

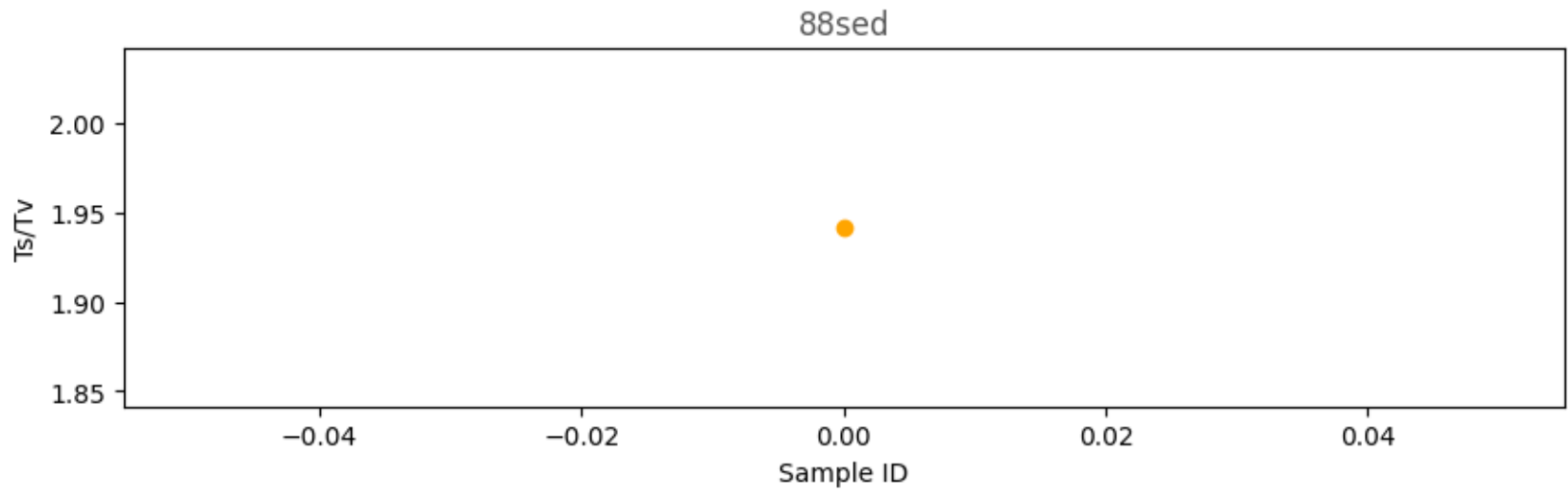
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
88sed	4,062,903	1.94	1.95	941,096	–	0	0
* frameshift ratio: out/(out+in)							

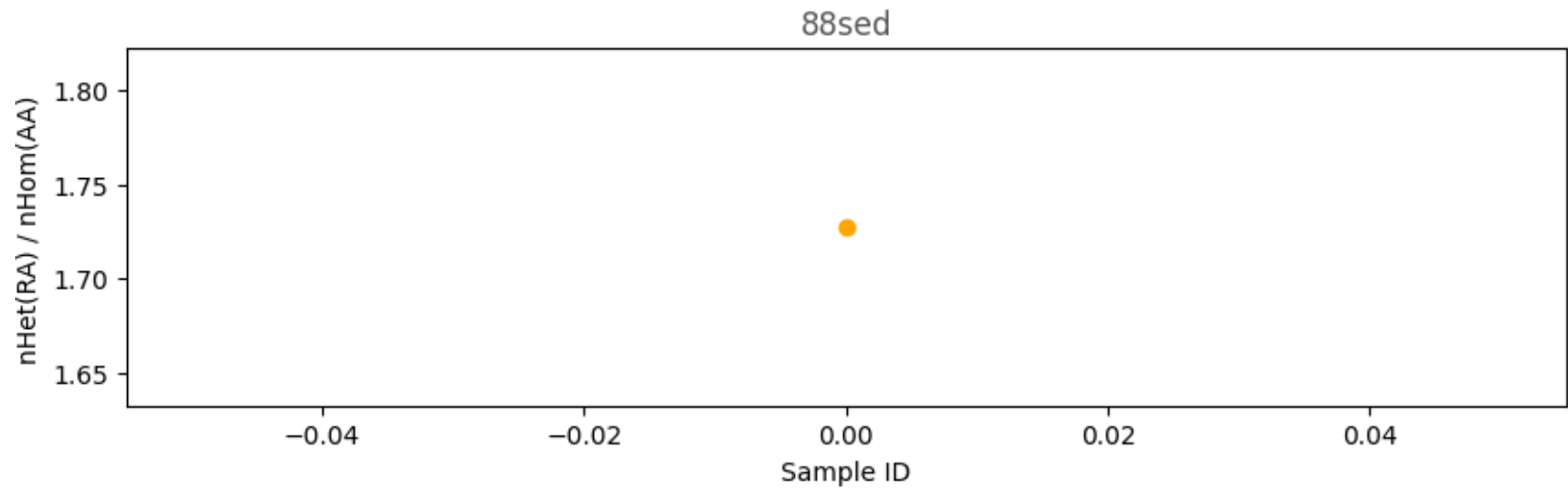
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
88sed	63.3%	1.92	67.8%	95,518	1,943

- 88sed .. /ngc/projects2/gm/data/archive/2022/variants/snv/88sedewsf-103825446004-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00501-220119_A01176_AH2CTLDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS03674_21RKG030588x01_snv_germline_raw.haplotype_caller.vcf.gz

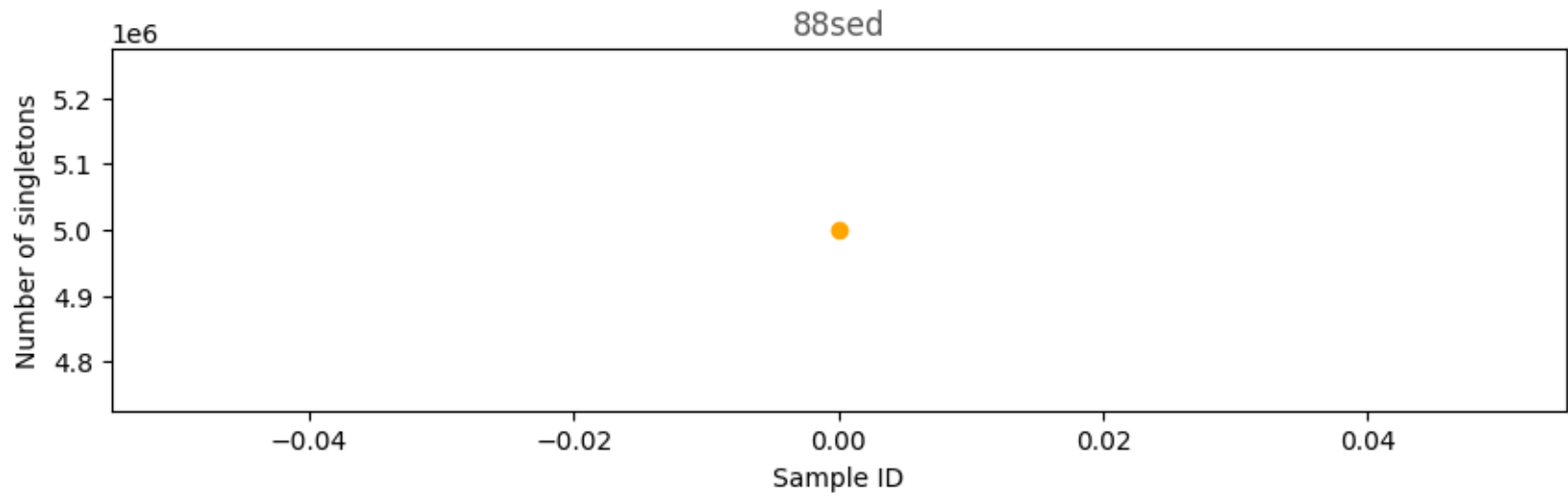
Ts/Tv by sample



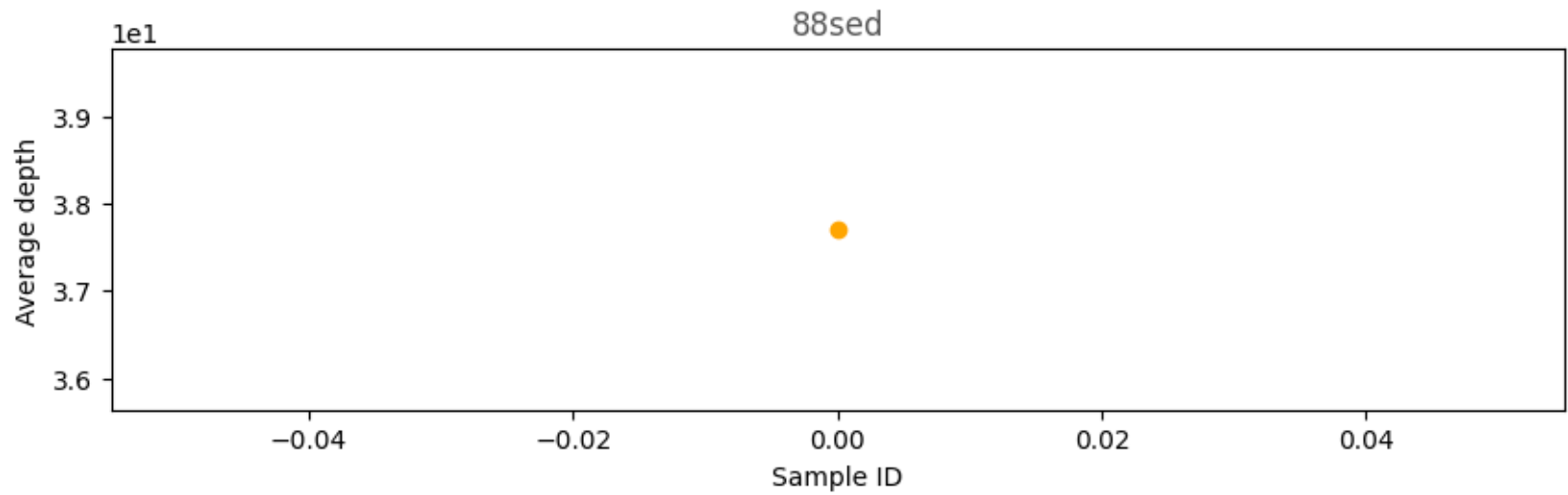
Hets vs non-ref Homs by sample



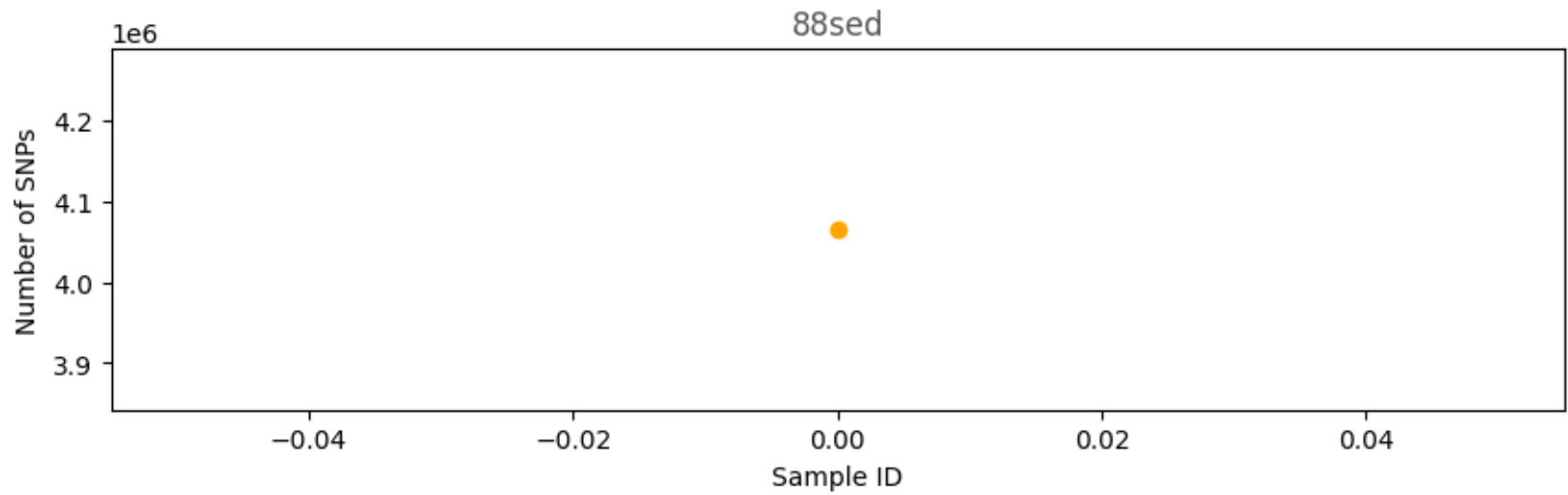
Singletons by sample (hets and homs)



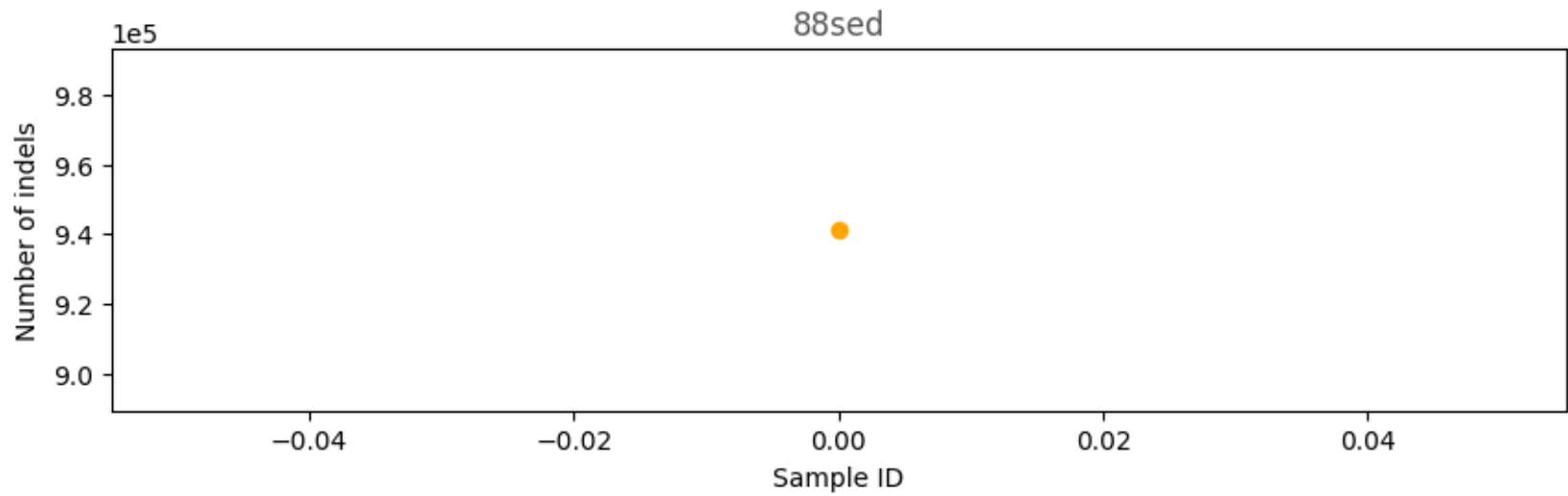
Average depth by sample



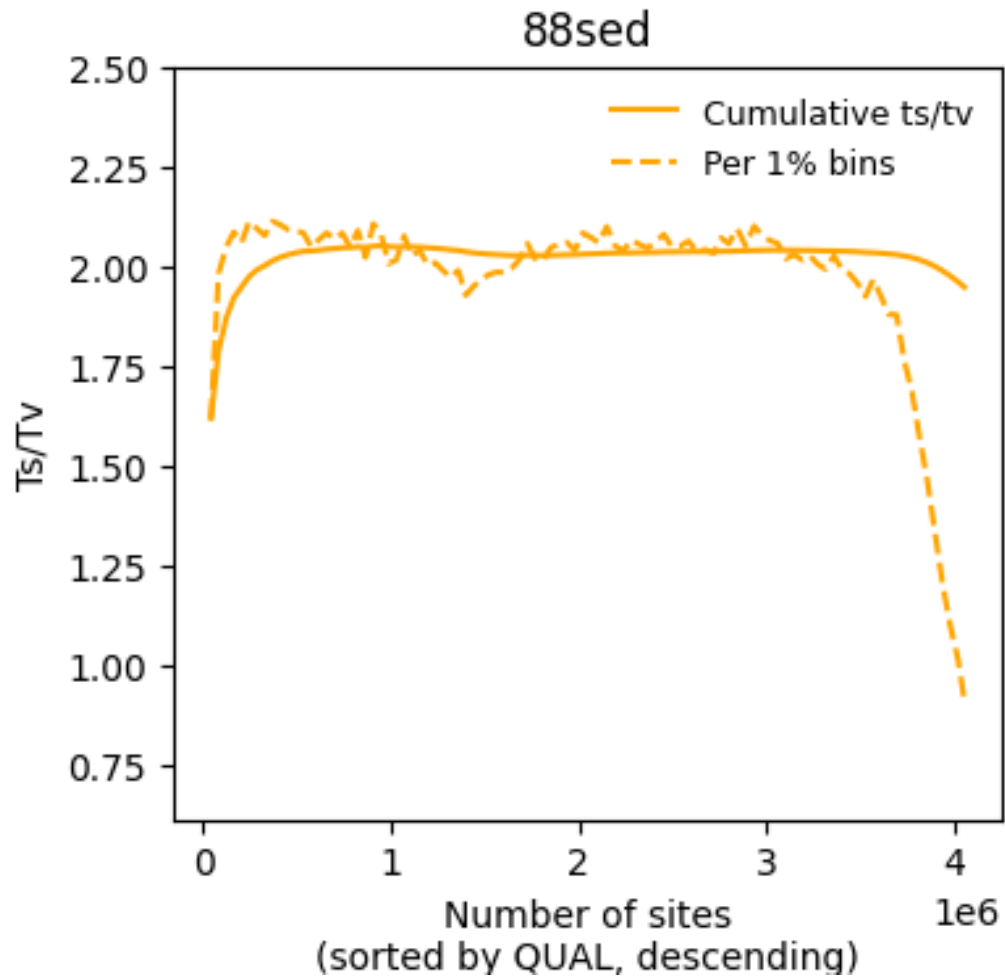
Number of SNPs by sample



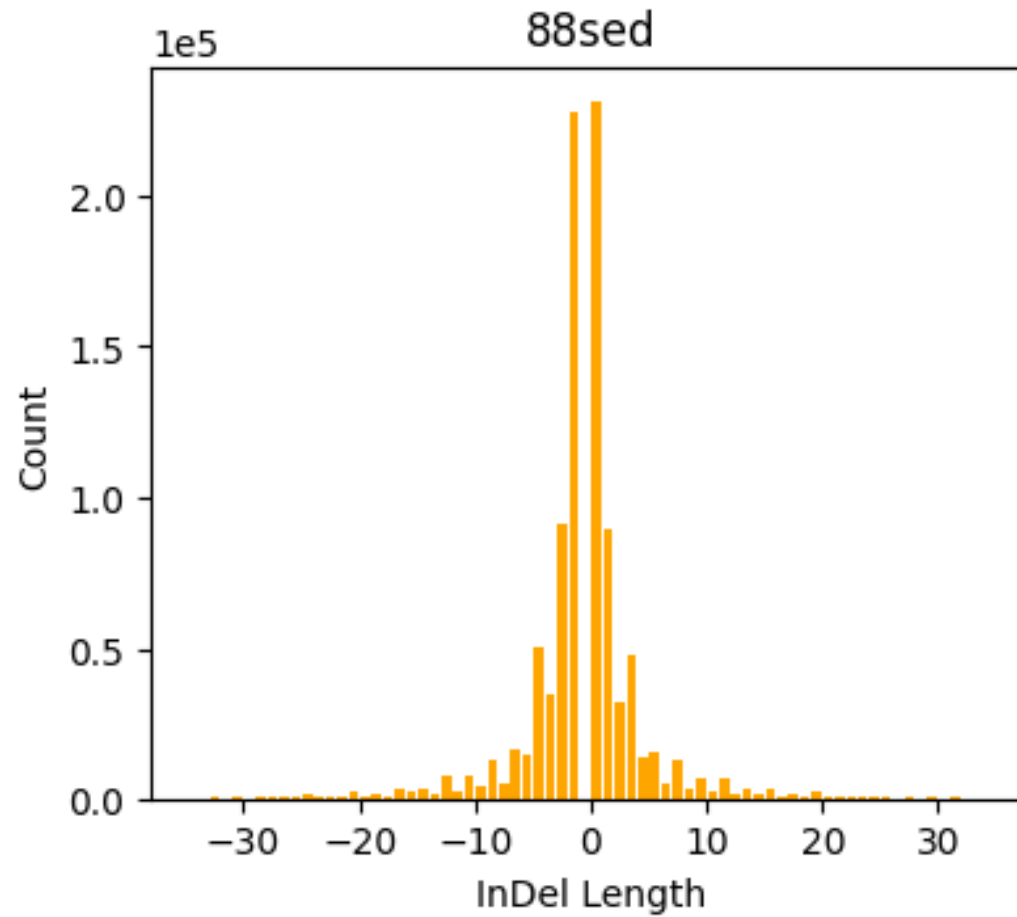
Number of indels by sample



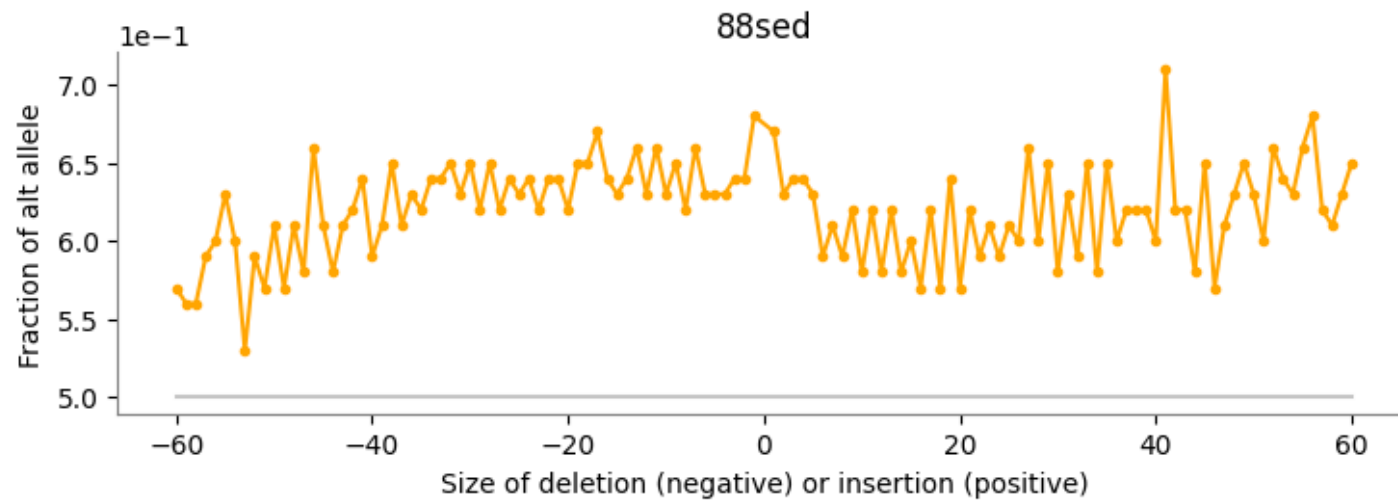
Ts/Tv stratified by QUAL



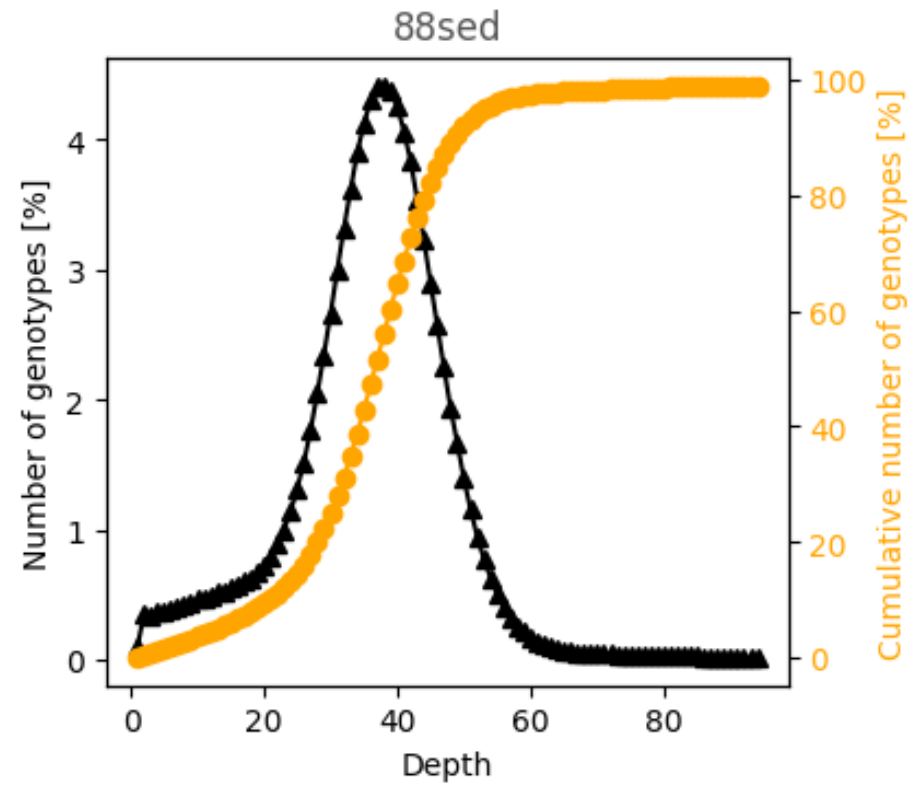
Indel distribution



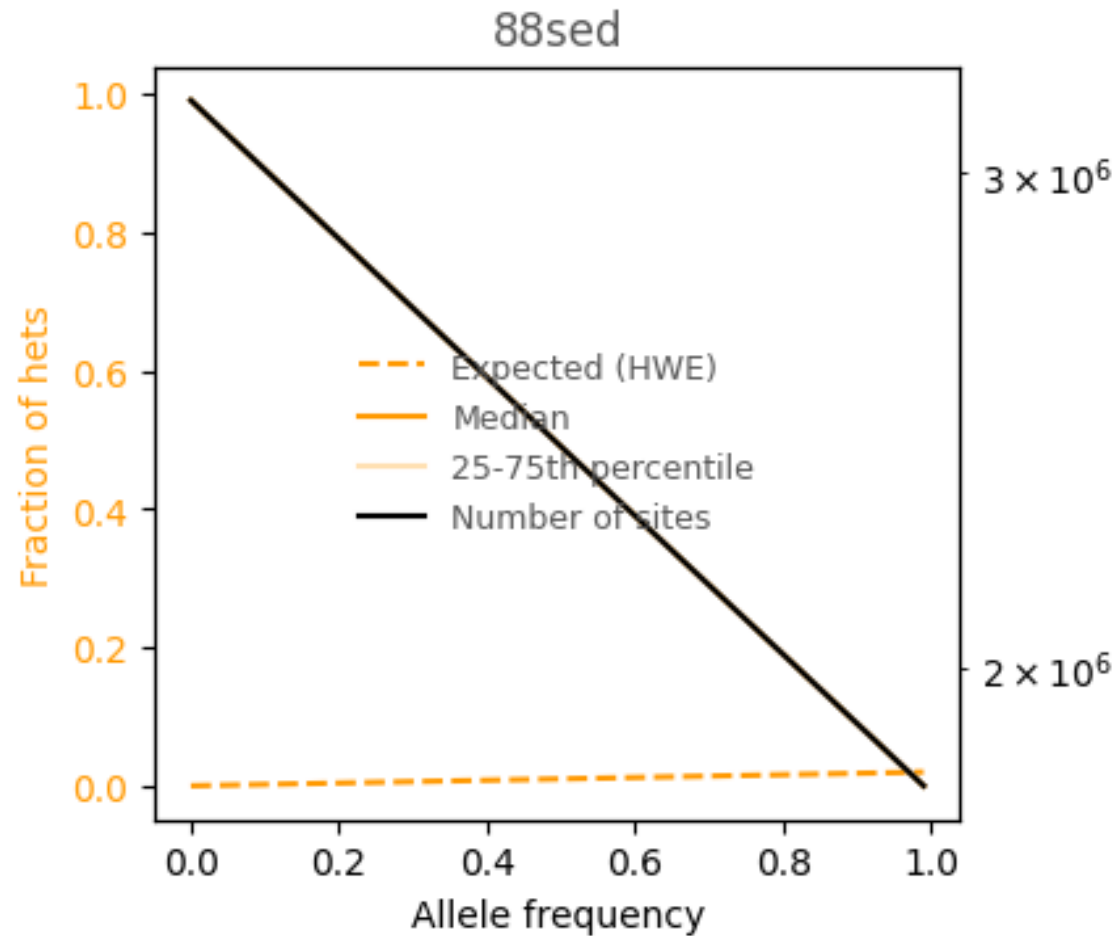
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

