

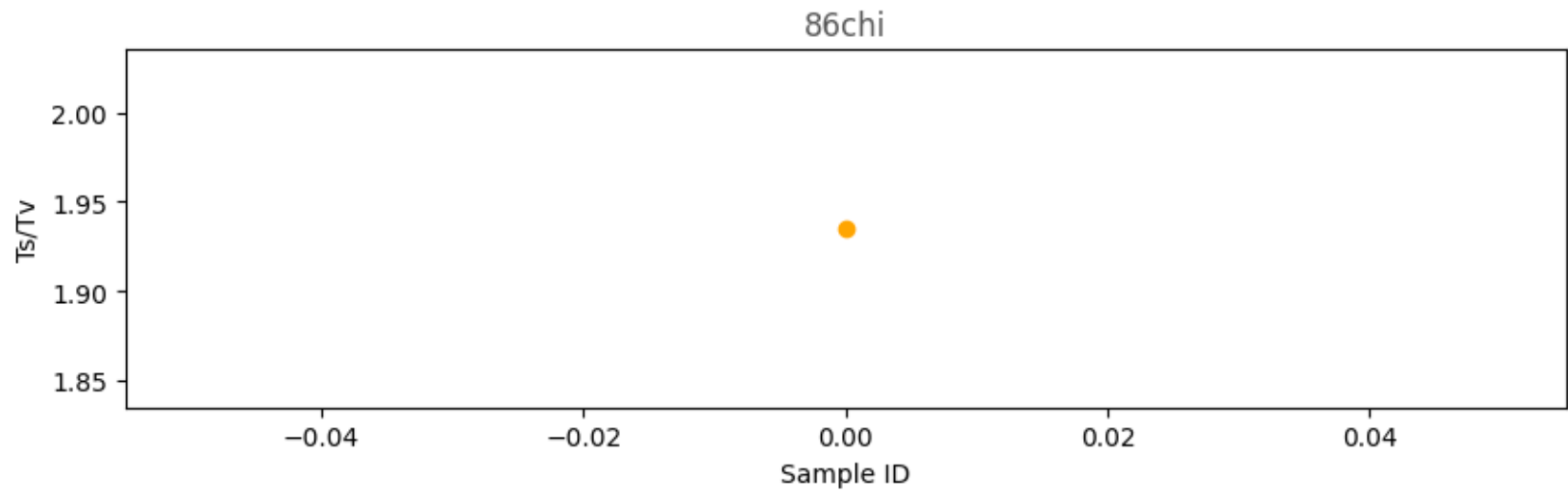
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
86chi	4,107,183	1.93	1.94	951,053	–	0	0
* frameshift ratio: out/(out+in)							

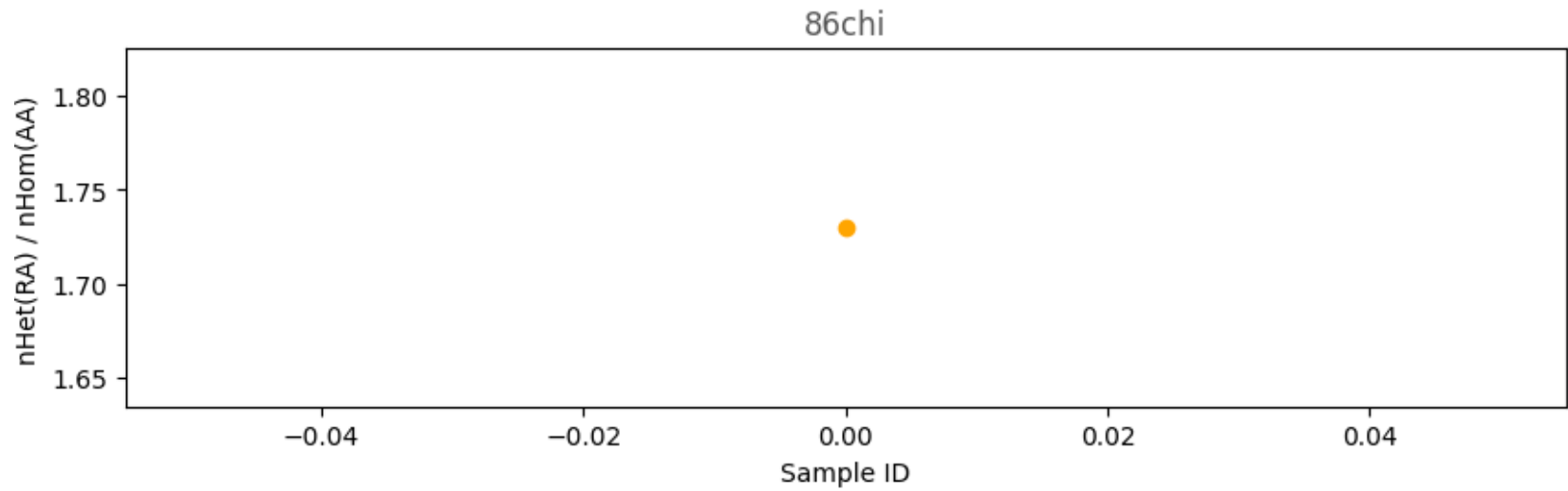
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
86chi	63.4%	1.91	68.3%	100,494	2,087

- 86chi .. /ngc/projects2/gm/data/archive/2022/variants/snv/86chiorif-110307304825-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-221116_A00559_BHF53WDSX5-RHGM_LABKA_WGSA
 KUT-WGSAKUT06229_22RKG027198x01_snv_germline_raw.haplotype_caller.vcf.gz

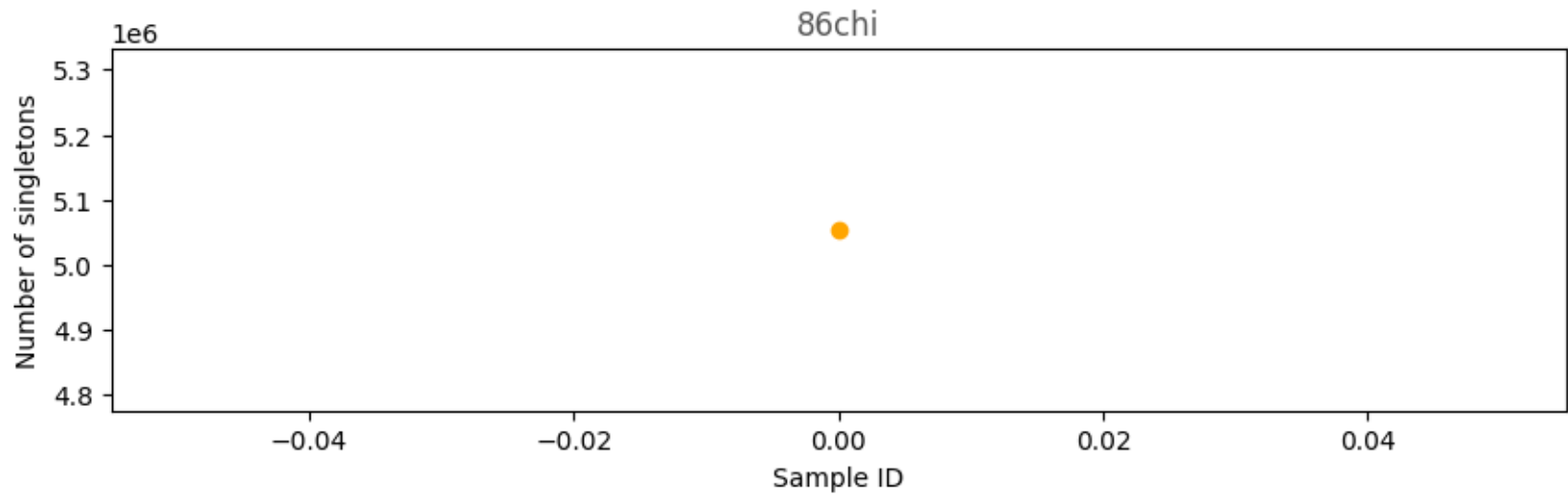
Ts/Tv by sample



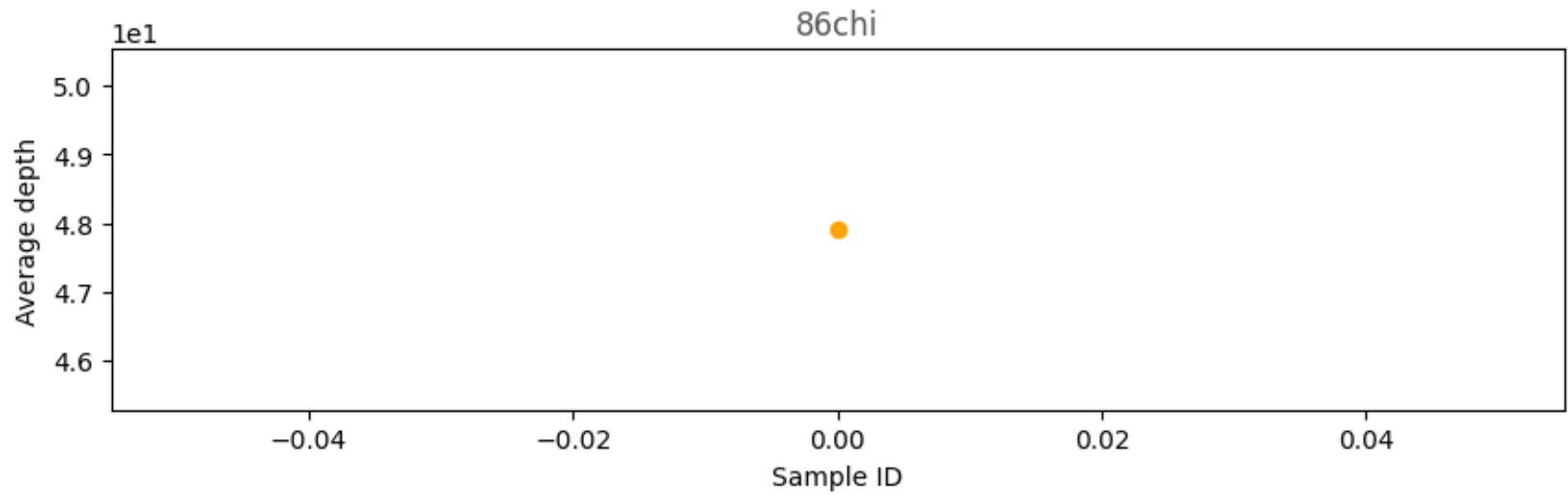
Hets vs non-ref Homs by sample



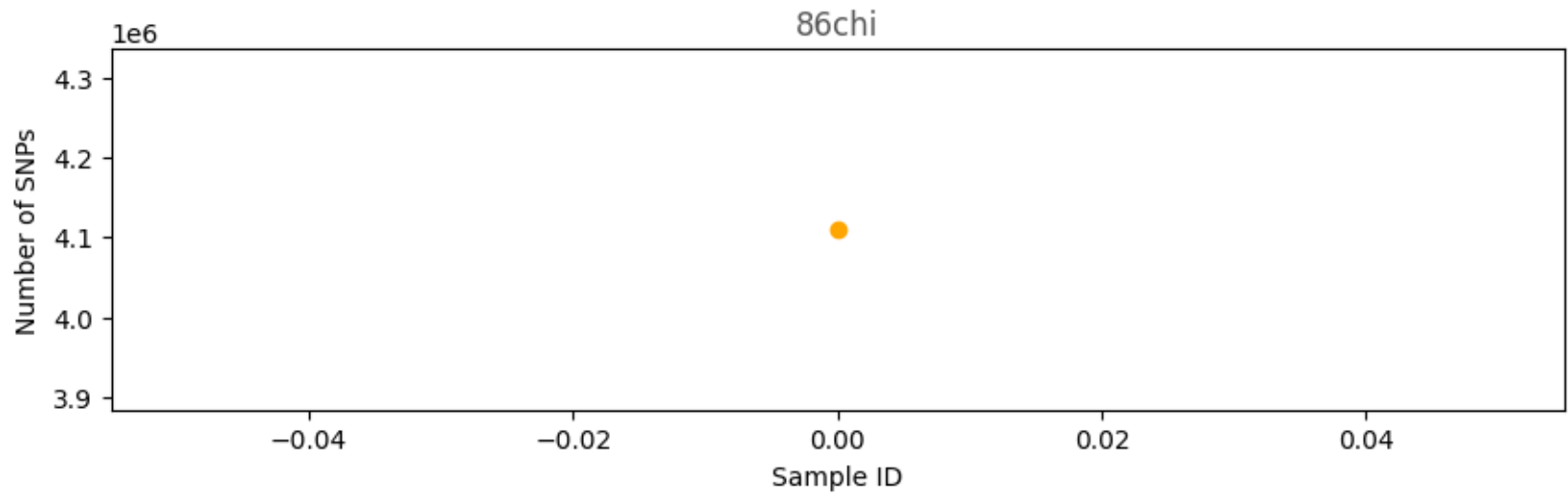
Singletons by sample (hets and homs)



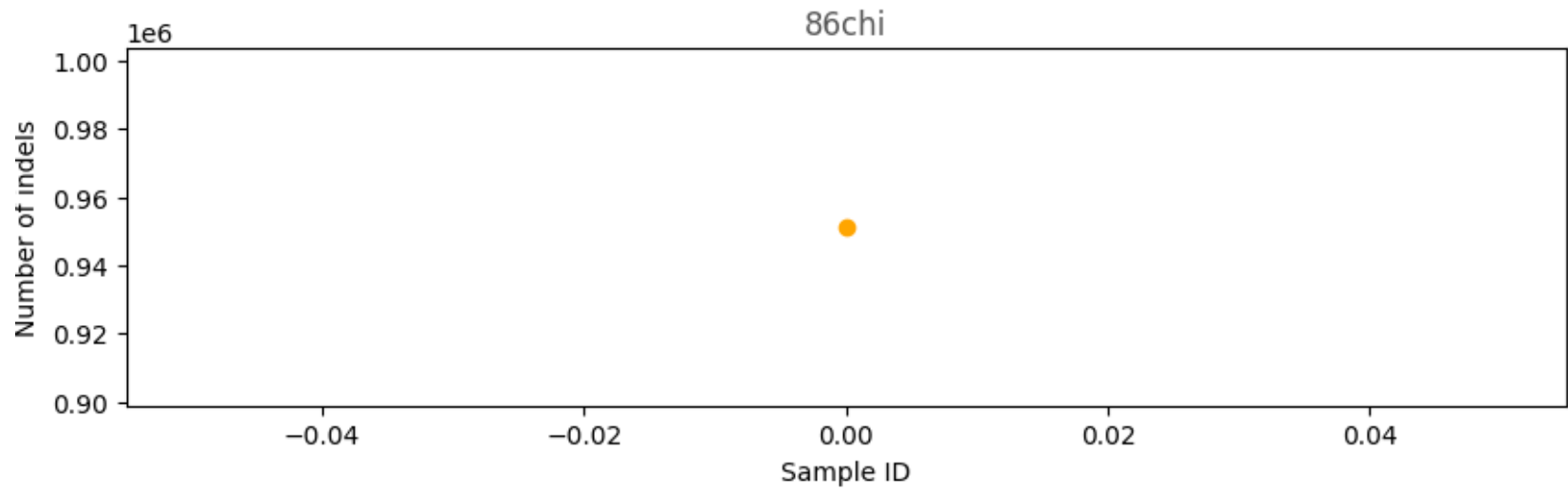
Average depth by sample



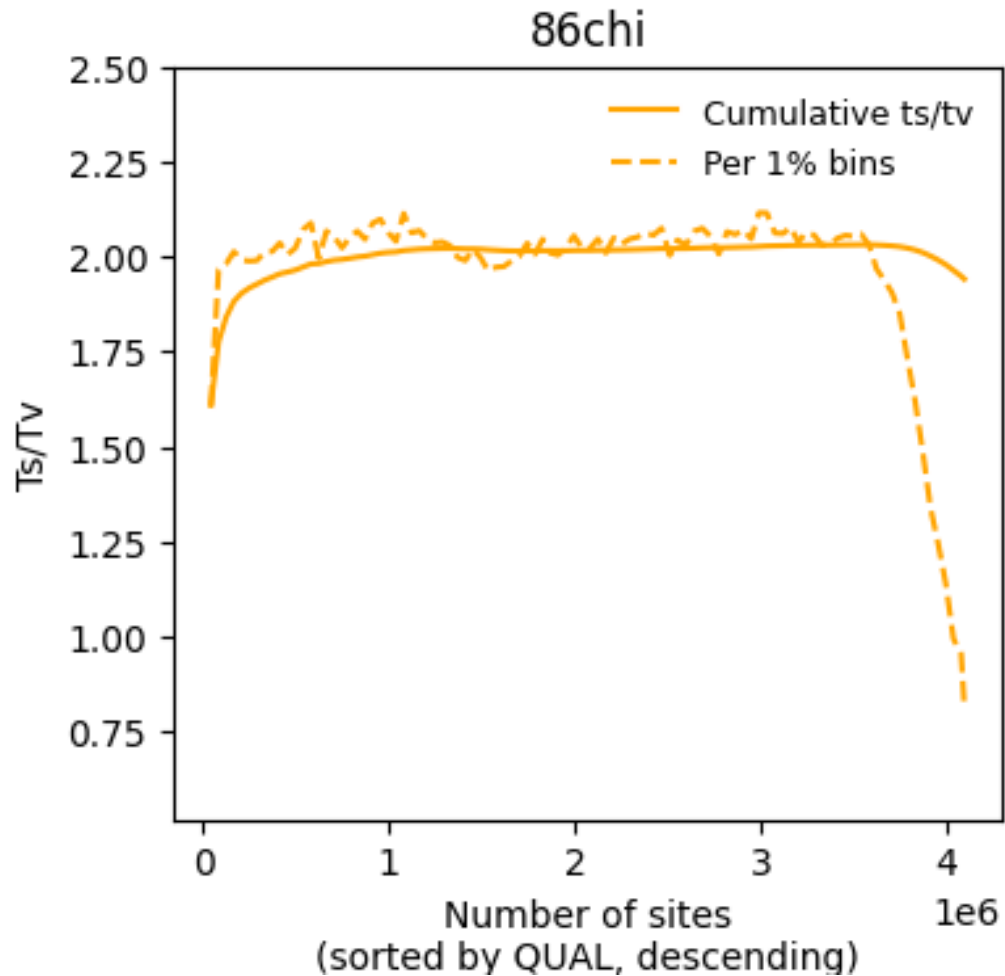
Number of SNPs by sample



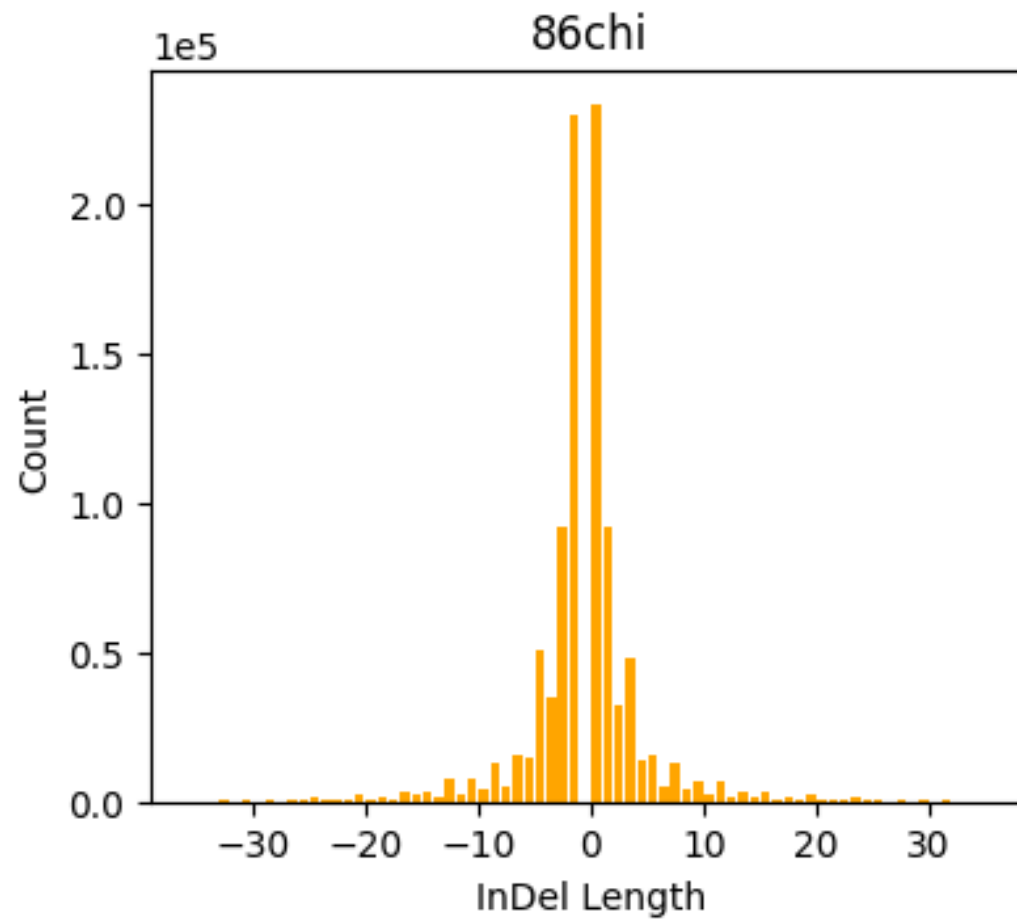
Number of indels by sample



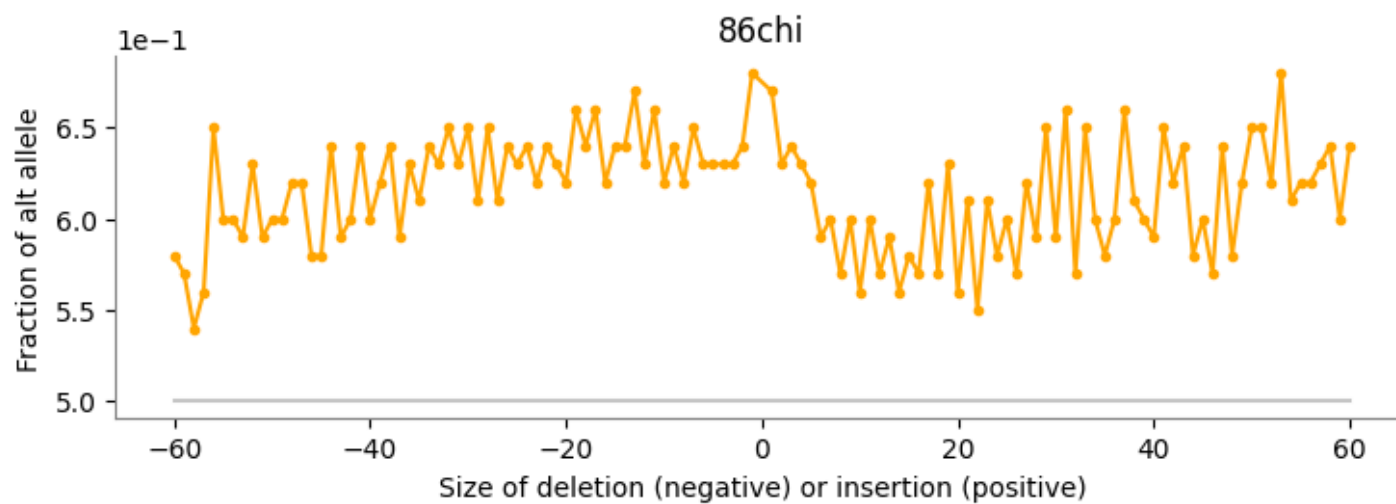
Ts/Tv stratified by QUAL



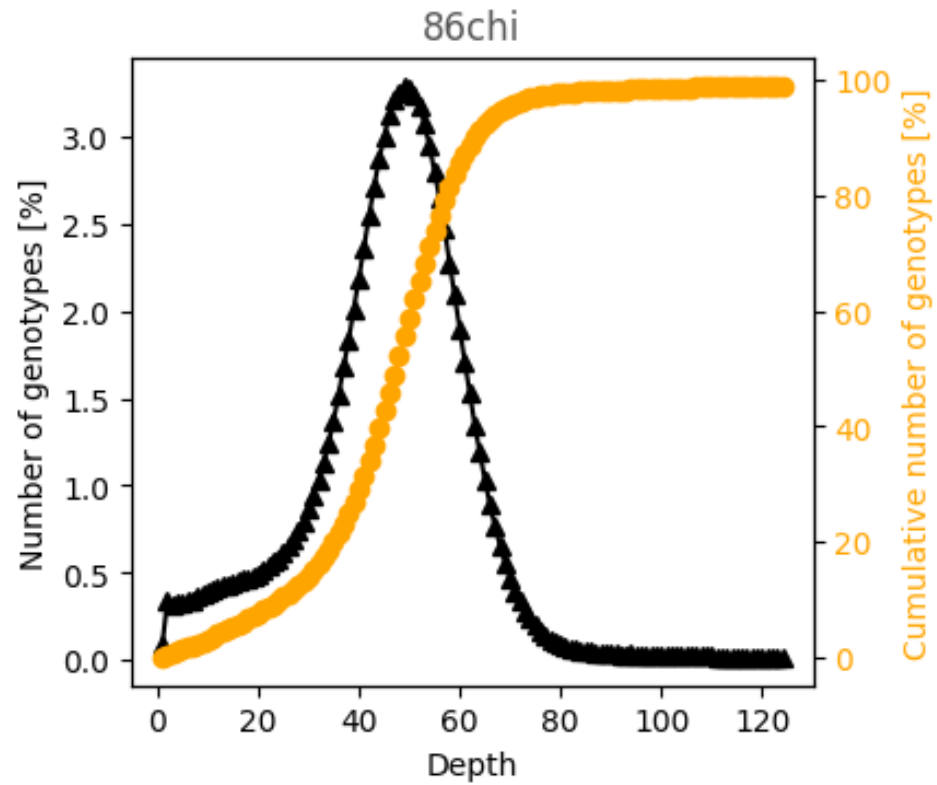
Indel distribution



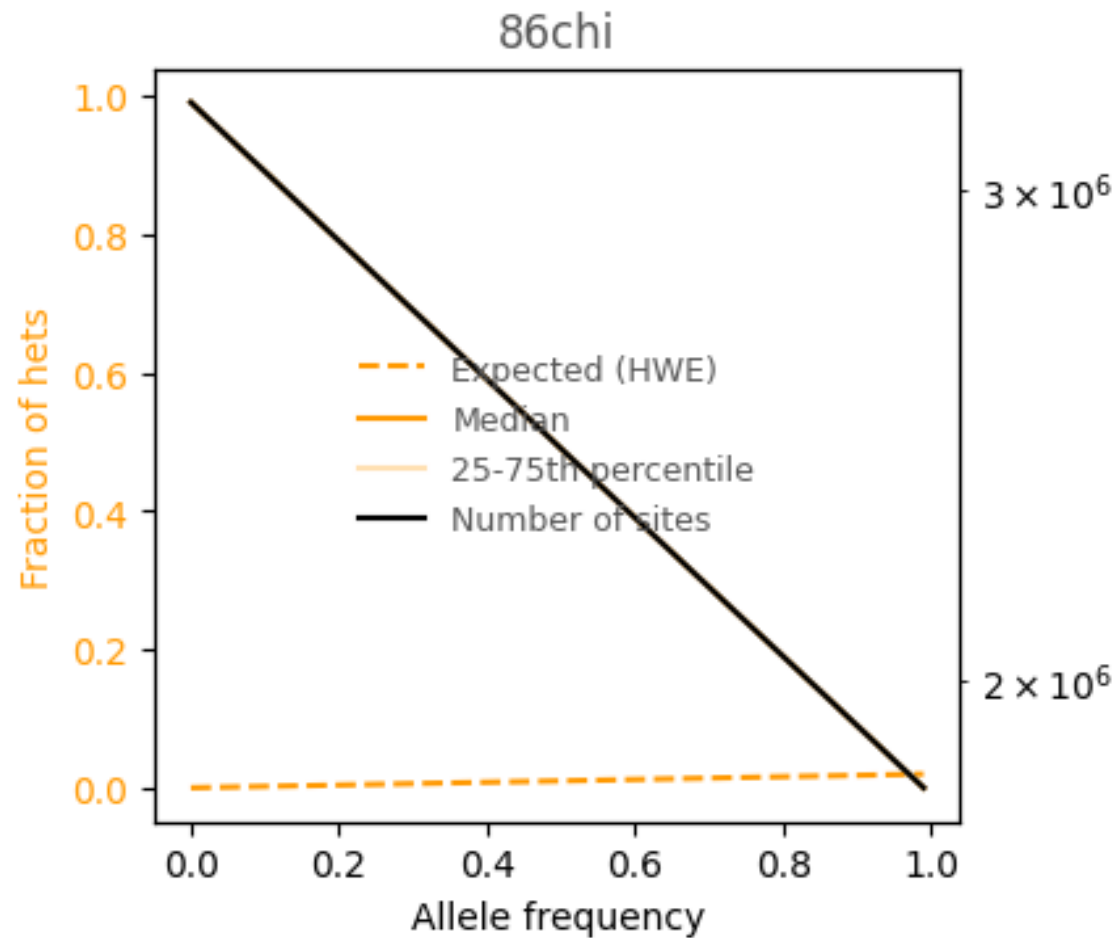
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

