

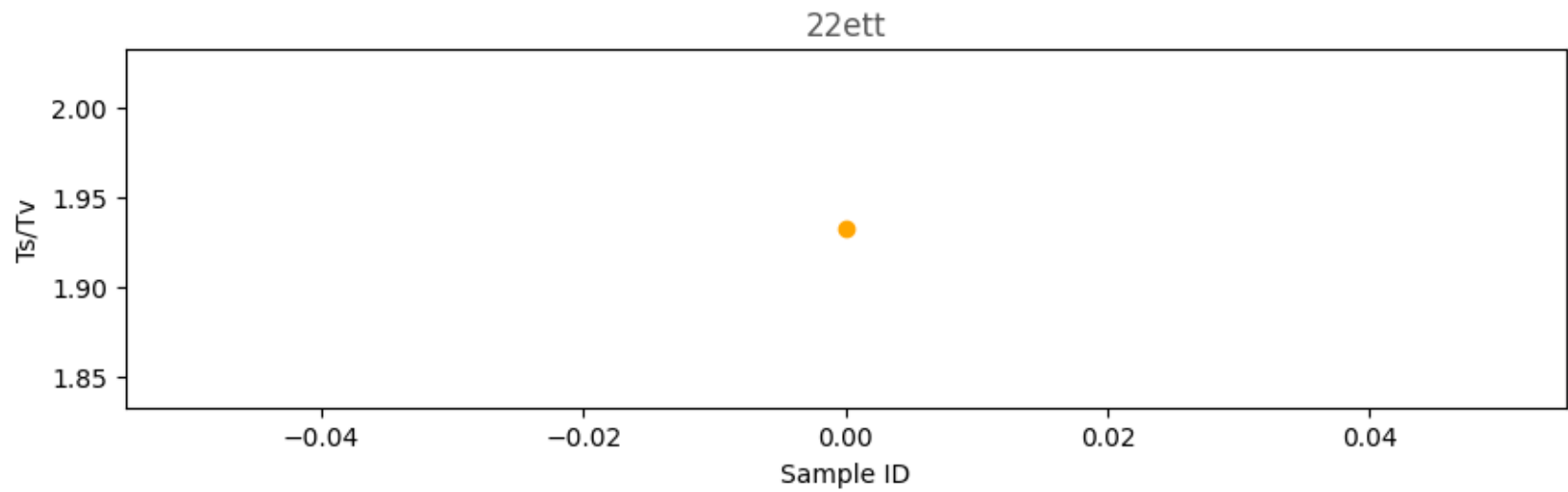
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
22ett	4,078,450	1.93	1.94	949,374	–	0	0
* frameshift ratio: out/(out+in)							

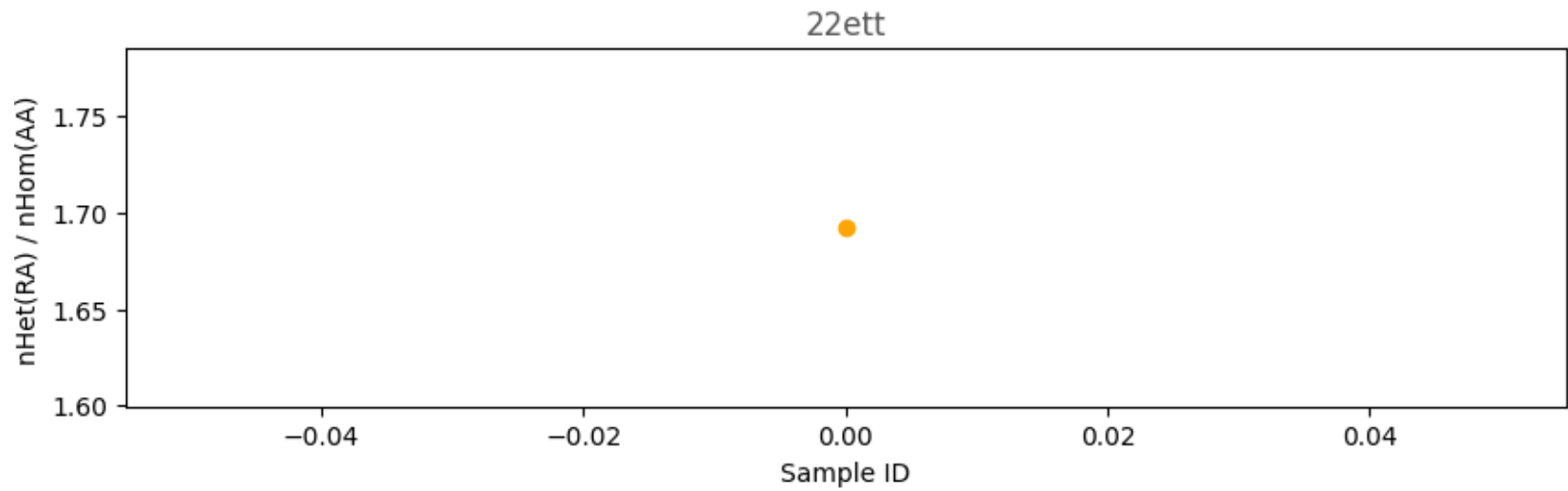
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
22ett	62.9%	1.91	67.8%	99,666	2,020

- 22ett .. /ngc/projects2/gm/data/archive/2022/variants/snv/22etthanf-110298125530-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-221019_A01411_AH3CYYDSX5-RHGM_LABKA_WGSA
 KUT-WGSAKUT05915_22RKG024855x01_snv_germline_raw.haplotype_caller.vcf.gz

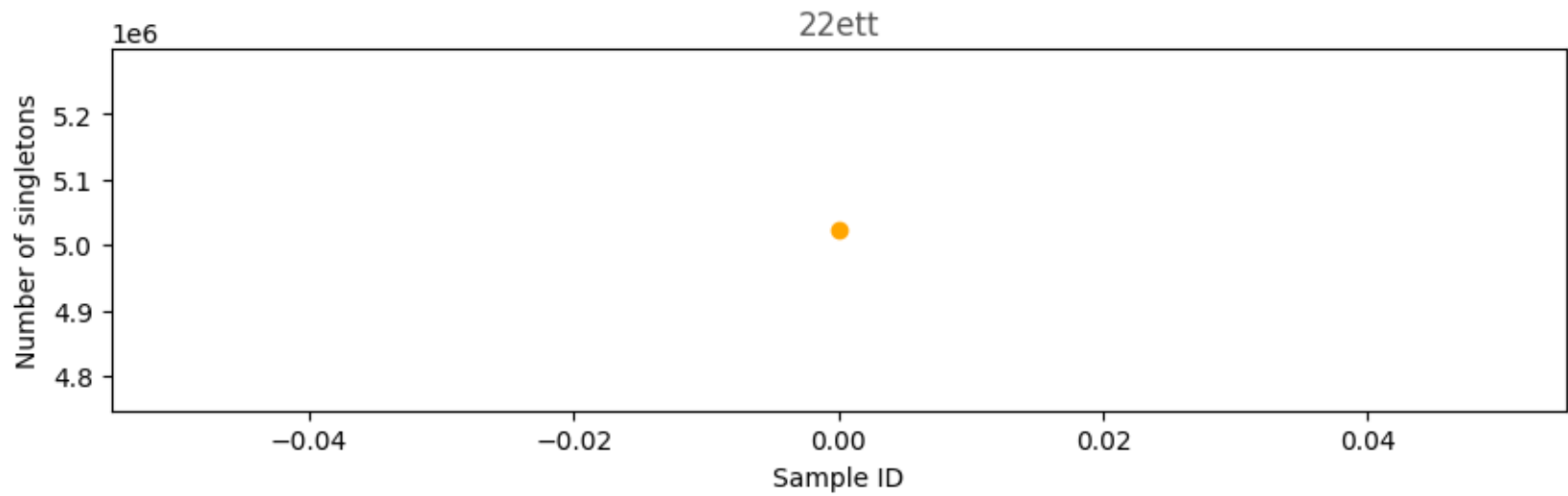
Ts/Tv by sample



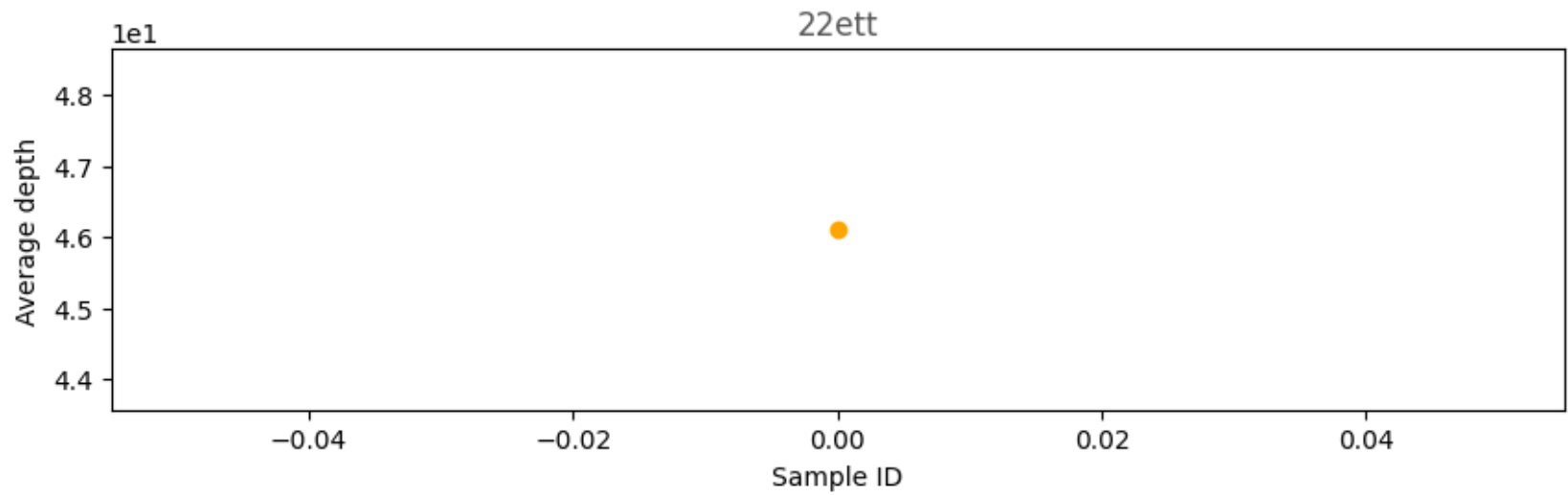
Hets vs non-ref Homs by sample



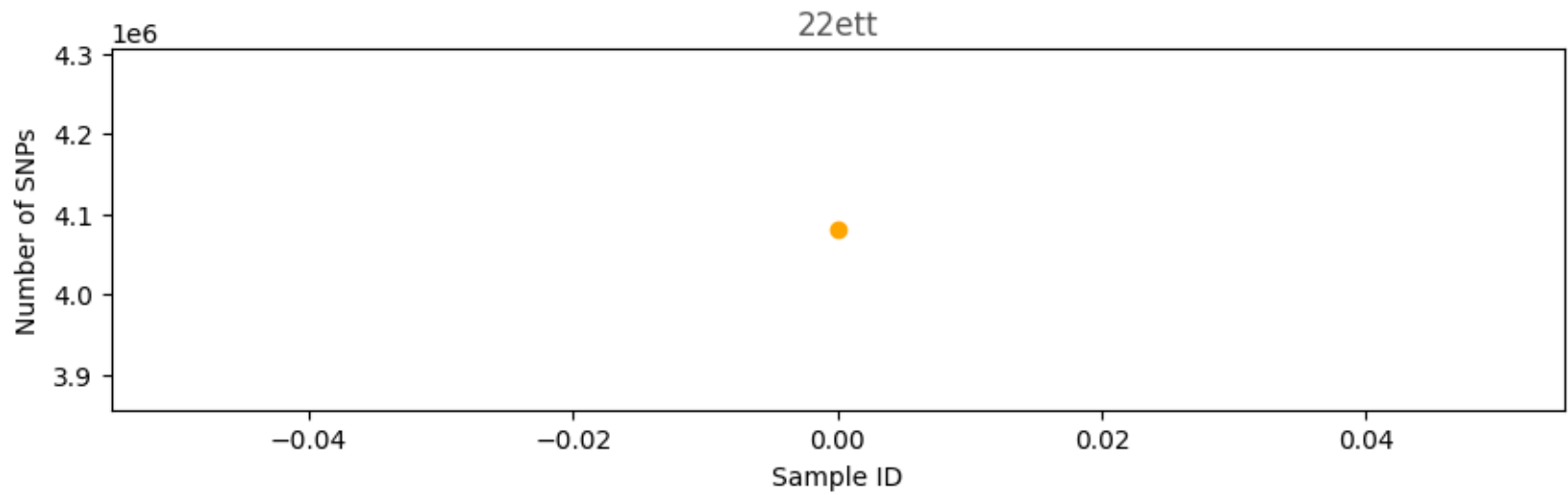
Singletons by sample (hets and homs)



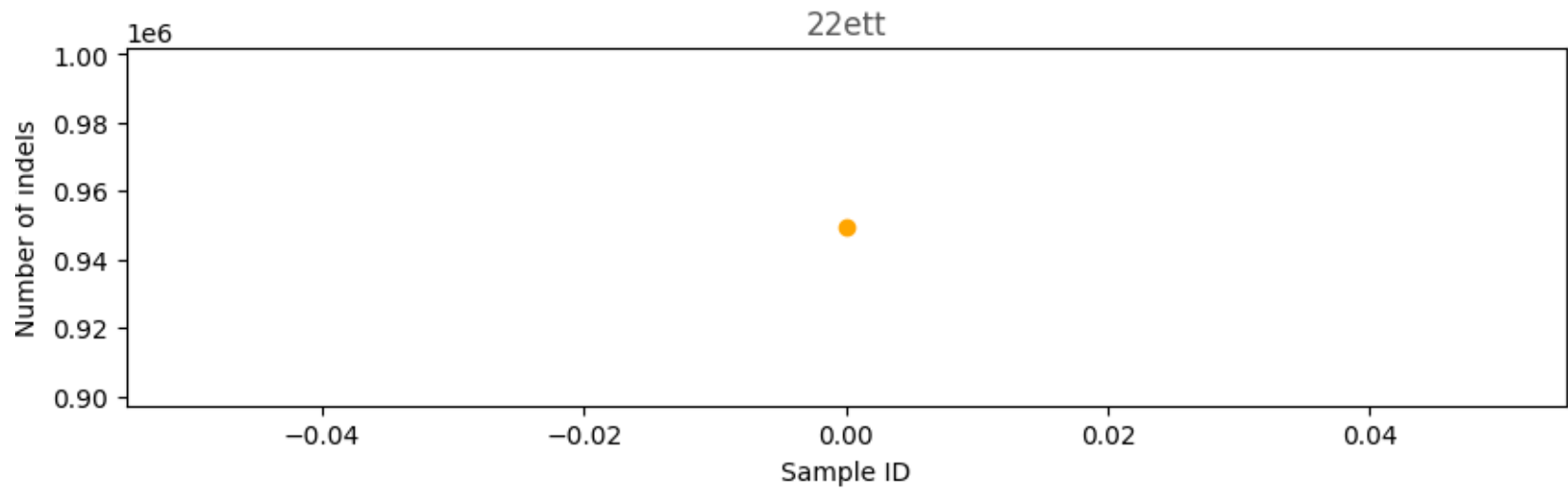
Average depth by sample



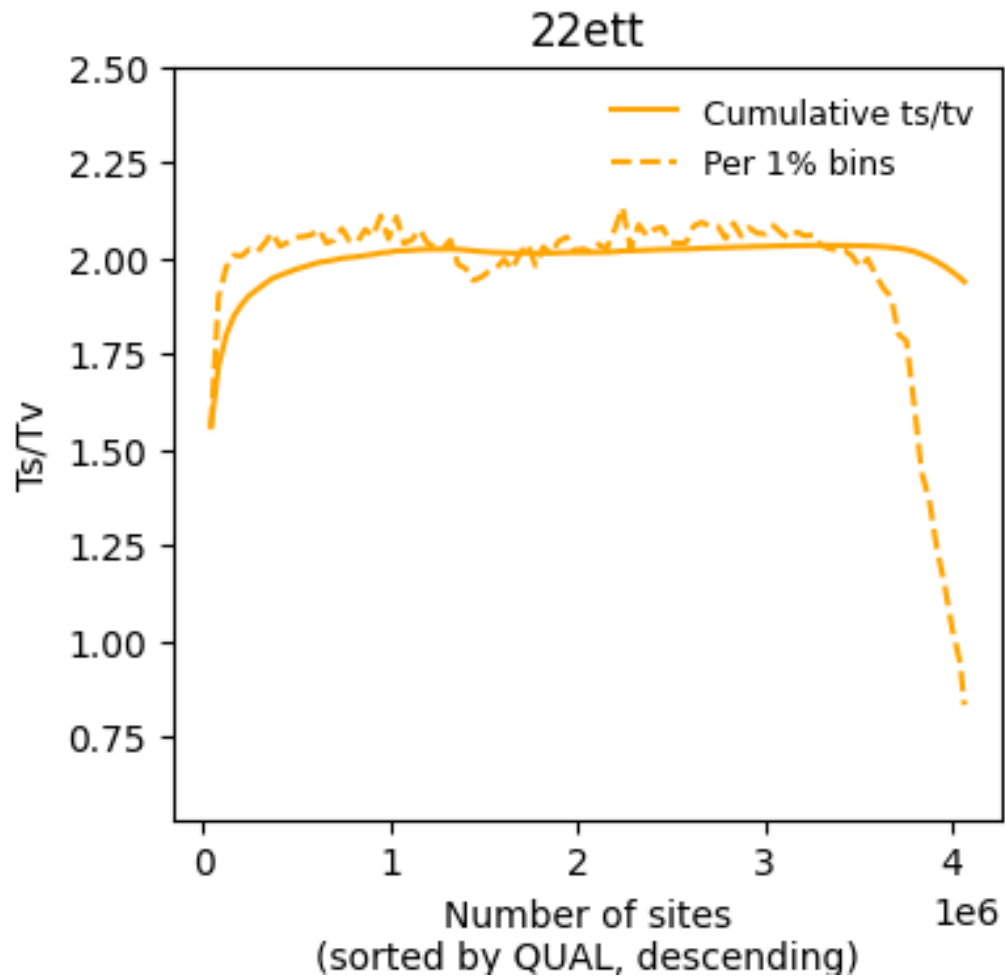
Number of SNPs by sample



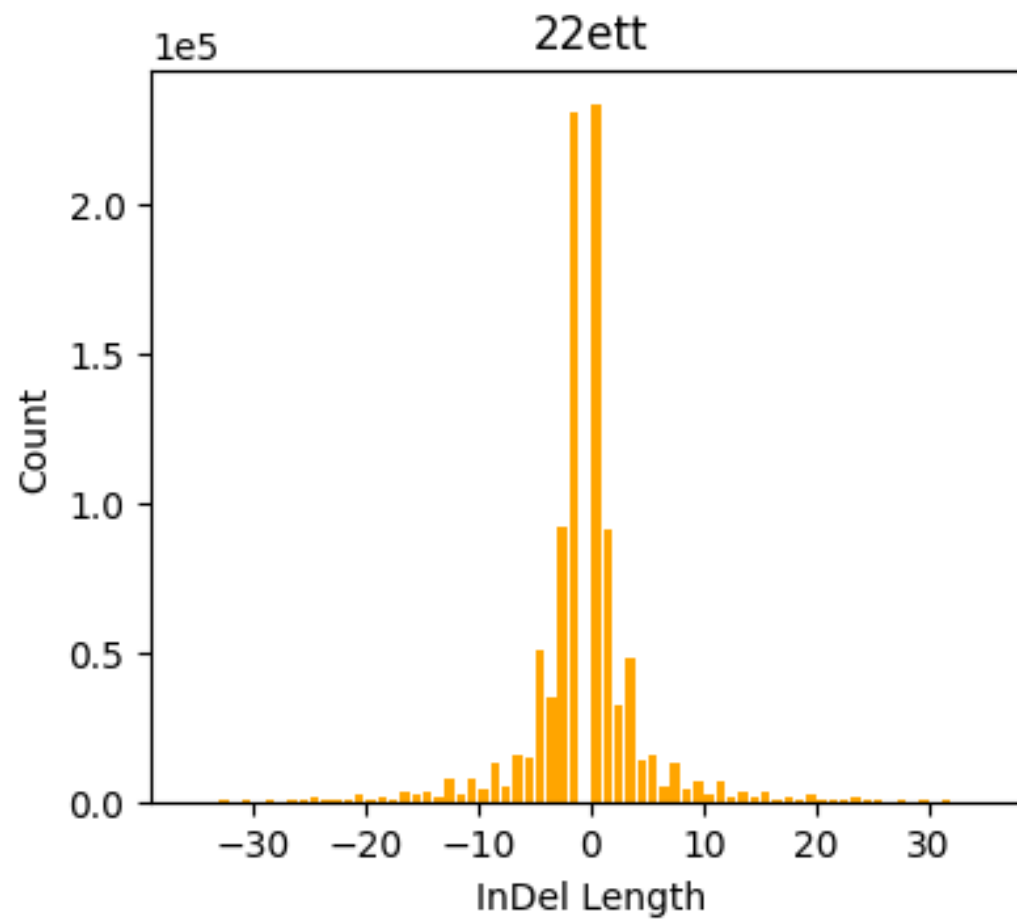
Number of indels by sample



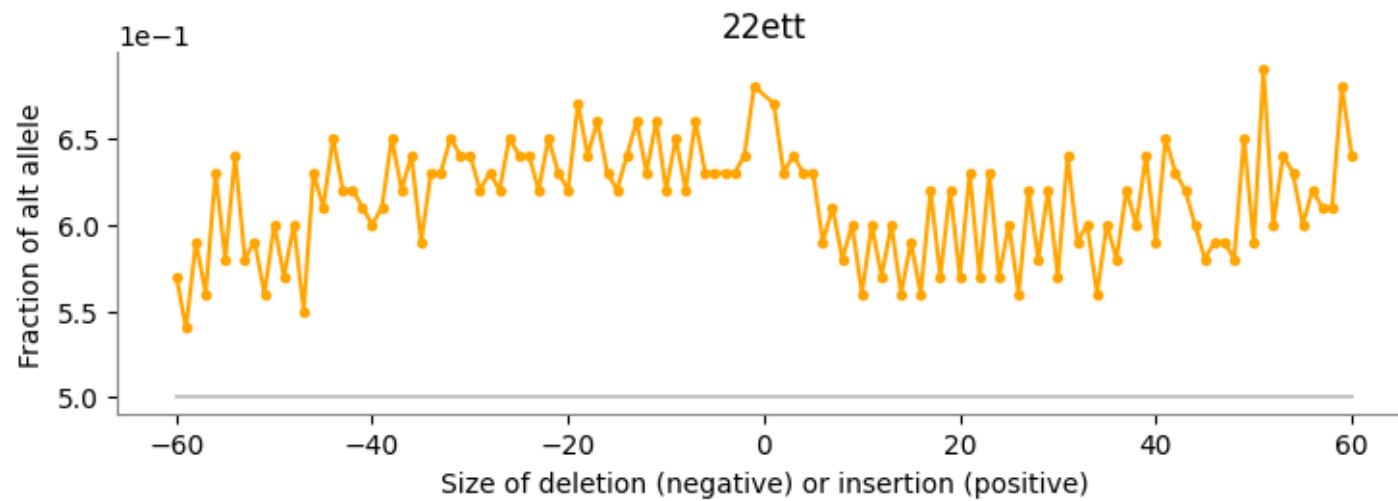
Ts/Tv stratified by QUAL



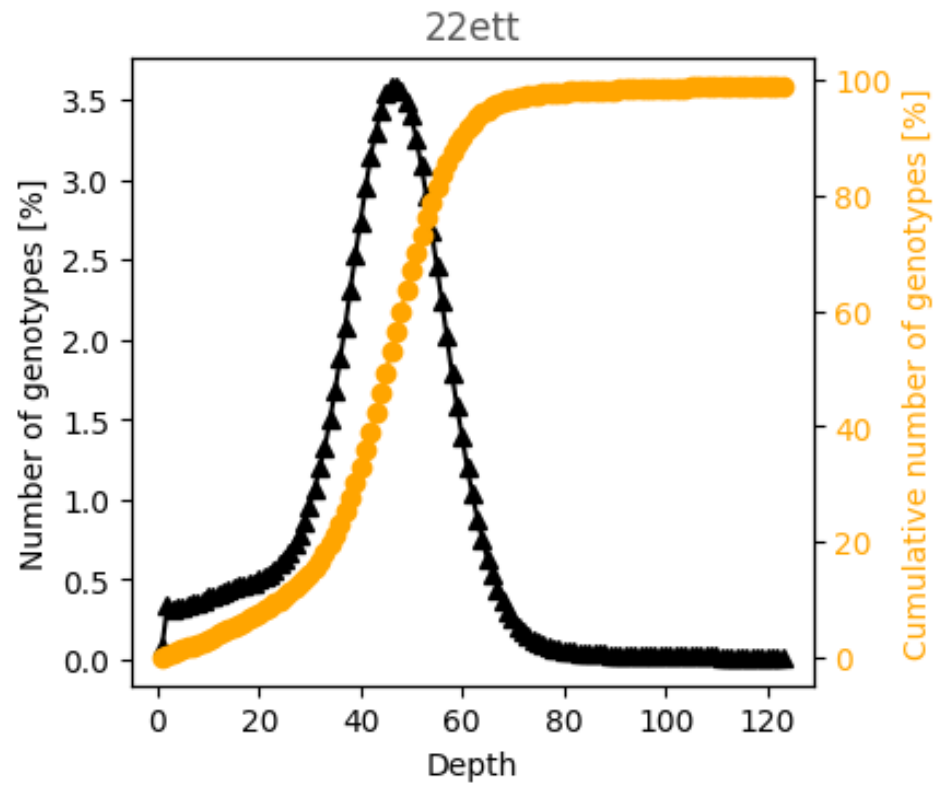
Indel distribution



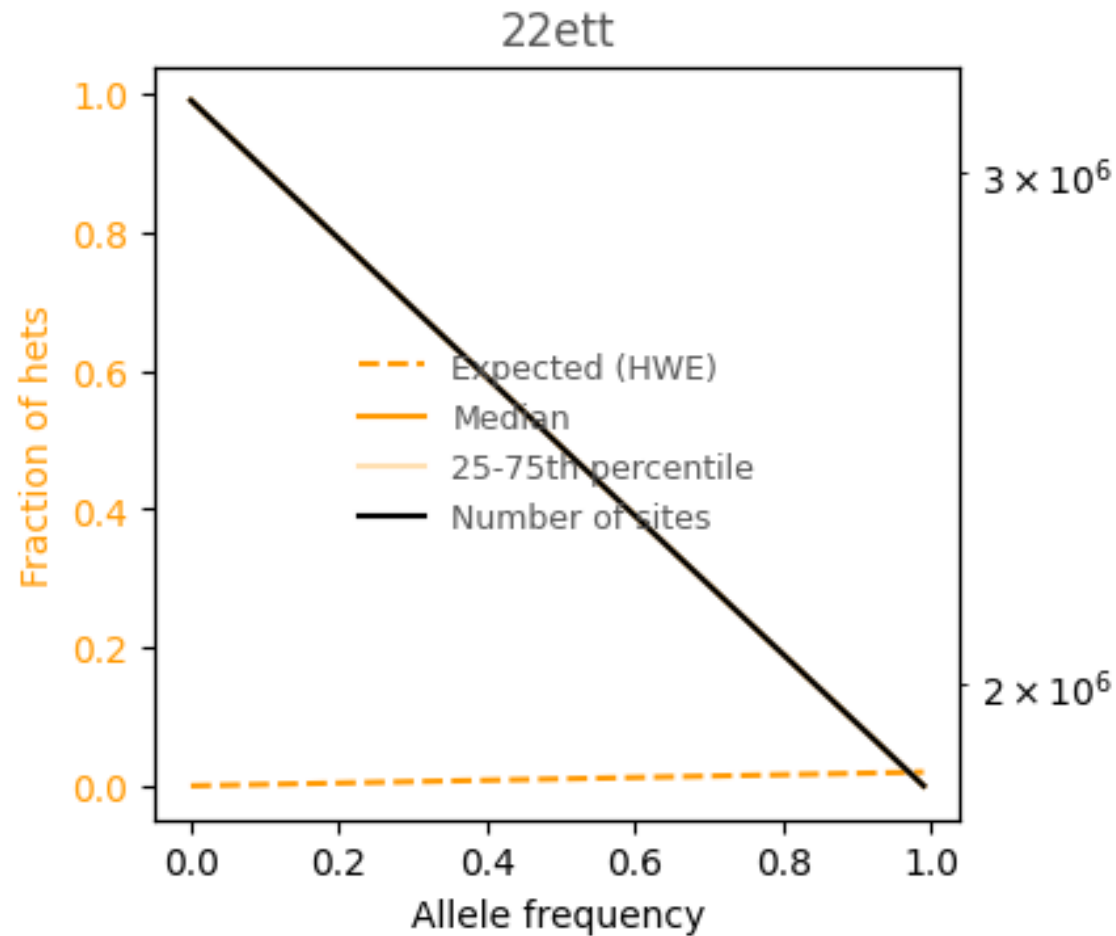
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

