

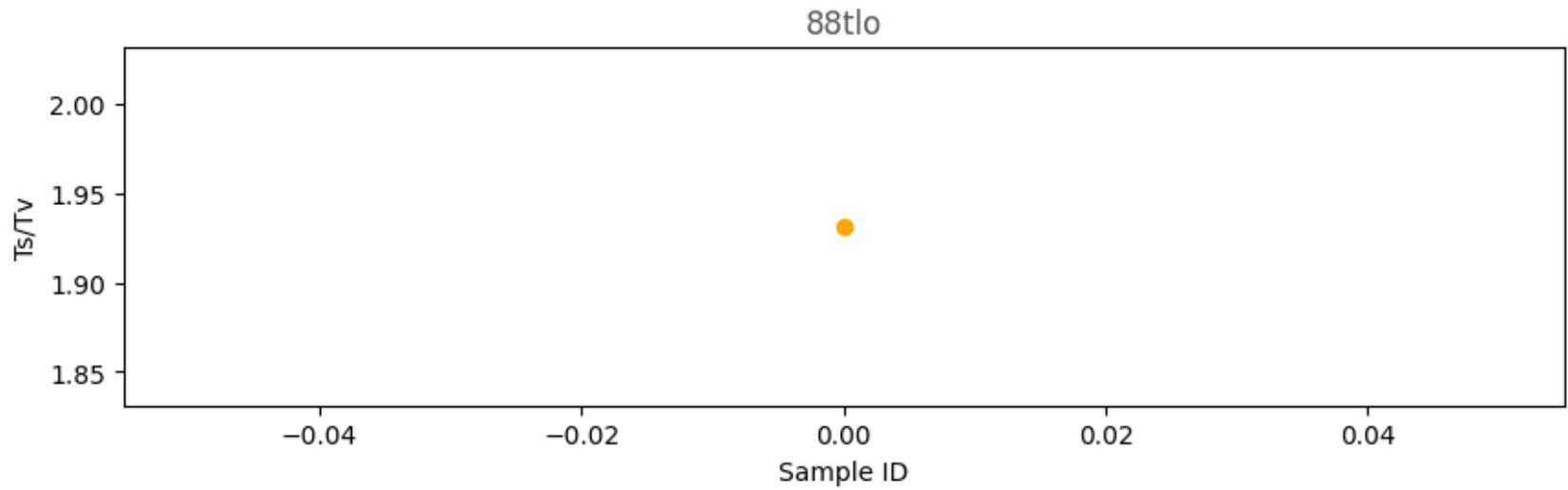
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
88tlo	4,075,344	1.93	1.94	944,188	–	0	0
* frameshift ratio: out/(out+in)							

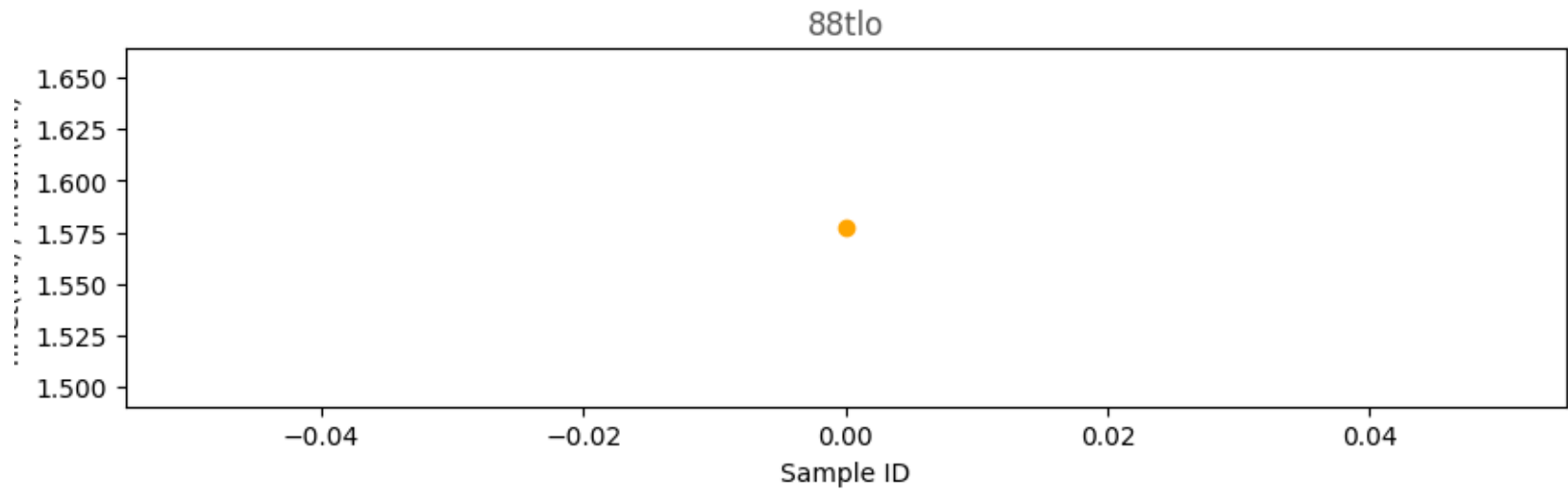
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
88tlo	61.2%	1.90	65.4%	94,914	1,945

- 88tlo .. /ngc/projects2/gm/data/archive/2022/variants/snv/88tloclim-103902945207-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01150-220824_A01961_AHMW7KDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS04901_snv_germline_raw.haplotype_caller.vcf.gz

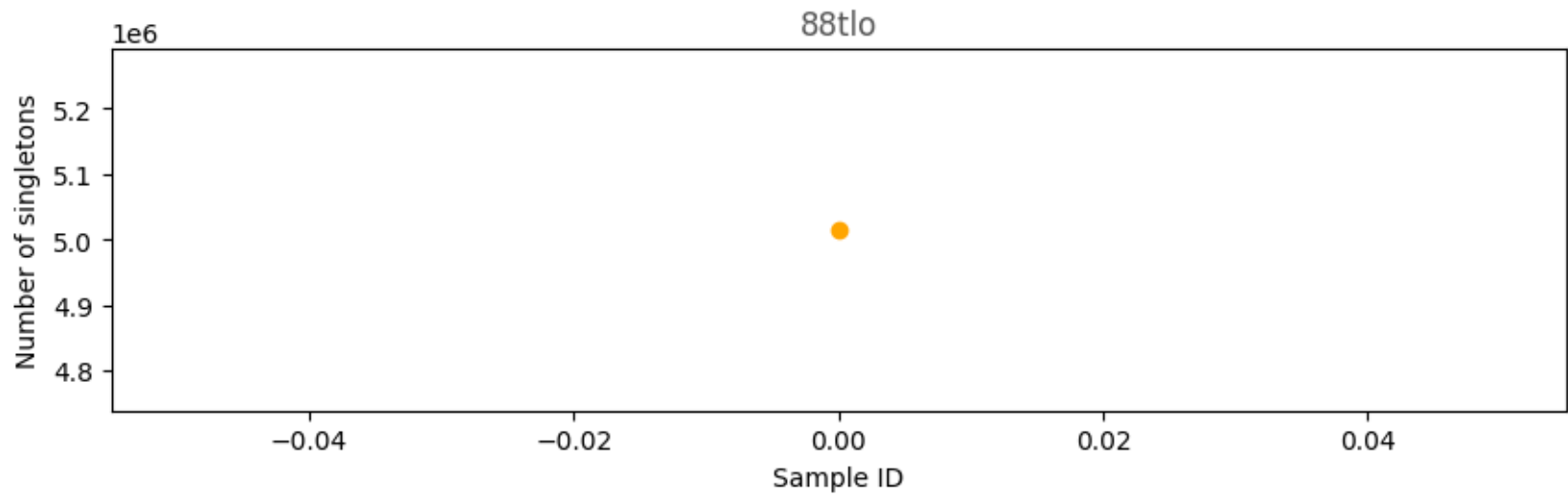
Ts/Tv by sample



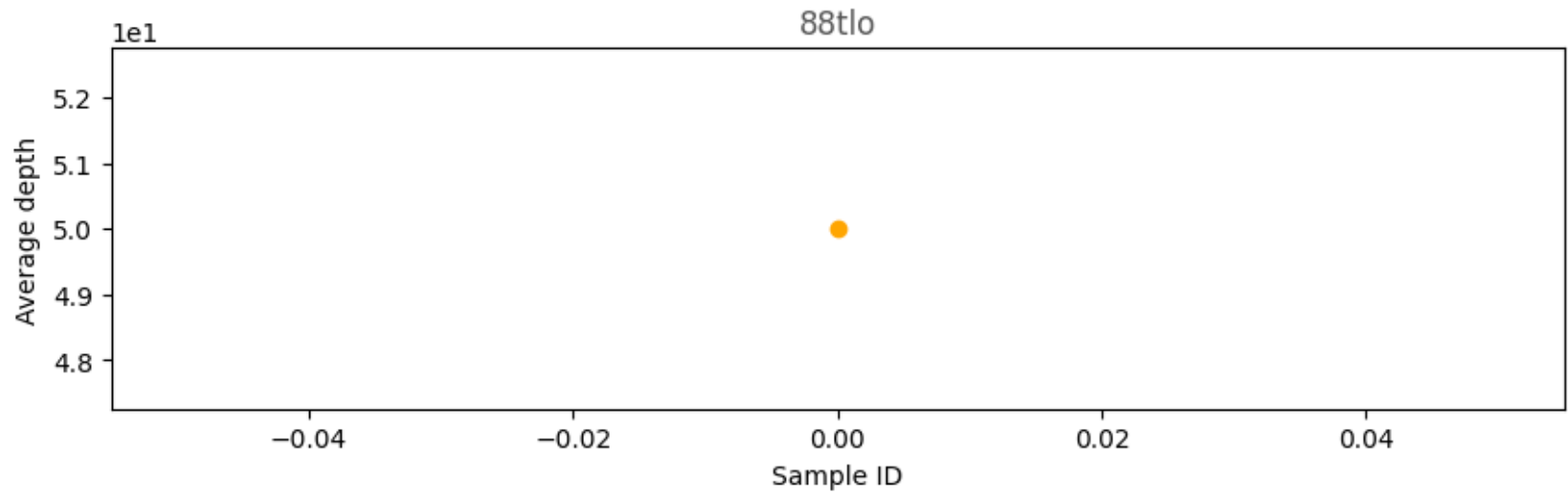
Hets vs non-ref Homs by sample



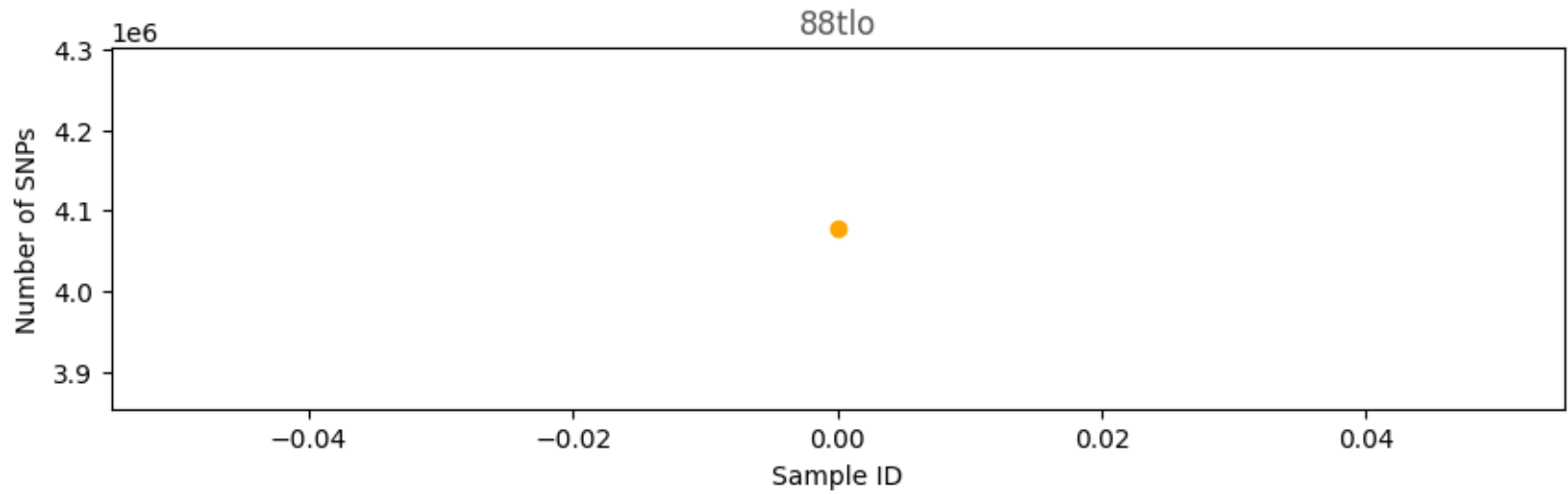
Singletons by sample (hets and homs)



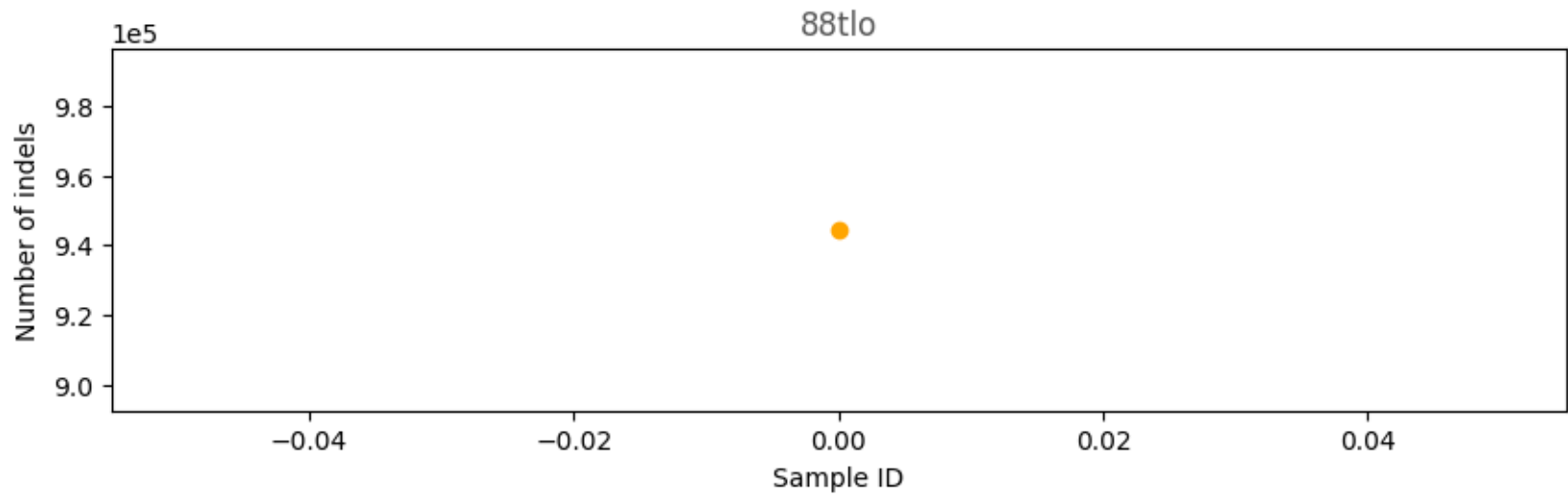
Average depth by sample



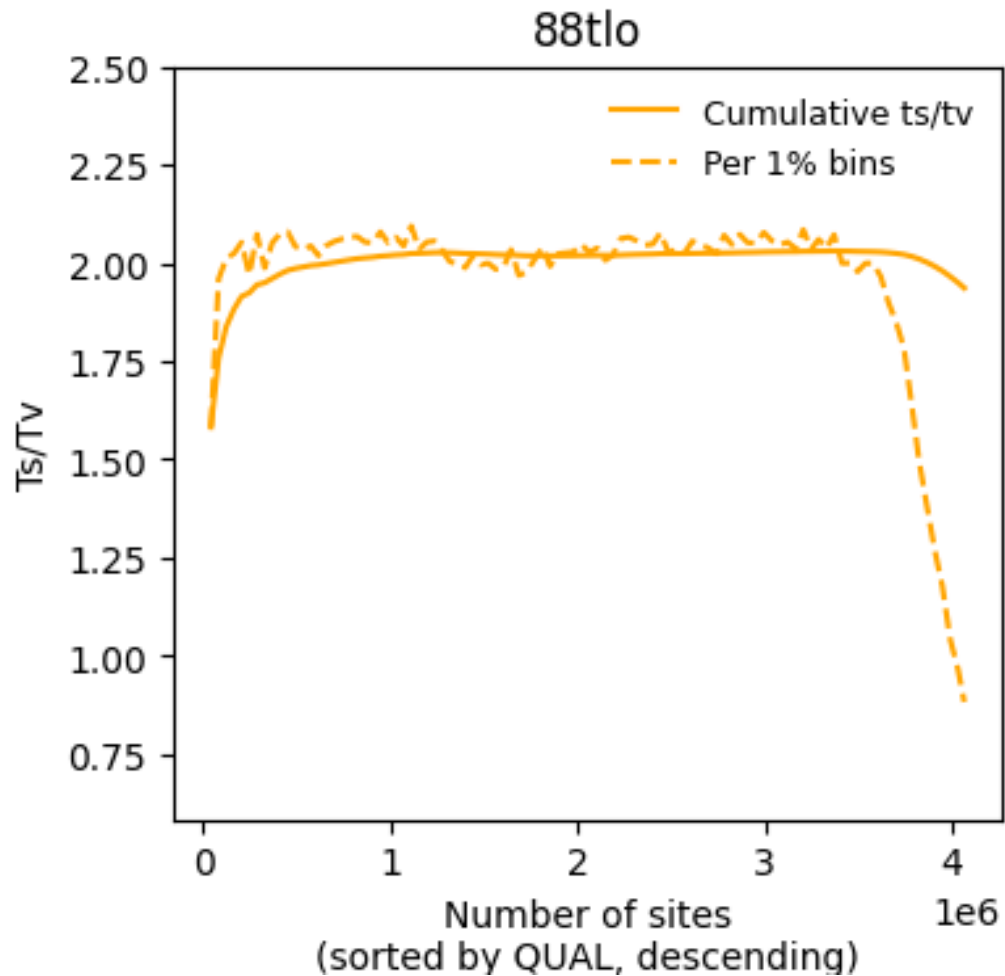
Number of SNPs by sample



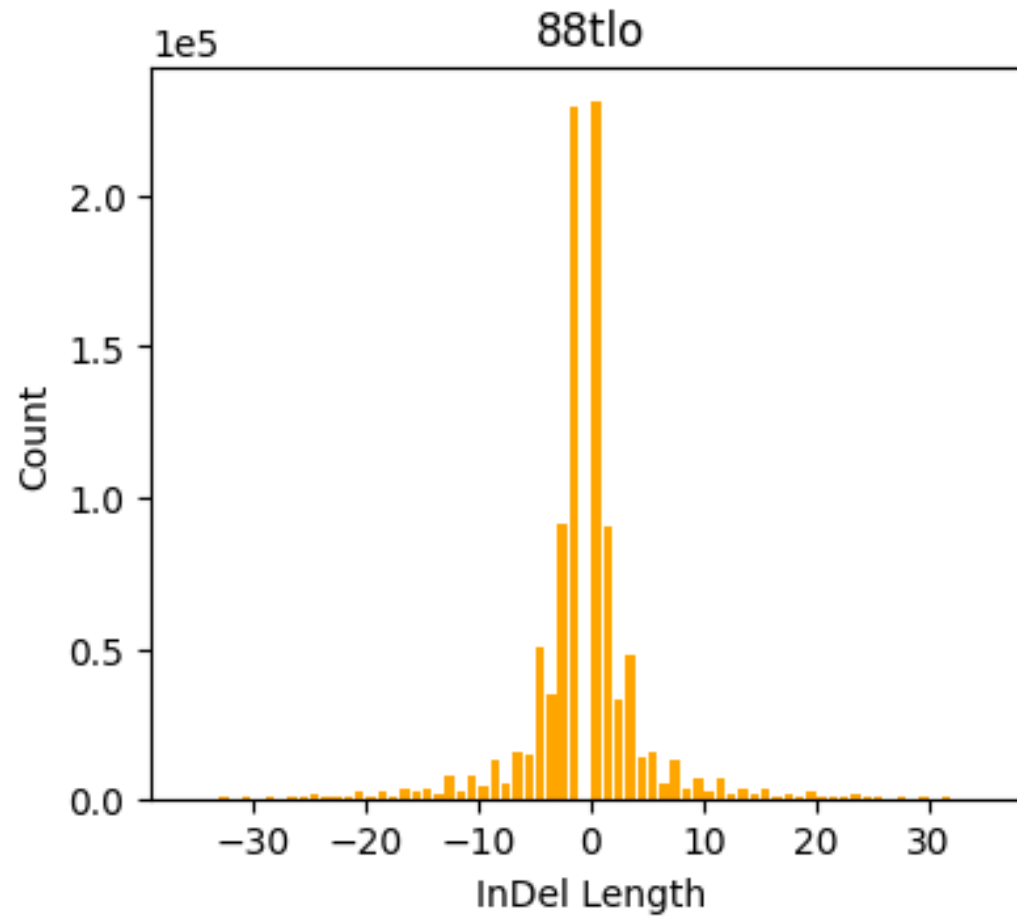
Number of indels by sample



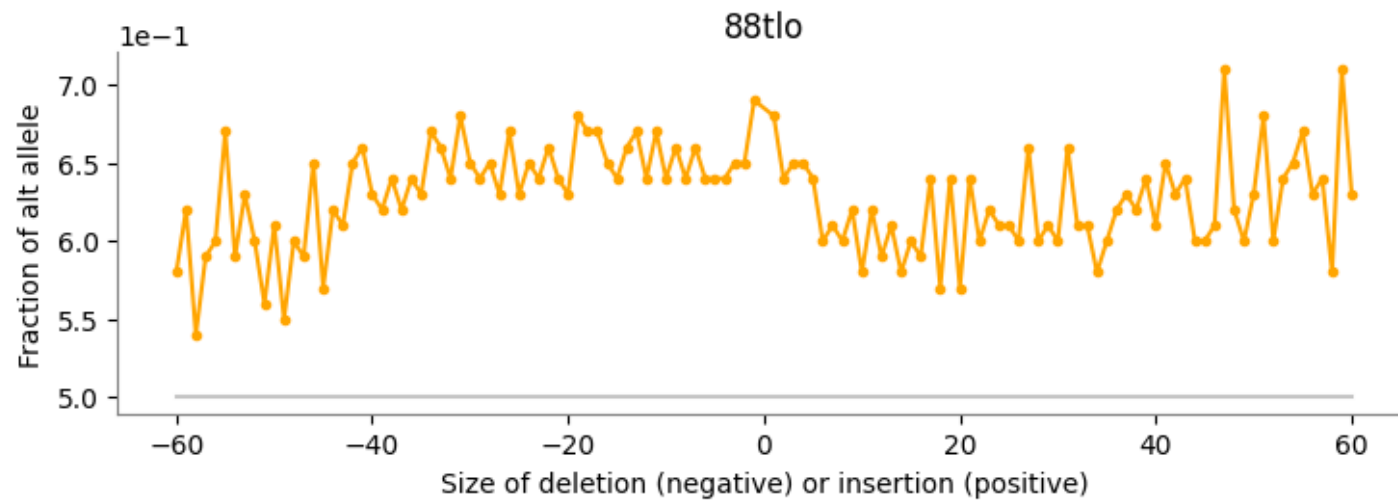
Ts/Tv stratified by QUAL



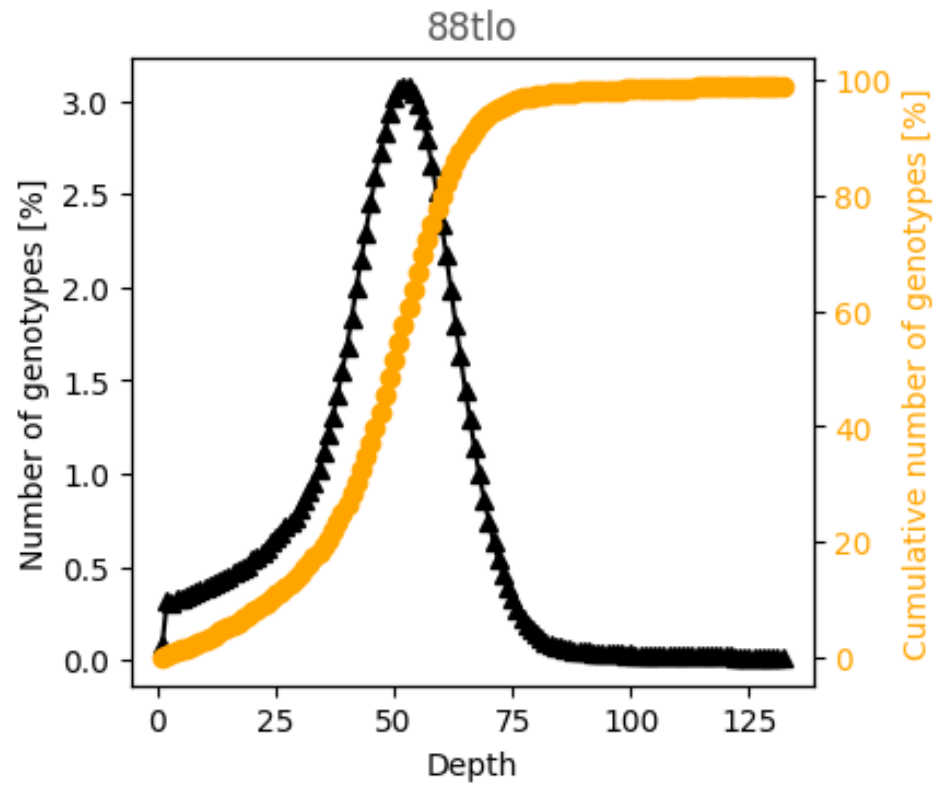
Indel distribution



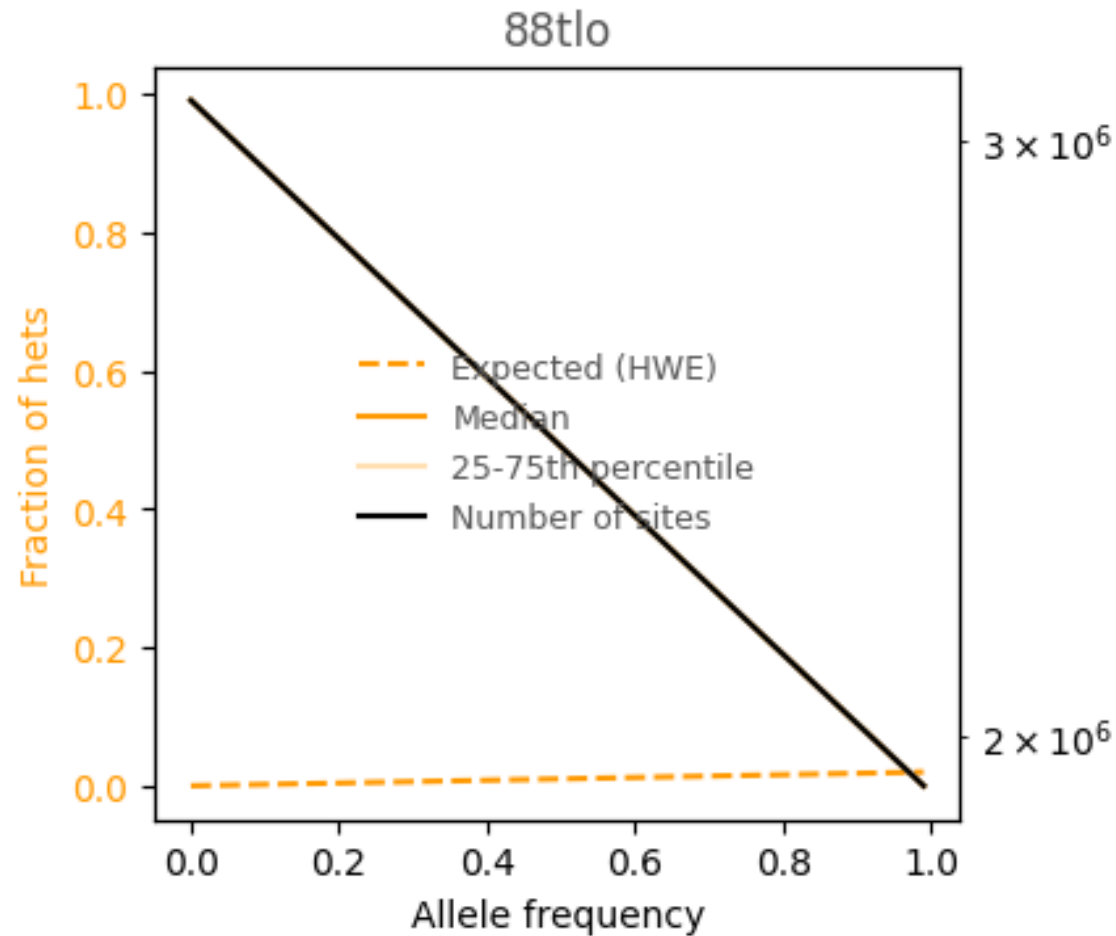
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

