

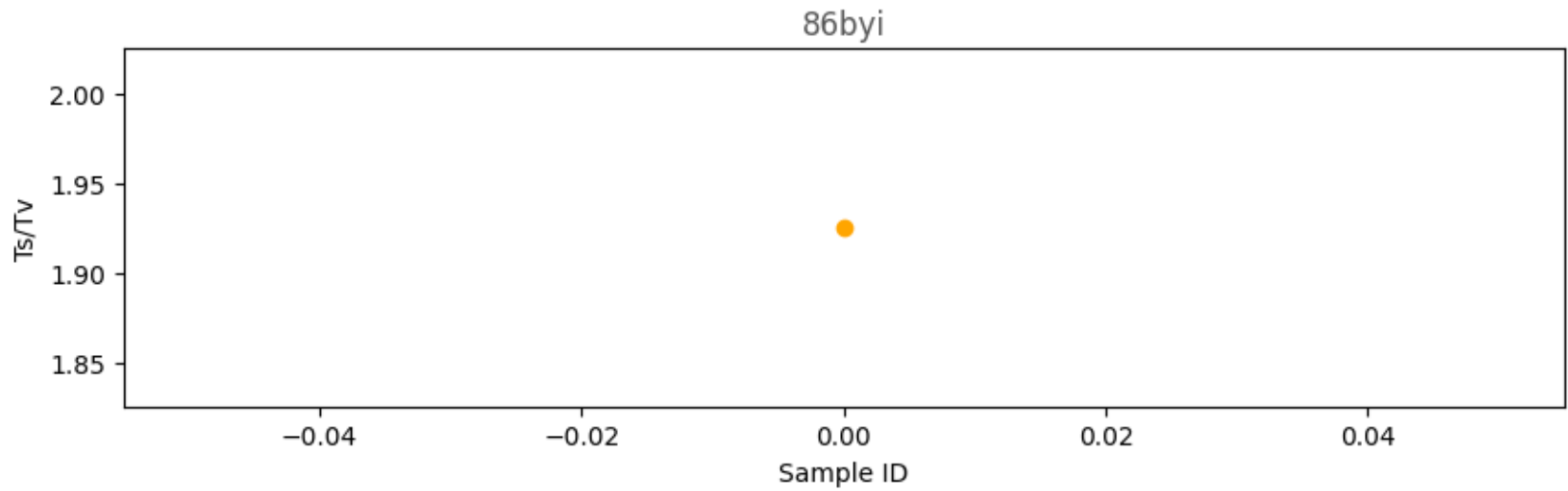
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
86byi	4,094,620	1.92	1.93	956,508	–	0	0
* frameshift ratio: out/(out+in)							

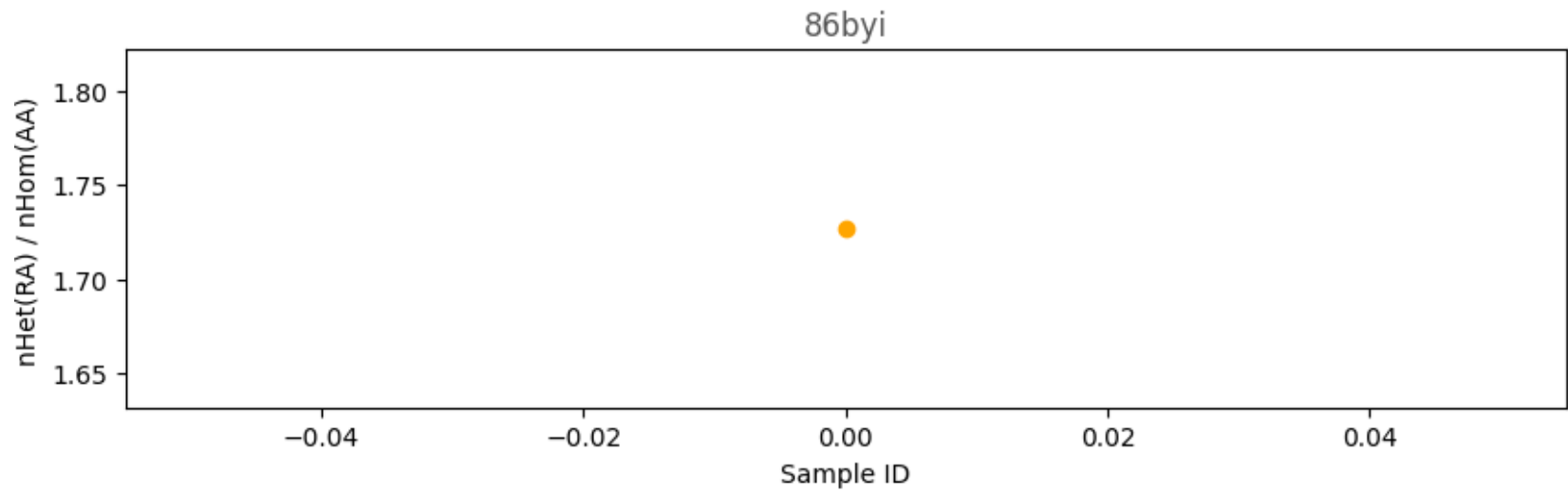
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
86byi	63.3%	1.89	68.5%	103,160	2,265

- 86byi .. /ngc/projects2/gm/data/archive/2022/variants/snv/86byinoif-110295486485-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-221012_A00559_BH2CMWDSX5-RHGM_LABKA_WGSA_KUT-WGSAKUT05804_22RKG023936x01_snv_germline_raw.haplotype_caller.vcf.gz

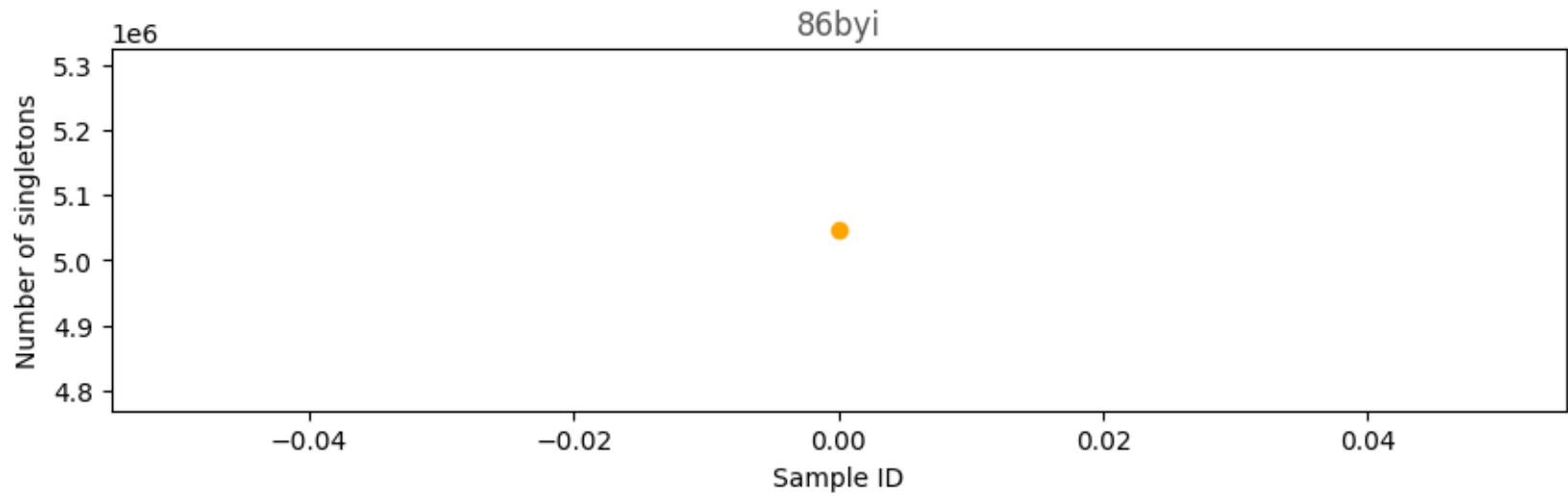
Ts/Tv by sample



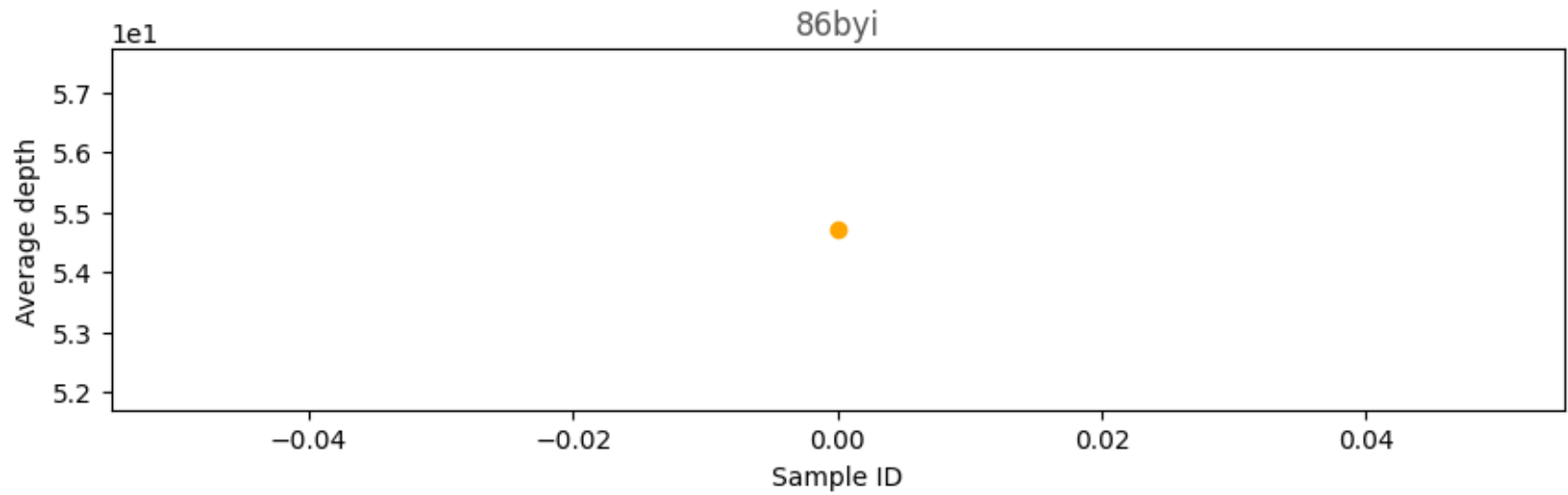
Hets vs non-ref Homs by sample



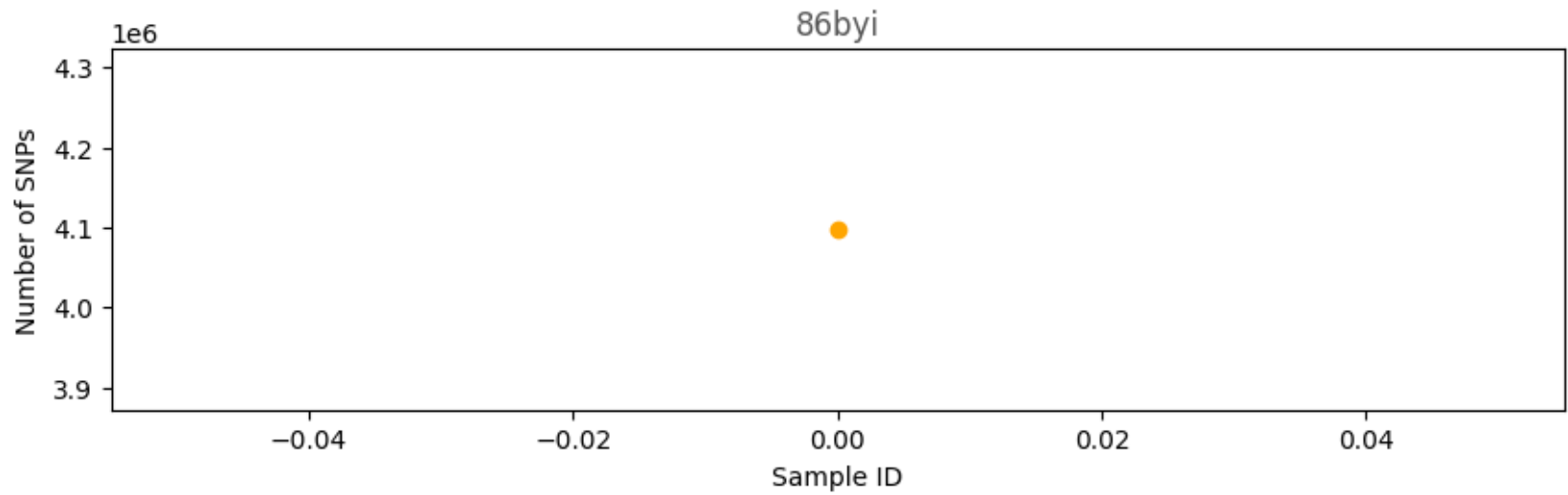
Singletons by sample (hets and homs)



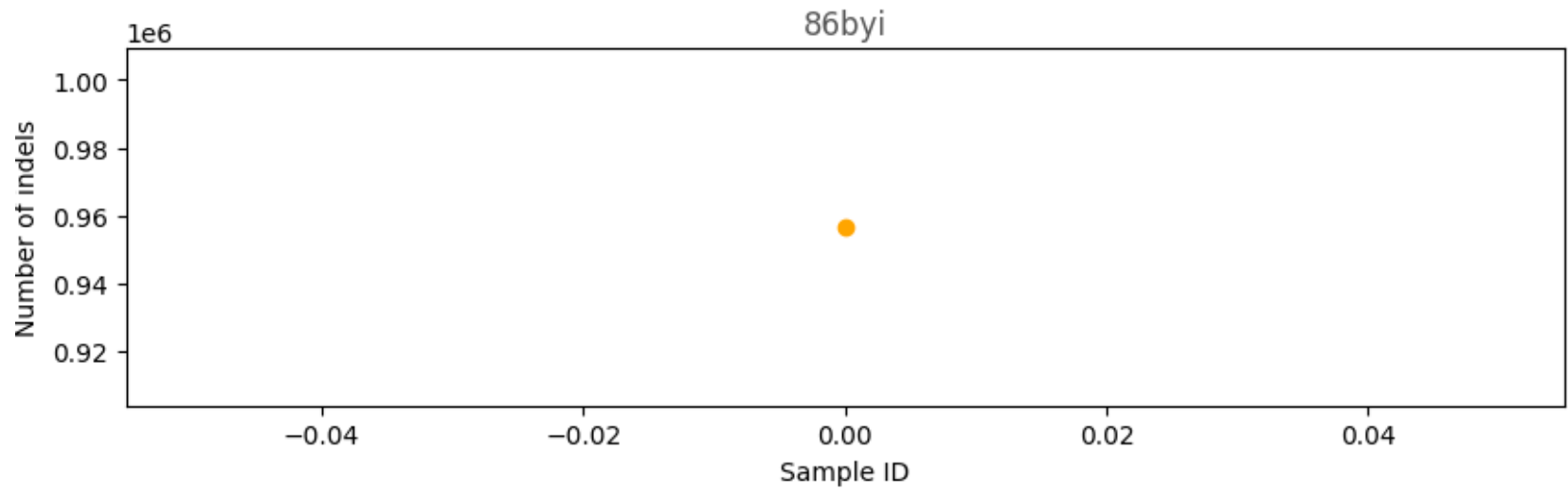
Average depth by sample



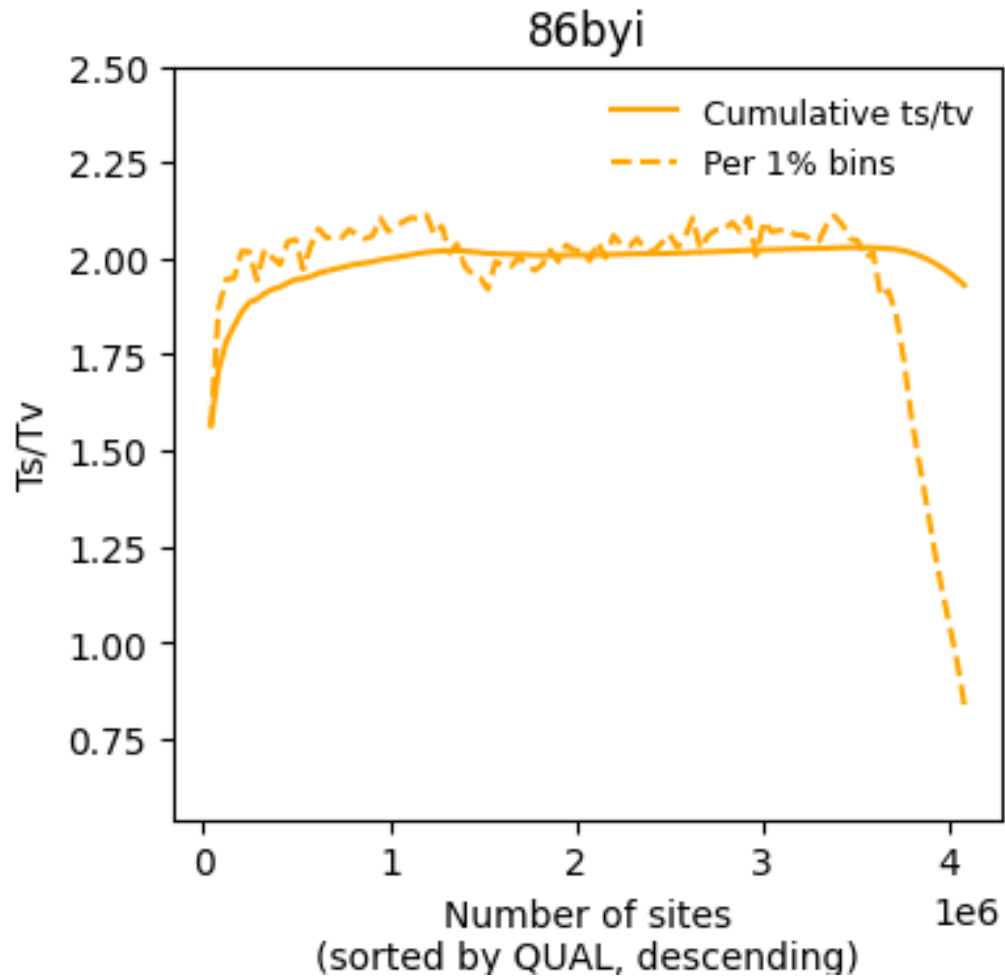
Number of SNPs by sample



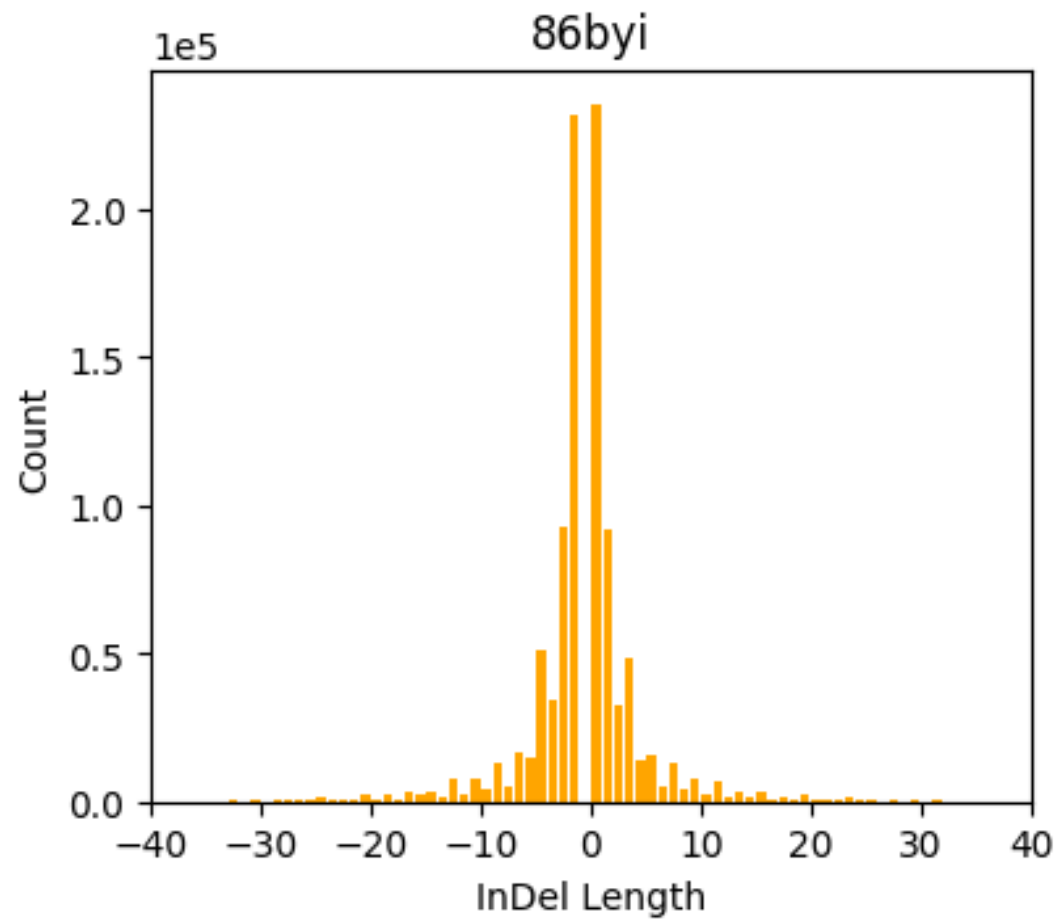
Number of indels by sample



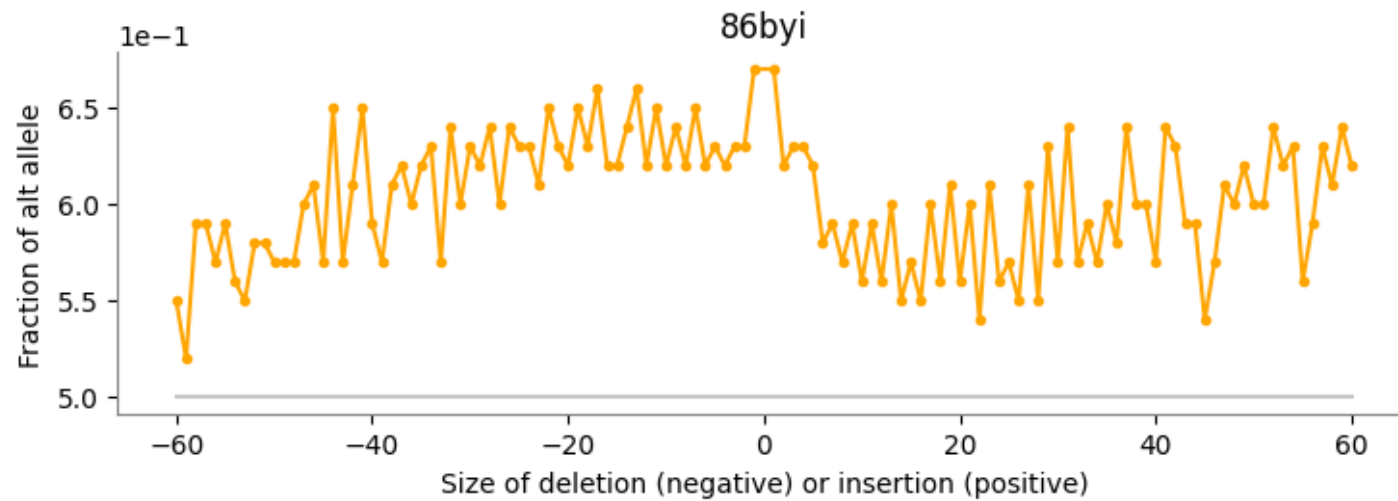
Ts/Tv stratified by QUAL



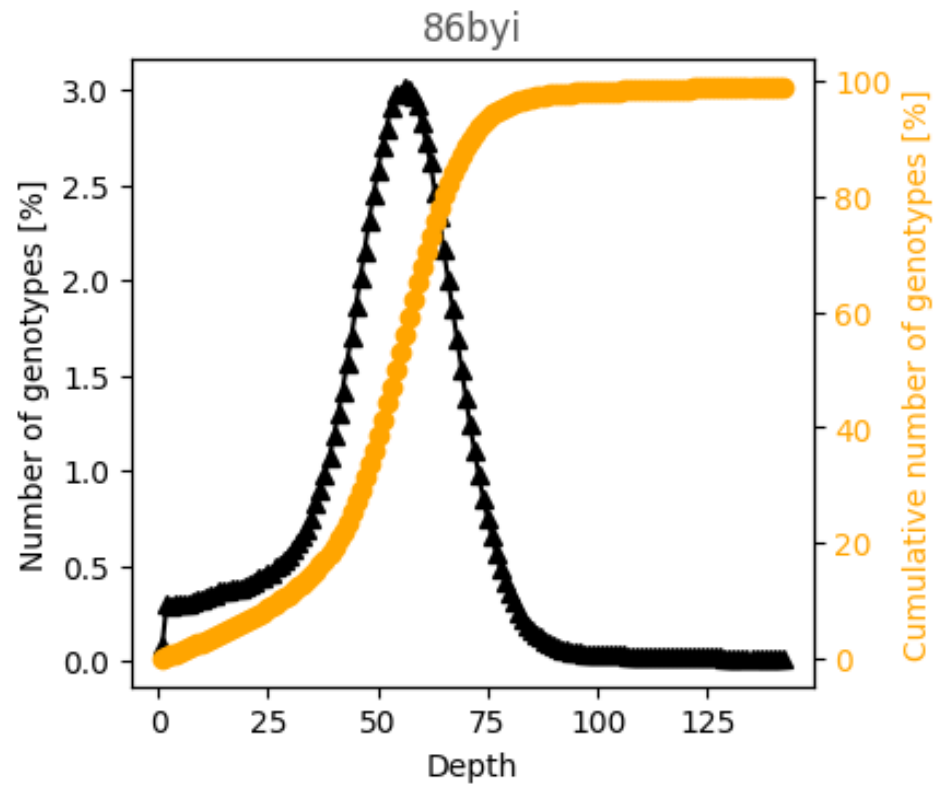
Indel distribution



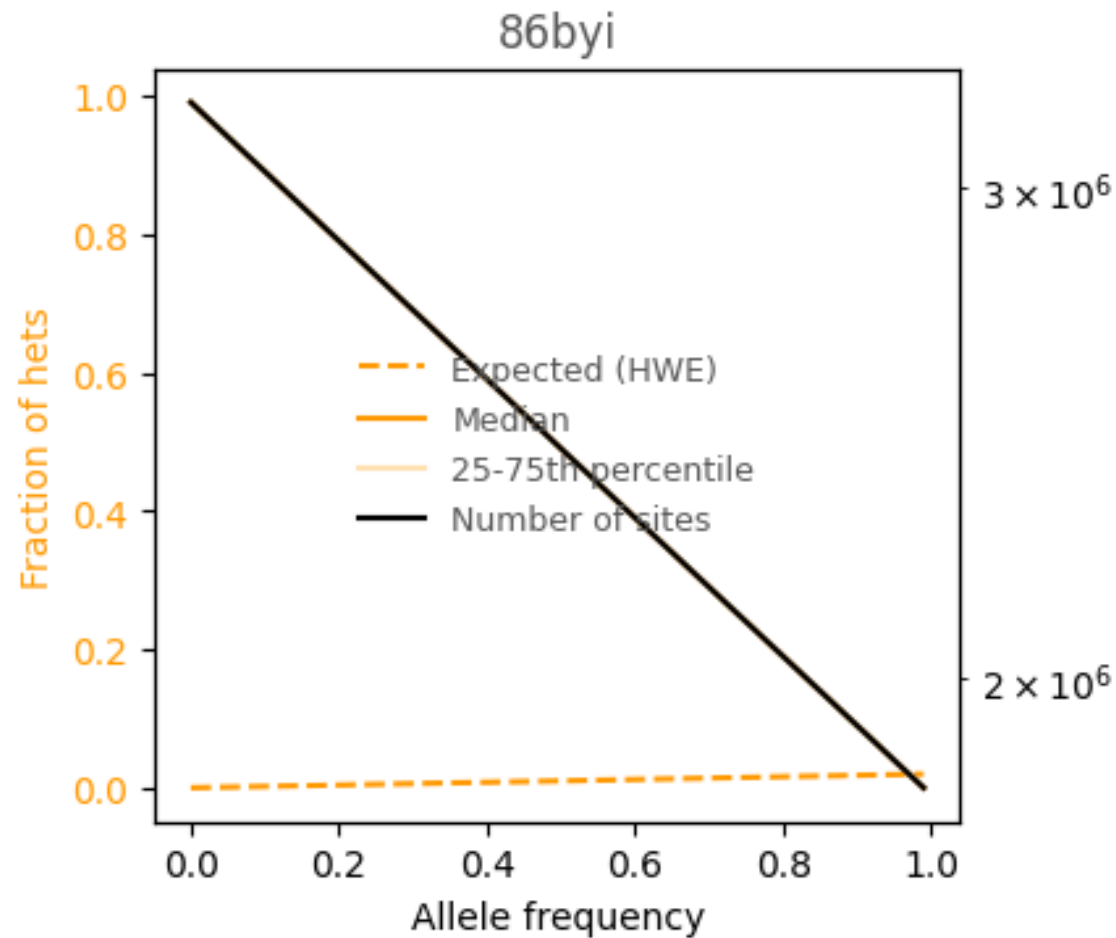
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

