

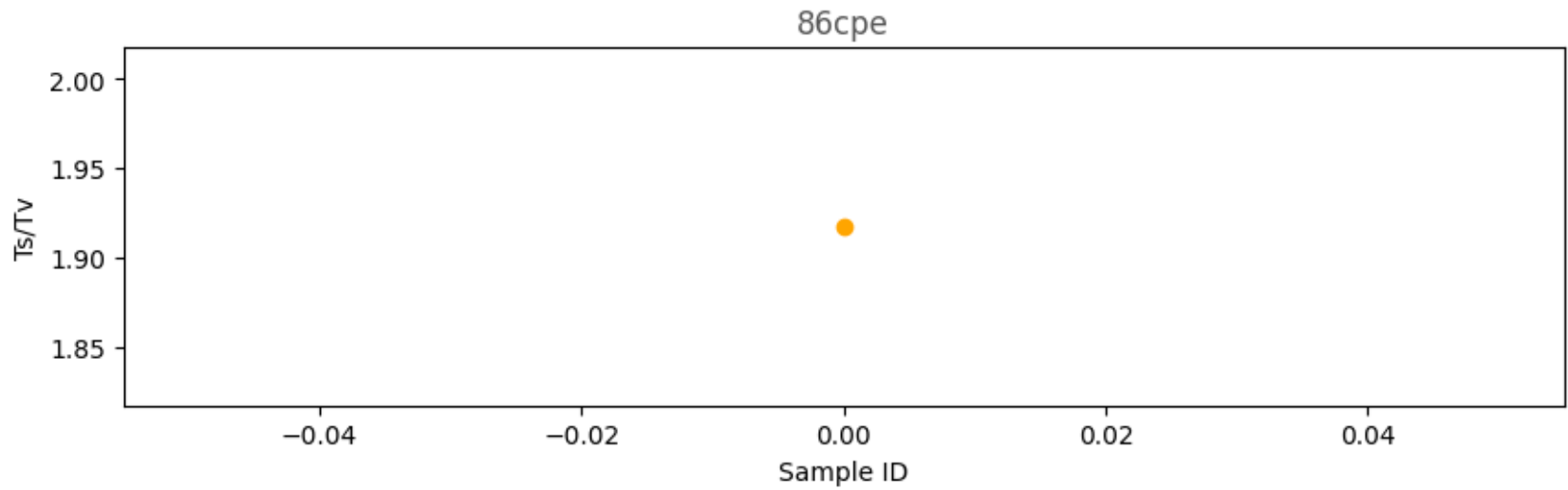
Summary Numbers

	SNPs			indels	MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	
86cpe	4,257,881	1.92	1.92	997,186	–	0
* frameshift ratio: out/(out+in)						

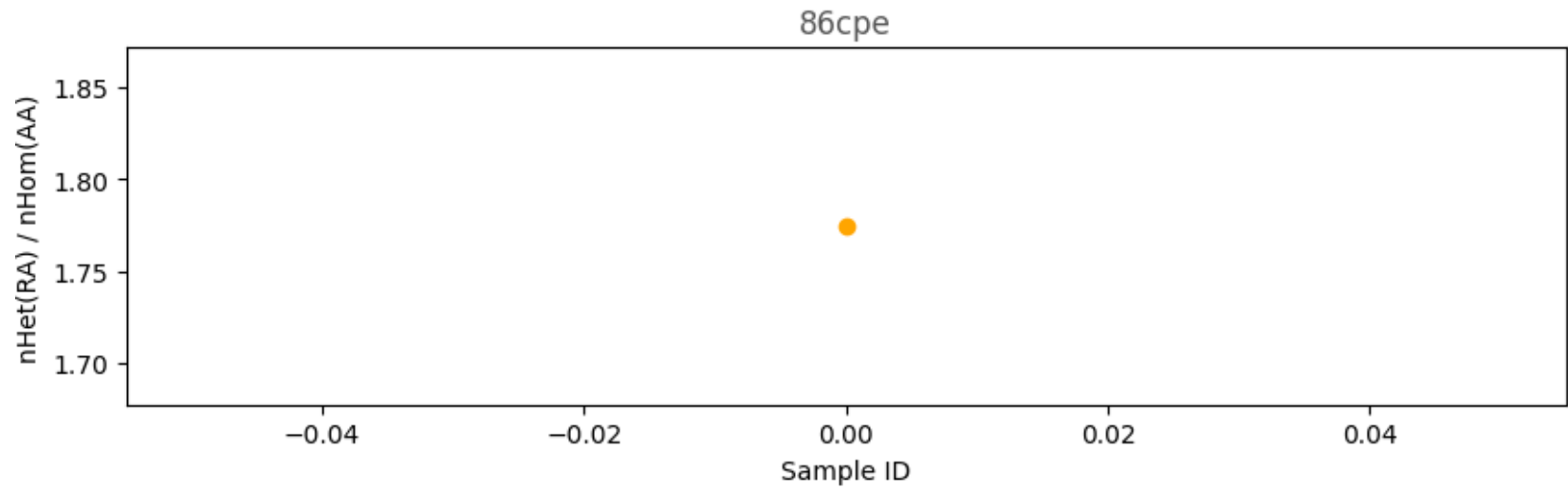
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
86cpe	64.0%	1.88	70.0%	117,232	2,207

- 86cpe .. /ngc/projects2/gm/data/archive/2022/variants/snv/86cpeandf-103745744606-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00183-210608_A01176_BHF27VDSX2-EXT_LAB
 KA_NGCWGS-NGCWGS00098_86krrwm30_snv_germline_raw.haplotype_caller.vcf.gz

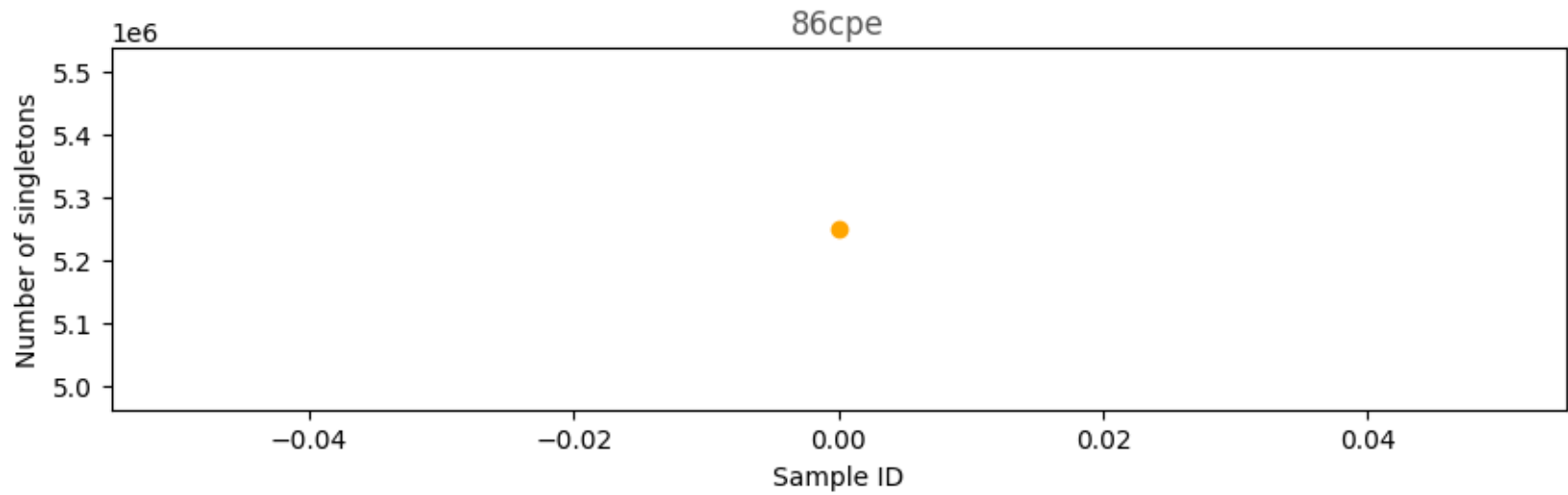
Ts/Tv by sample



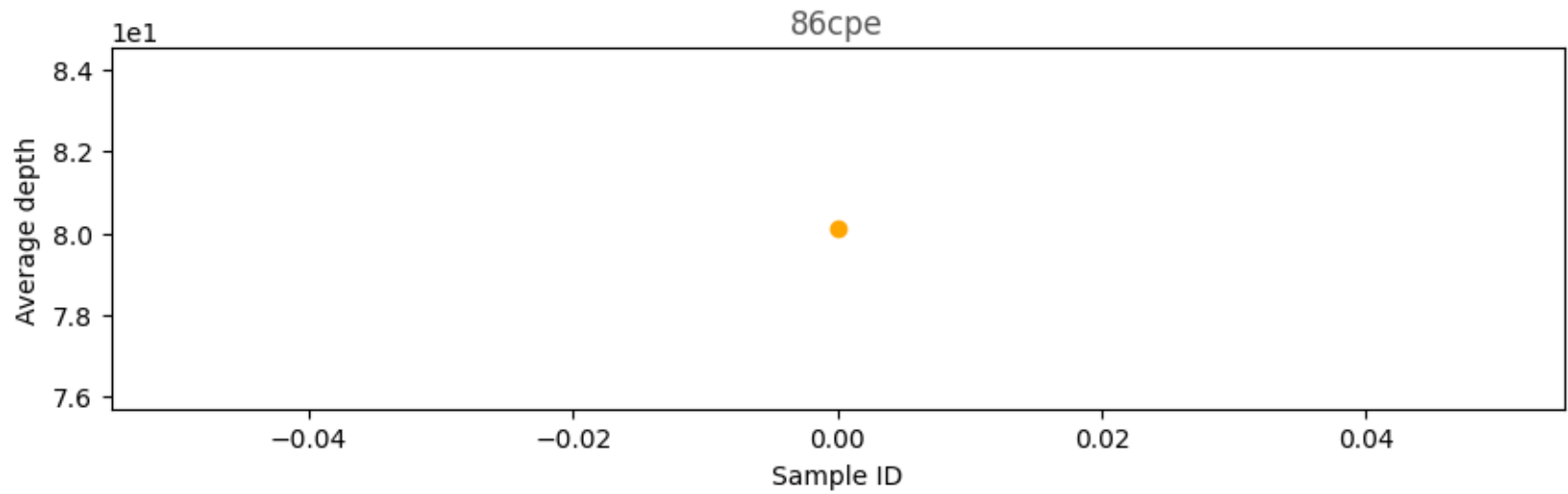
Hets vs non-ref Homs by sample



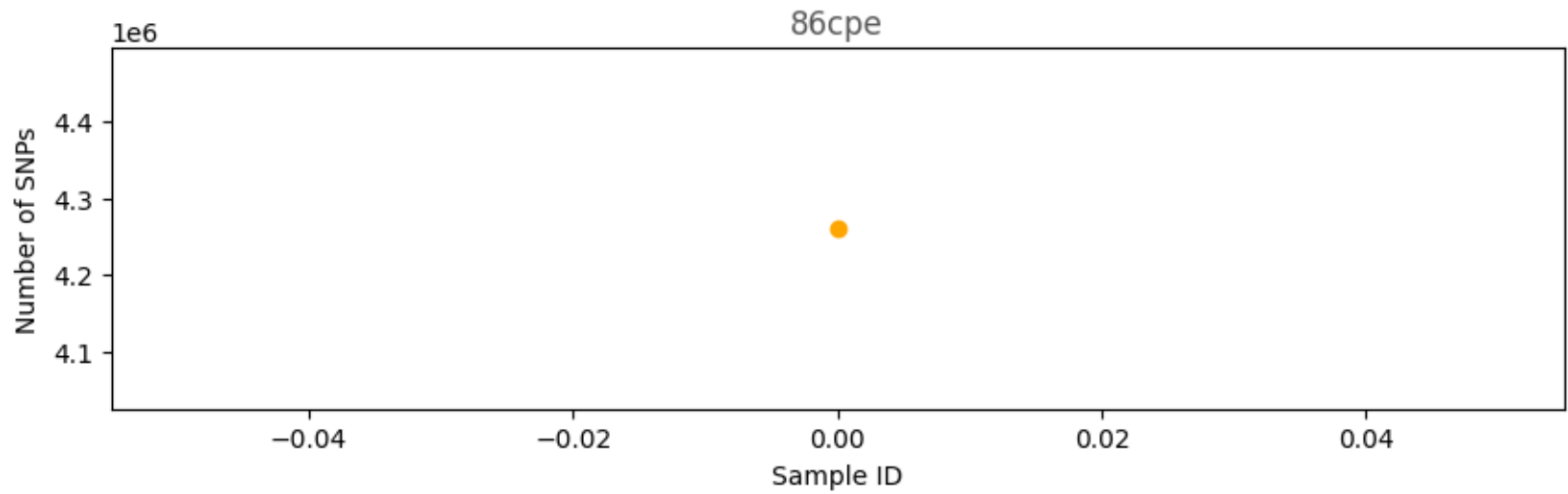
Singletons by sample (hets and homs)



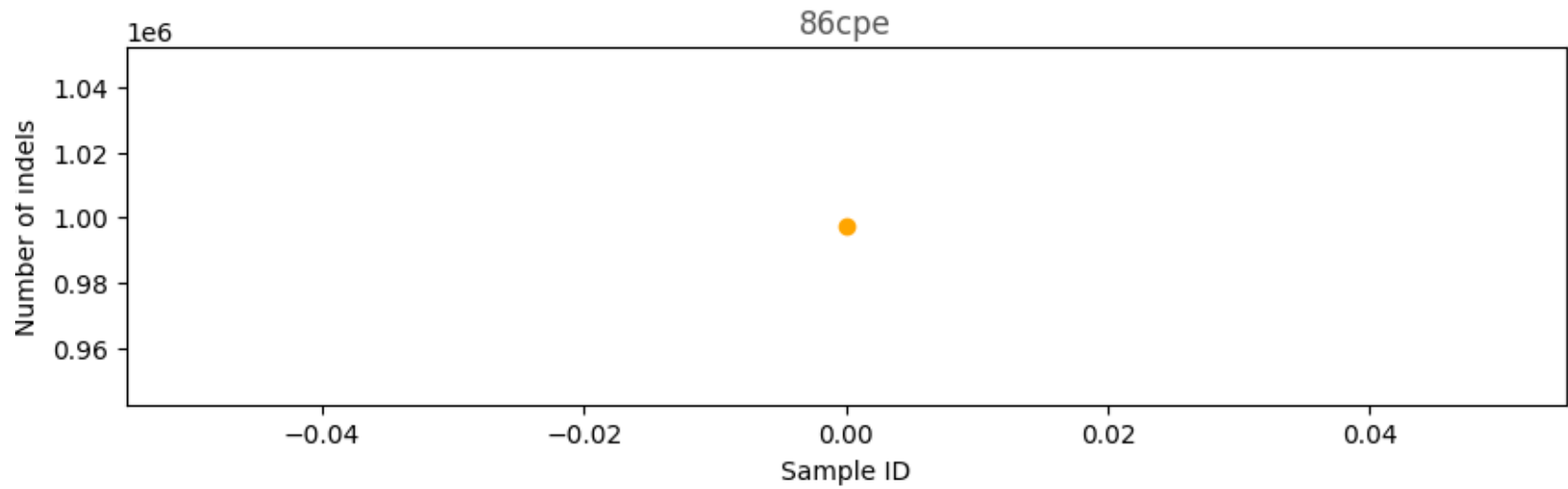
Average depth by sample



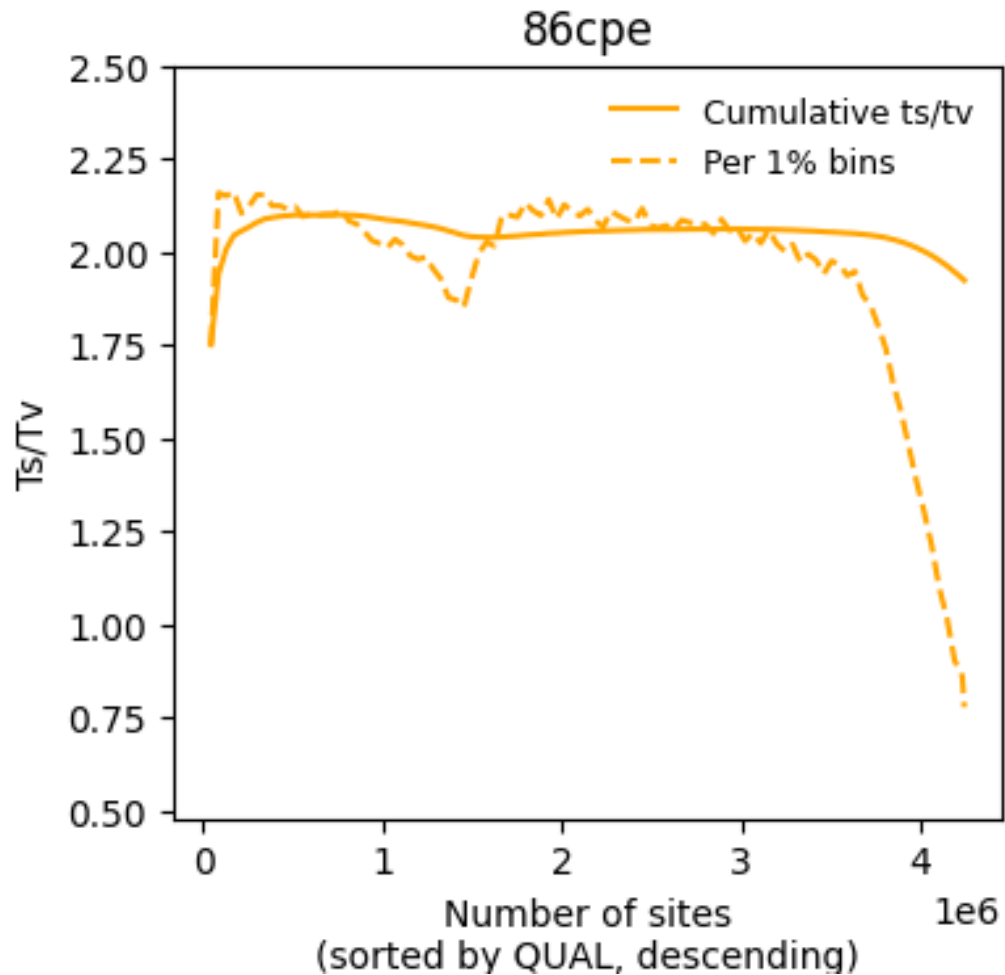
Number of SNPs by sample



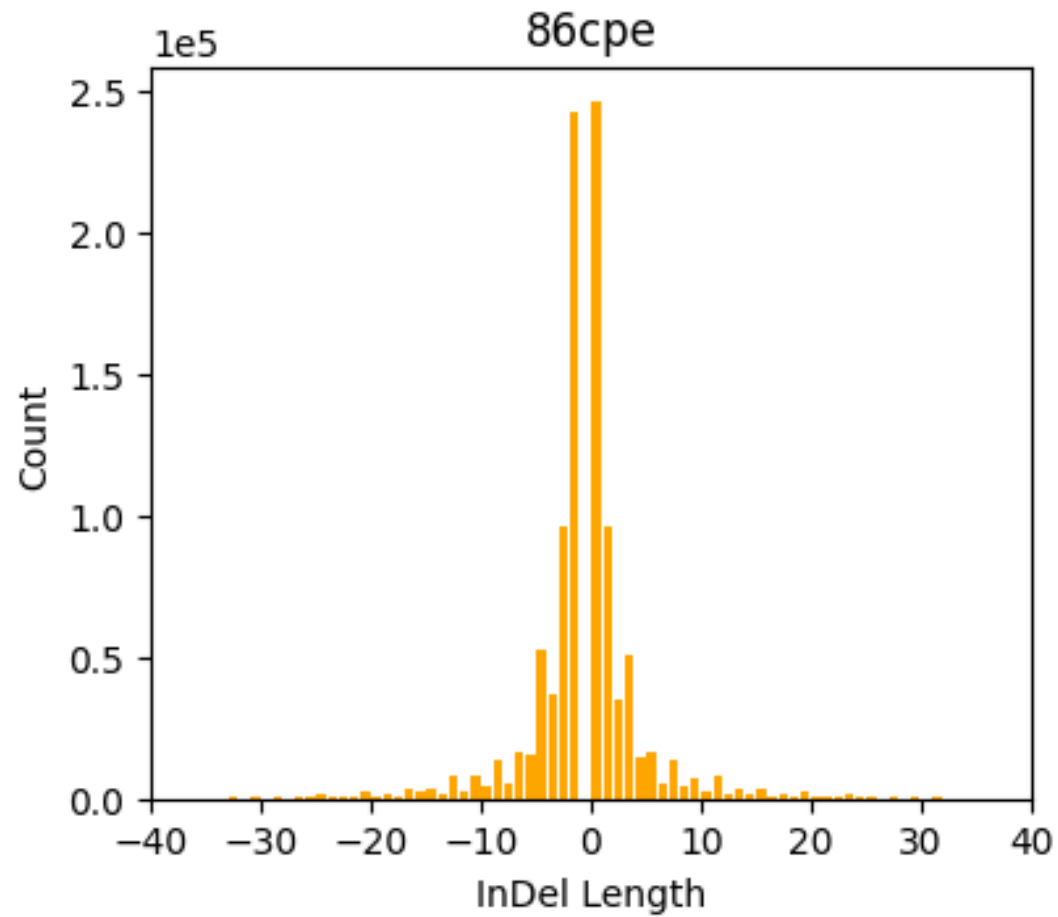
Number of indels by sample



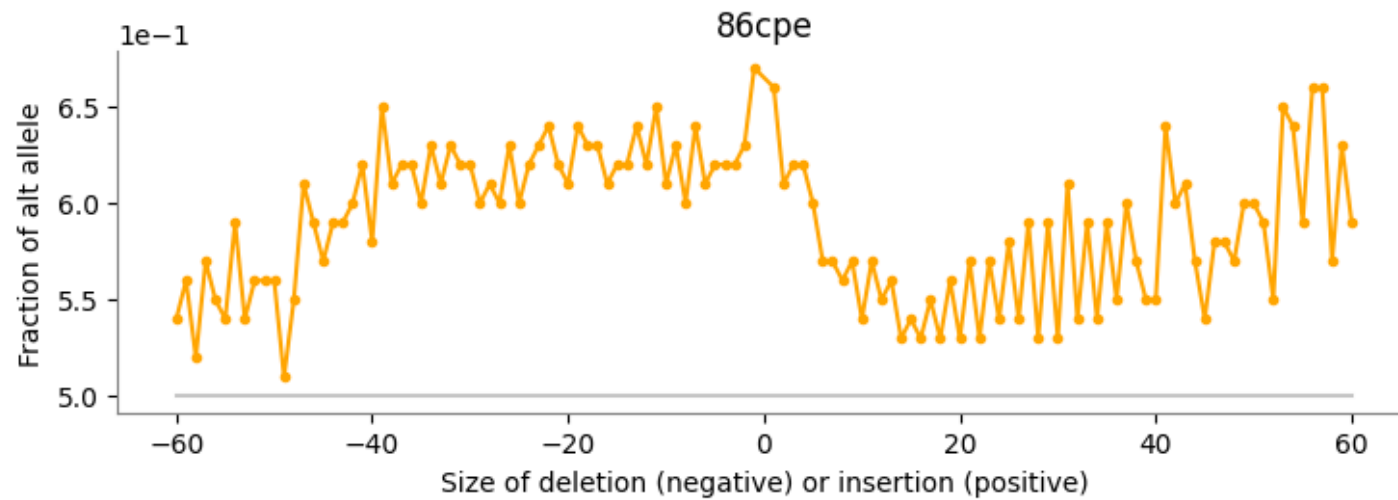
Ts/Tv stratified by QUAL



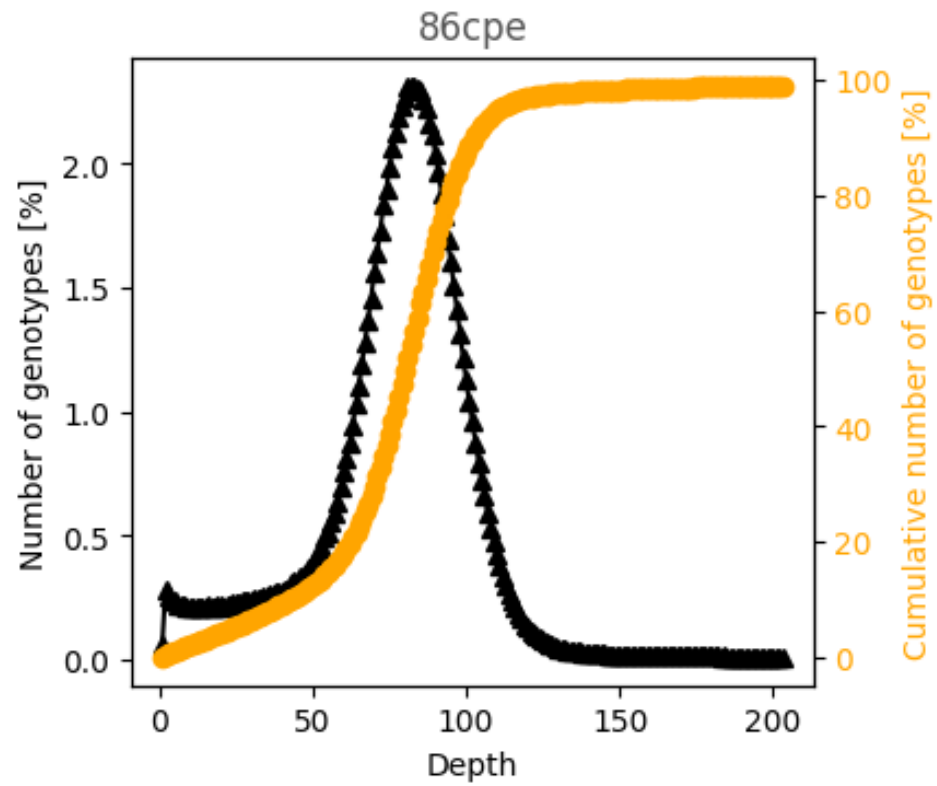
Indel distribution



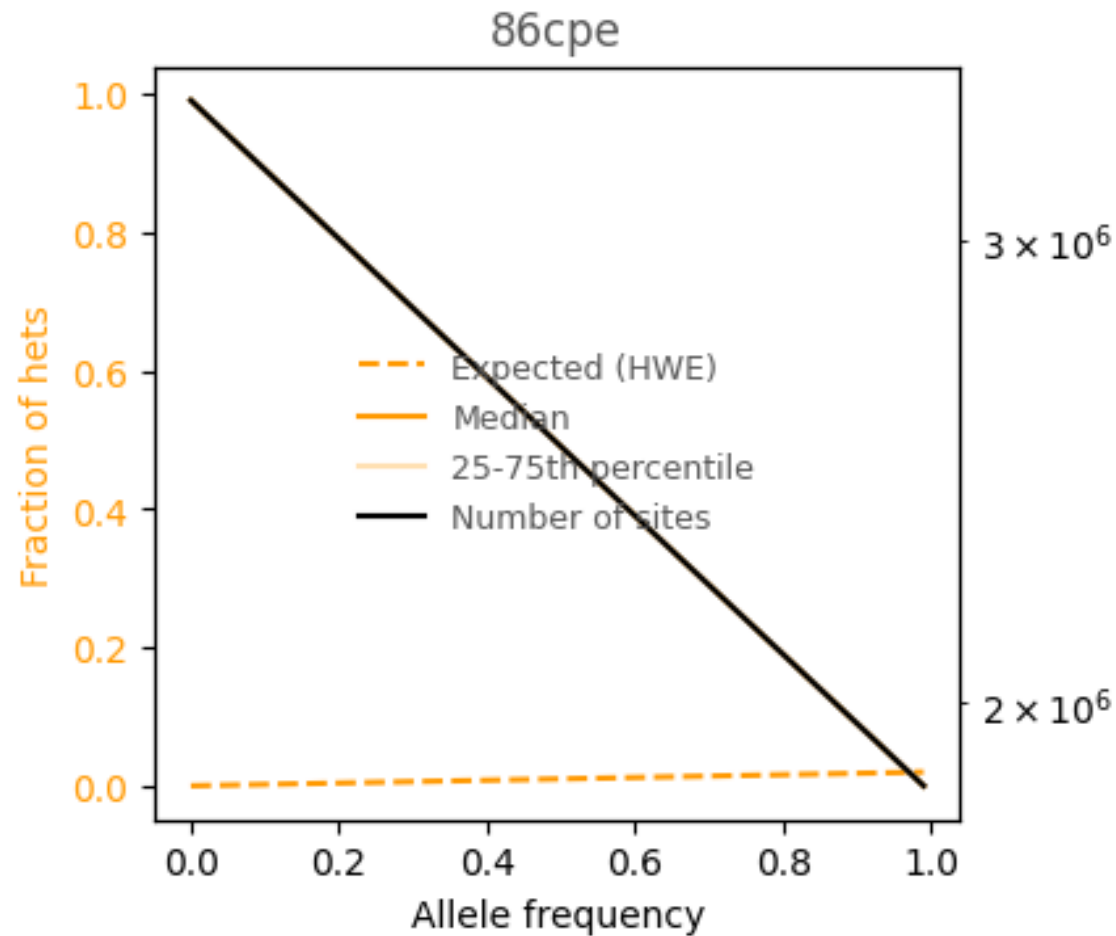
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

