

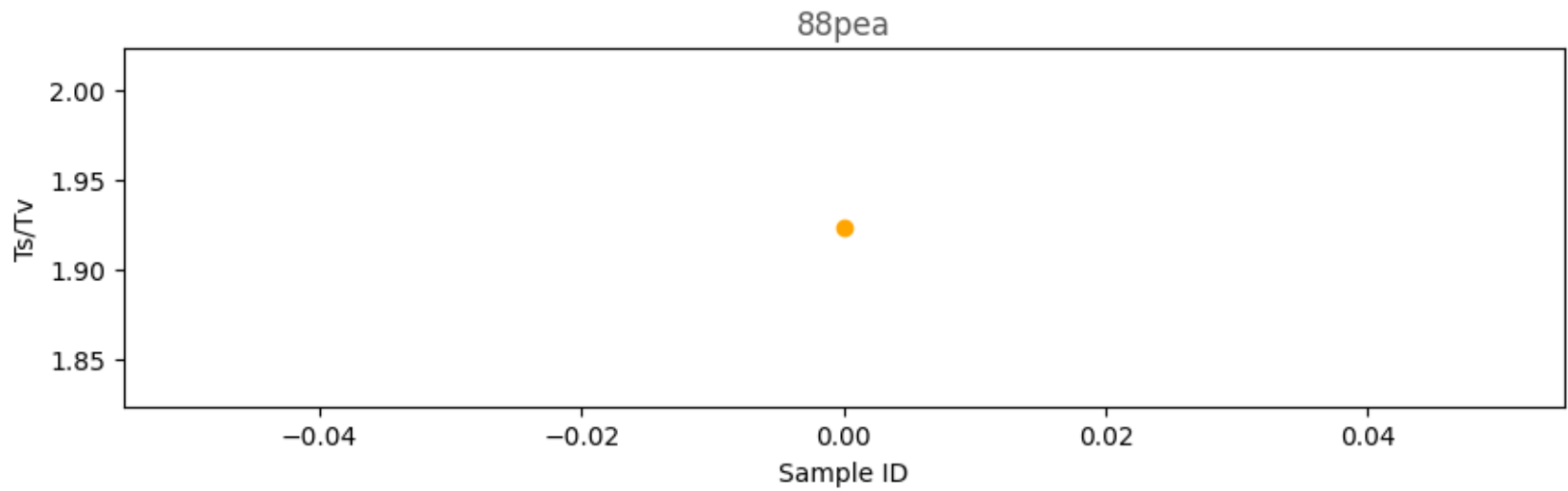
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
88pea	4,132,264	1.92	1.93	960,485	–	0	0
* frameshift ratio: out/(out+in)							

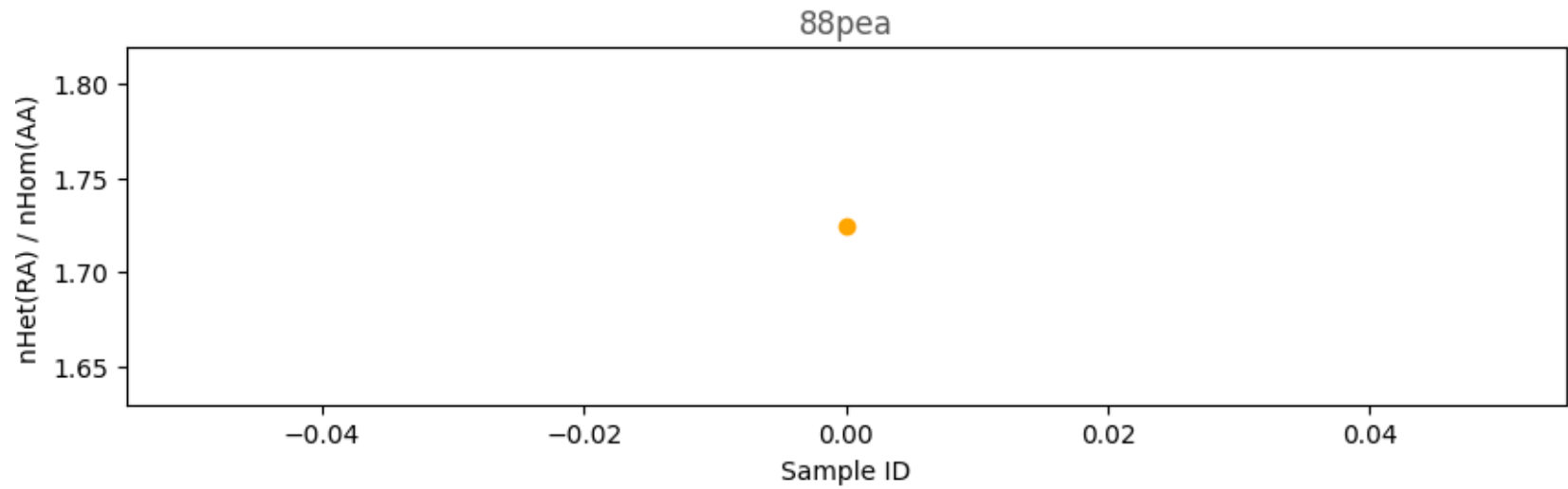
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
88pea	63.3%	1.89	68.4%	103,275	2,181

- 88pea .. /ngc/projects2/gm/data/archive/2022/variants/snv/88pearadf-103872029694-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00813-220525_A00559_BHJ5JJDSX3-EXT_LAB
KA_NGCWGS-NGCWGS04319_snv_germline_raw.haplotype_caller.vcf.gz

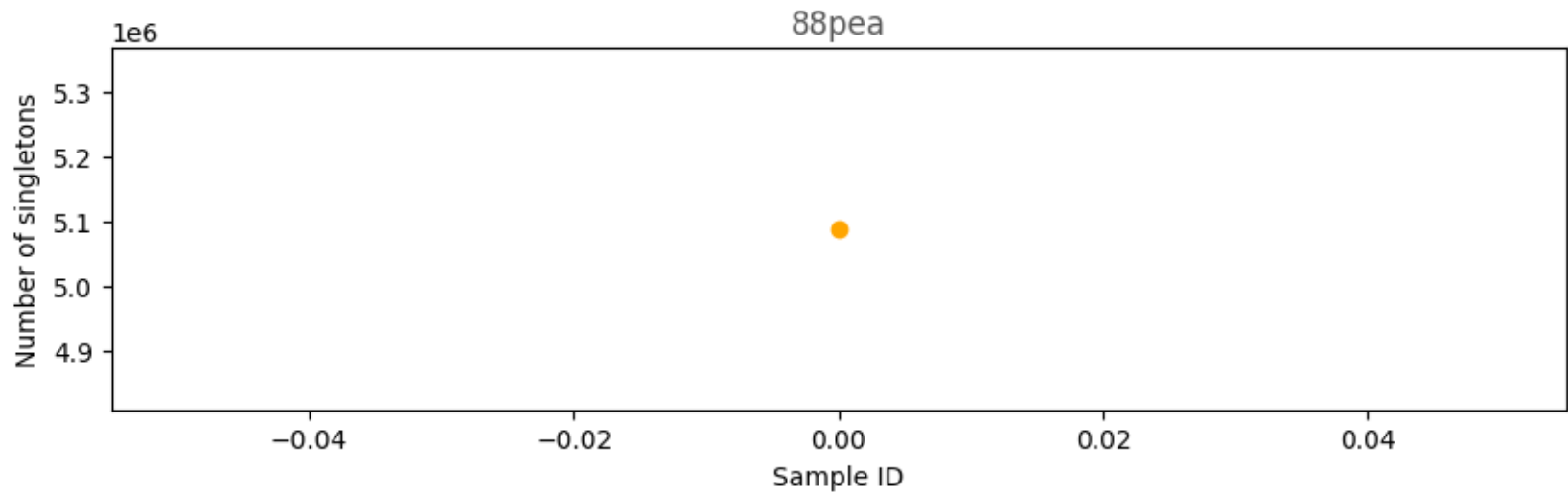
Ts/Tv by sample



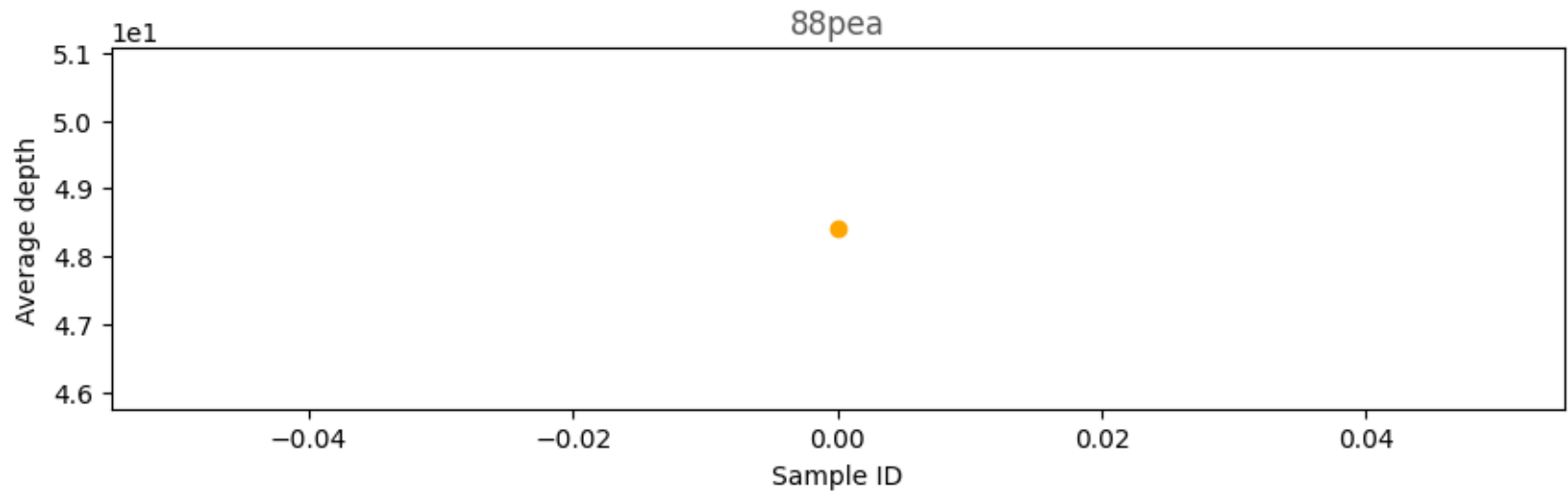
Hets vs non-ref Homs by sample



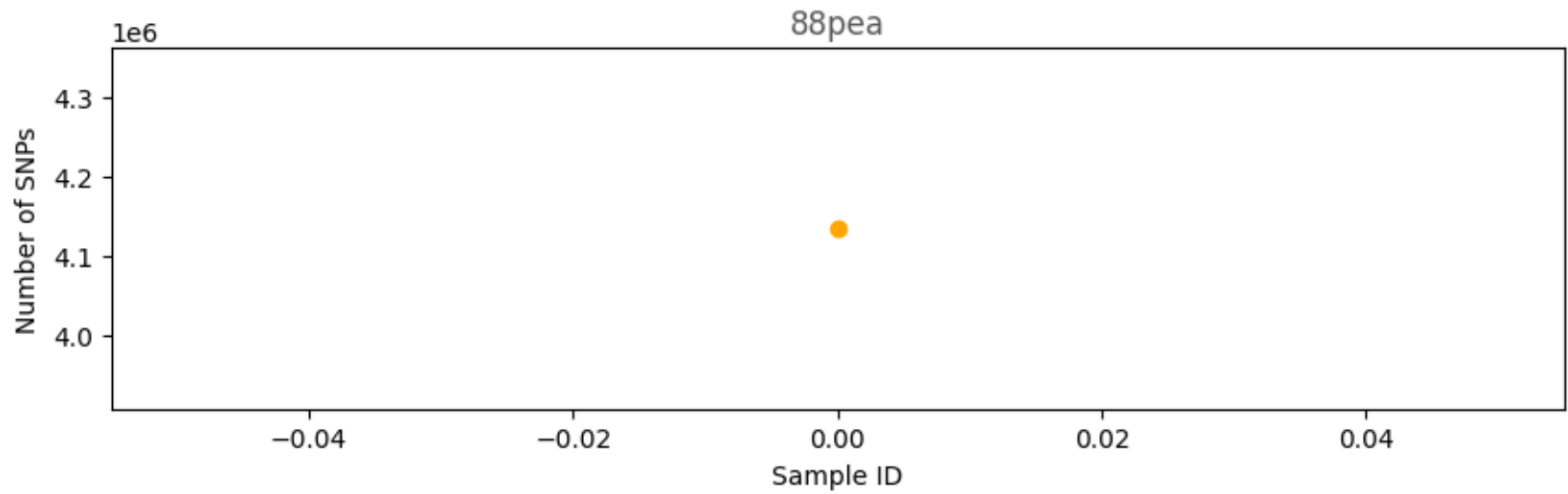
Singletons by sample (hets and homs)



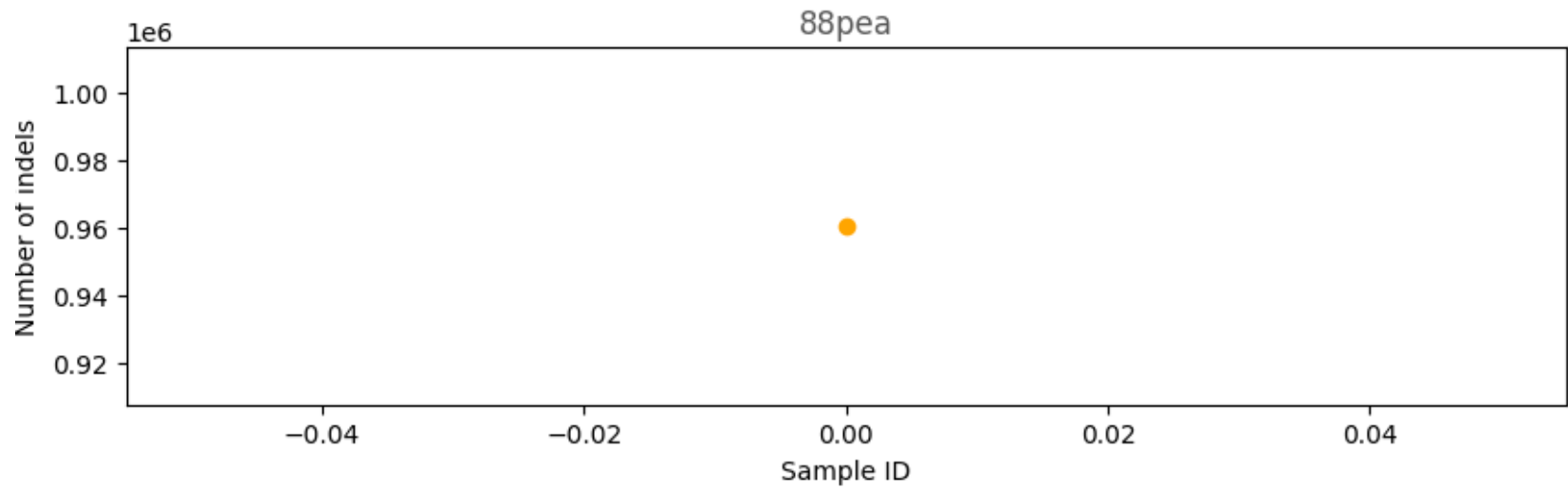
Average depth by sample



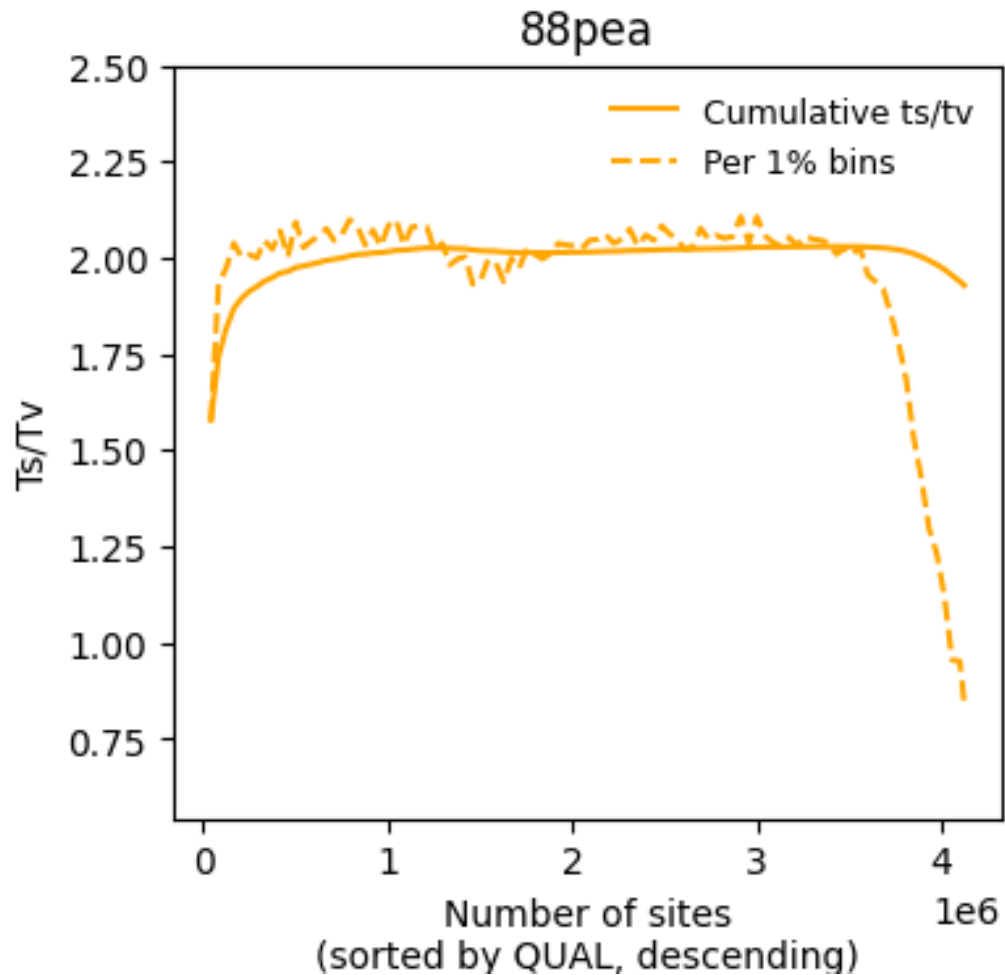
Number of SNPs by sample



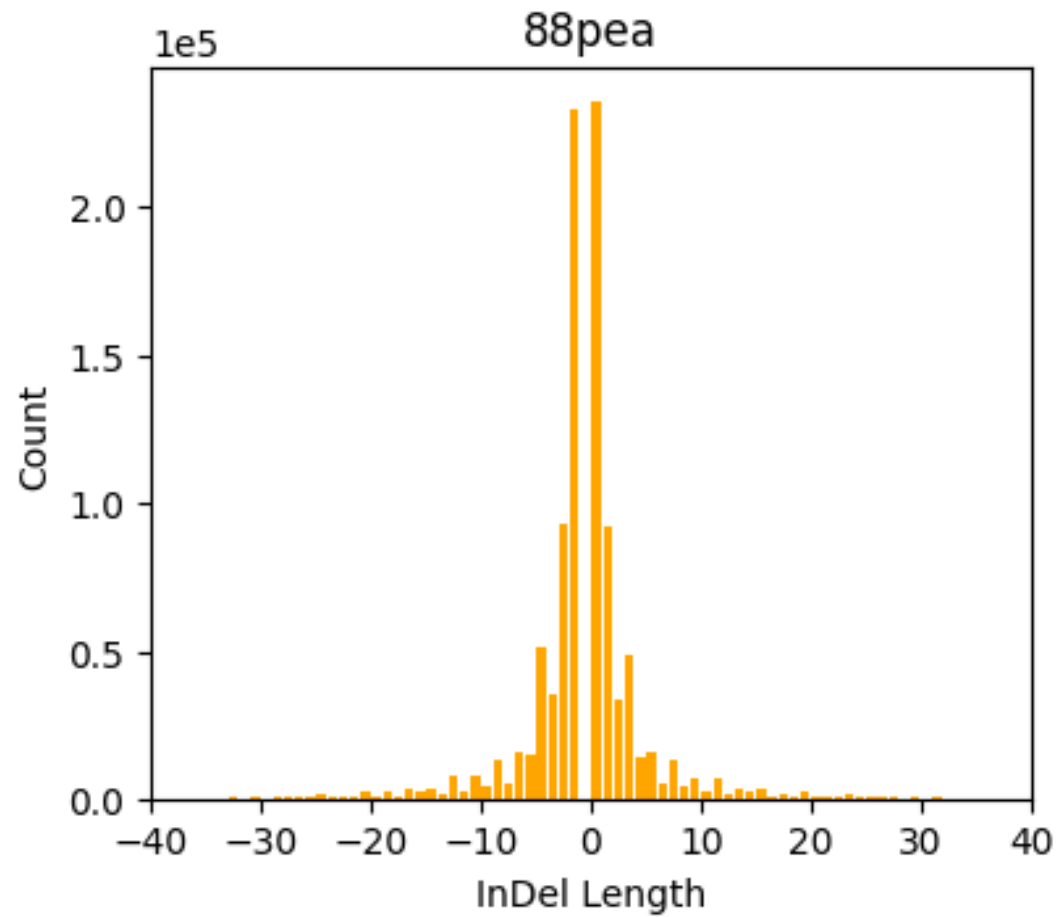
Number of indels by sample



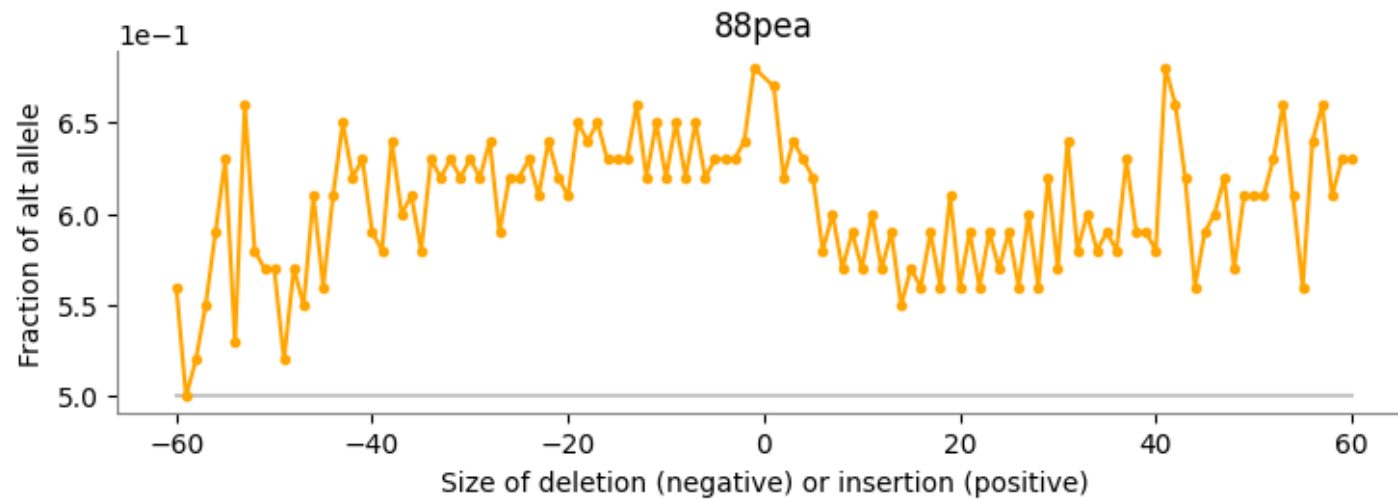
Ts/Tv stratified by QUAL



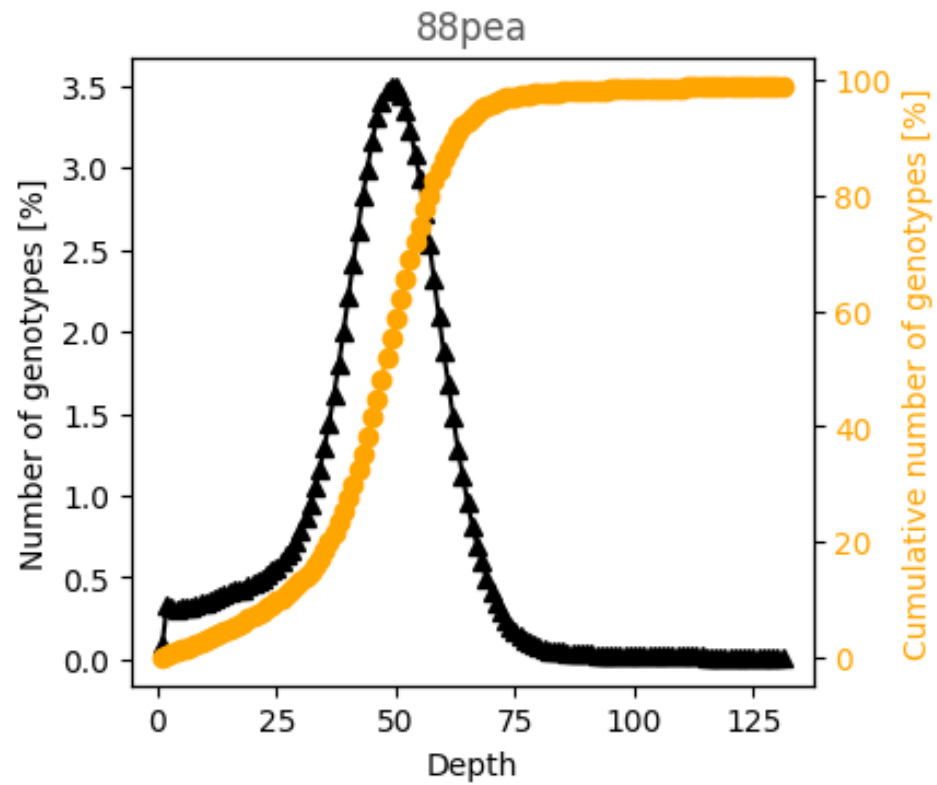
Indel distribution



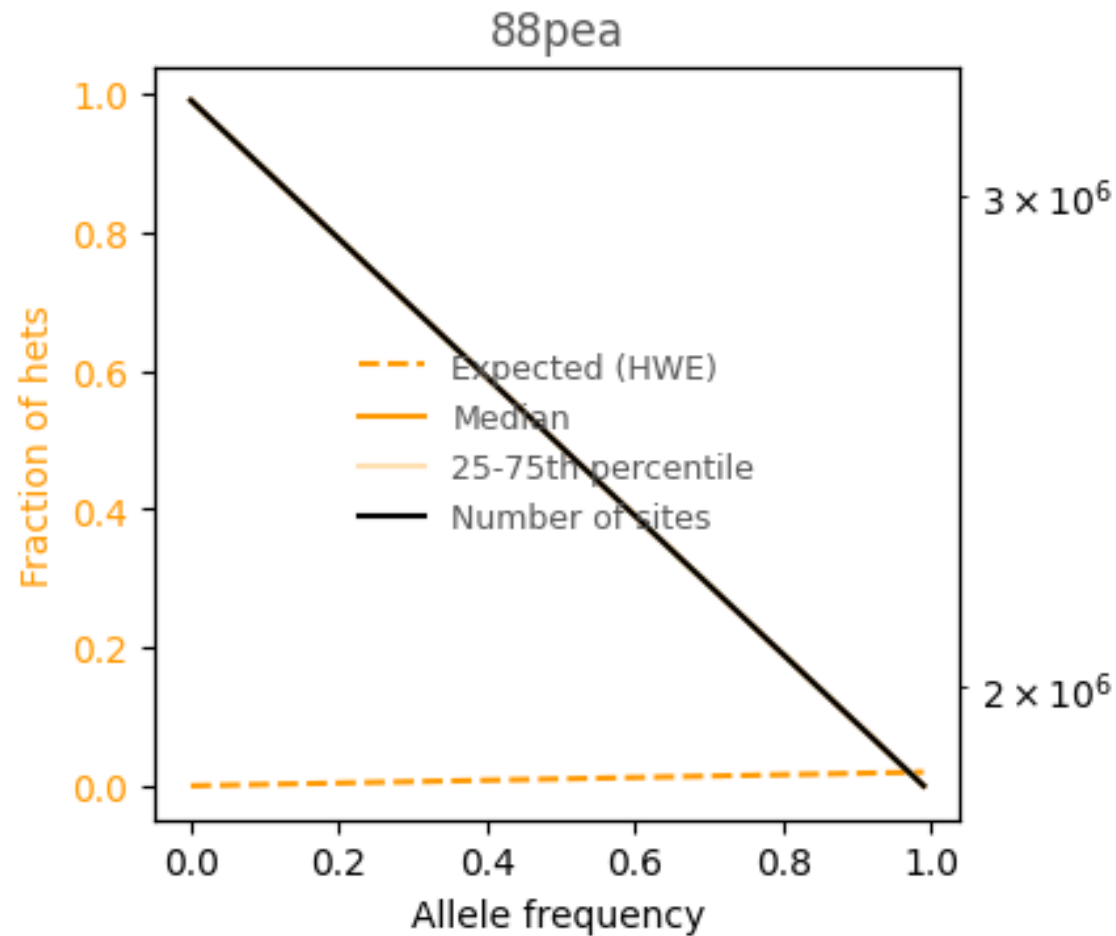
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

