

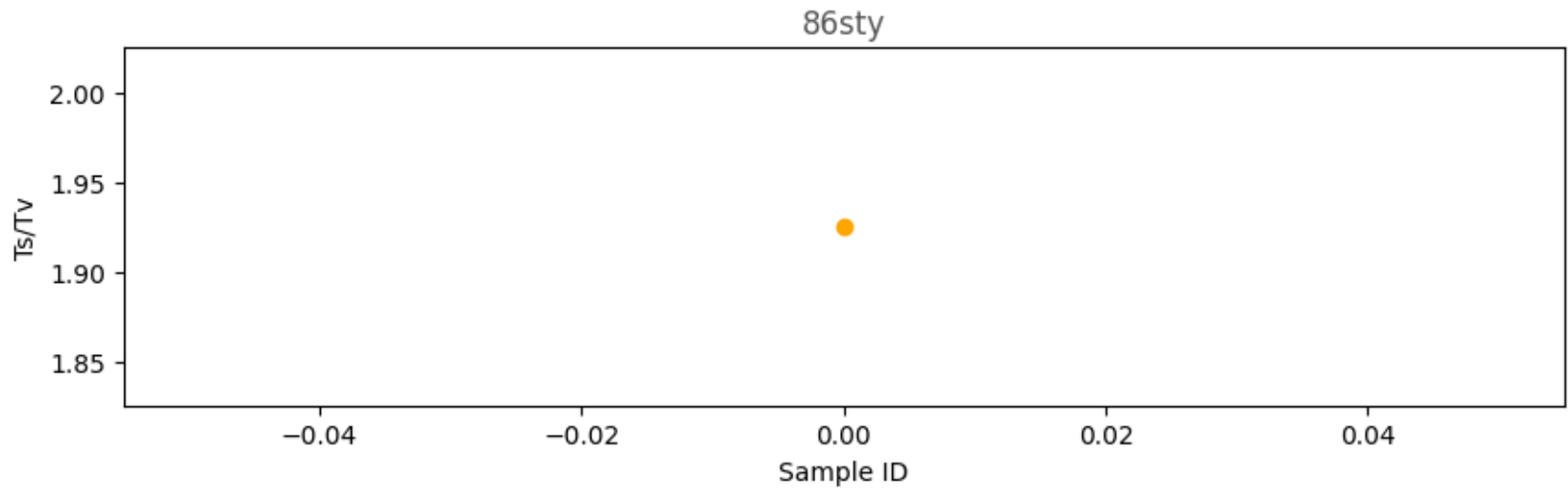
Summary Numbers

	SNPs			indels	MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	
86sty	4,102,394	1.93	1.93	956,545	–	0
* frameshift ratio: out/(out+in)						

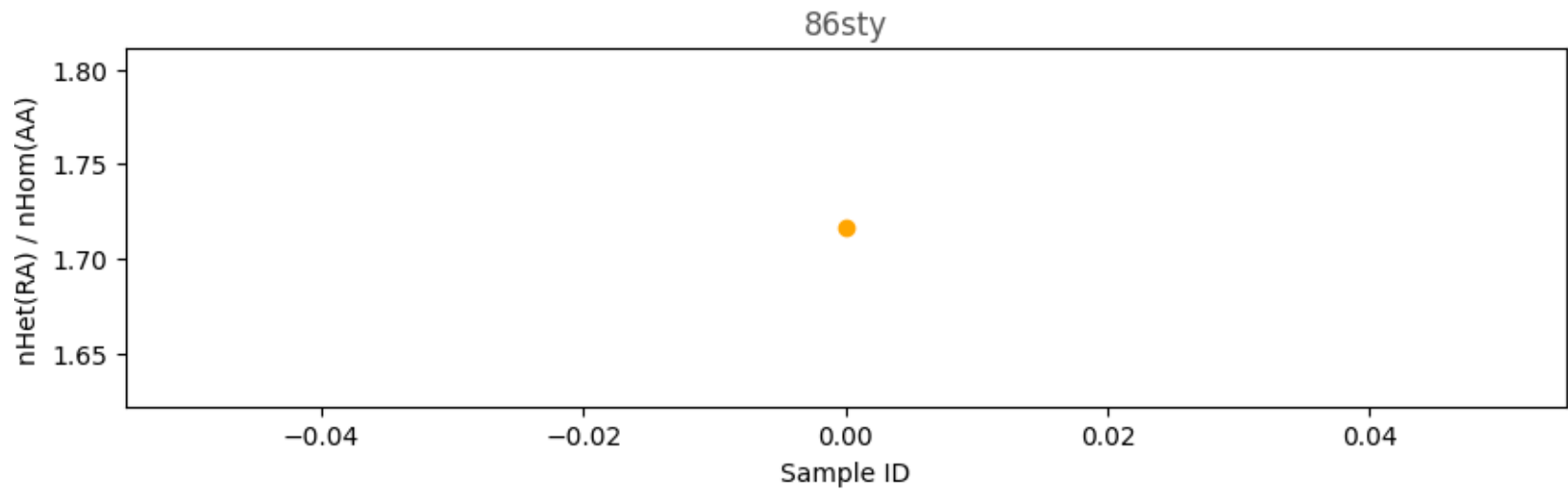
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
86sty	63.2%	1.90	68.5%	103,140	2,216

- 86sty .. /ngc/projects2/gm/data/archive/2022/variants/snv/86styataf-103812852866-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211203_A01411_BHVMTHDSX2-RHGM_LABKA_WGSA
 KUT-WGSAKUT03469_21RKG029698x01_snv_germline_raw.haplotype_caller.vcf.gz

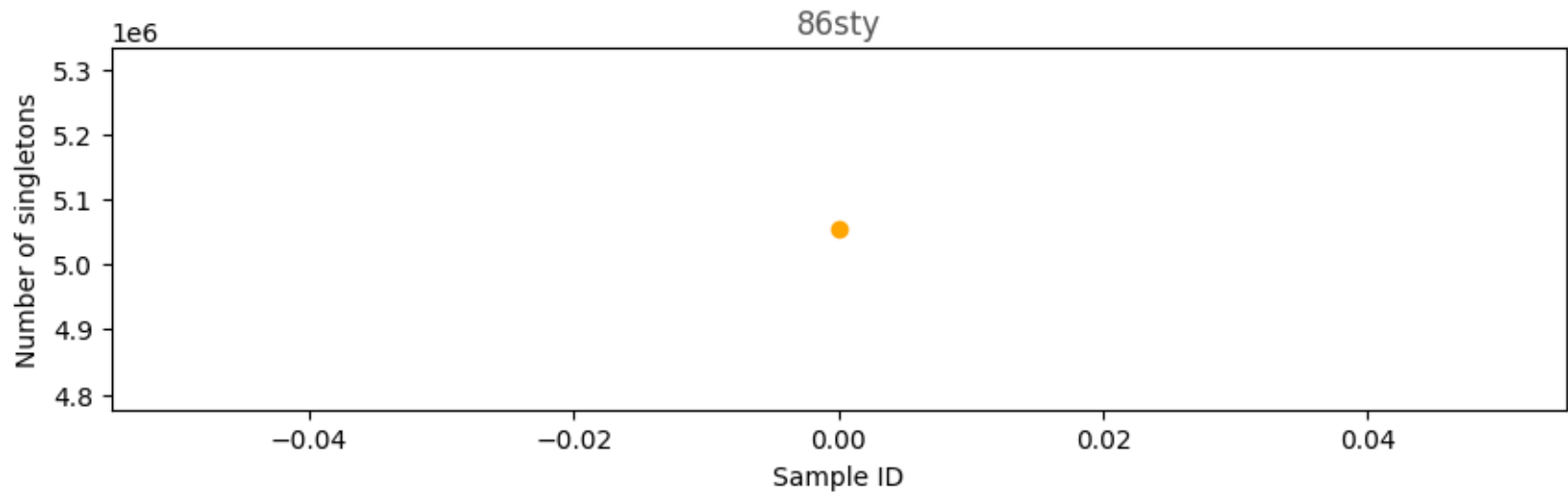
Ts/Tv by sample



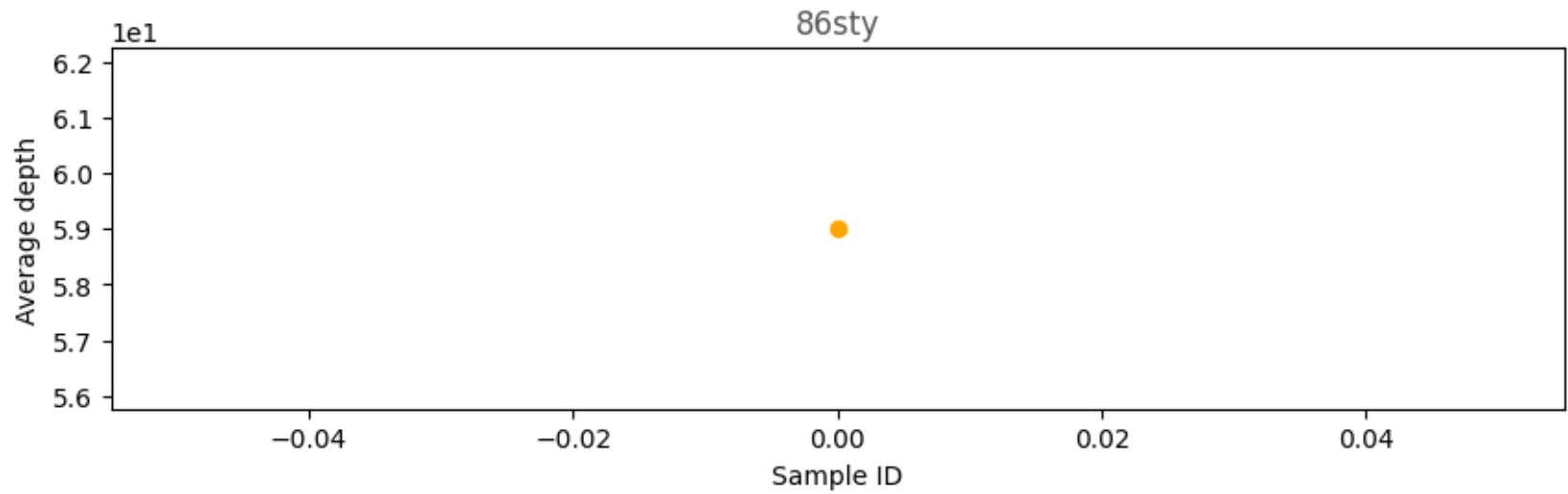
Hets vs non-ref Homs by sample



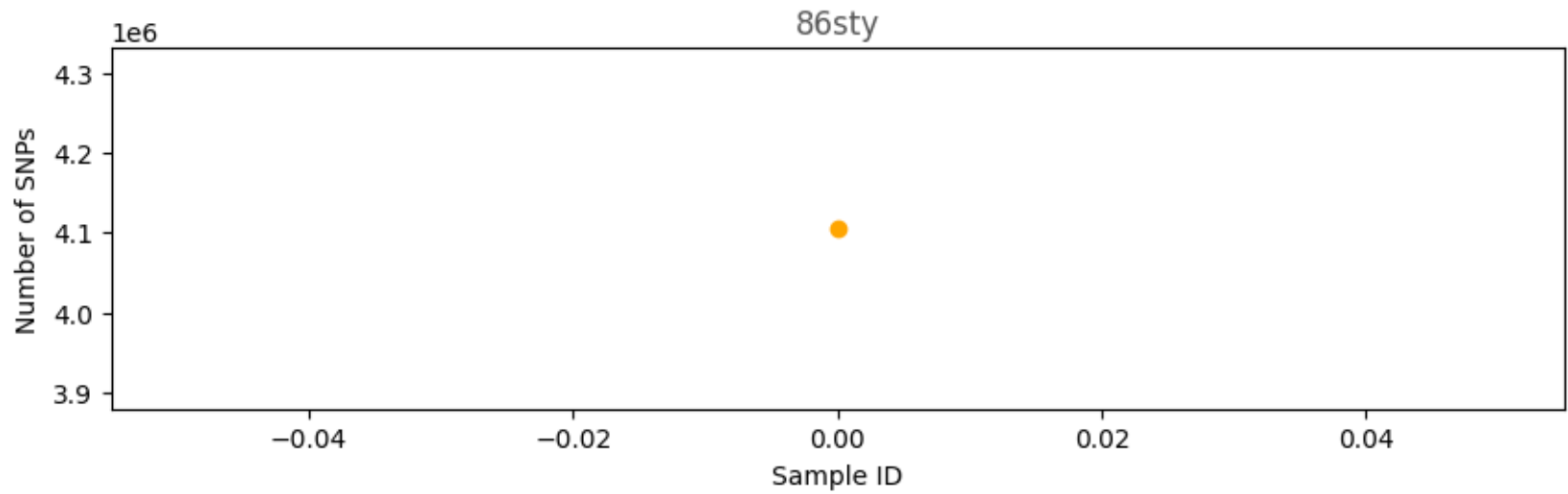
Singletons by sample (hets and homs)



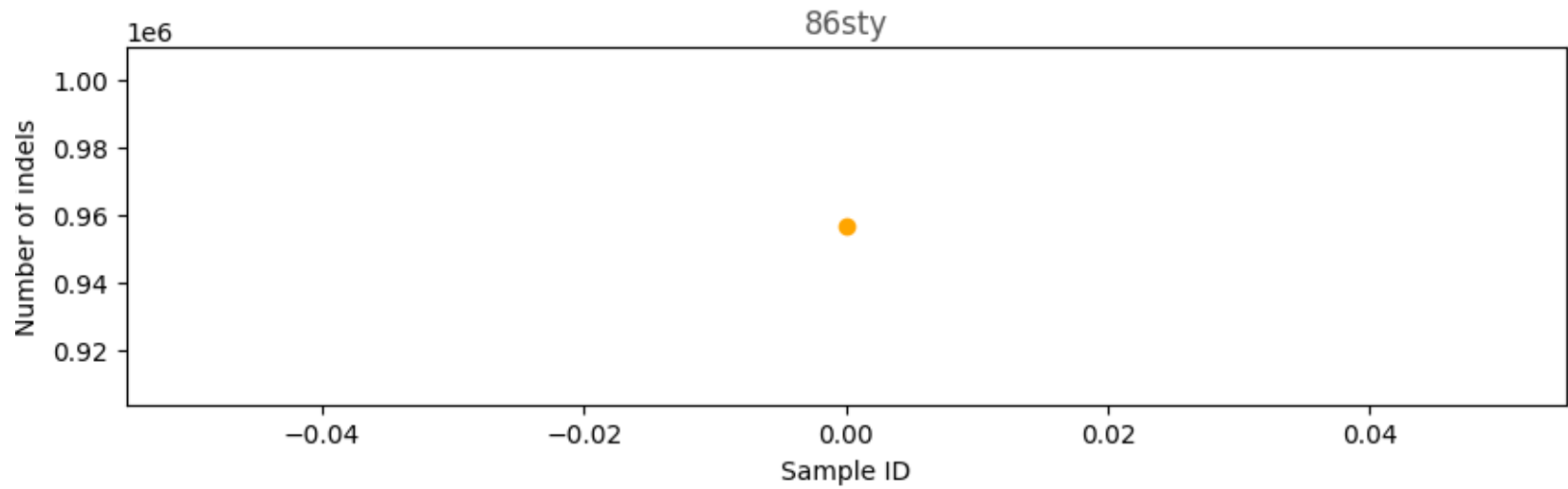
Average depth by sample



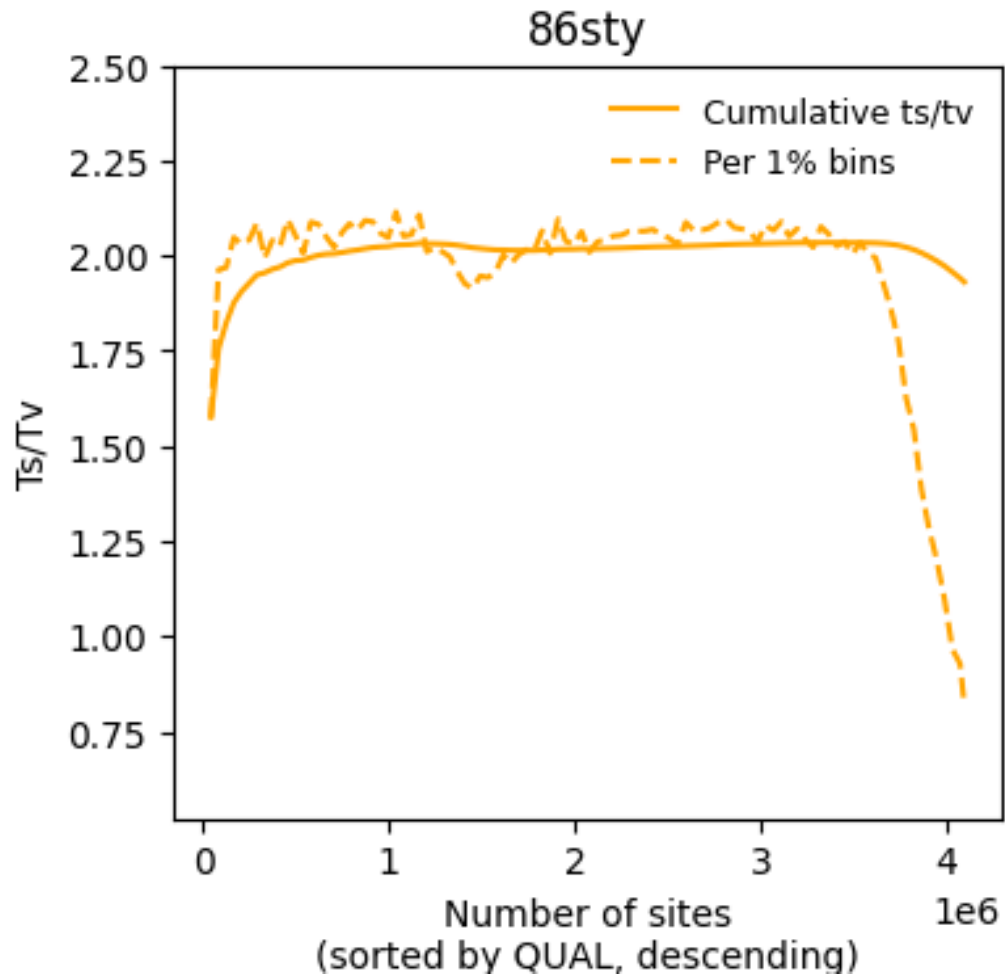
Number of SNPs by sample



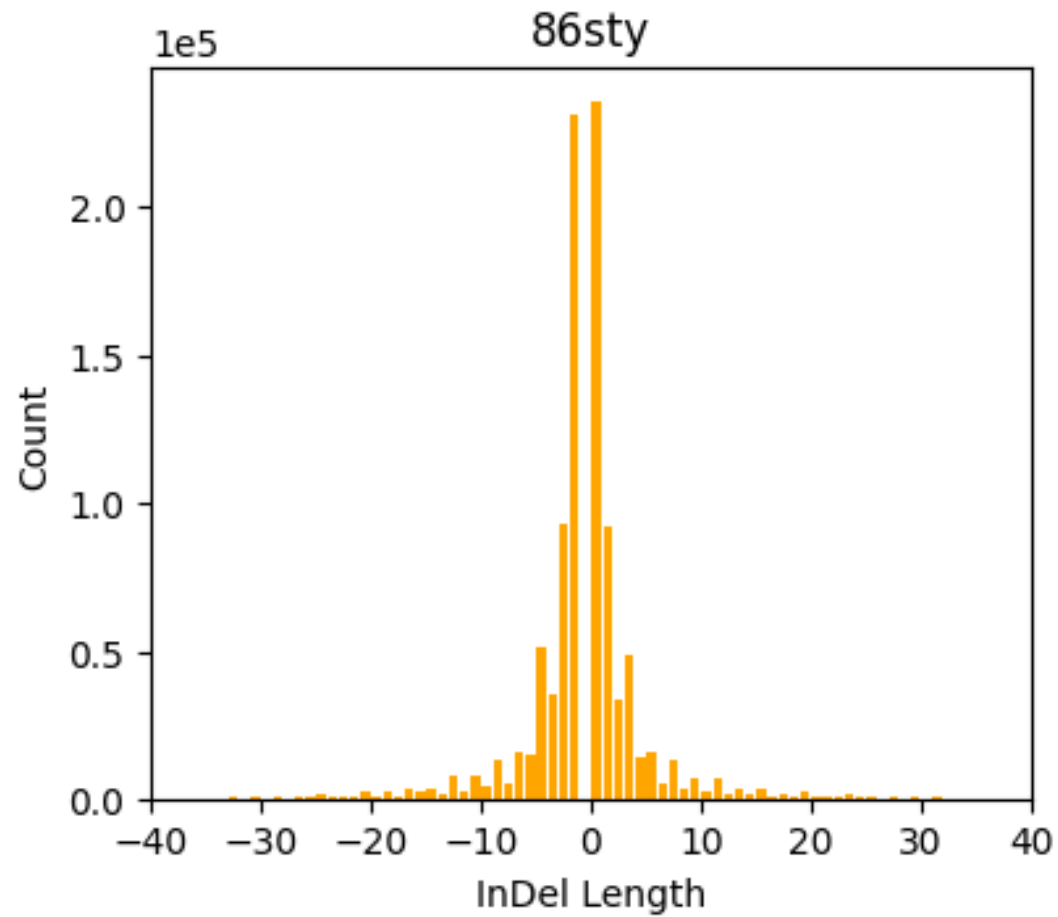
Number of indels by sample



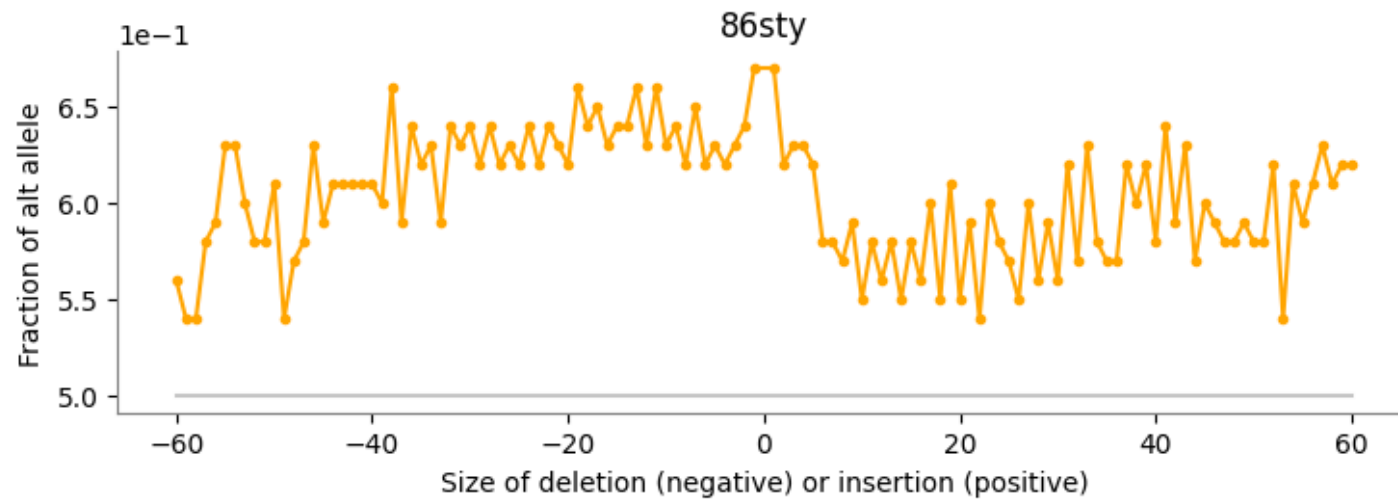
Ts/Tv stratified by QUAL



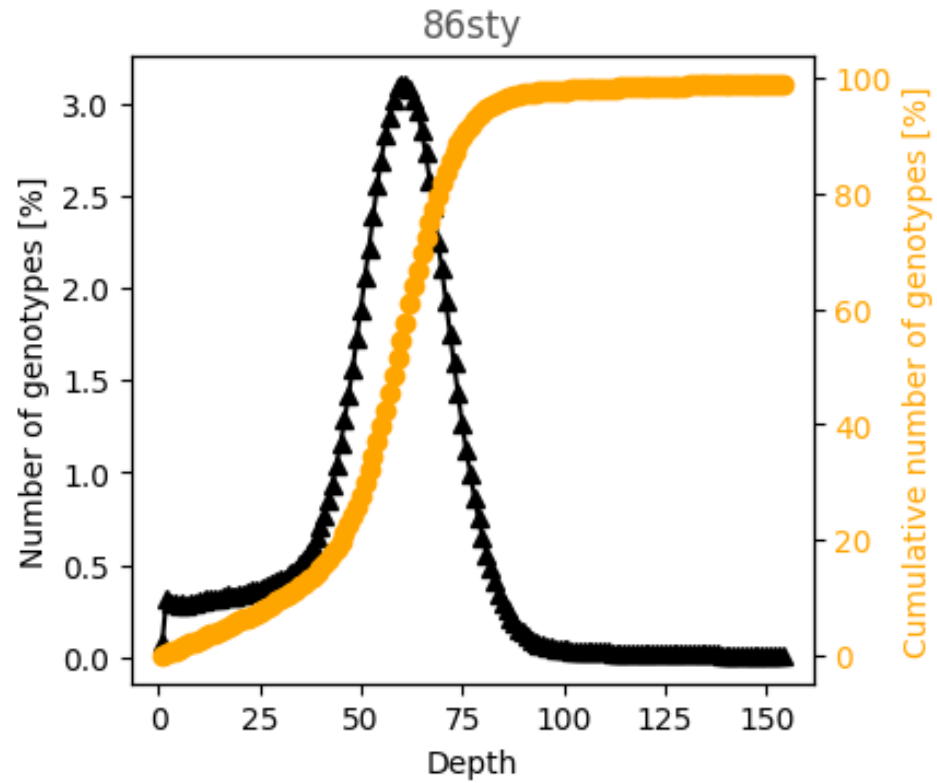
Indel distribution



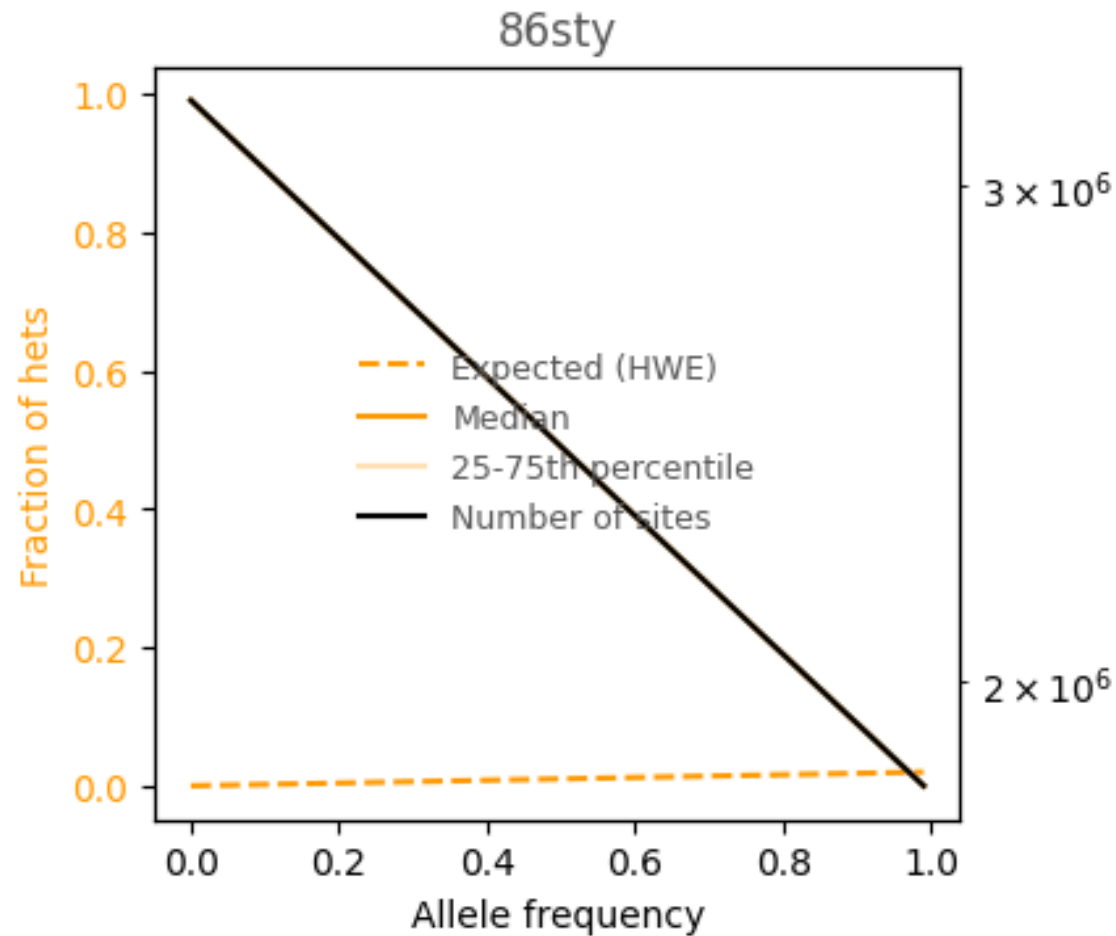
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

