

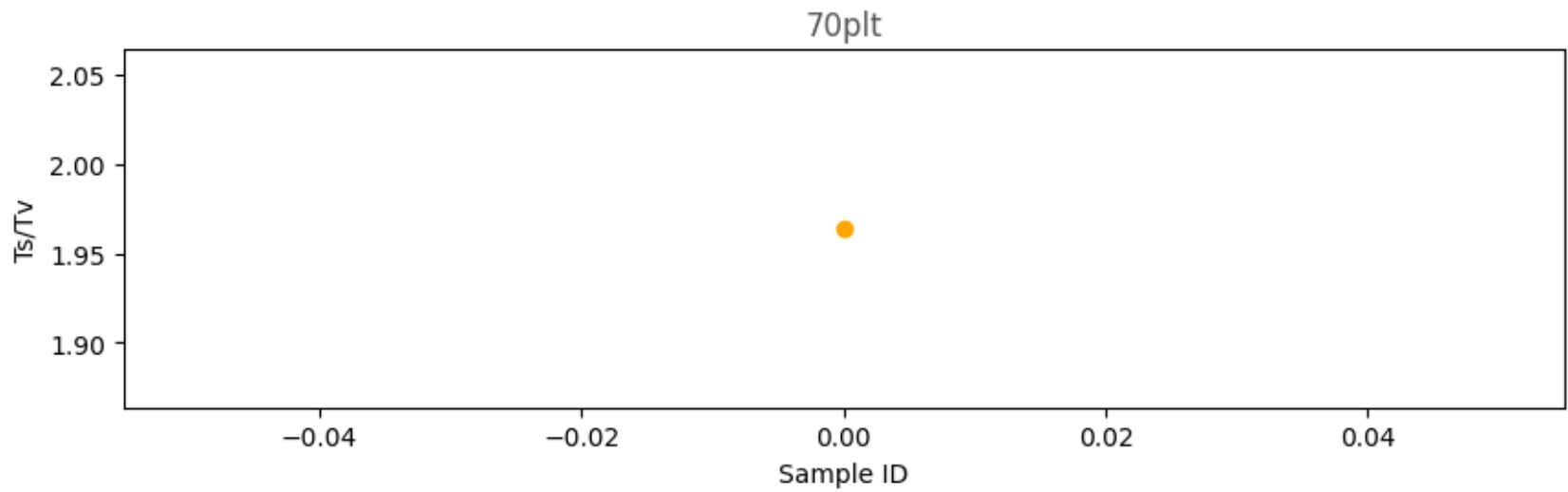
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
70plt	3,995,610	1.96	1.97	935,861	–	0	0
* frameshift ratio: out/(out+in)							

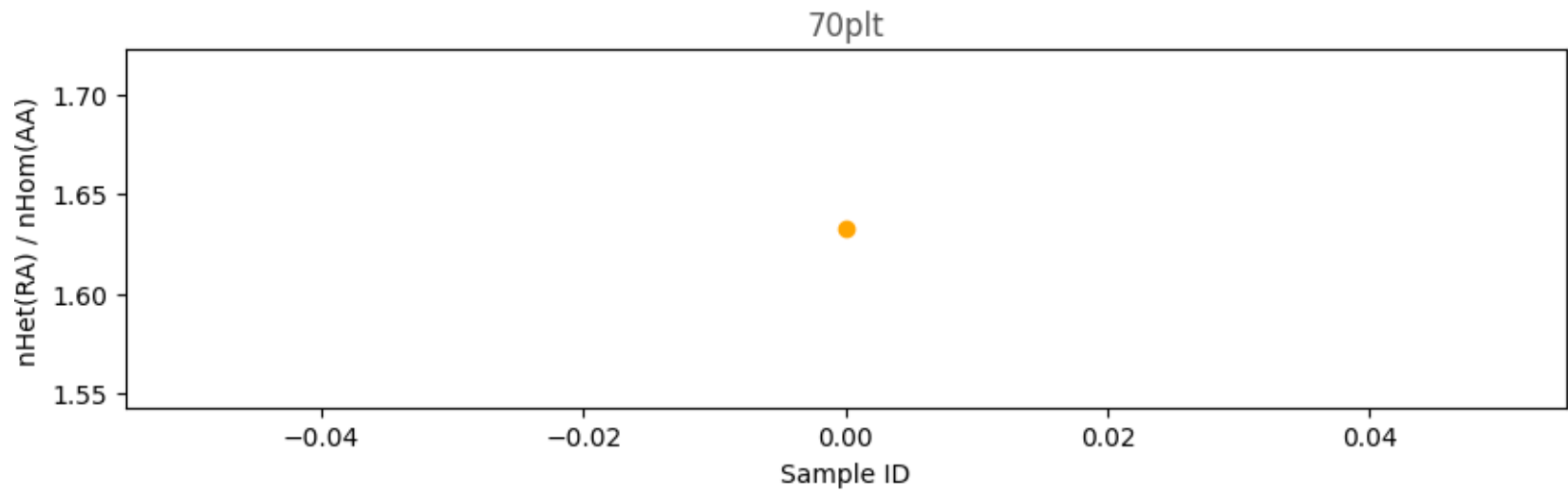
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
70plt	62.0%	1.94	66.3%	91,475	1,596

- 70plt .. /ngc/projects2/gm/data/archive/2022/variants/snv/70pltbsim-103821305554-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220105_A01466_AH5T7TDSX3-RHGM_LABKA_WGS-
 WGS03620_snv_germline_raw.haplotype_caller.vcf.gz

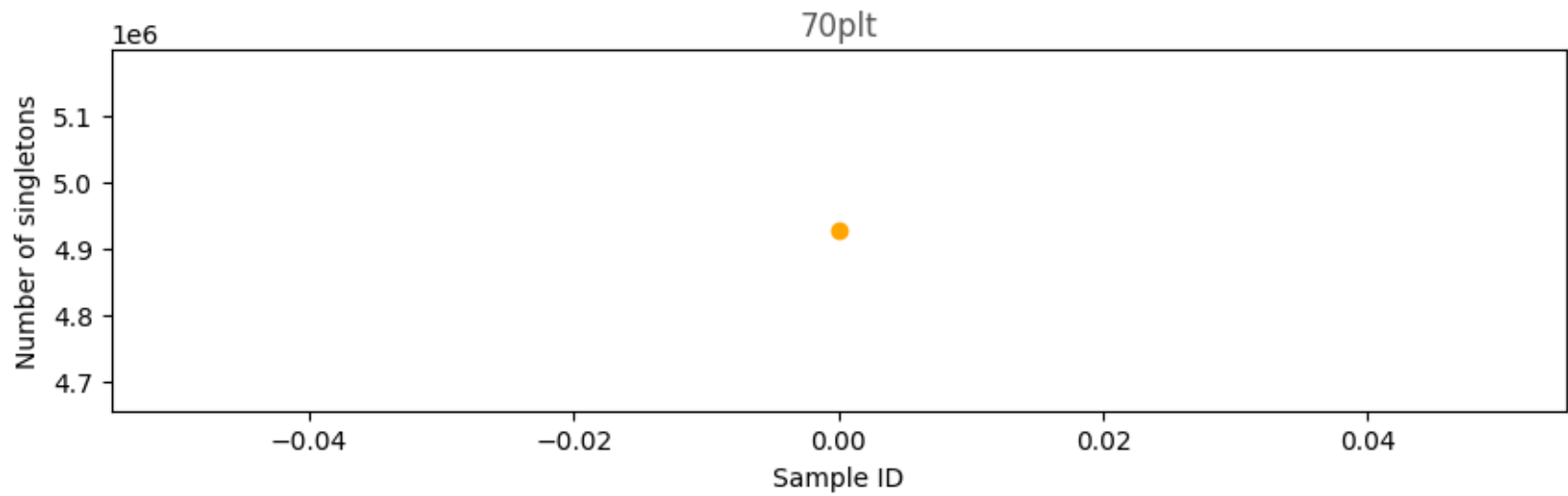
Ts/Tv by sample



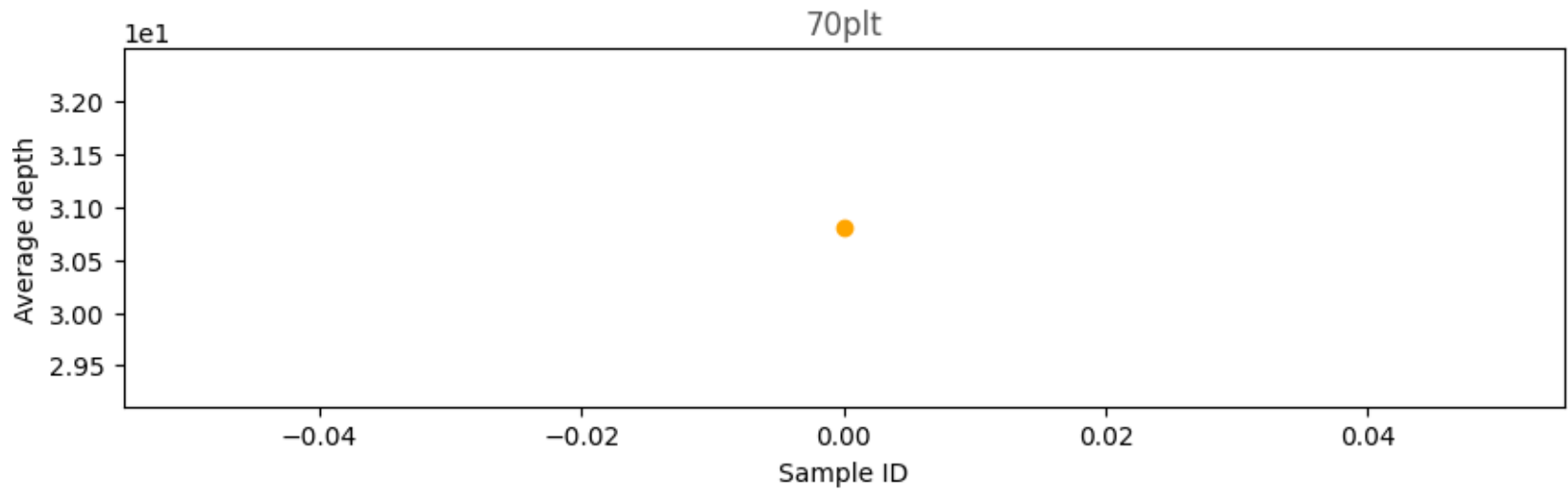
Hets vs non-ref Homs by sample



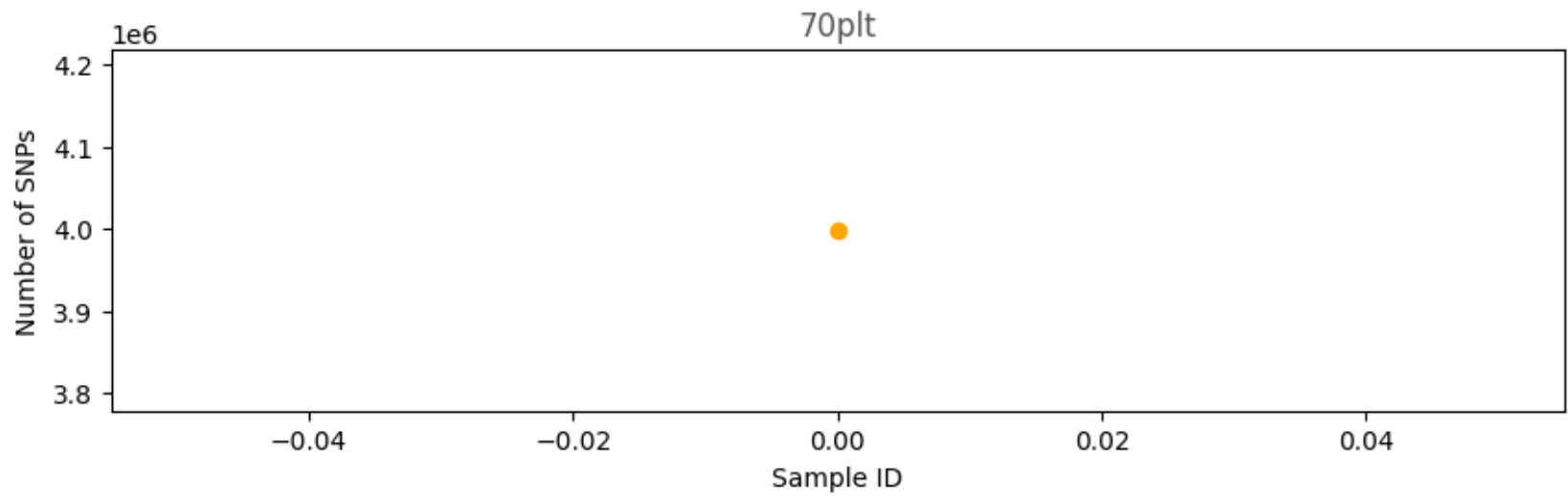
Singletons by sample (hets and homs)



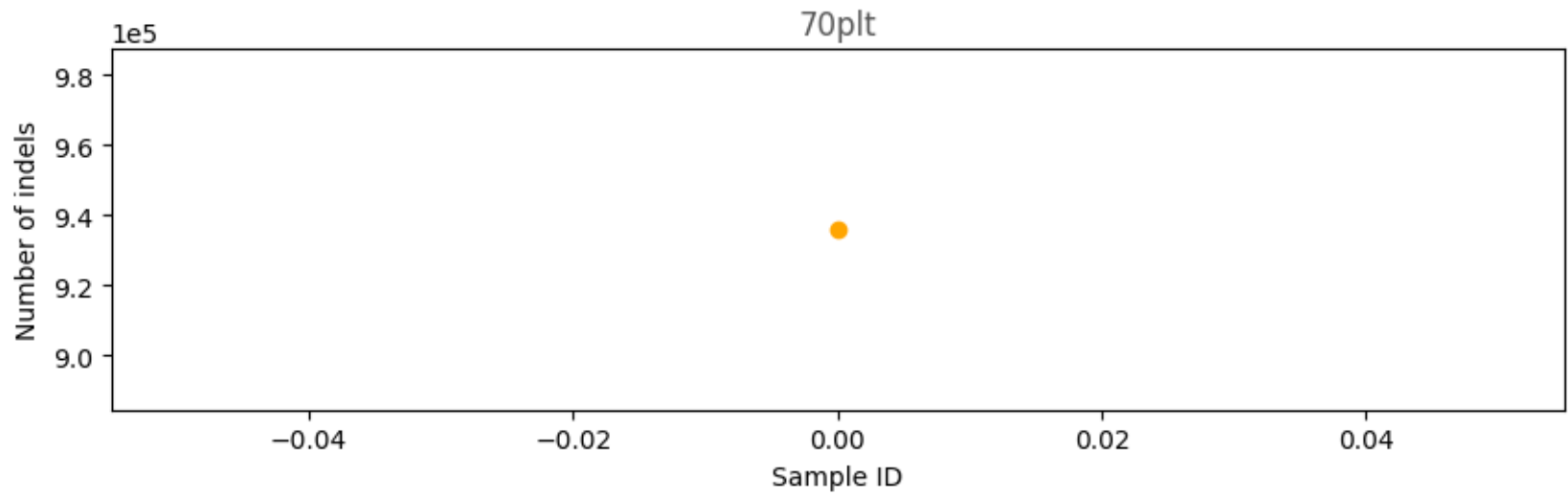
Average depth by sample



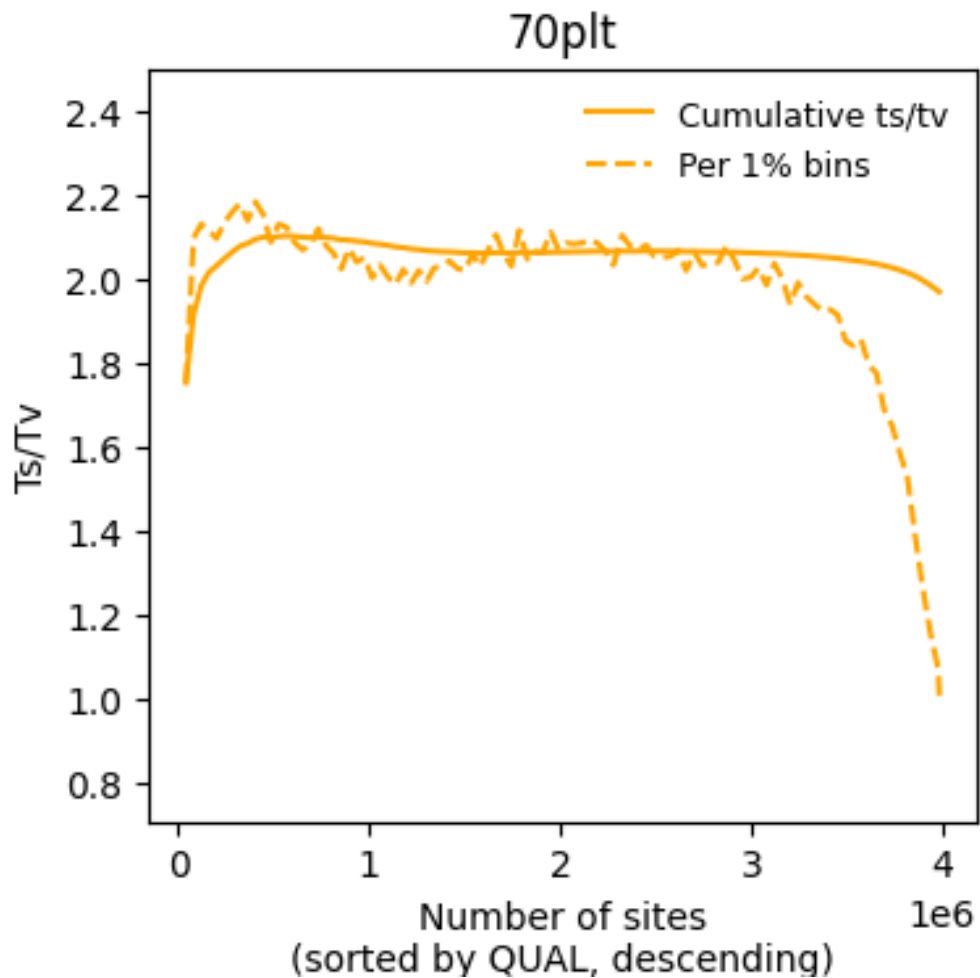
Number of SNPs by sample



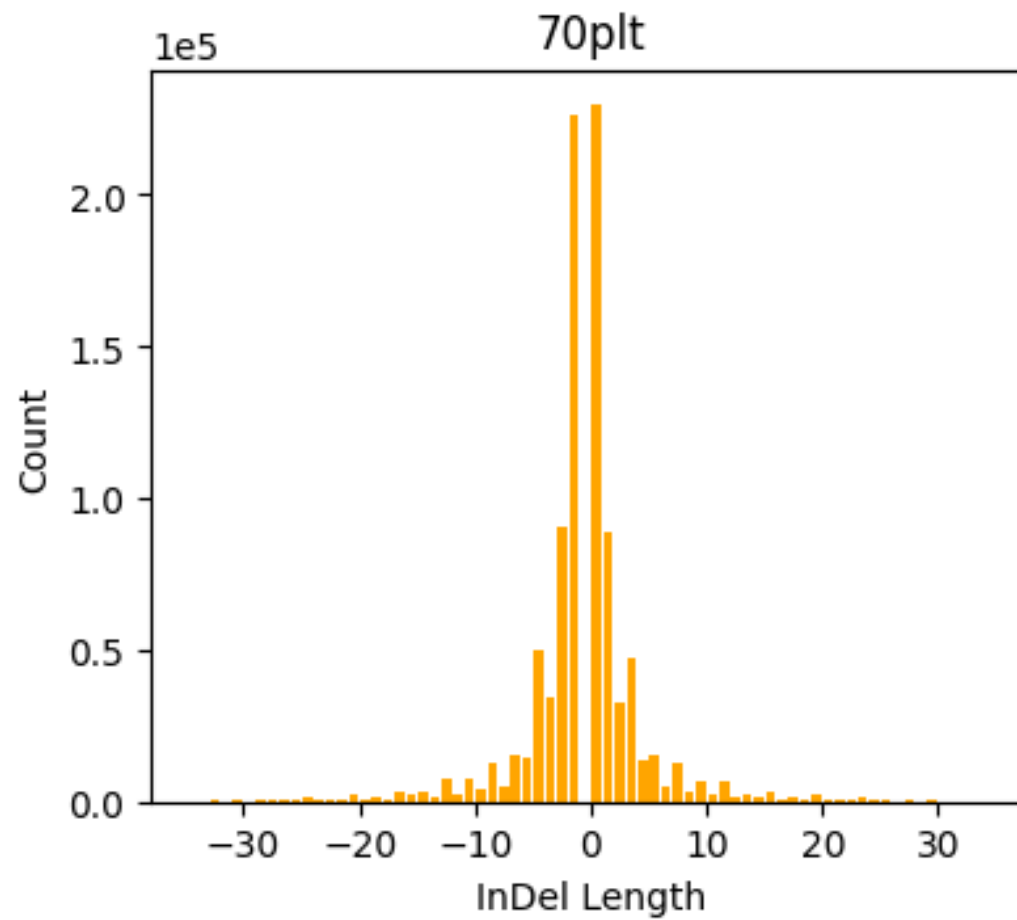
Number of indels by sample



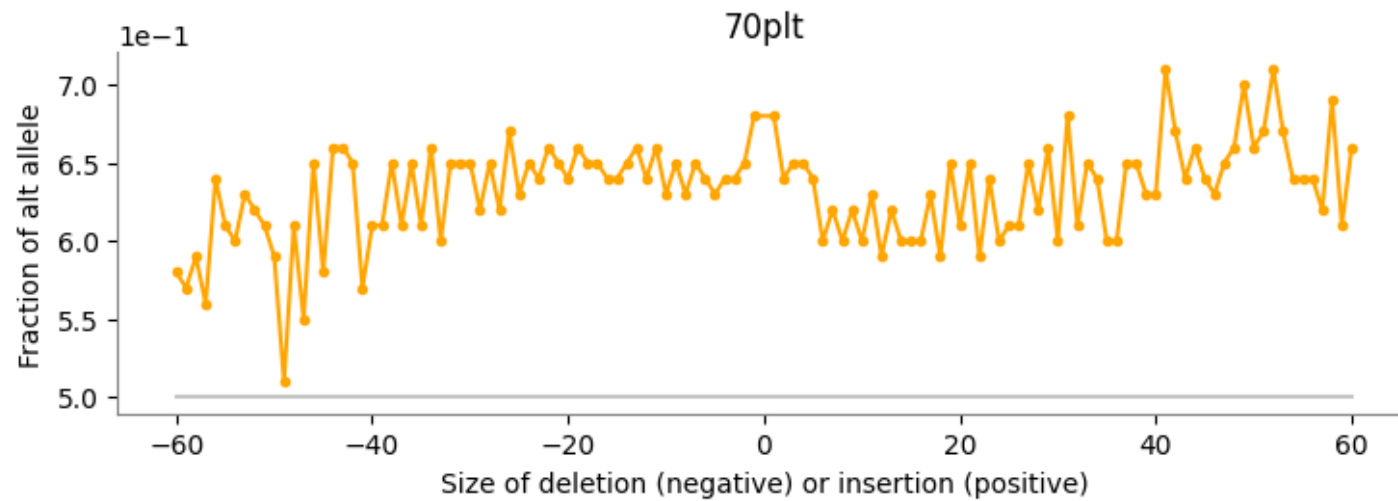
Ts/Tv stratified by QUAL



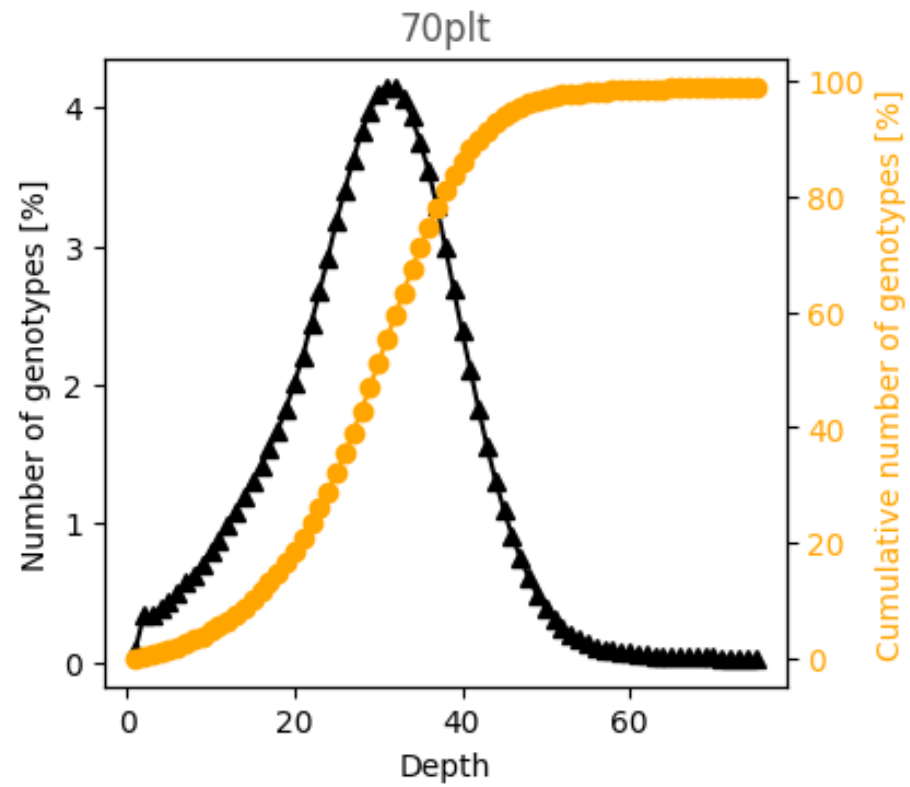
Indel distribution



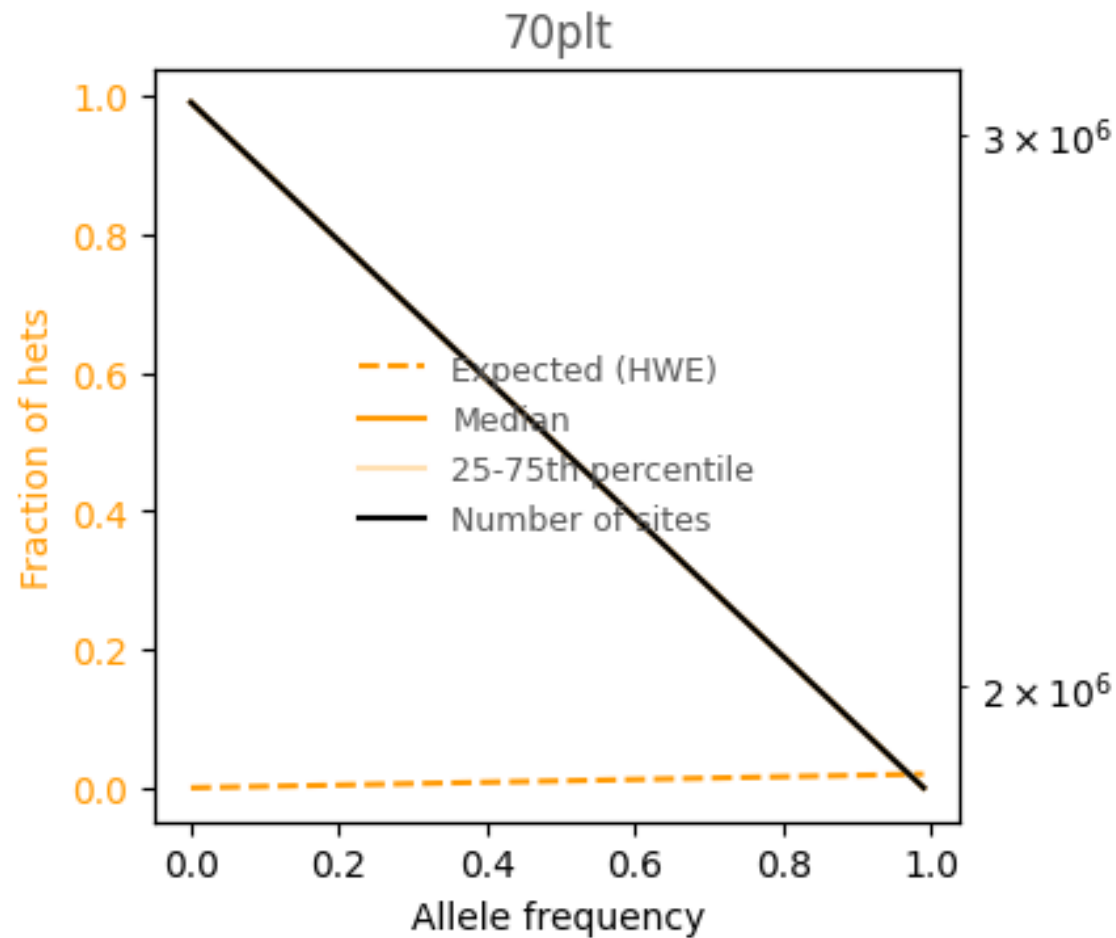
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

