

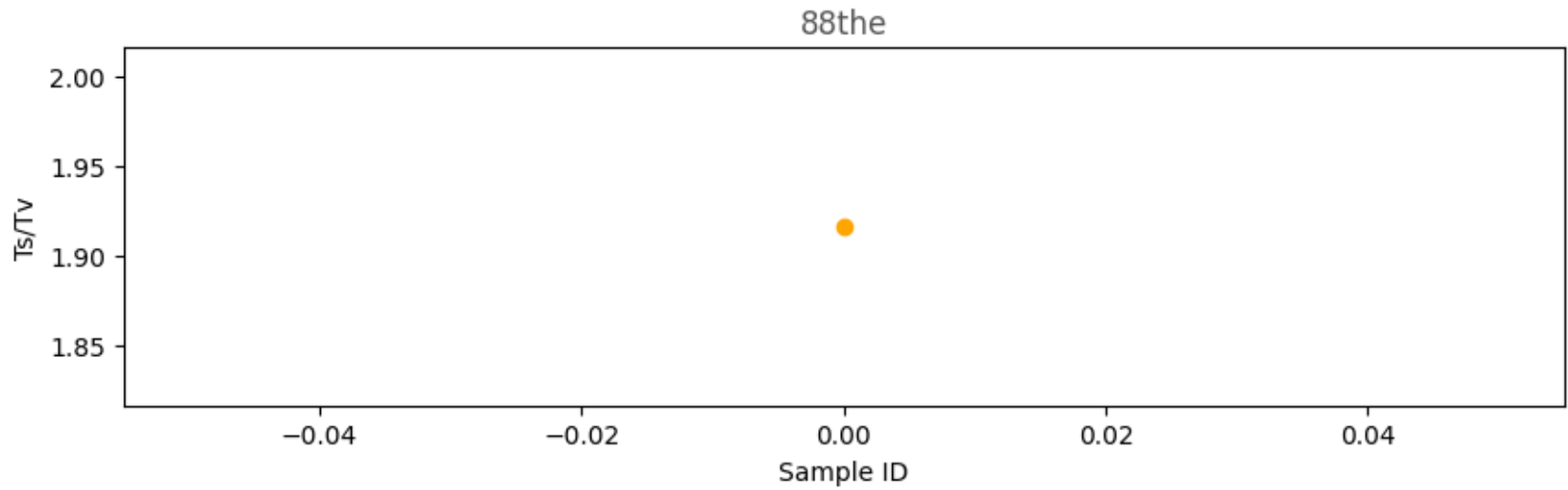
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
88the	4,080,346	1.92	1.92	955,078	–	0	0
* frameshift ratio: out/(out+in)							

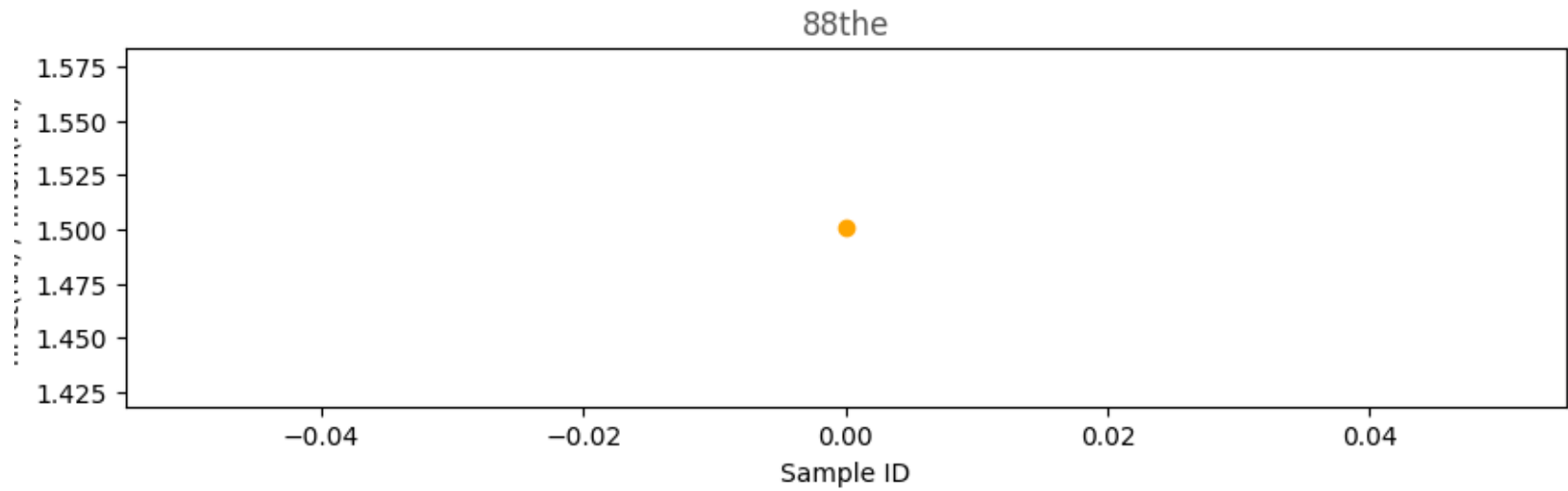
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
88the	60.0%	1.87	65.1%	101,899	2,123

- 88the .. /ngc/projects2/gm/data/archive/2022/variants/snv/88thesrtm-103897532926-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220803_A00559_AHNW22DSX3-RHGM_LABKA_WGSA_KUT-WGSAKUT04752_22RKG018398x01_snv_germline_raw.haplotype_caller.vcf.gz

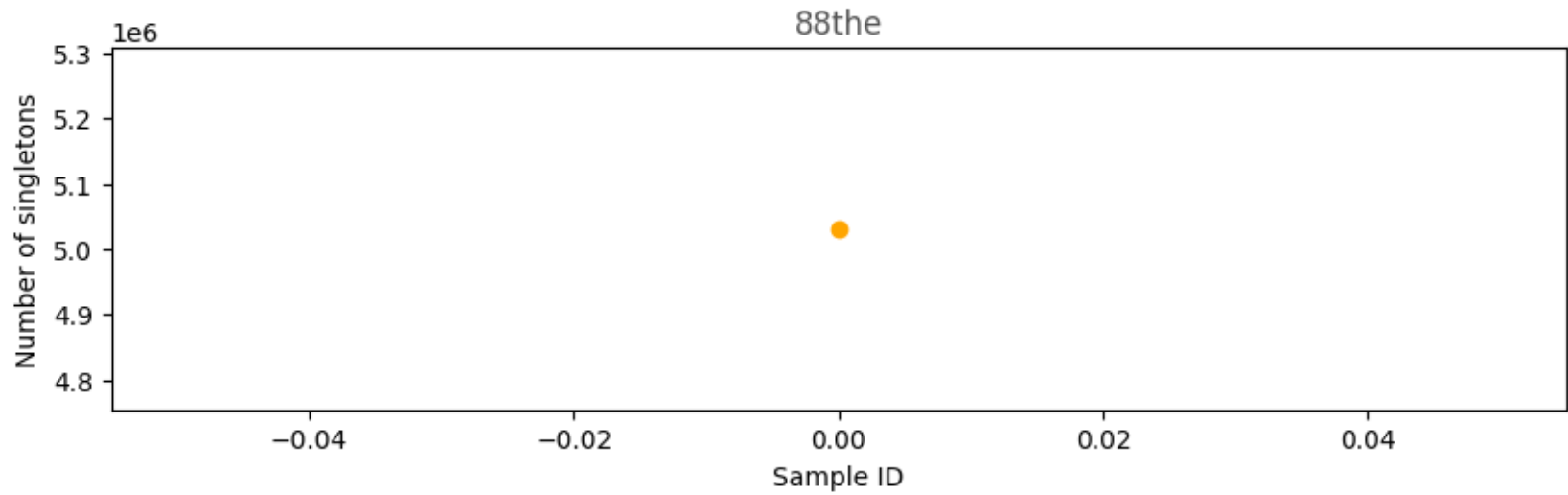
Ts/Tv by sample



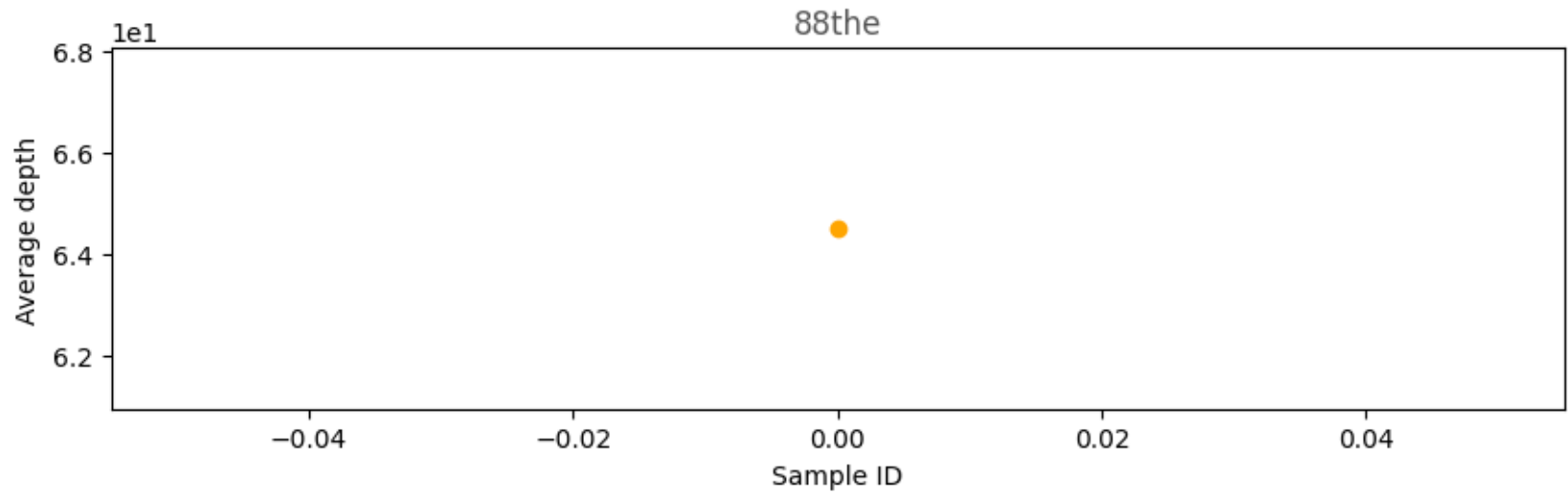
Hets vs non-ref Homs by sample



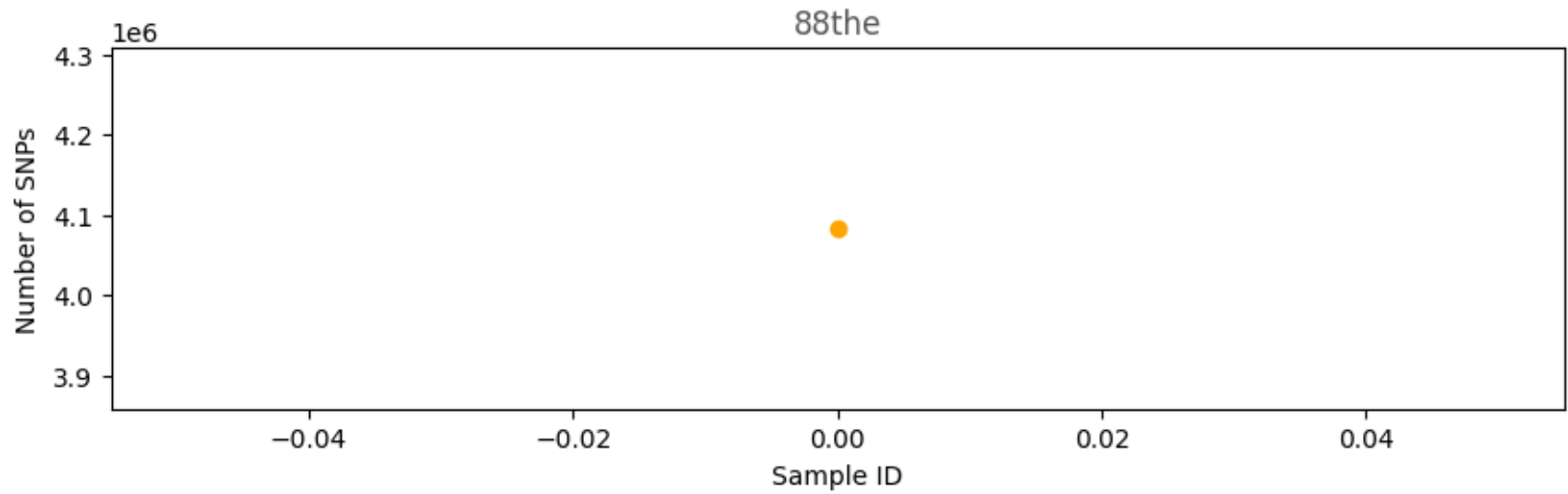
Singletons by sample (hets and homs)



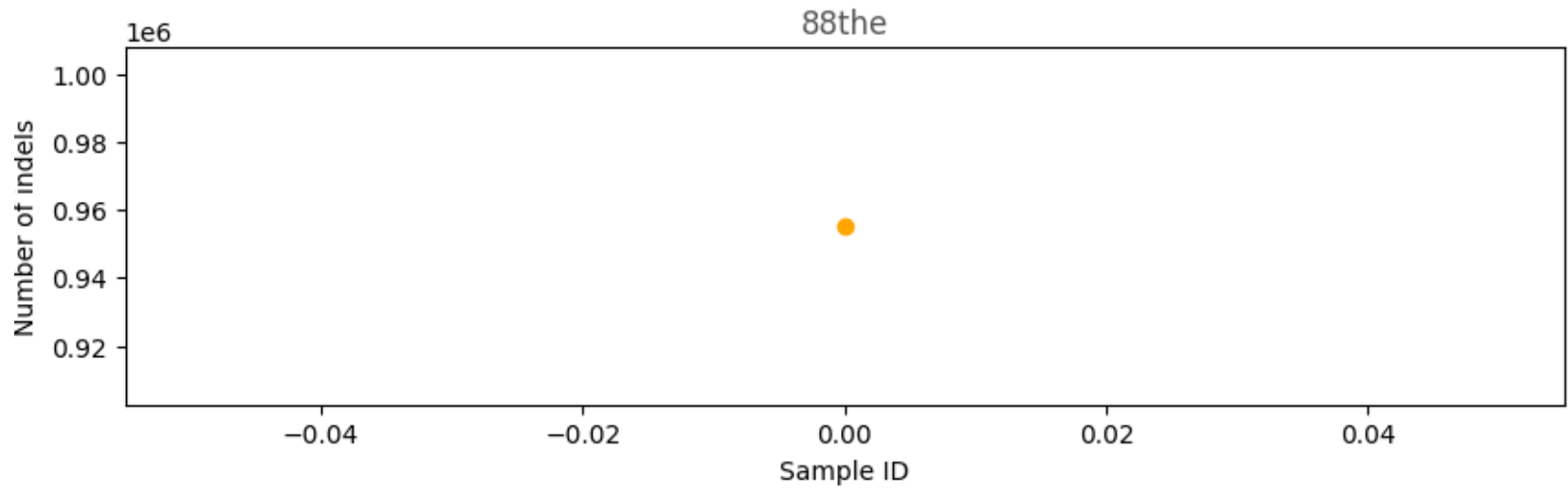
Average depth by sample



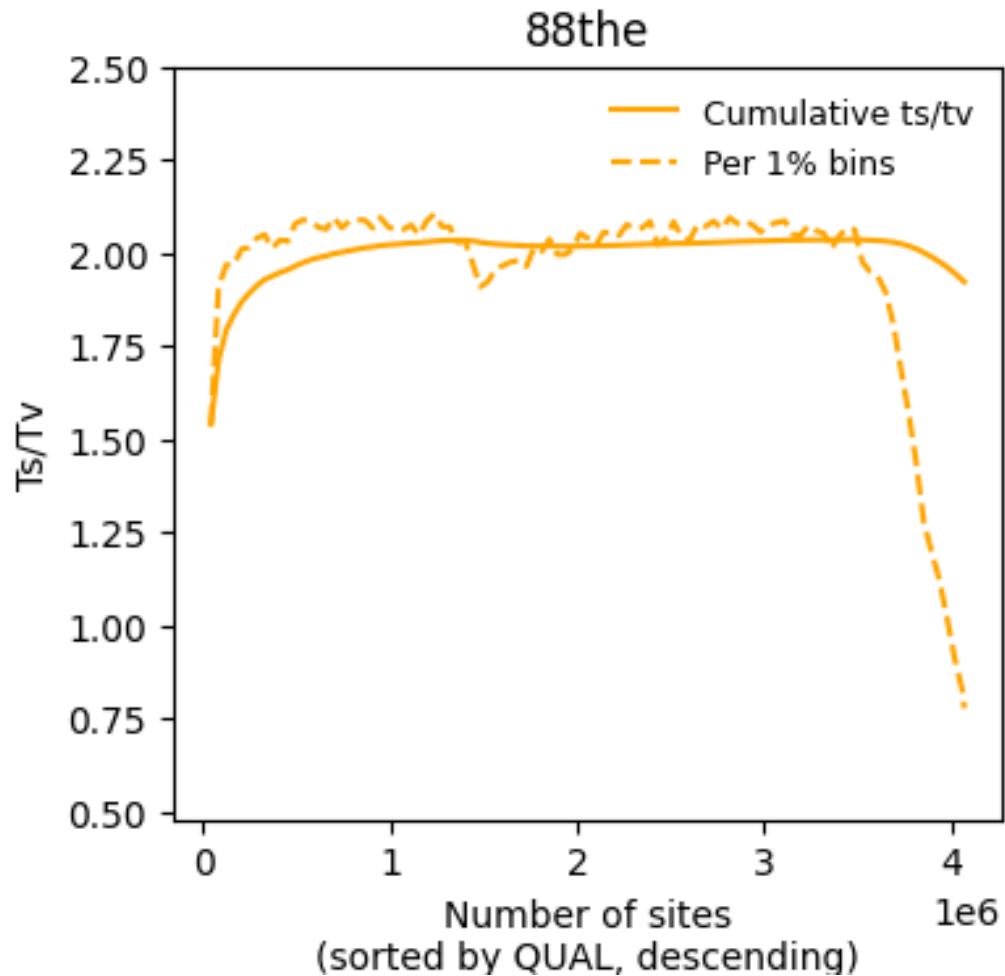
Number of SNPs by sample



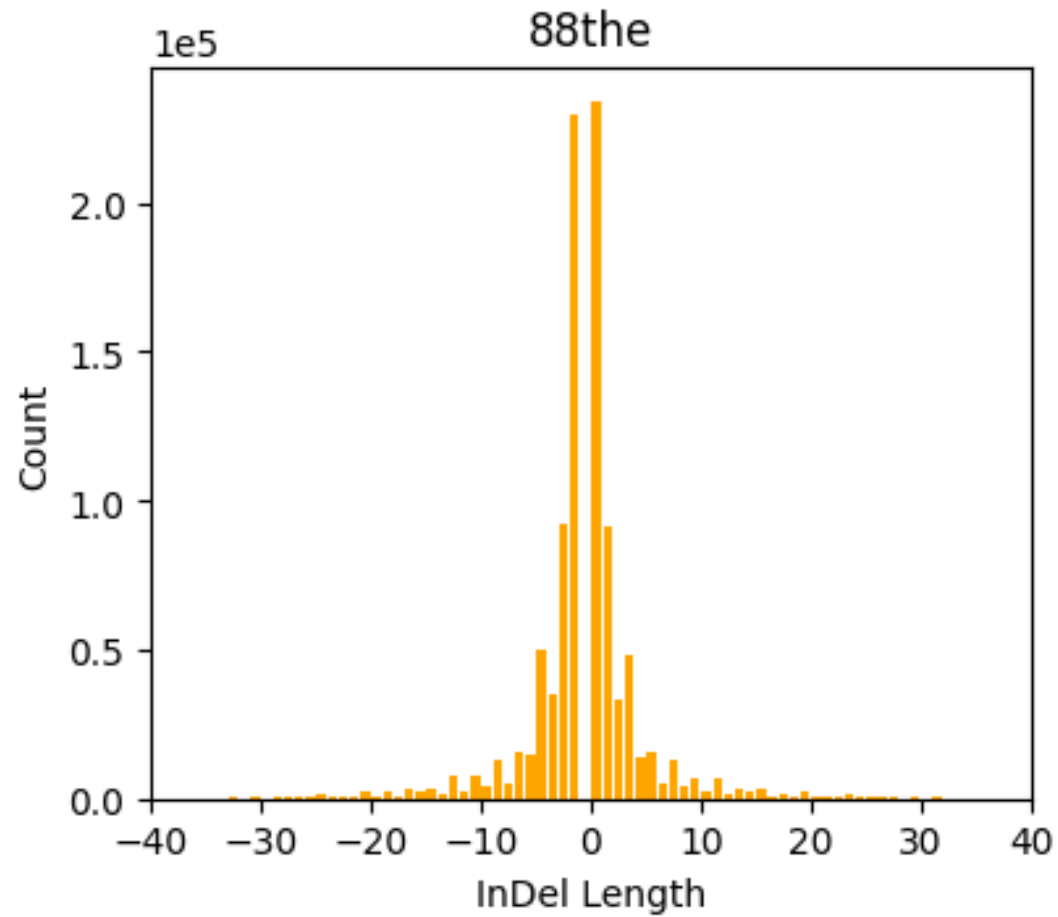
Number of indels by sample



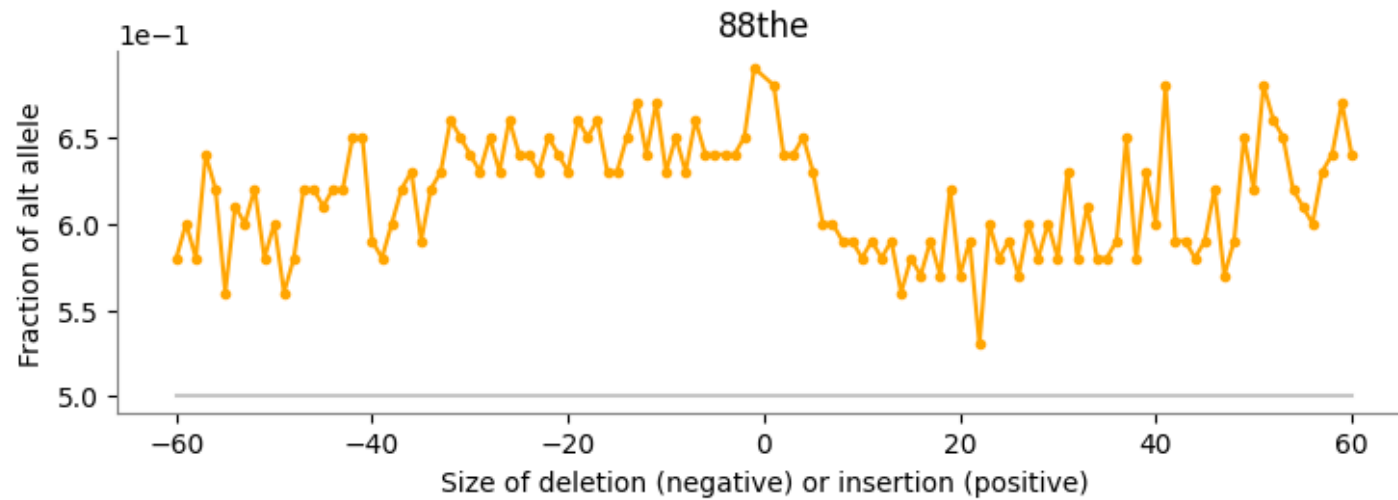
Ts/Tv stratified by QUAL



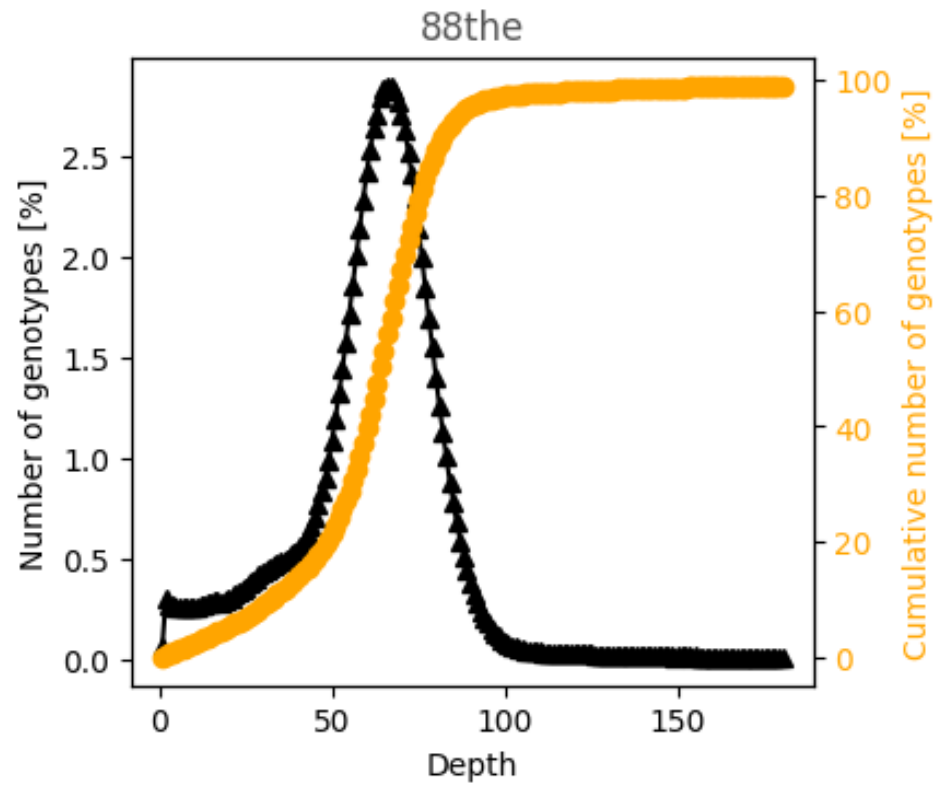
Indel distribution



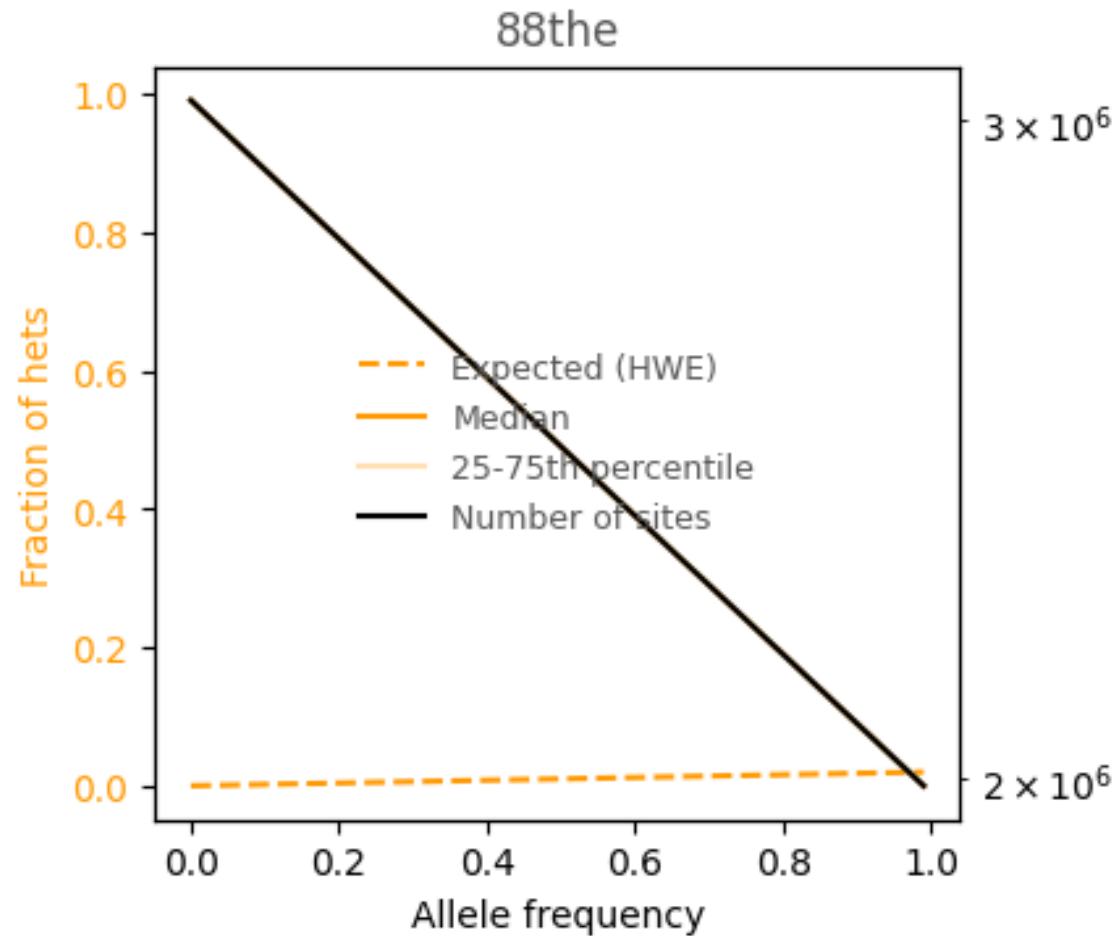
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

