

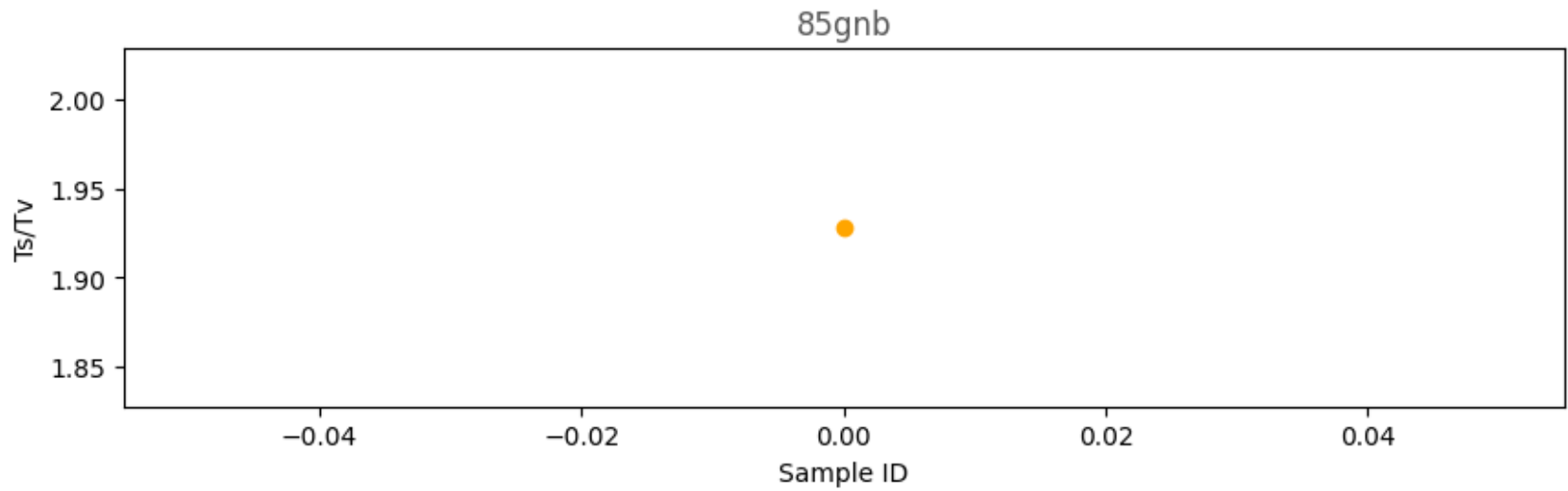
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
85gnb	4,108,583	1.93	1.93	960,275	–	0	0
* frameshift ratio: out/(out+in)							

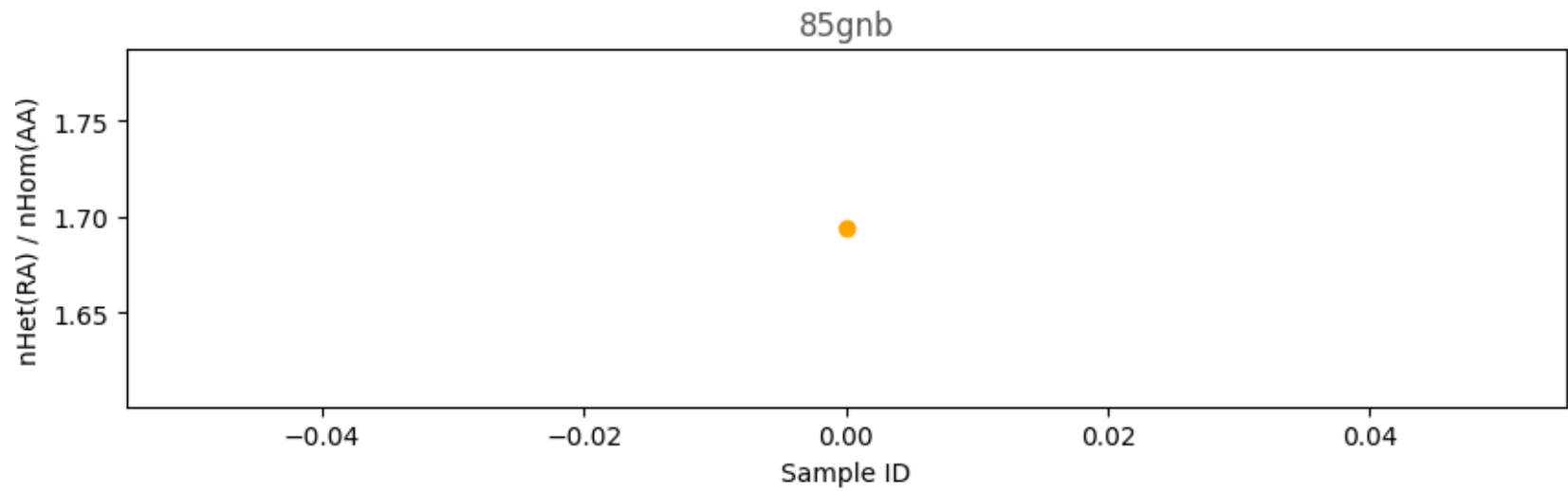
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
85gnb	62.9%	1.90	68.4%	103,344	2,292

- 85gnb .. /ngc/projects2/gm/data/archive/2022/variants/snv/85gnb0878-NA12878-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02307-221111_A01961_BH3TCFDSX5-EXT_LABKA_NG
CWGS-NGCWGS06195_snv_germline_raw.haplotype_caller.vcf.gz

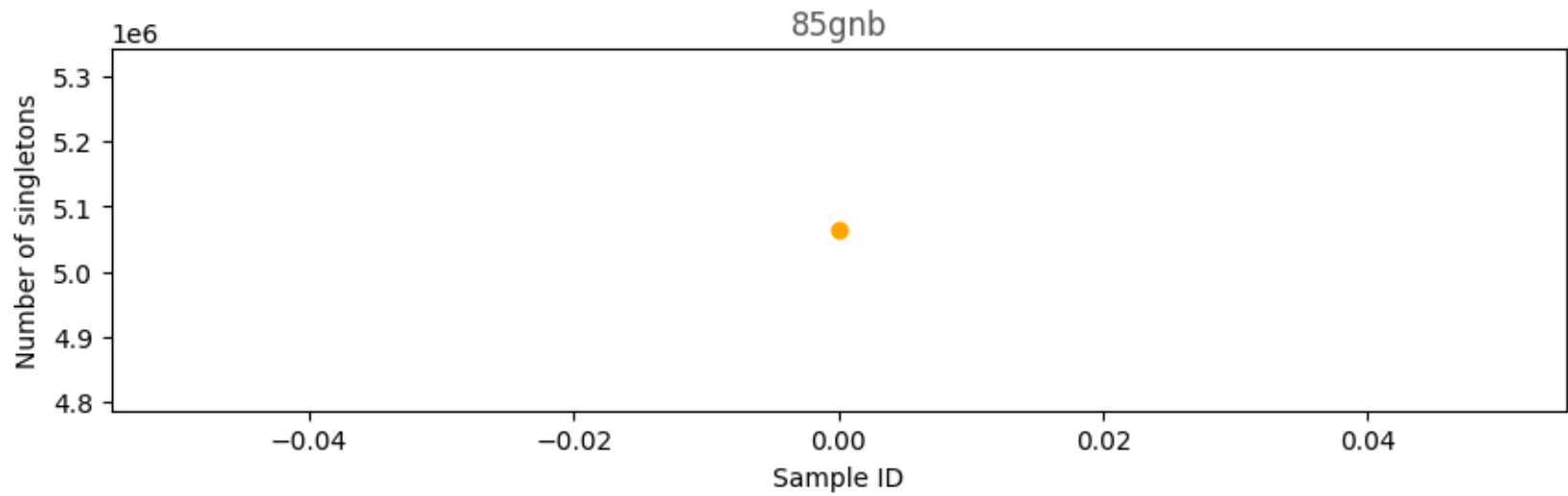
Ts/Tv by sample



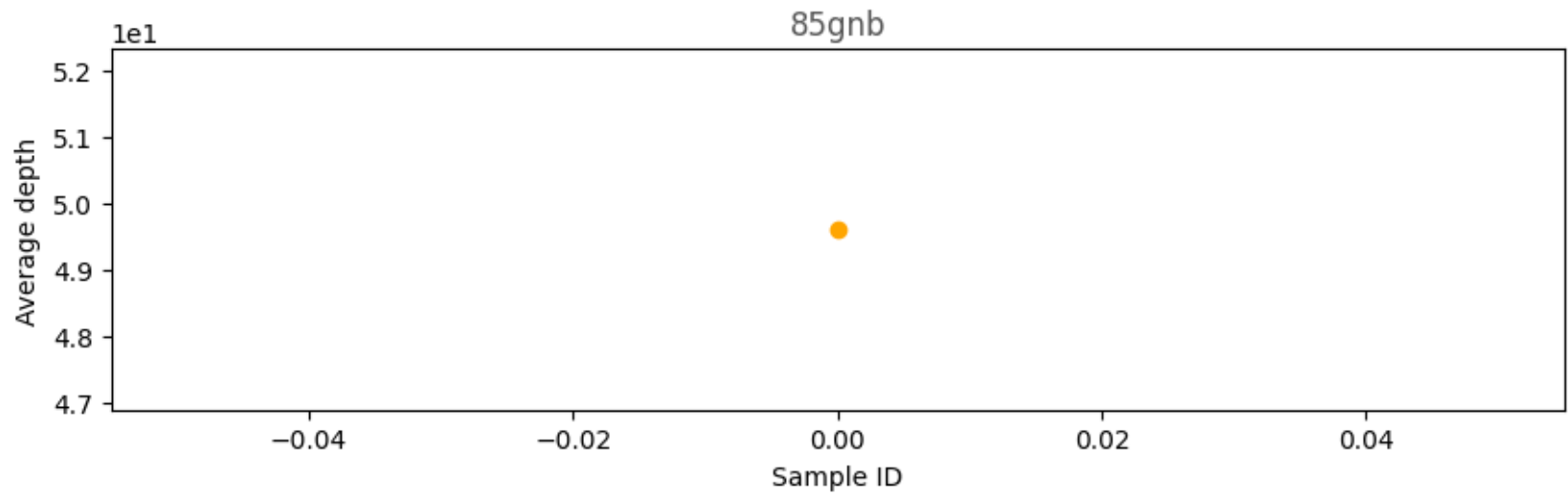
Hets vs non-ref Homs by sample



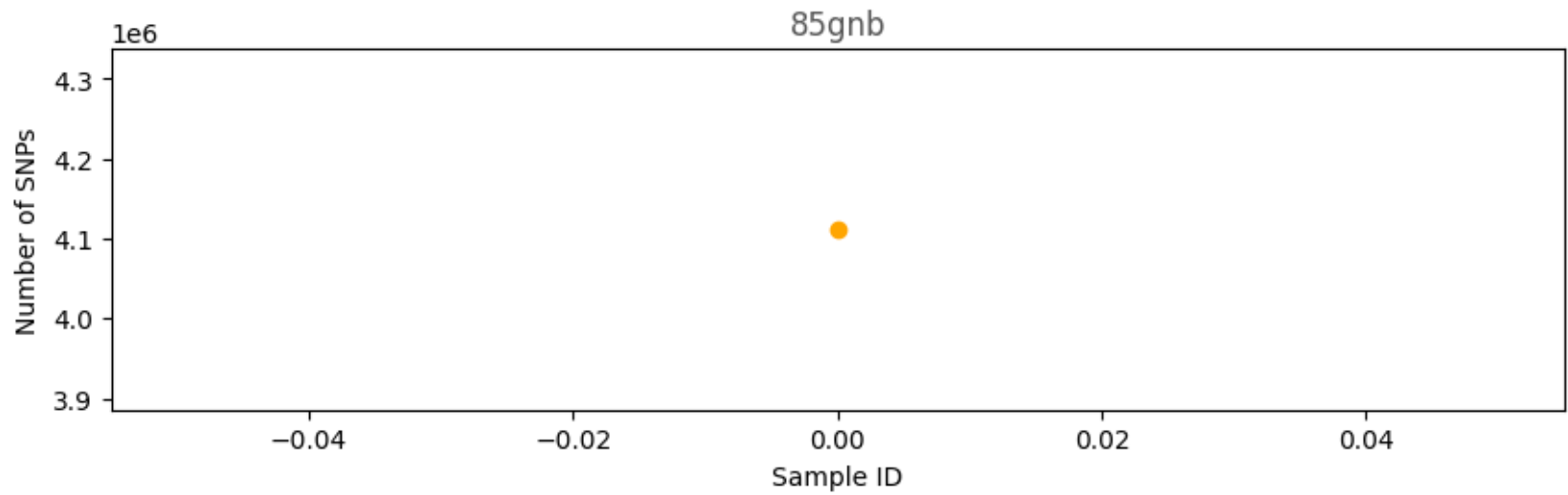
Singletons by sample (hets and homs)



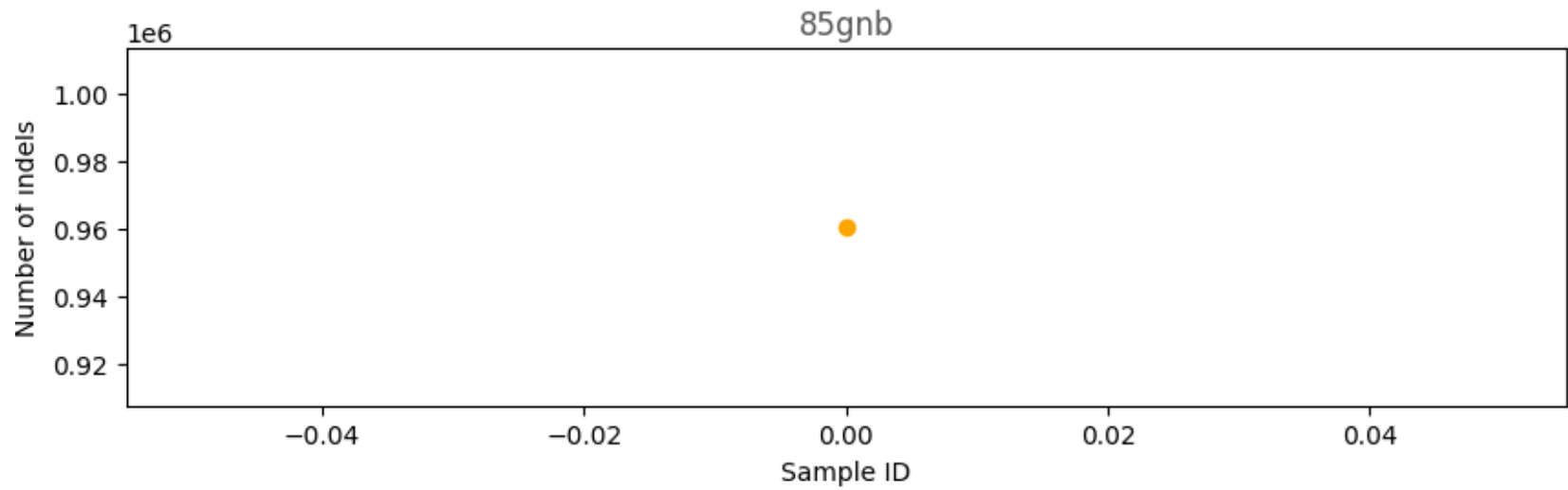
Average depth by sample



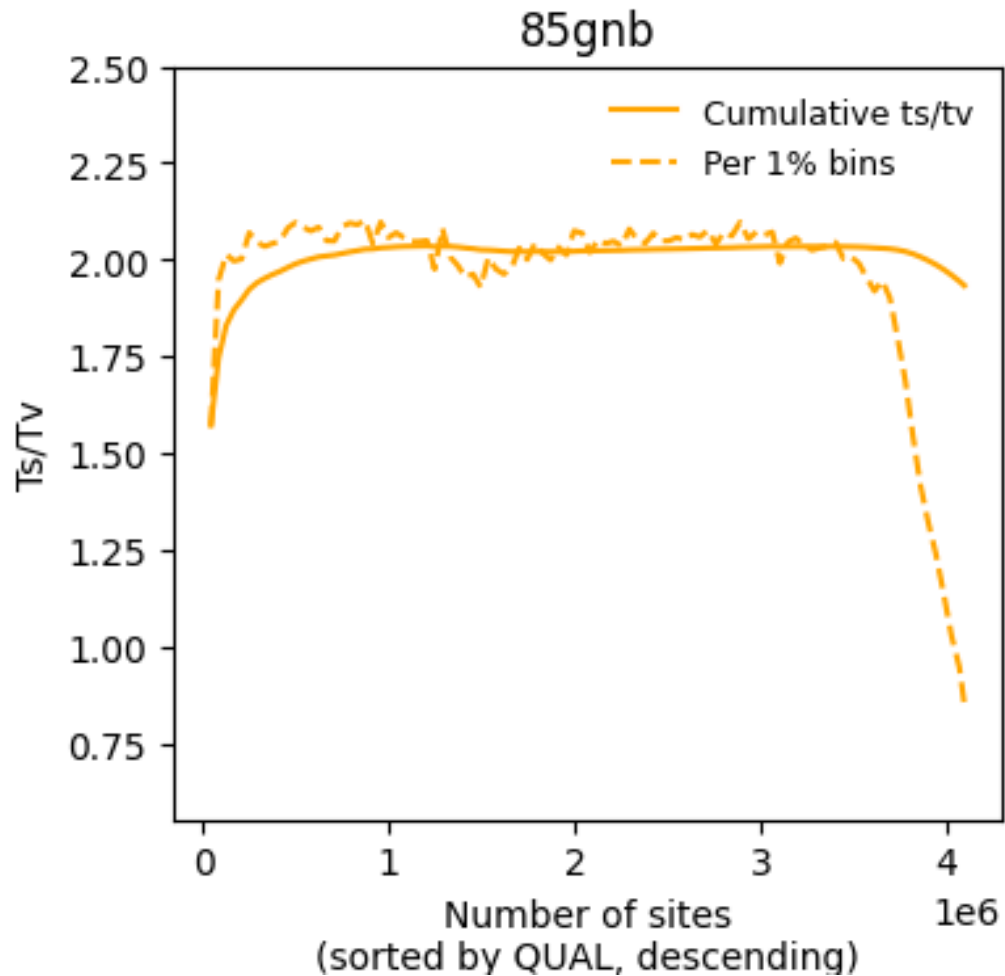
Number of SNPs by sample



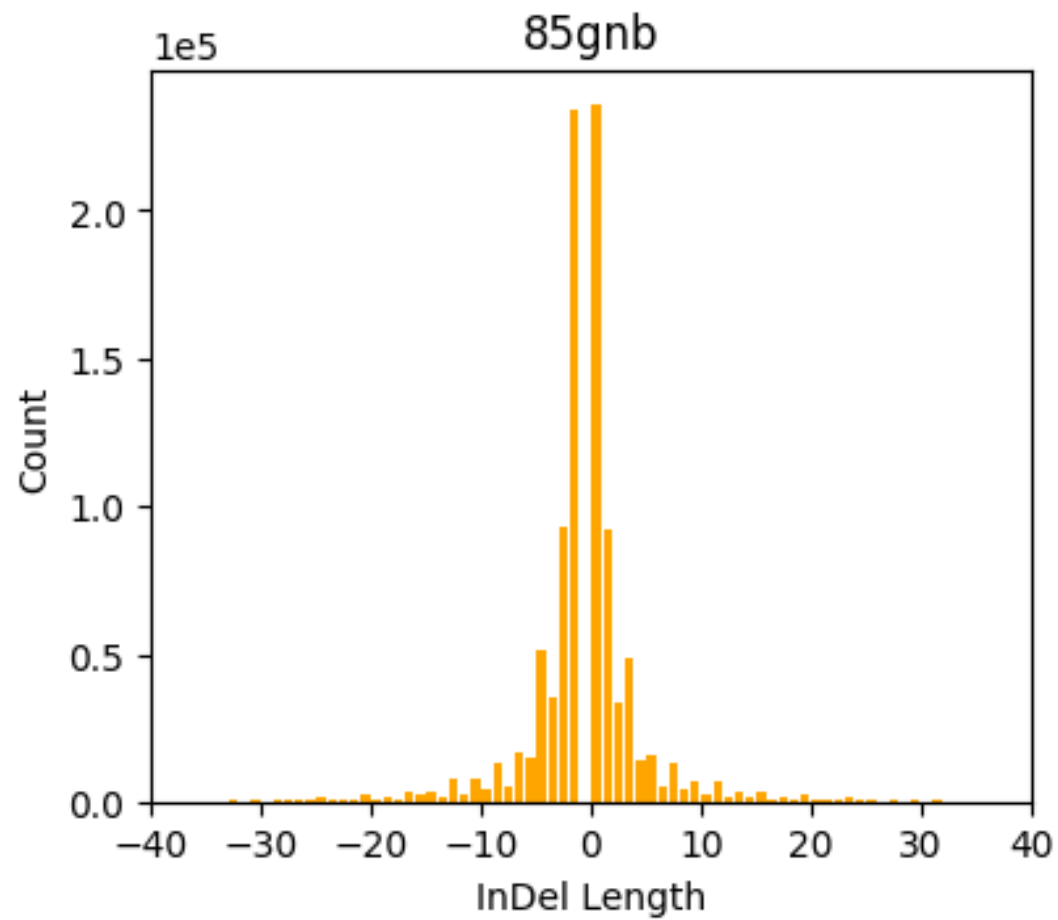
Number of indels by sample



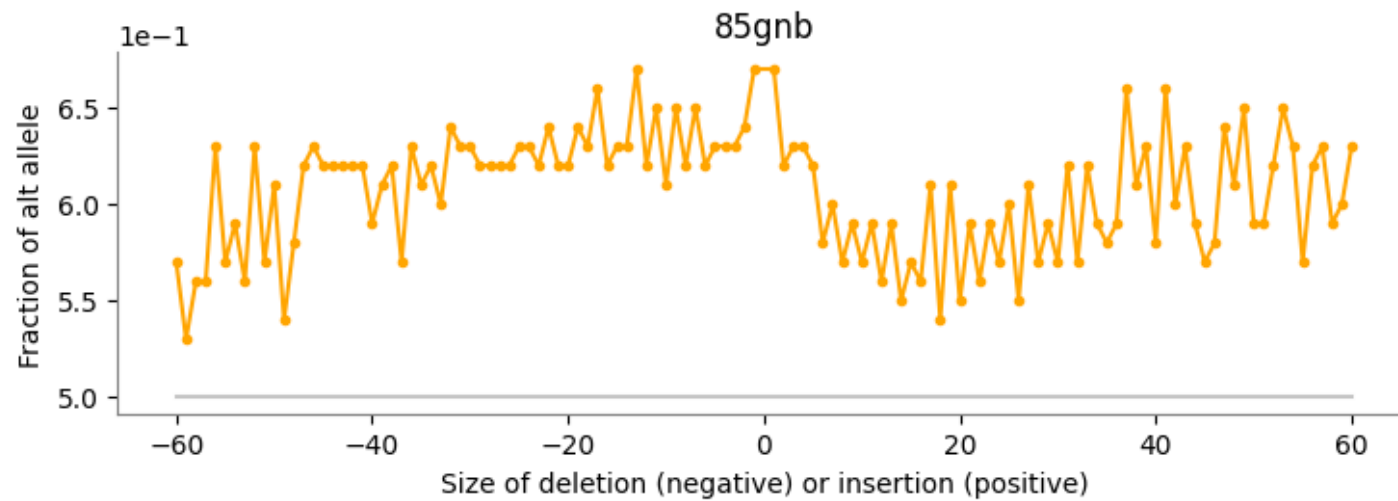
Ts/Tv stratified by QUAL



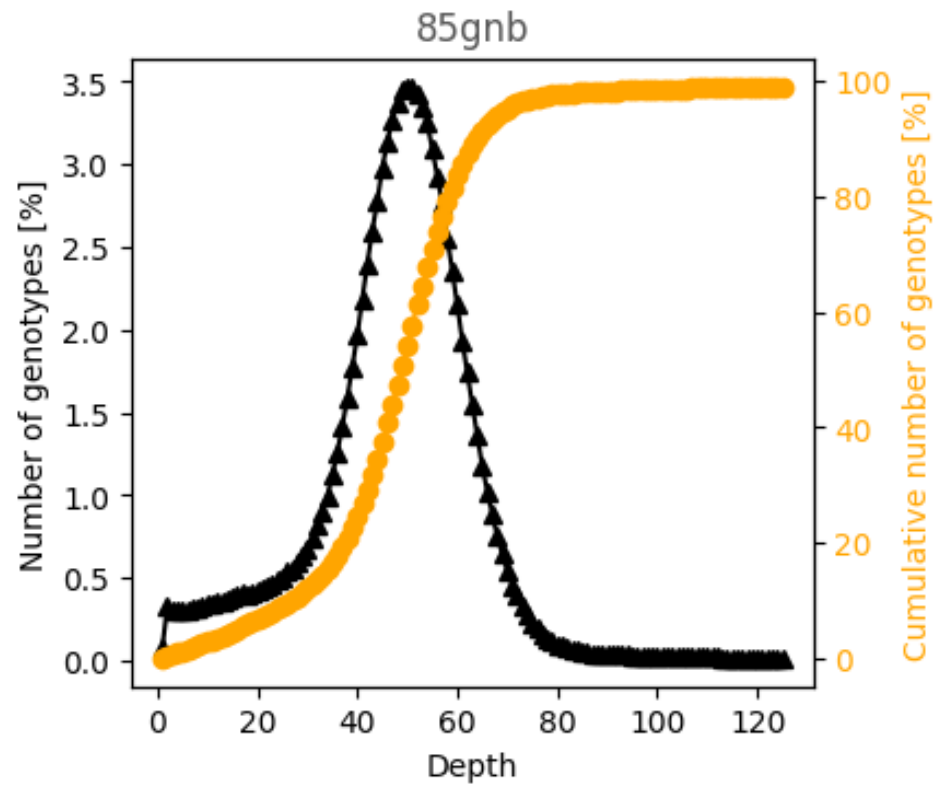
Indel distribution



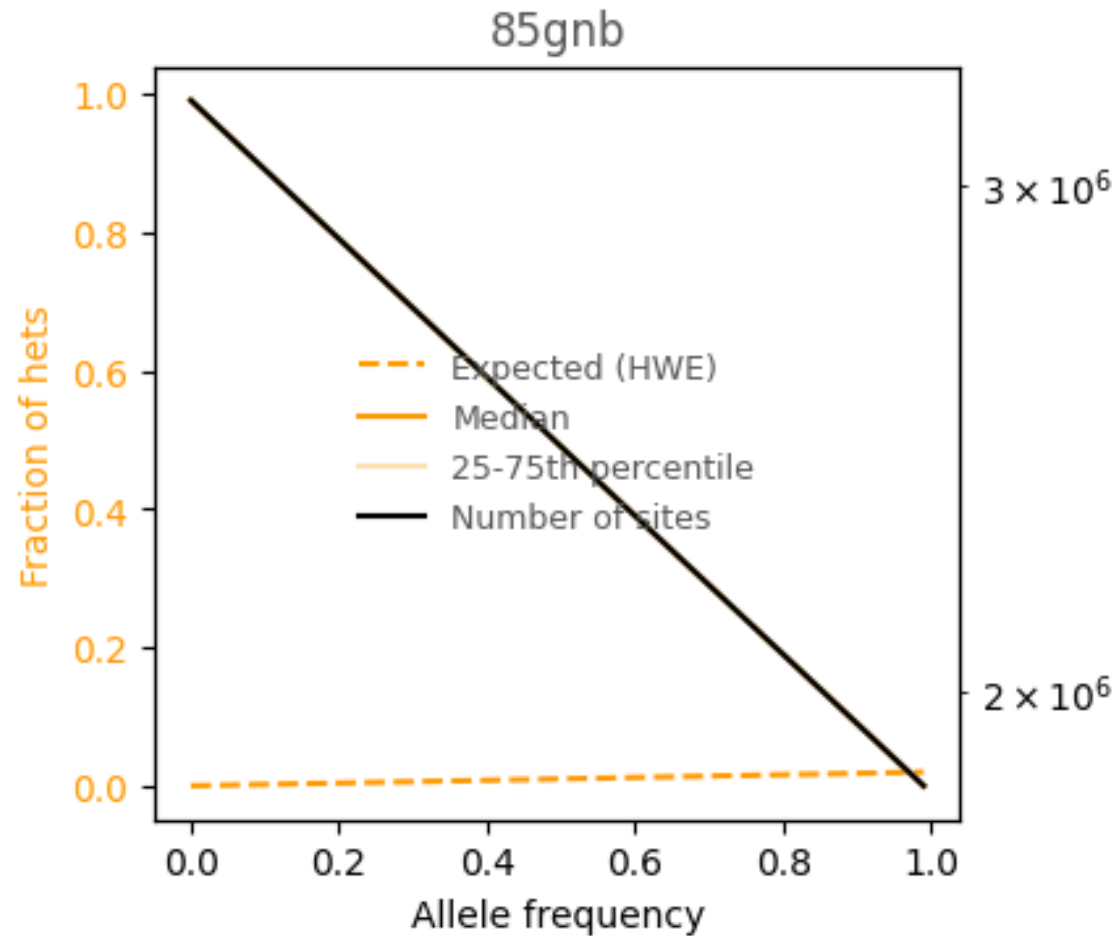
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

