

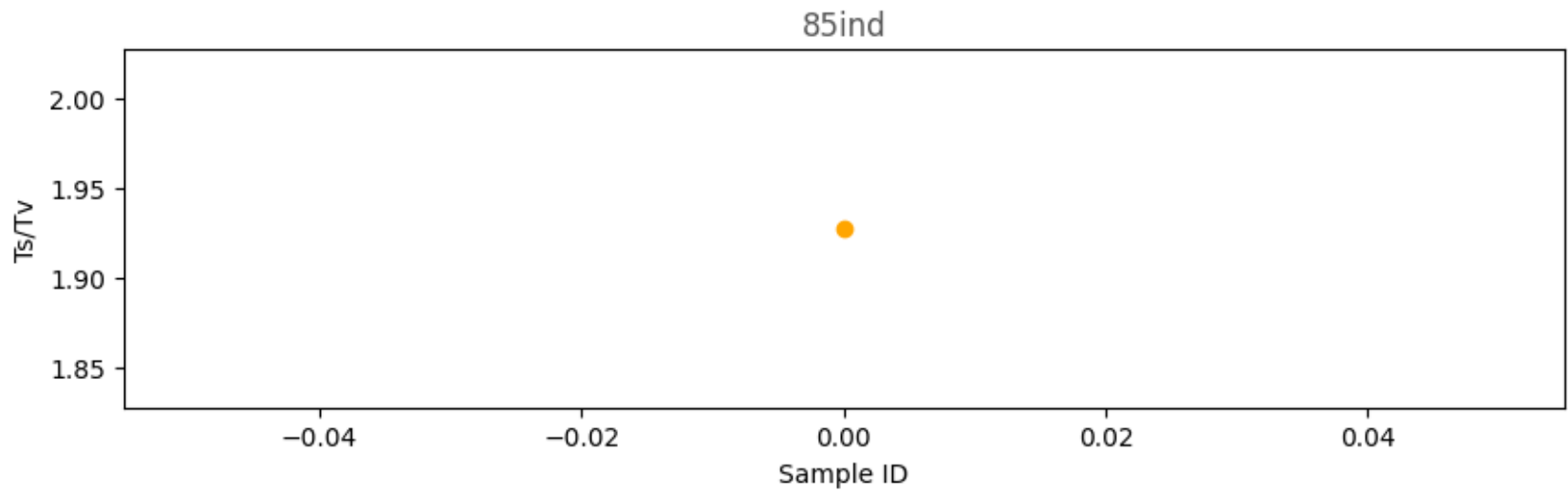
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
85ind	4,064,573	1.93	1.93	940,521	–	0	0
* frameshift ratio: out/(out+in)							

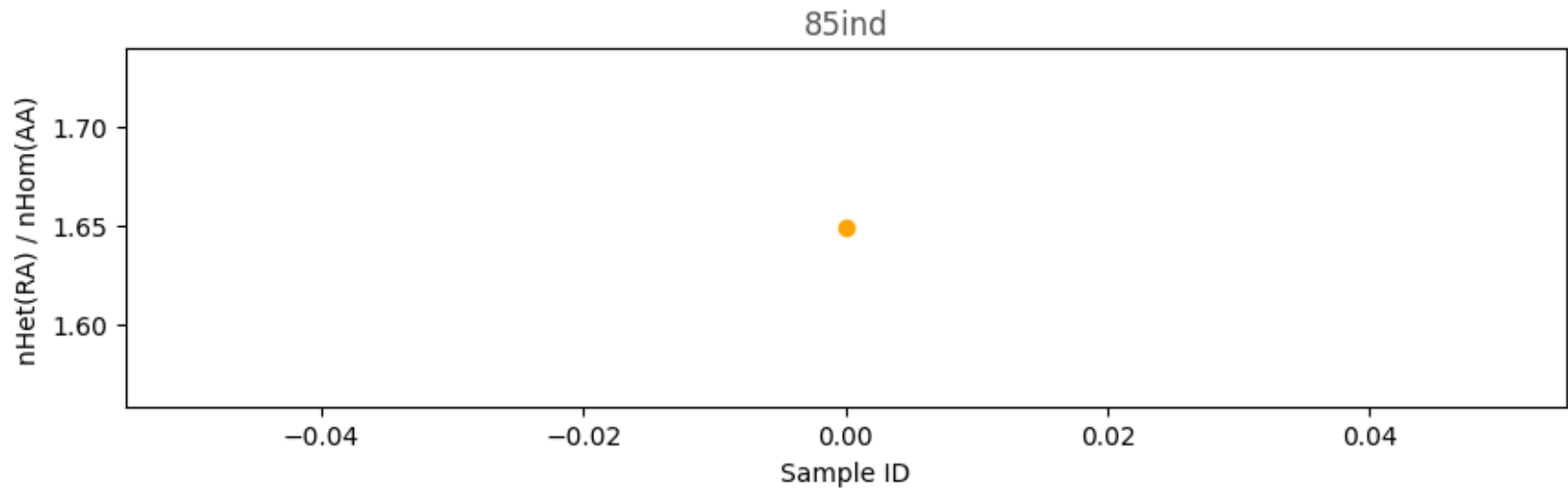
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
85ind	62.3%	1.89	66.6%	95,760	1,893

- 85ind .. /ngc/projects2/gm/data/archive/2022/variants/snv/85indatom-103807518970-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00418-211203_A01466_BHN2G5DSX2-EXT_LAB
KA_NGCWGS-NGCWGS03440_21RKG026478x01_snv_germline_raw.haplotype_caller.vcf.gz

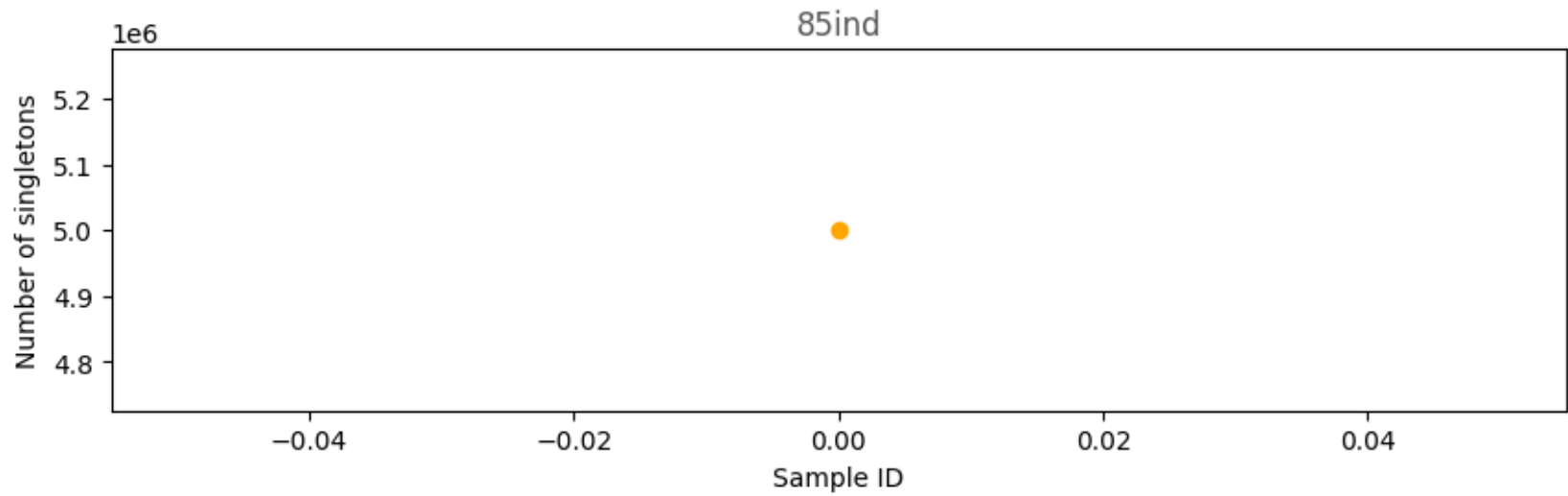
Ts/Tv by sample



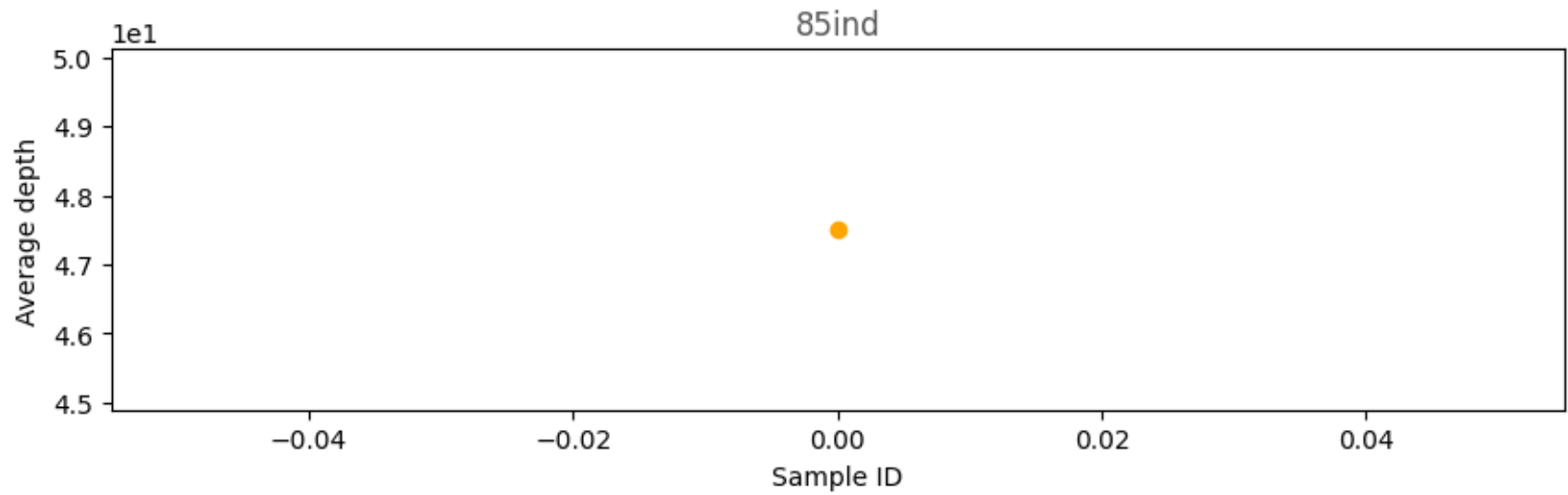
Hets vs non-ref Homs by sample



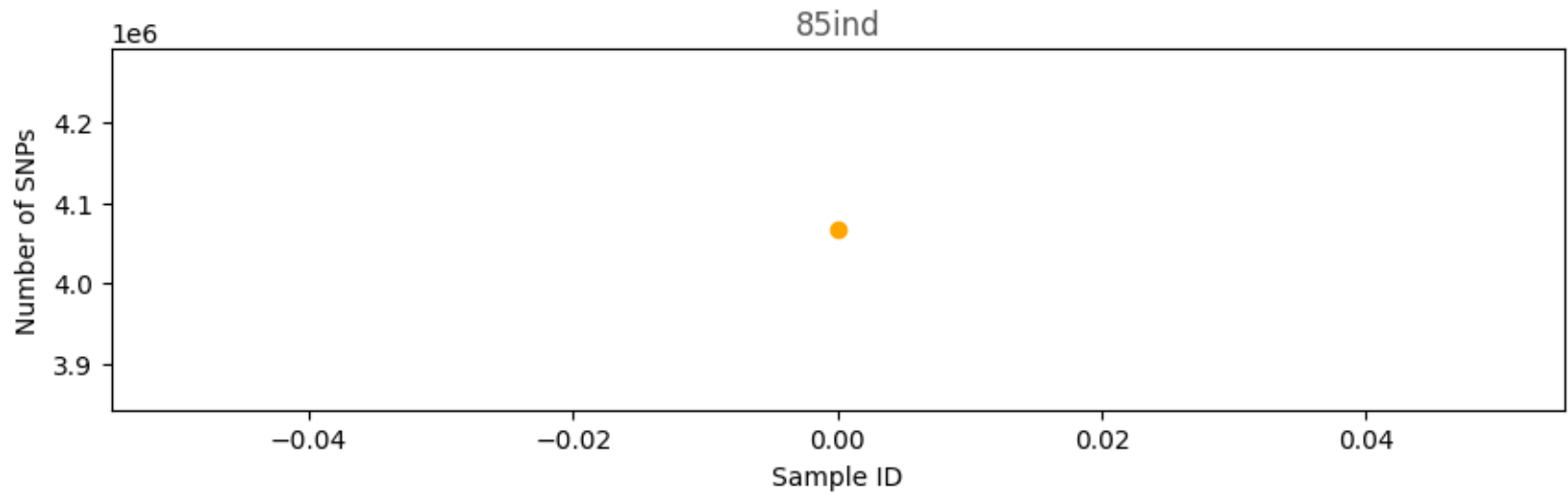
Singletons by sample (hets and homs)



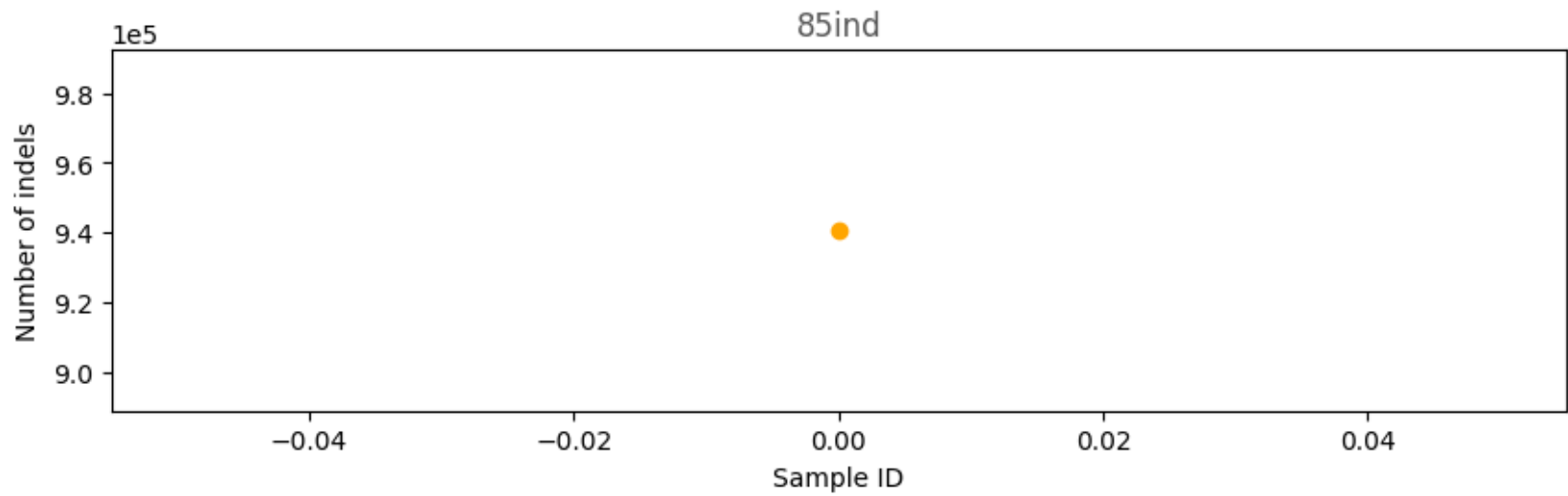
Average depth by sample



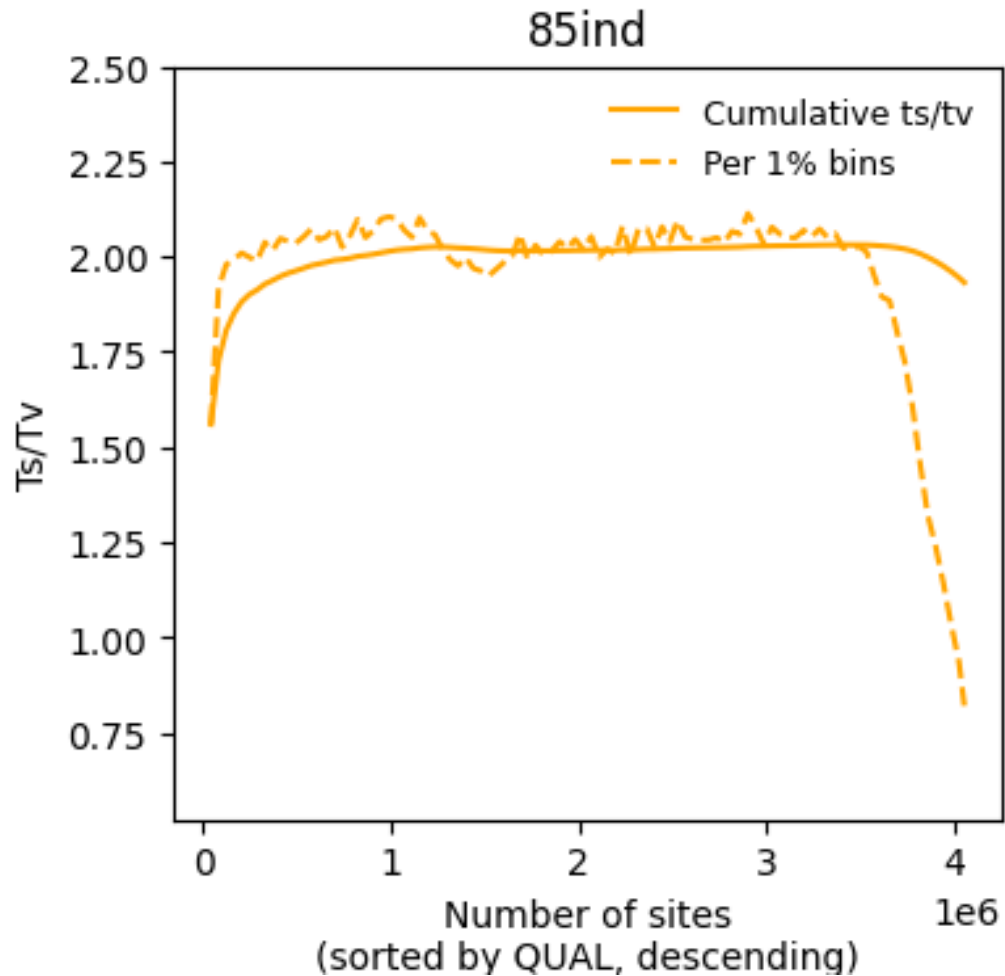
Number of SNPs by sample



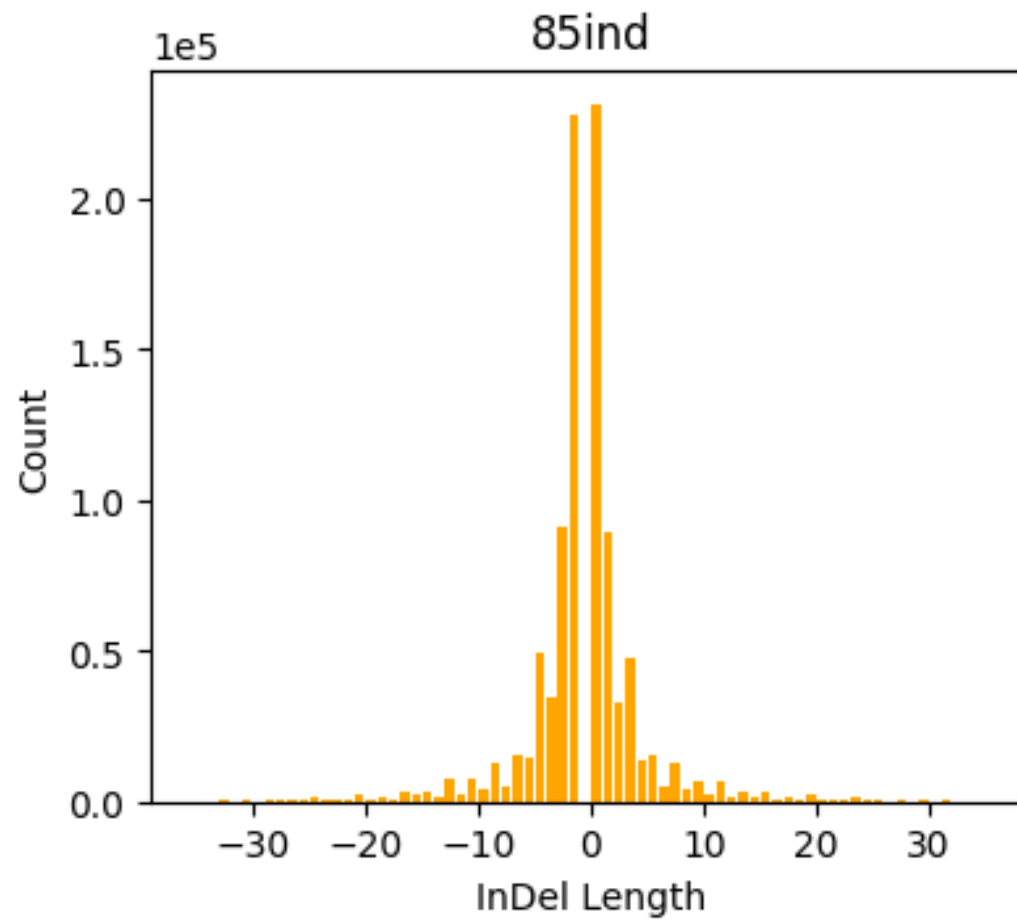
Number of indels by sample



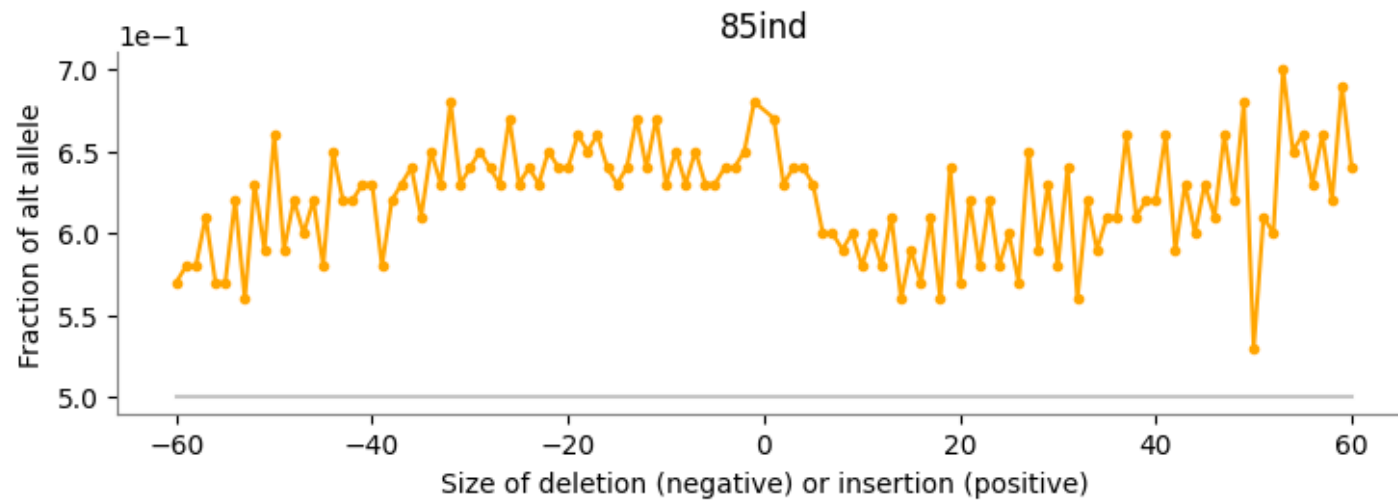
Ts/Tv stratified by QUAL



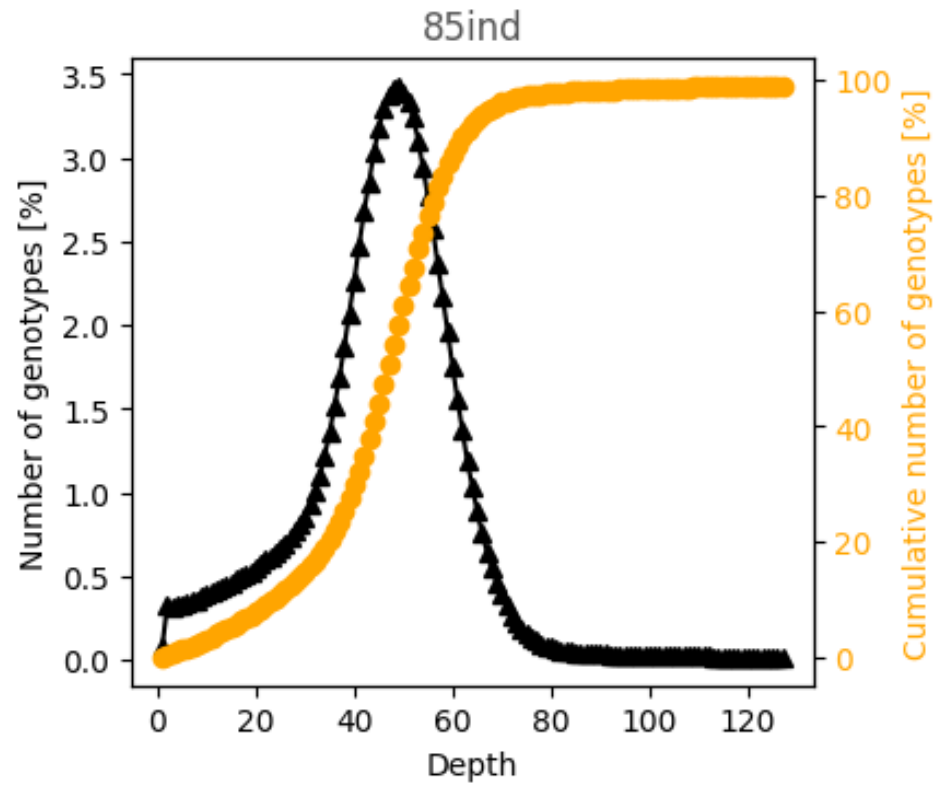
Indel distribution



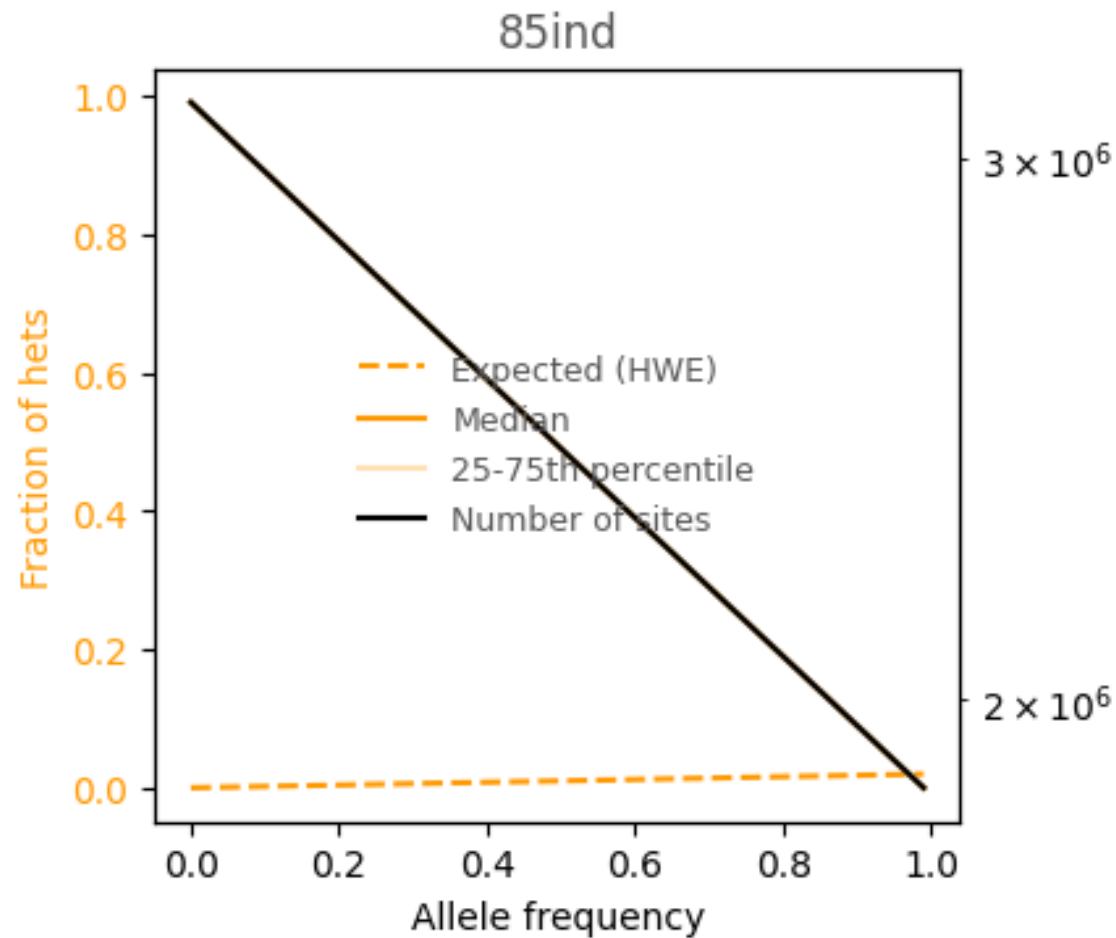
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

