

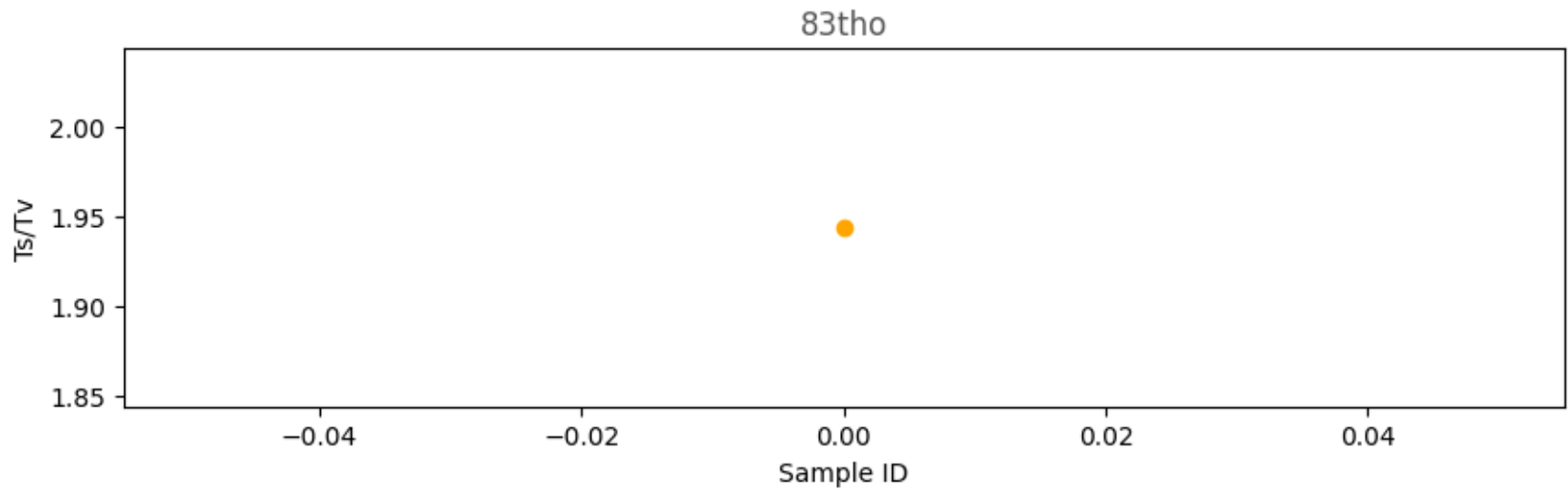
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
83tho	4,024,834	1.94	1.95	938,695	–	0	0
* frameshift ratio: out/(out+in)							

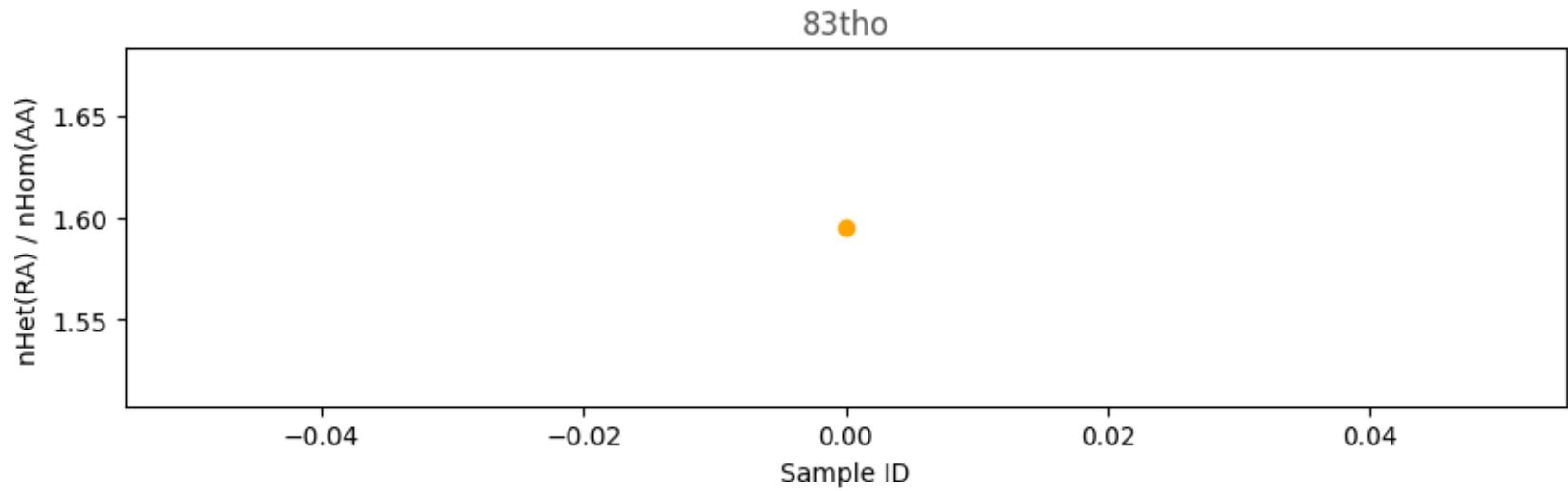
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
83tho	61.5%	1.92	65.9%	94,023	1,886

- 83tho .. /ngc/projects2/gm/data/archive/2022/variants/snv/83thoranm-103900719595-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01605-220916_A01176_BHYT22DSX3-EXT_LAB
 KA_NGCWGS-NGCWGS05353_22RKG019346_snv_germline_raw.haplotype_caller.vcf.gz

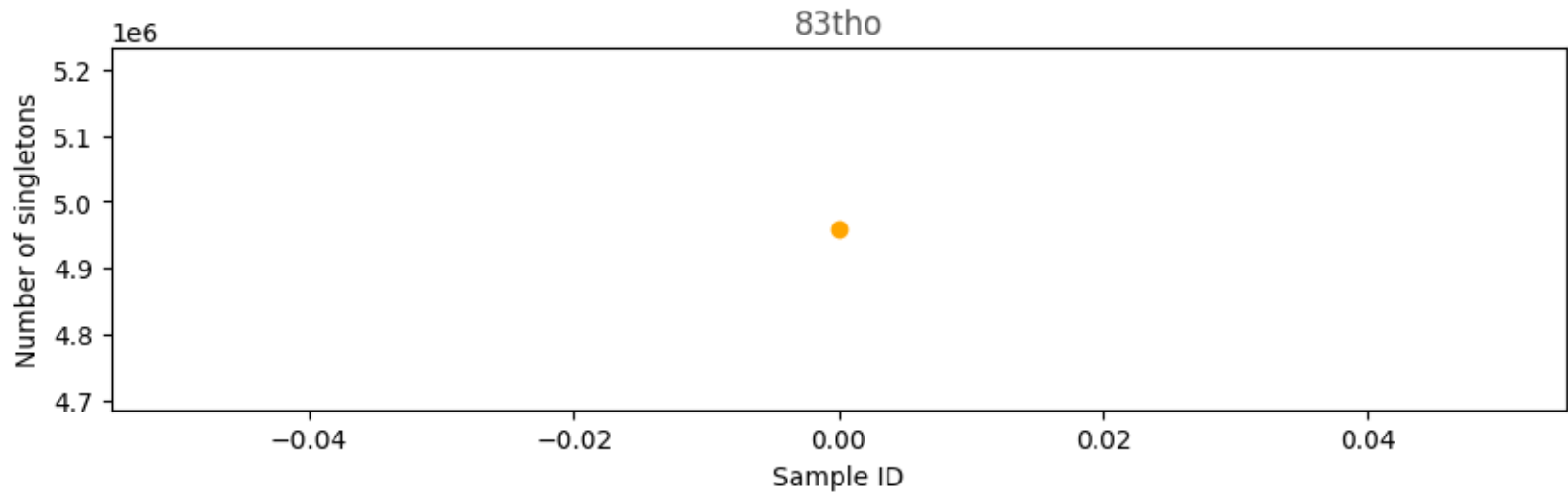
Ts/Tv by sample



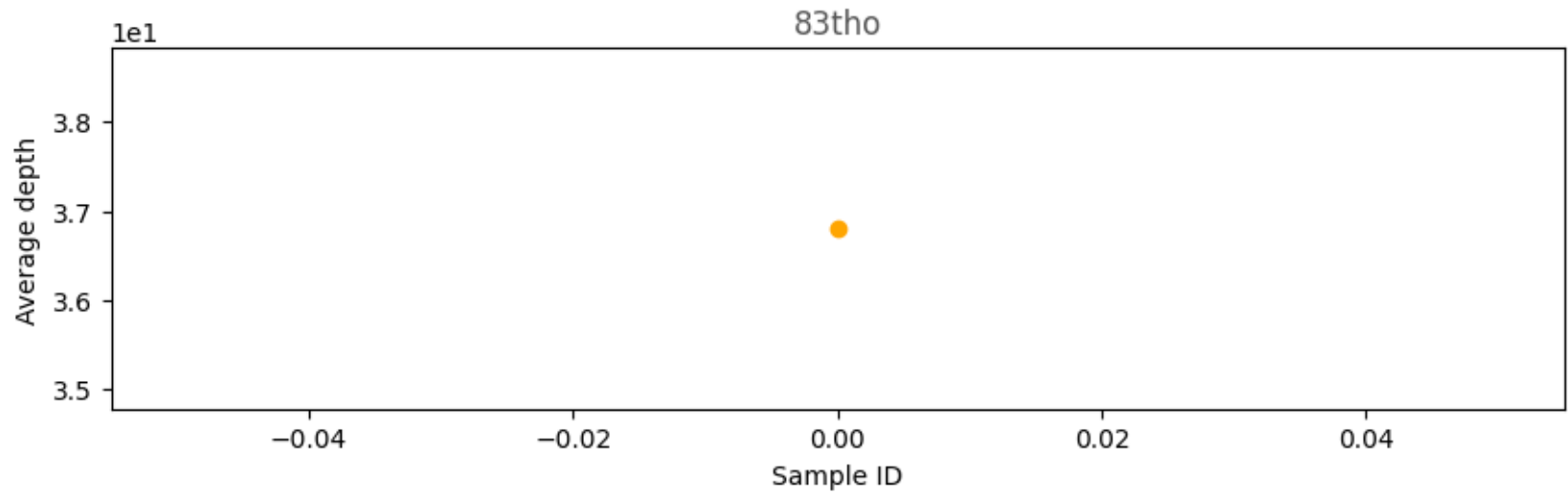
Hets vs non-ref Homs by sample



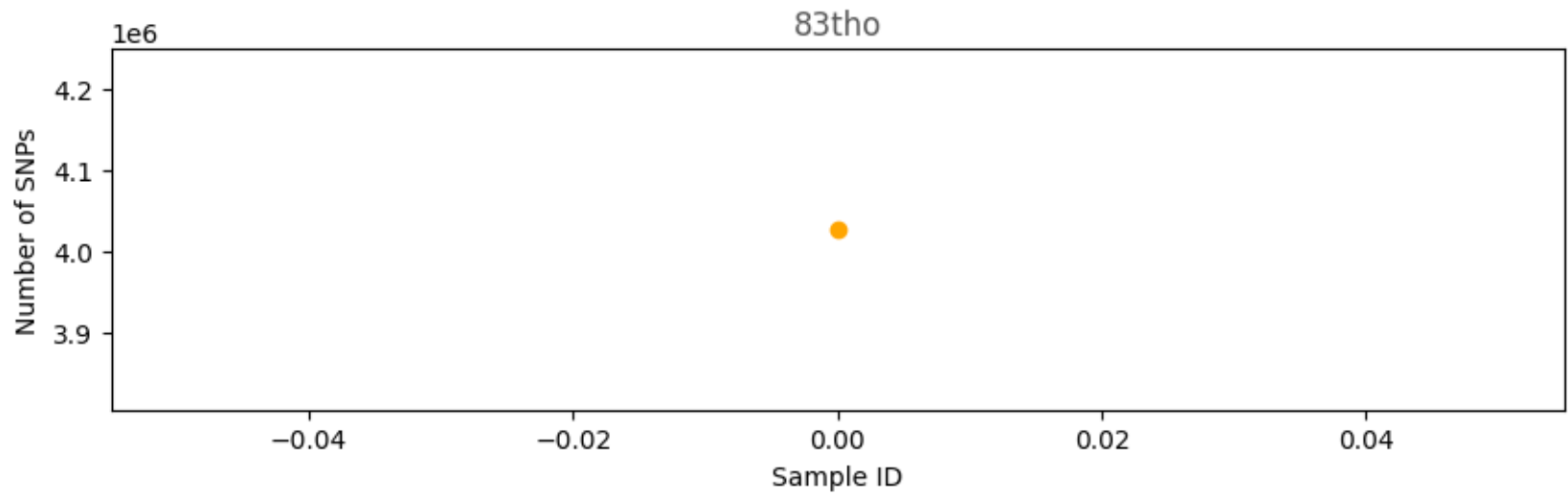
Singletons by sample (hets and homs)



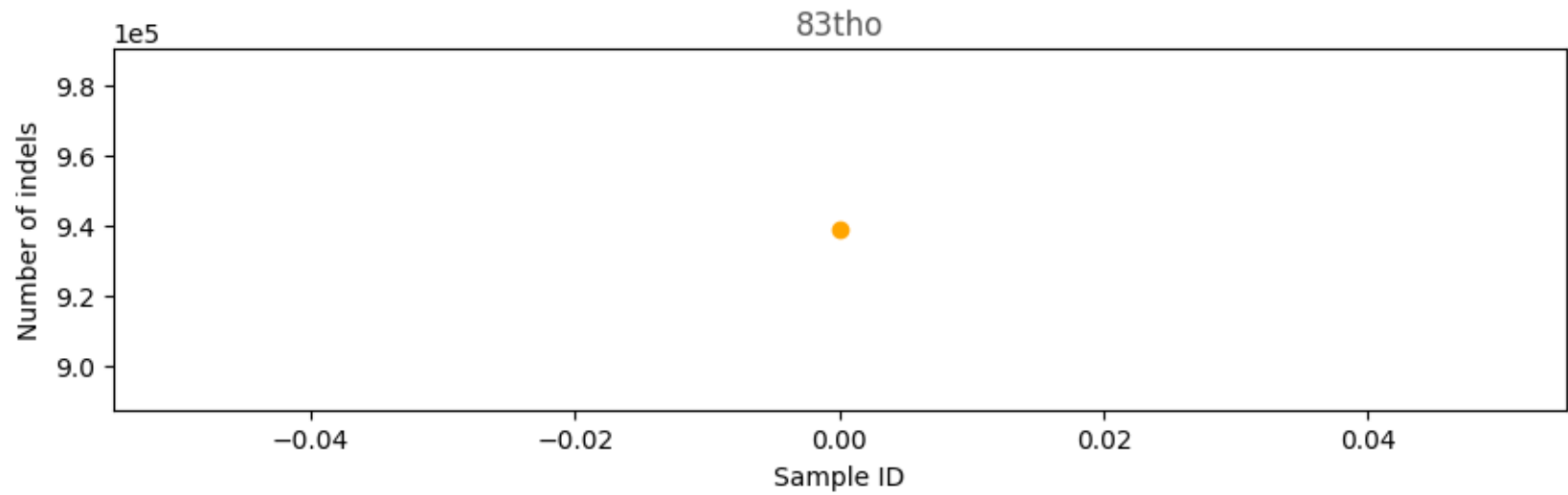
Average depth by sample



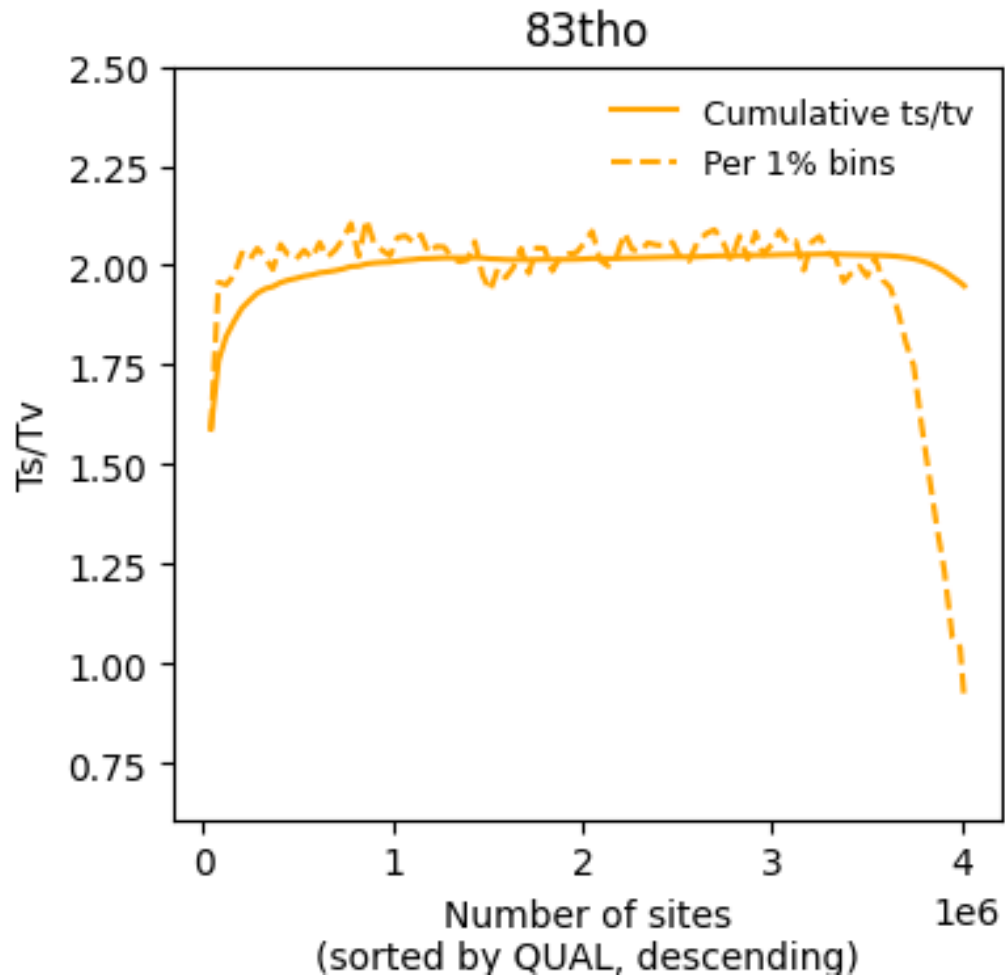
Number of SNPs by sample



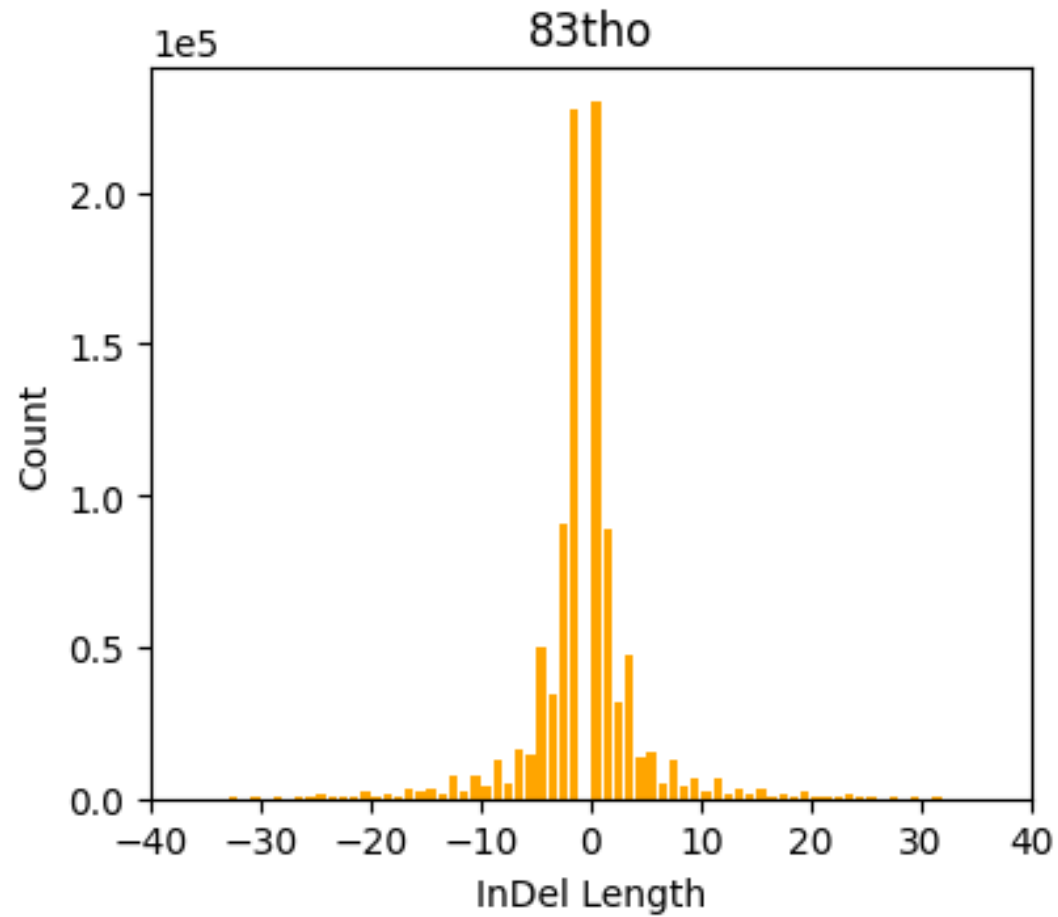
Number of indels by sample



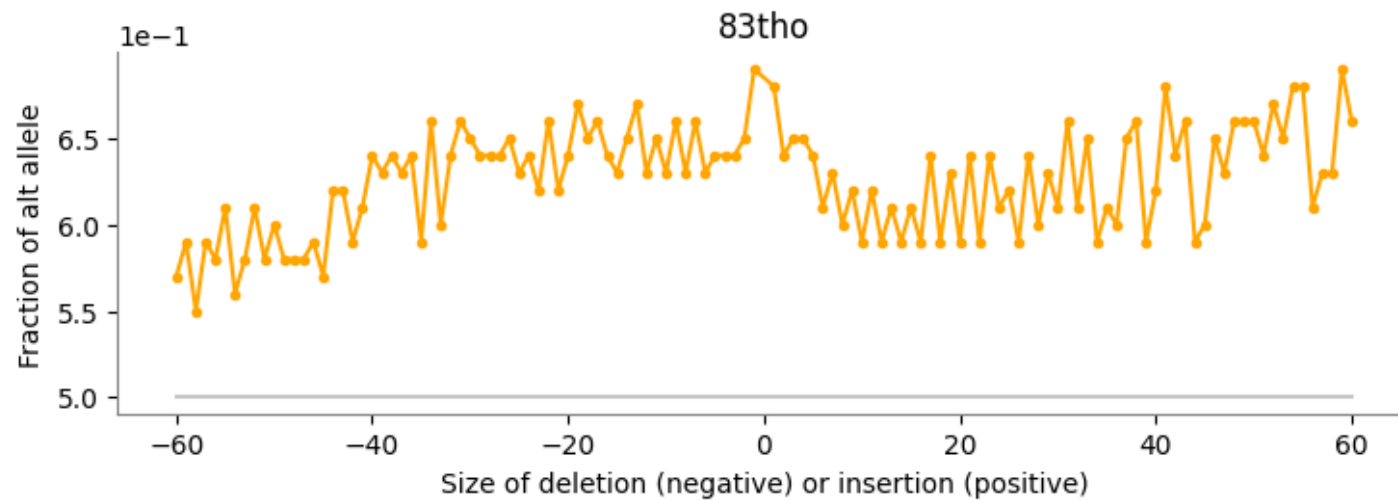
Ts/Tv stratified by QUAL



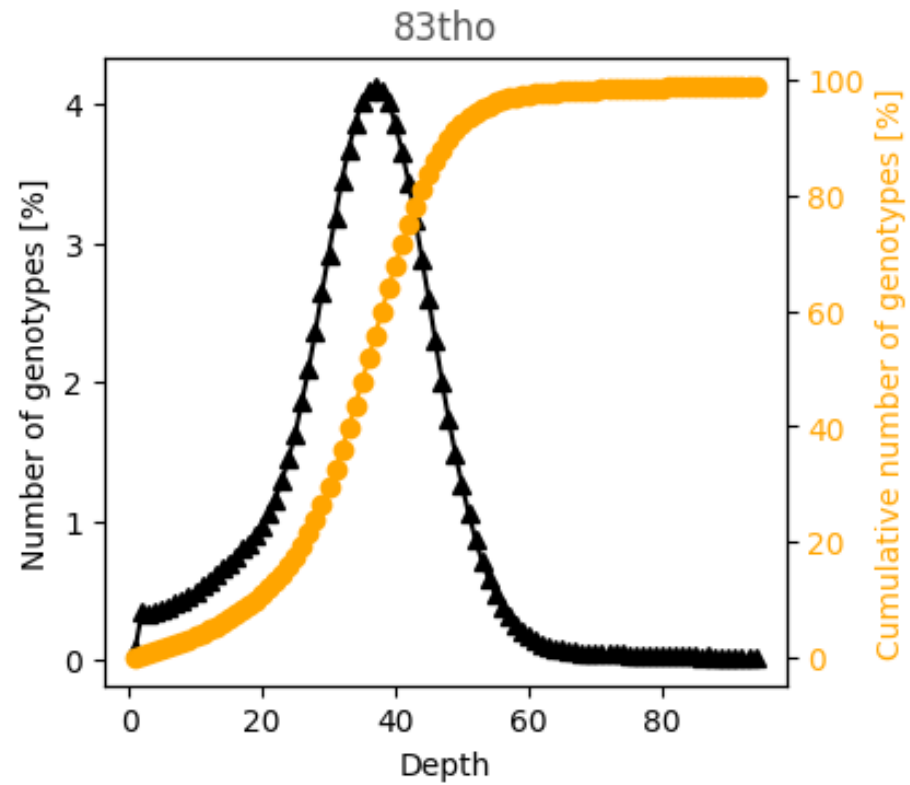
Indel distribution



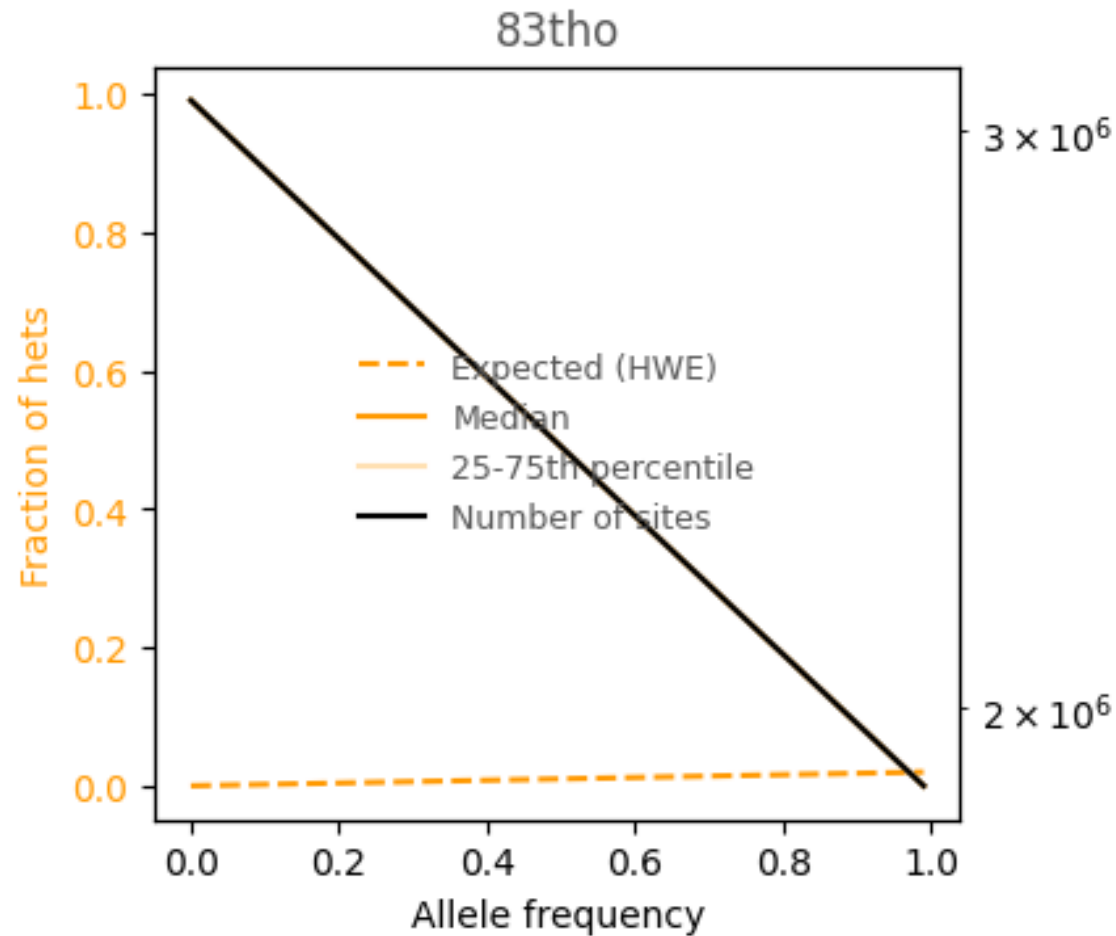
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

