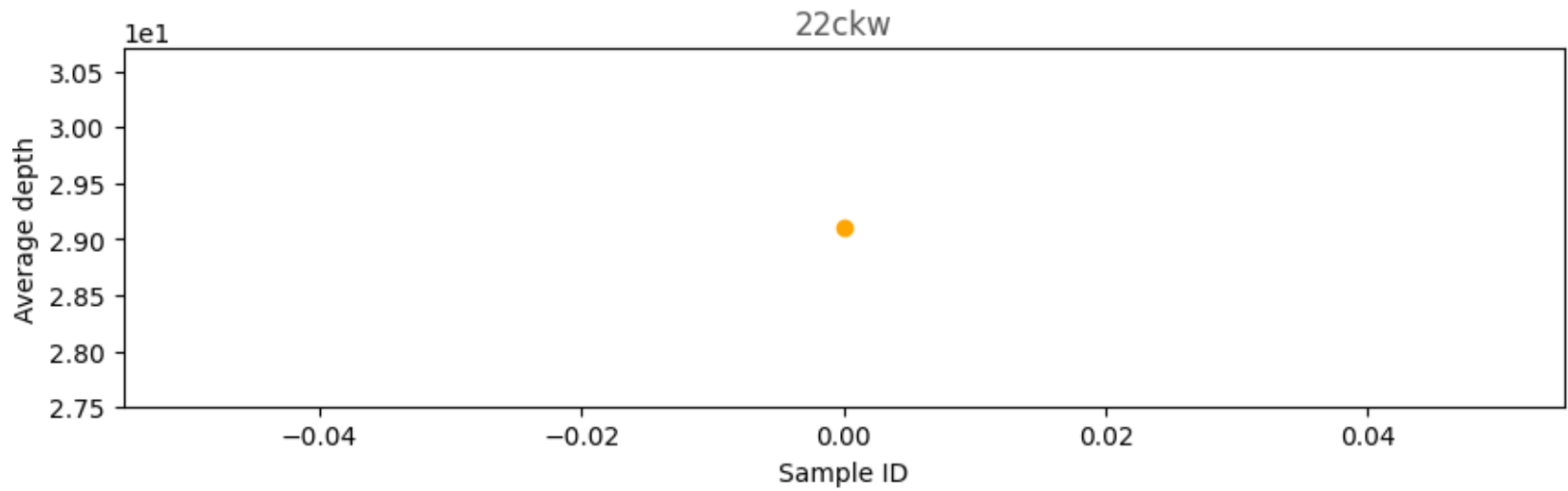
Hets vs non-ref Homs by sample



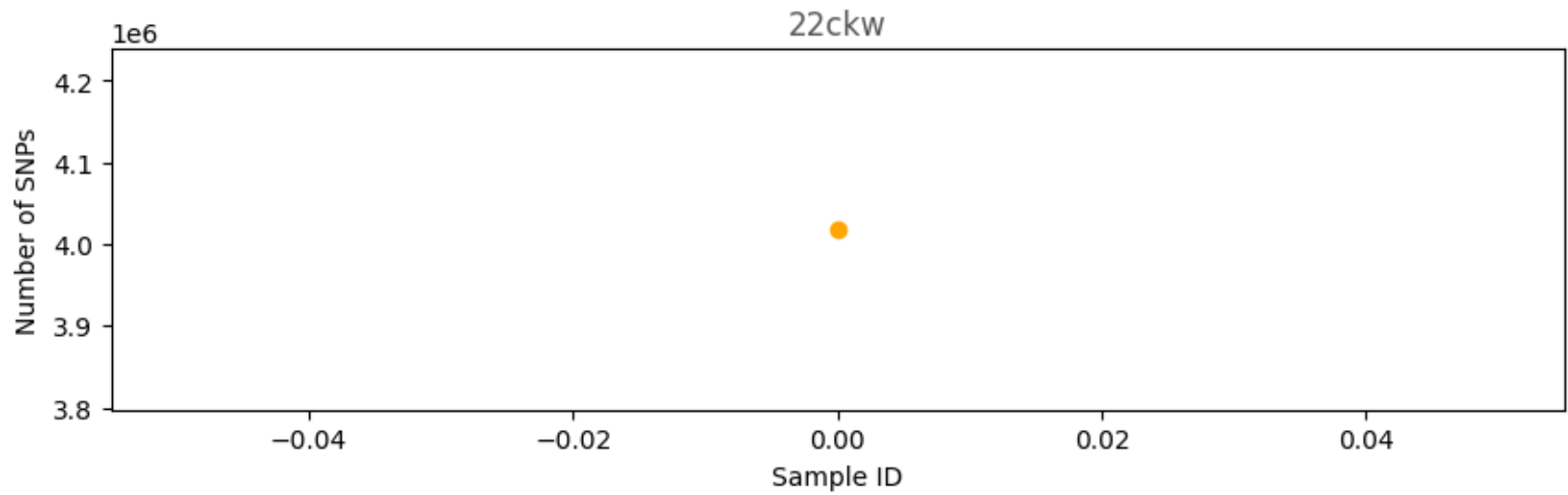
Singletons by sample (hets and homs)



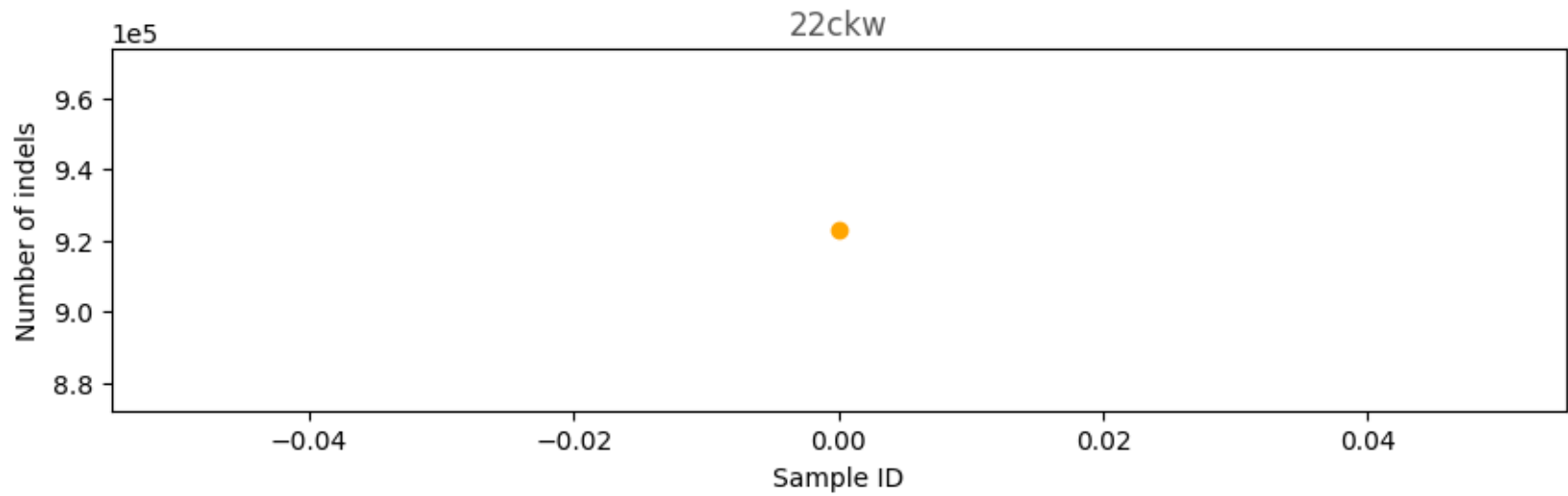
Average depth by sample



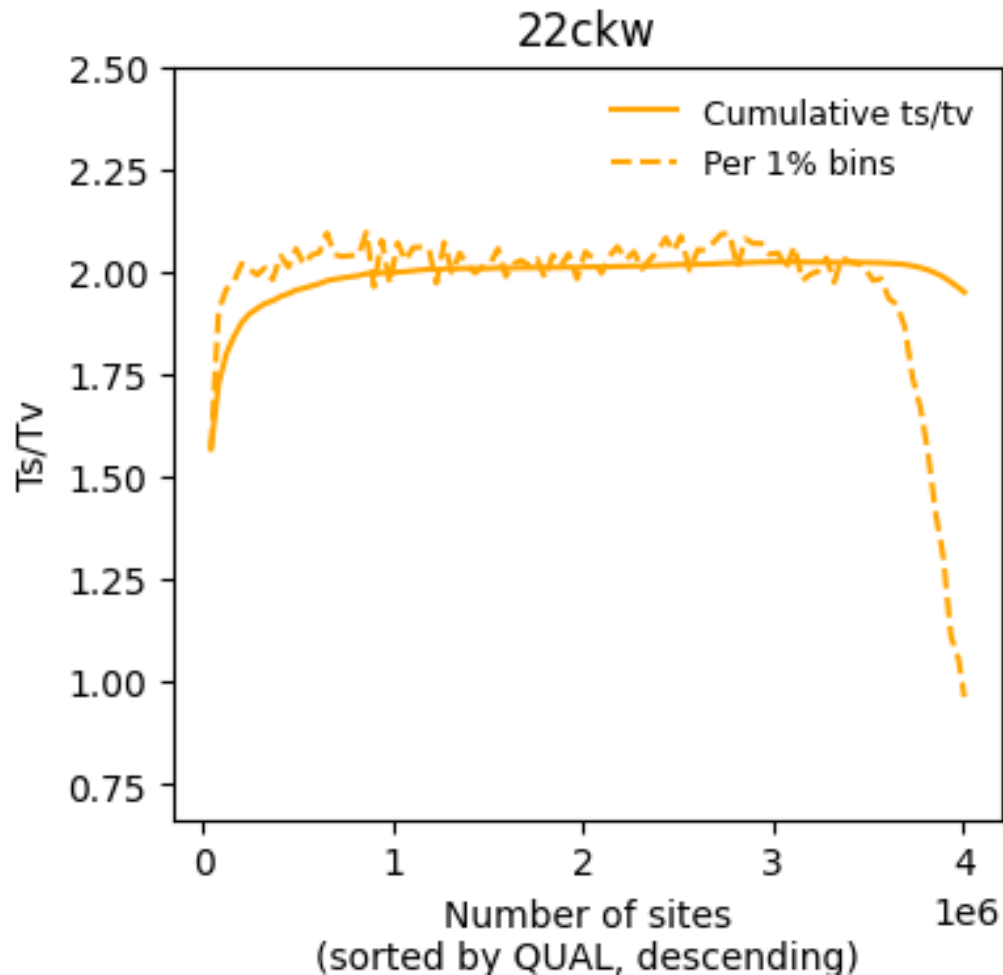
Number of SNPs by sample



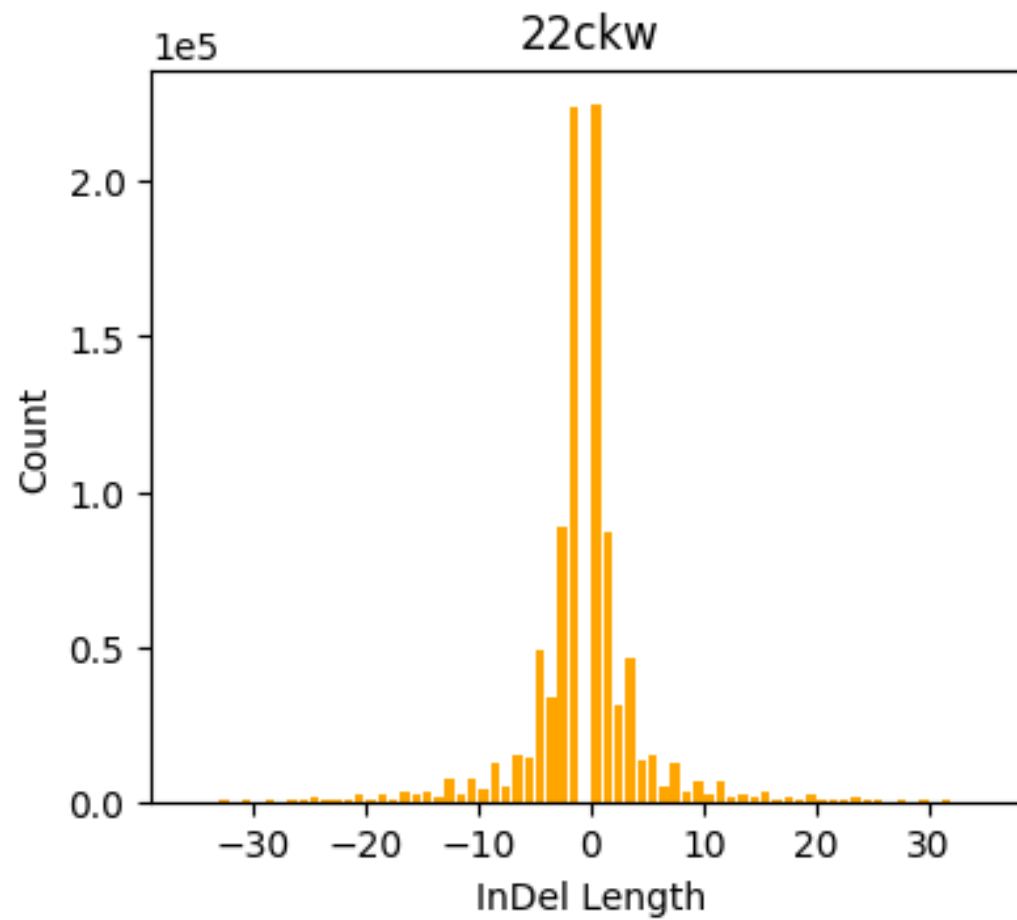
Number of indels by sample



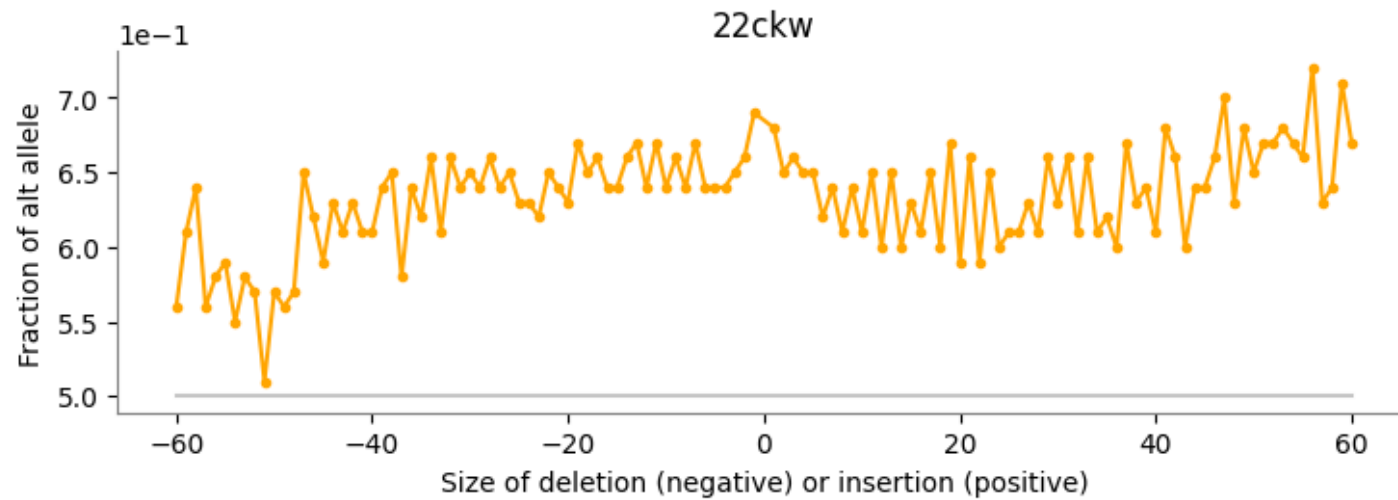
Ts/Tv stratified by QUAL



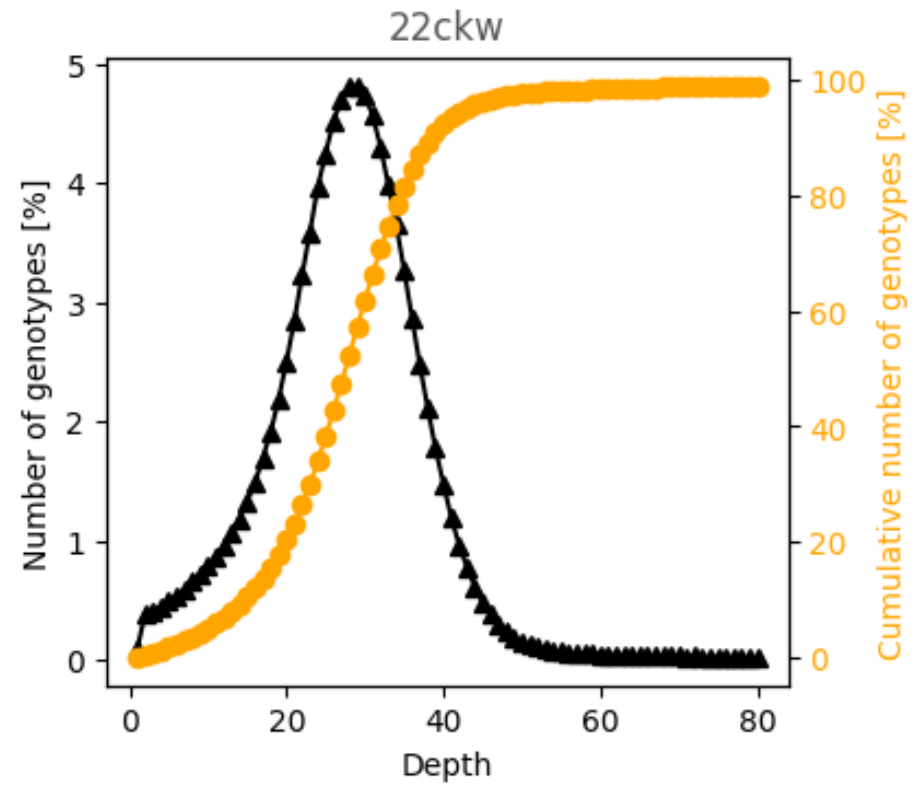
Indel distribution



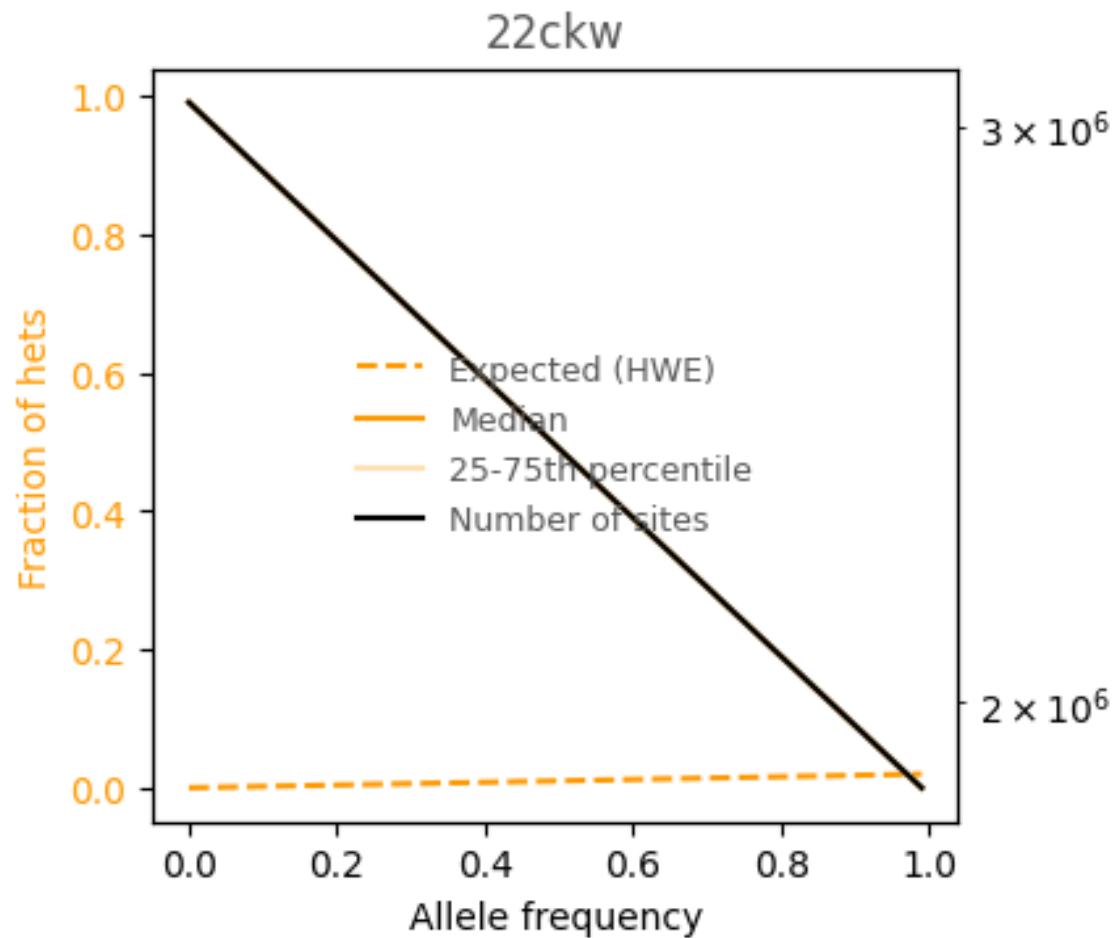
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

