

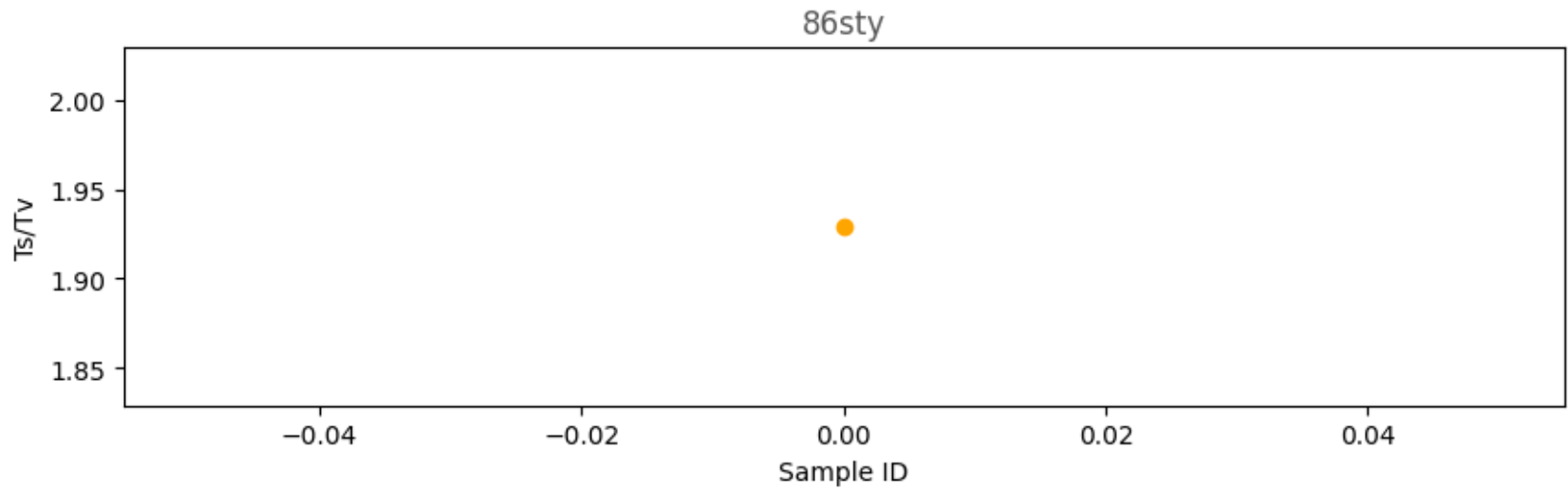
Summary Numbers

	SNPs			indels	MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	
86sty	4,065,020	1.93	1.93	941,977	–	0
* frameshift ratio: out/(out+in)						

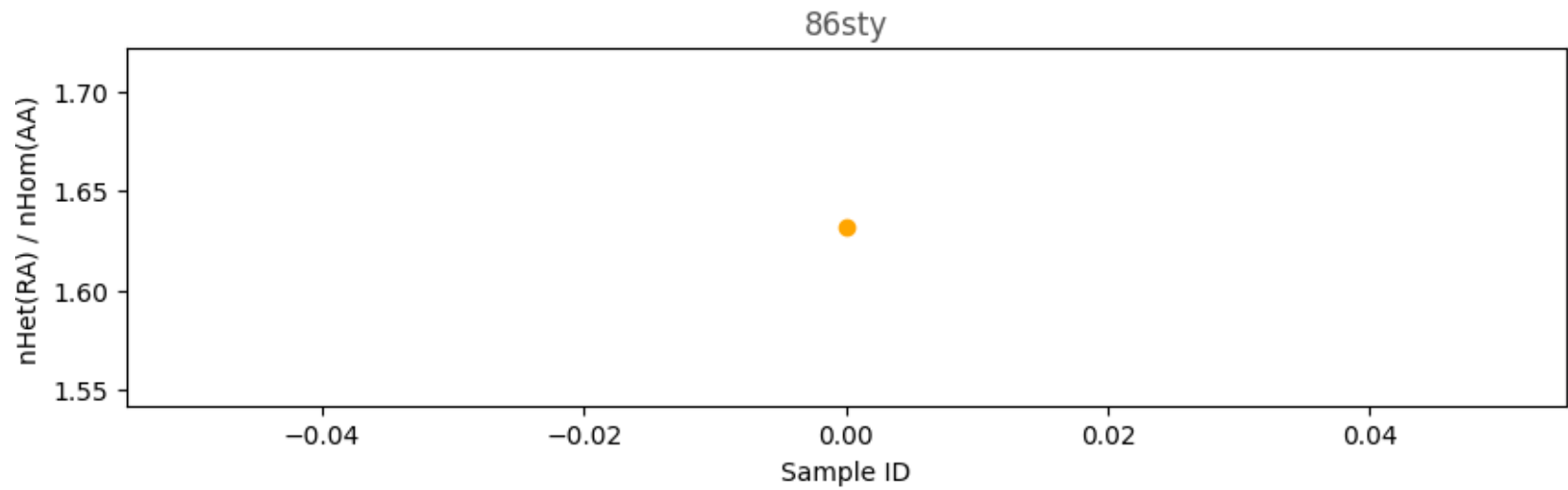
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
86sty	62.0%	1.90	66.6%	96,903	1,979

- 86sty .. /ngc/projects2/gm/data/archive/2022/variants/snv/86styehcm-103801038801-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211110_A01411_BHT75LDSX2-RHGM_LABKA_WGS-
 WGS03327_snv_germline_raw.haplotype_caller.vcf.gz

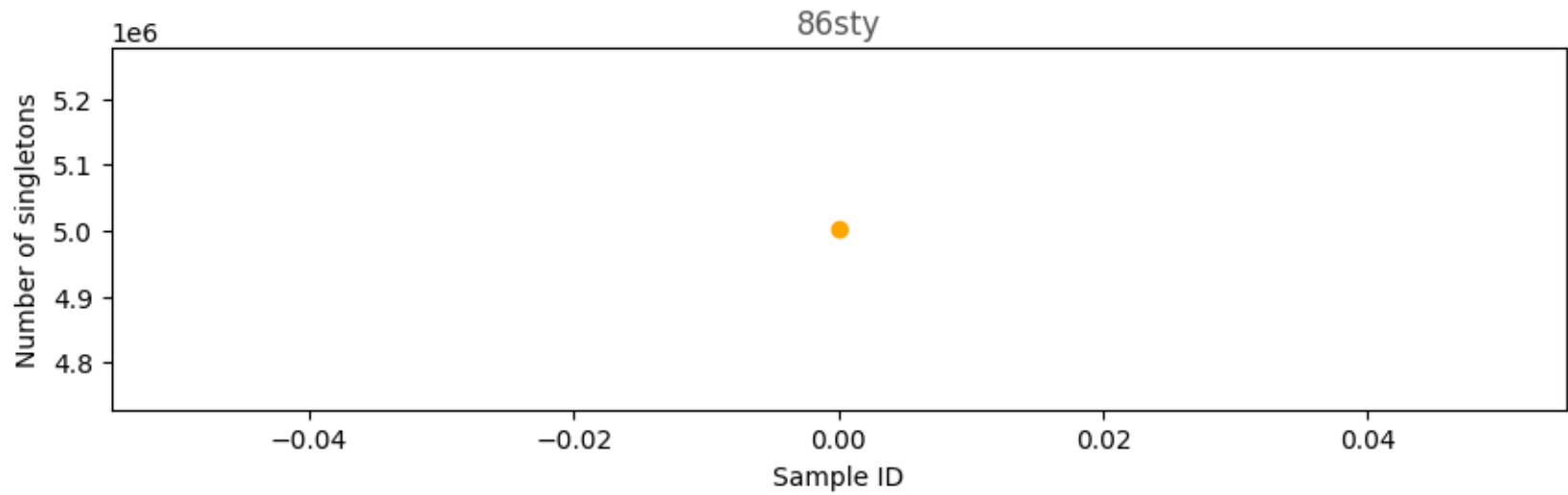
Ts/Tv by sample



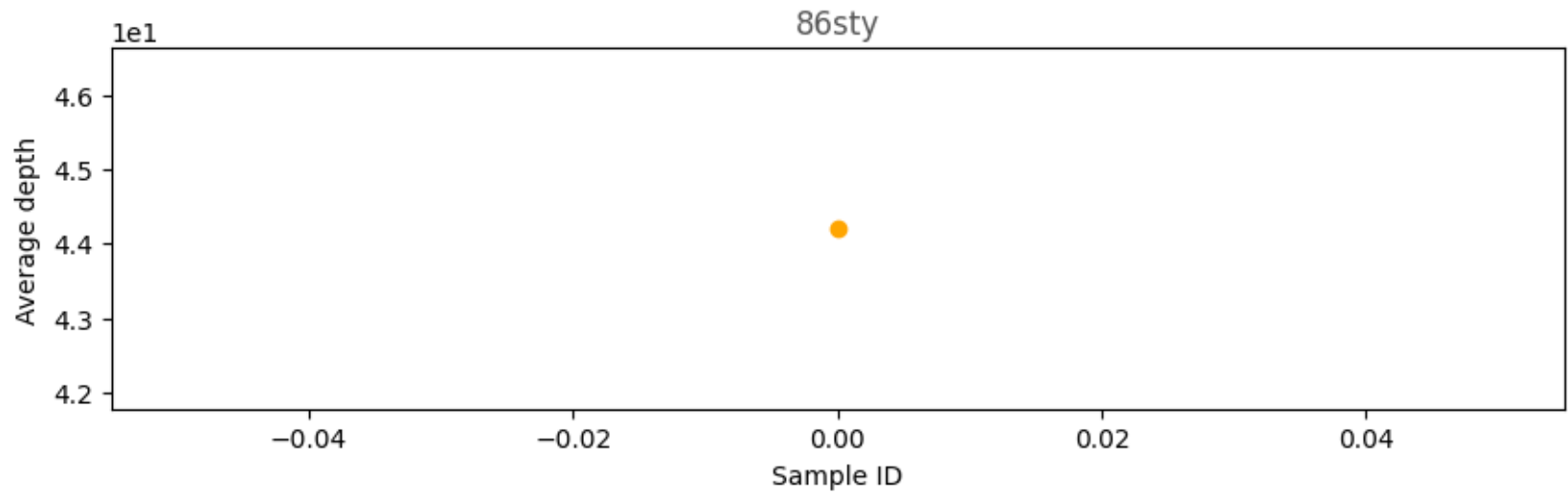
Hets vs non-ref Homs by sample



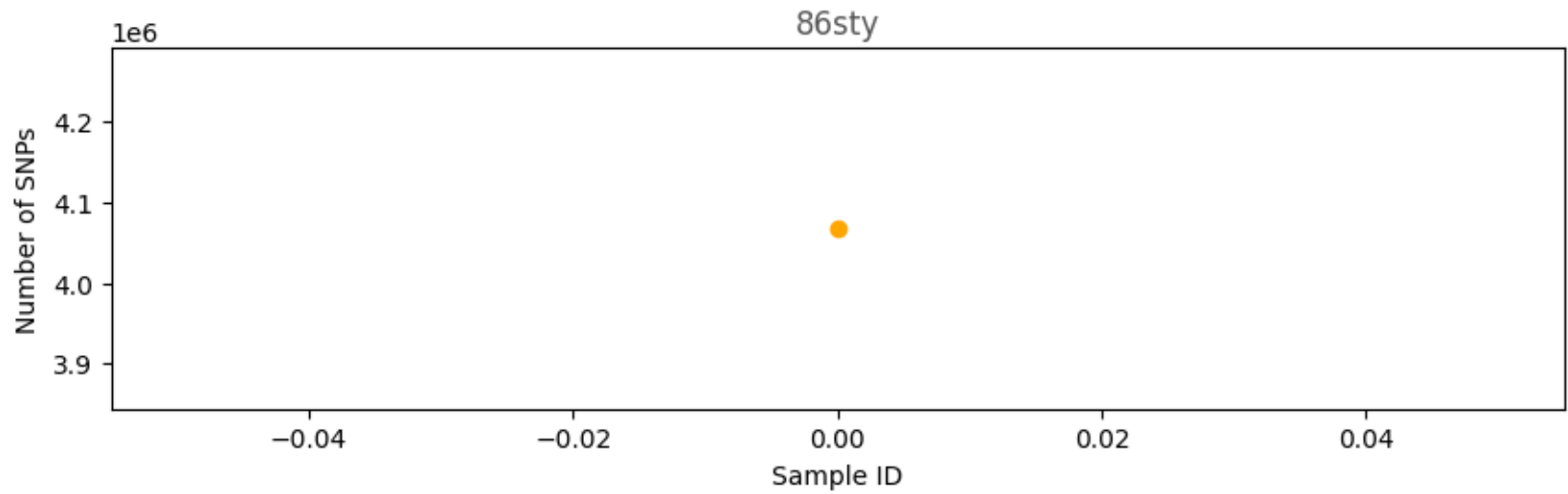
Singletons by sample (hets and homs)



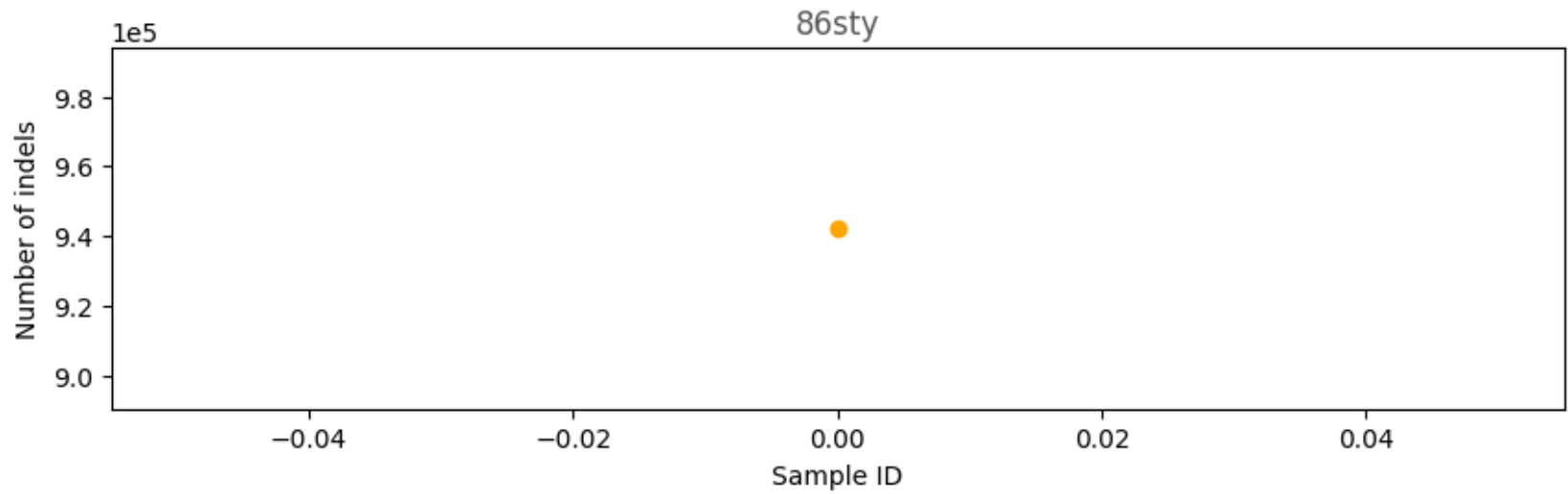
Average depth by sample



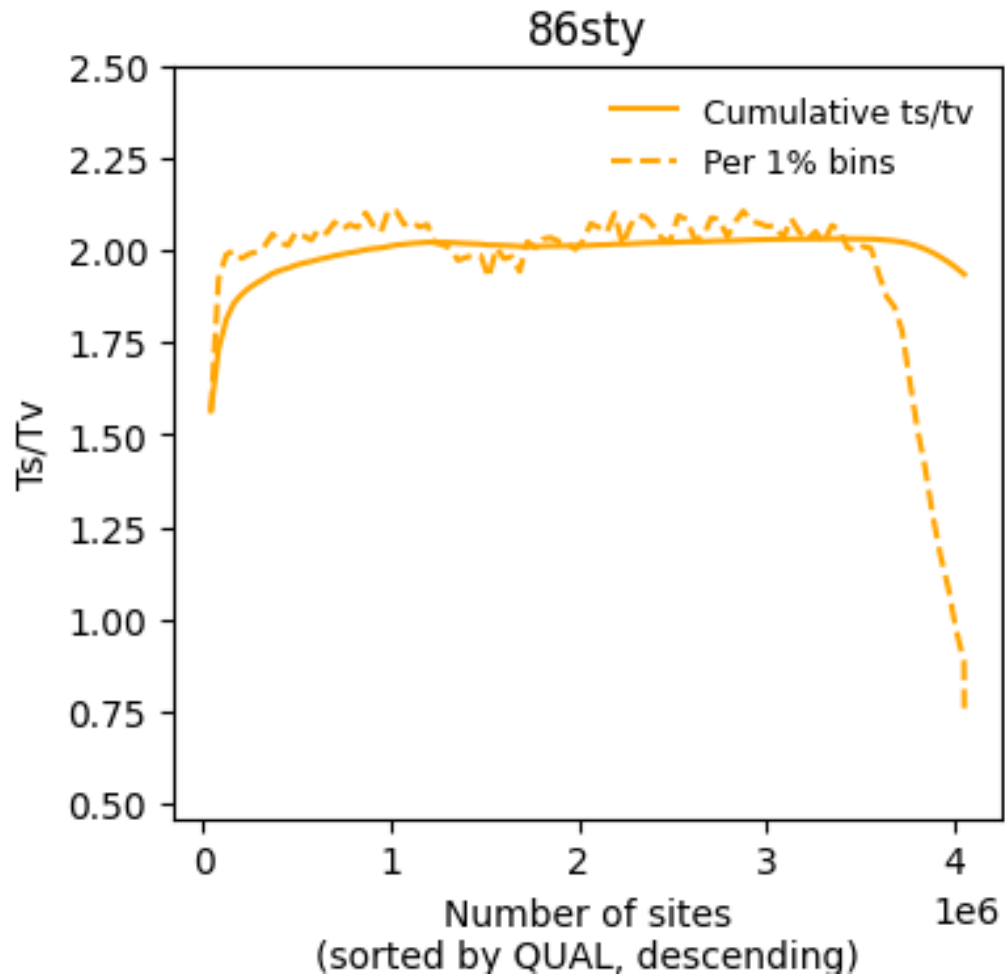
Number of SNPs by sample



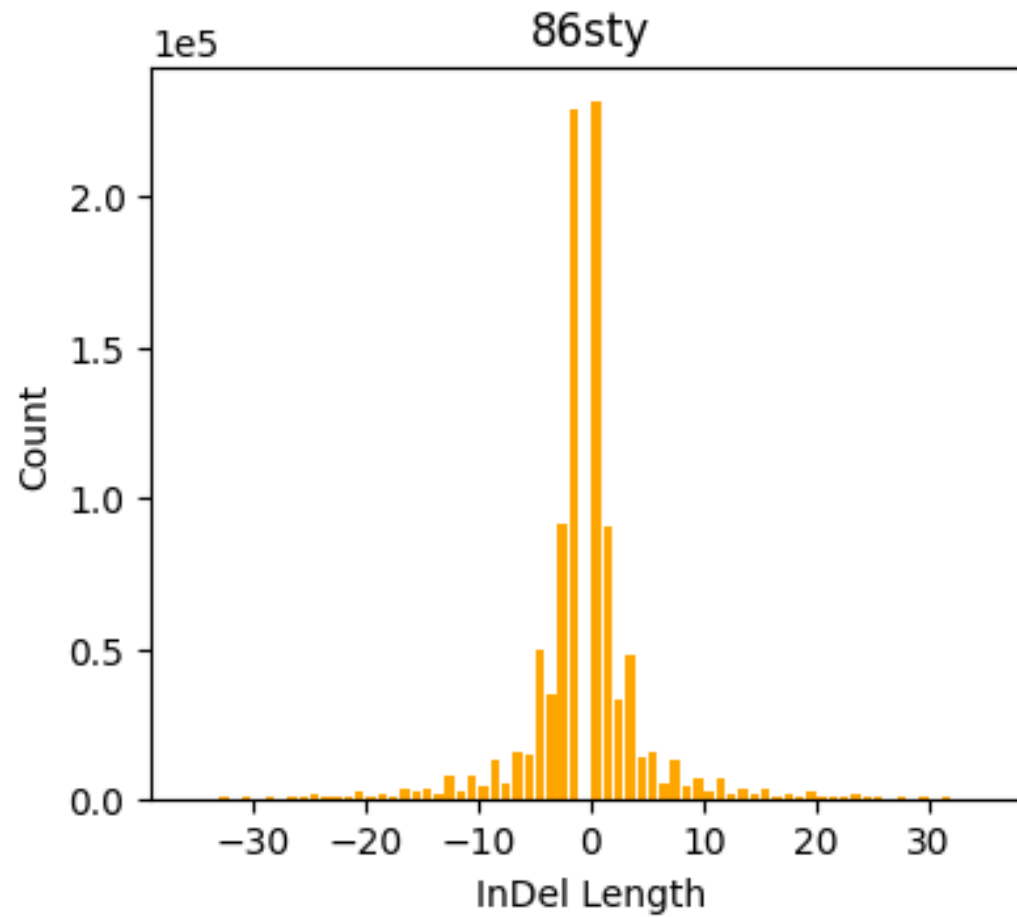
Number of indels by sample



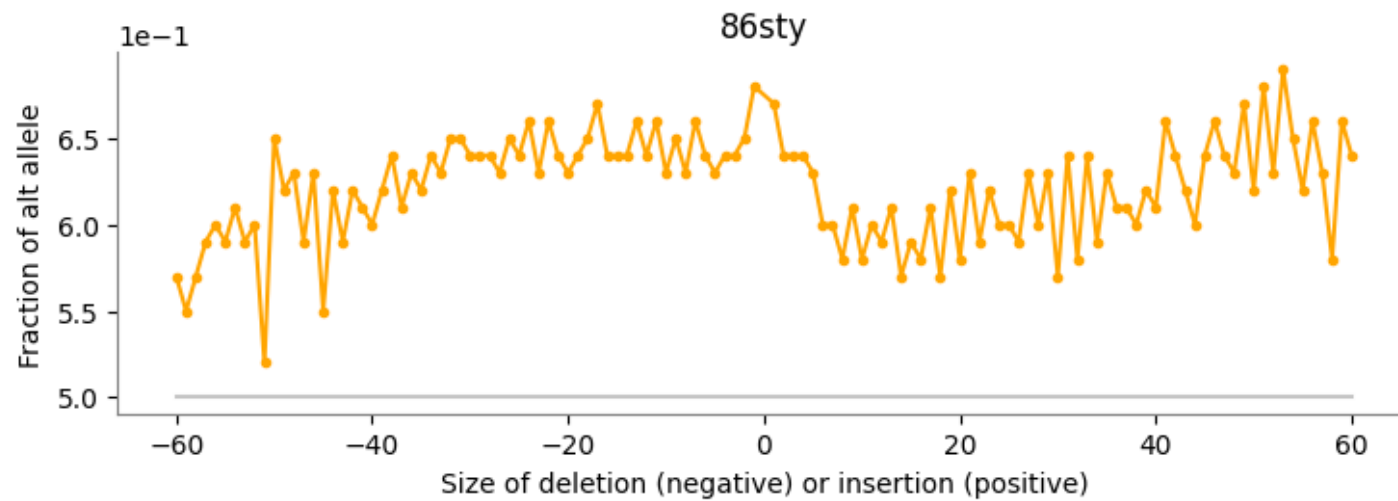
Ts/Tv stratified by QUAL



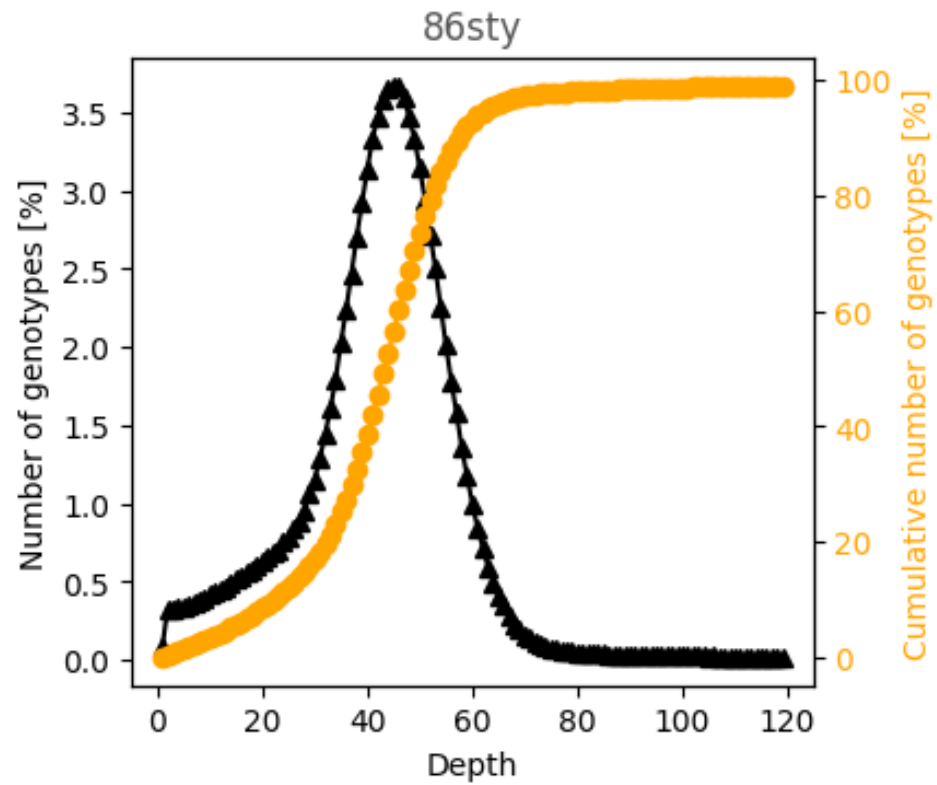
Indel distribution



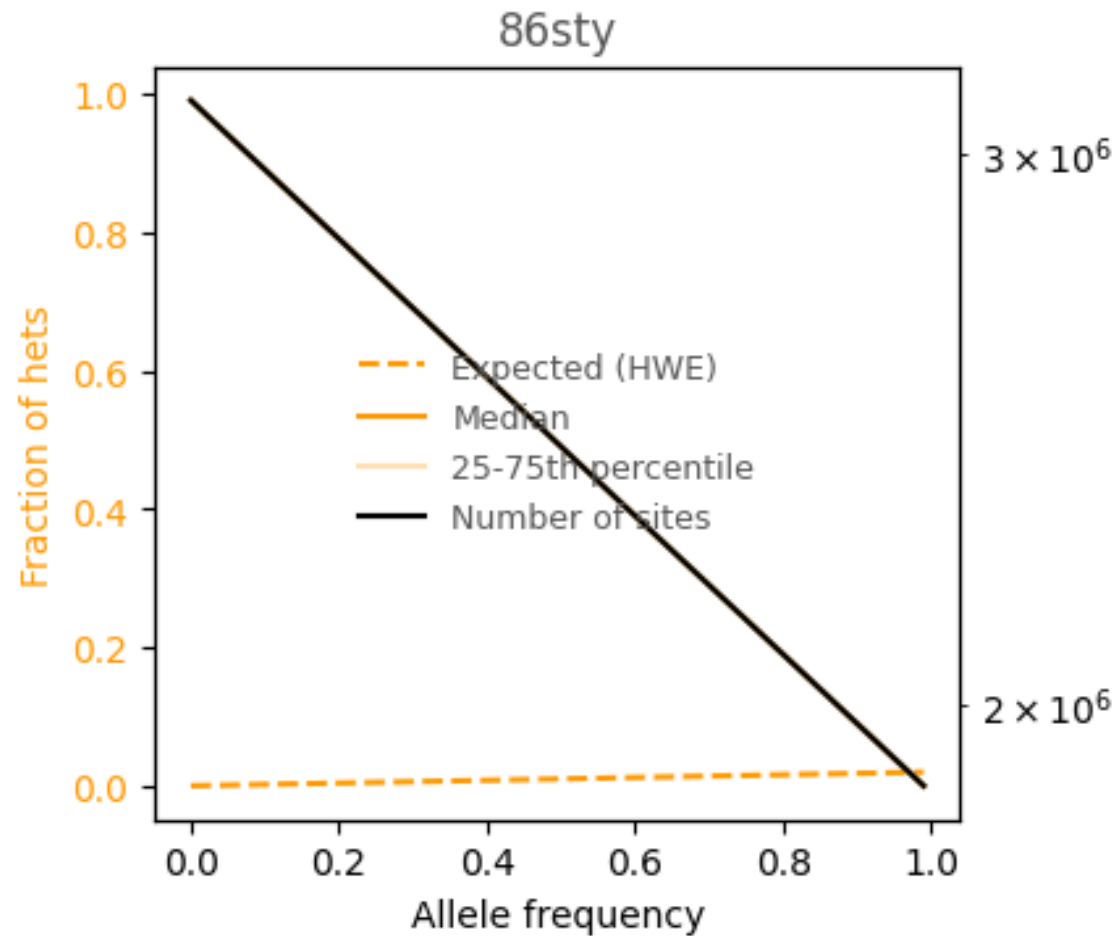
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

