

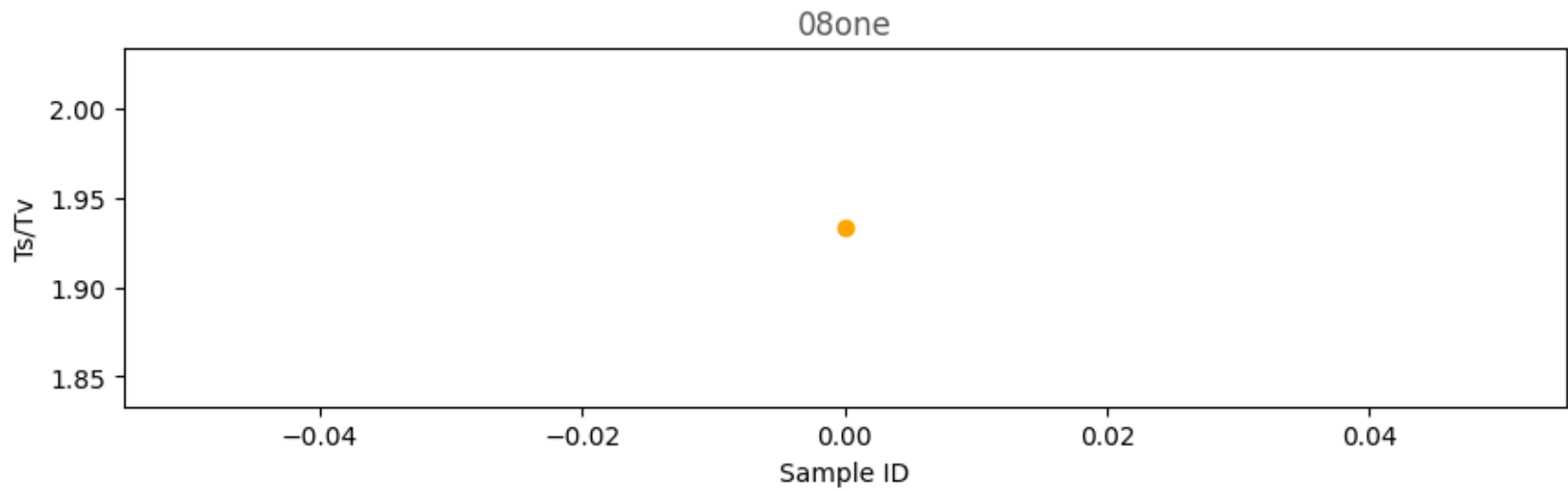
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
08one	3,983,335	1.93	1.94	932,401	–	0	0
* frameshift ratio: out/(out+in)							

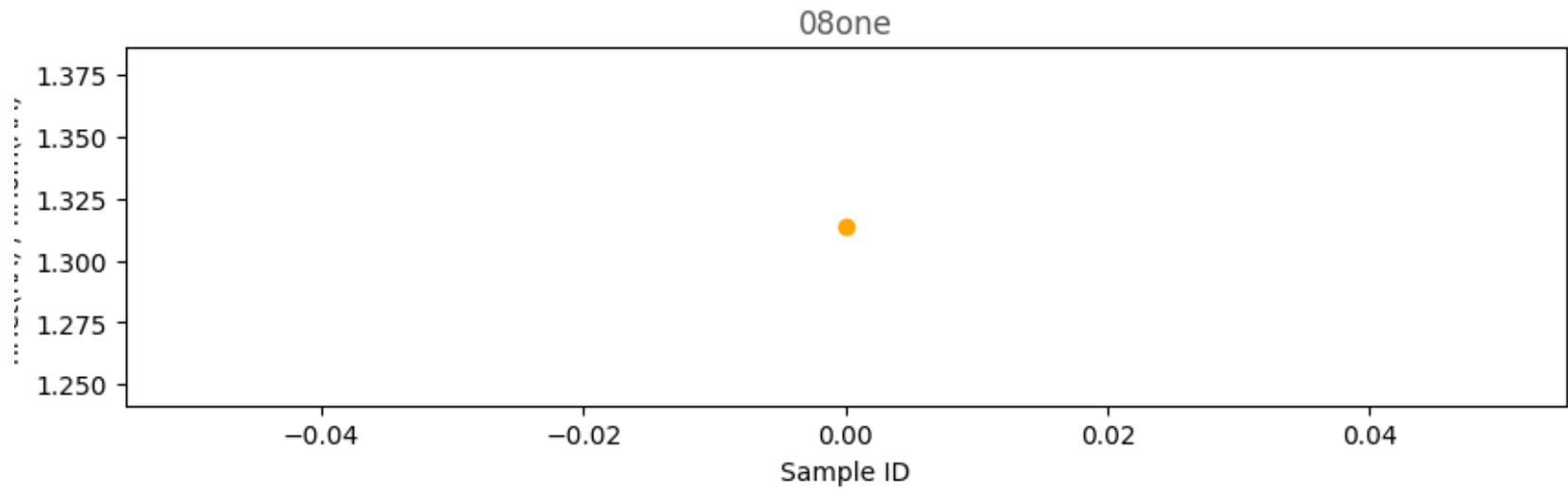
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
08one	56.8%	1.90	61.4%	89,312	1,876

- 08one .. /ngc/projects2/gm/data/archive/2022/variants/snv/08onertom-103825452497-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00513-220119_A01176_BH2CT5DSX3-EXT_LAB
KA_NGCWGS-NGCWGS03686_21RKG030137x01_snv_germline_raw.haplotype_caller.vcf.gz

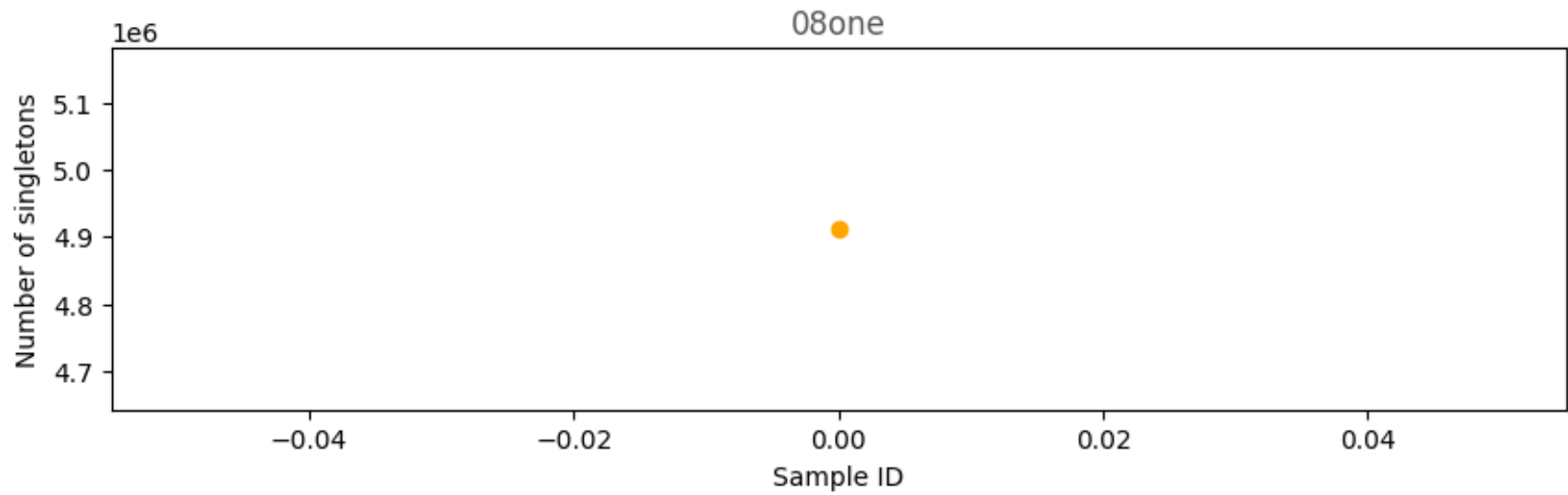
Ts/Tv by sample



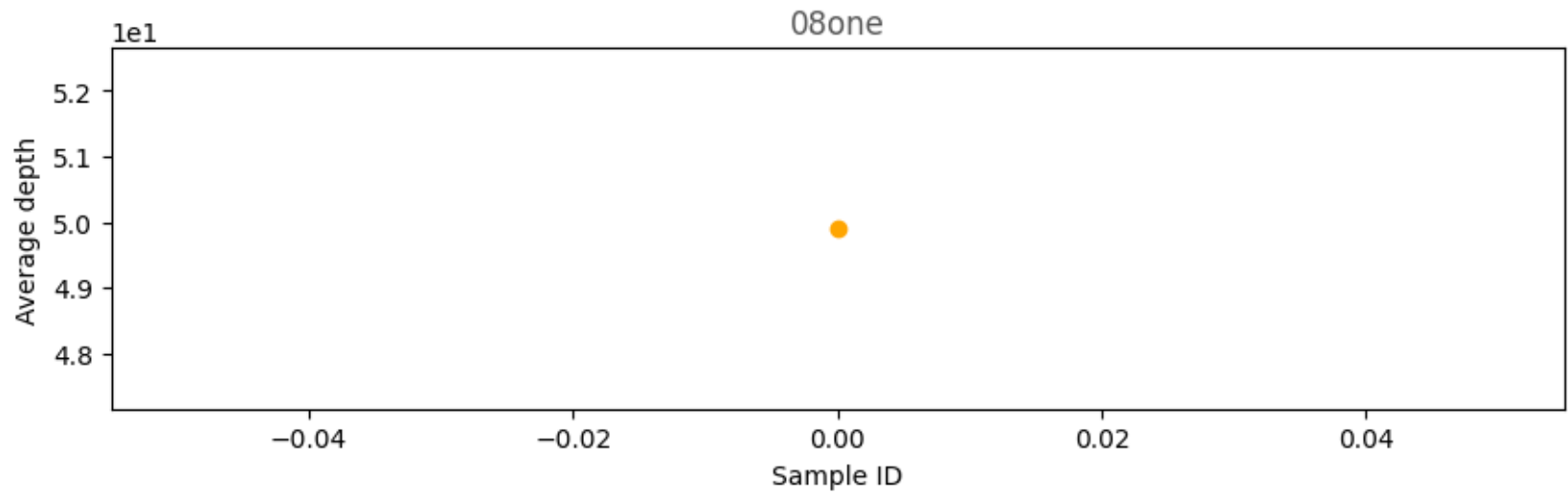
Hets vs non-ref Homs by sample



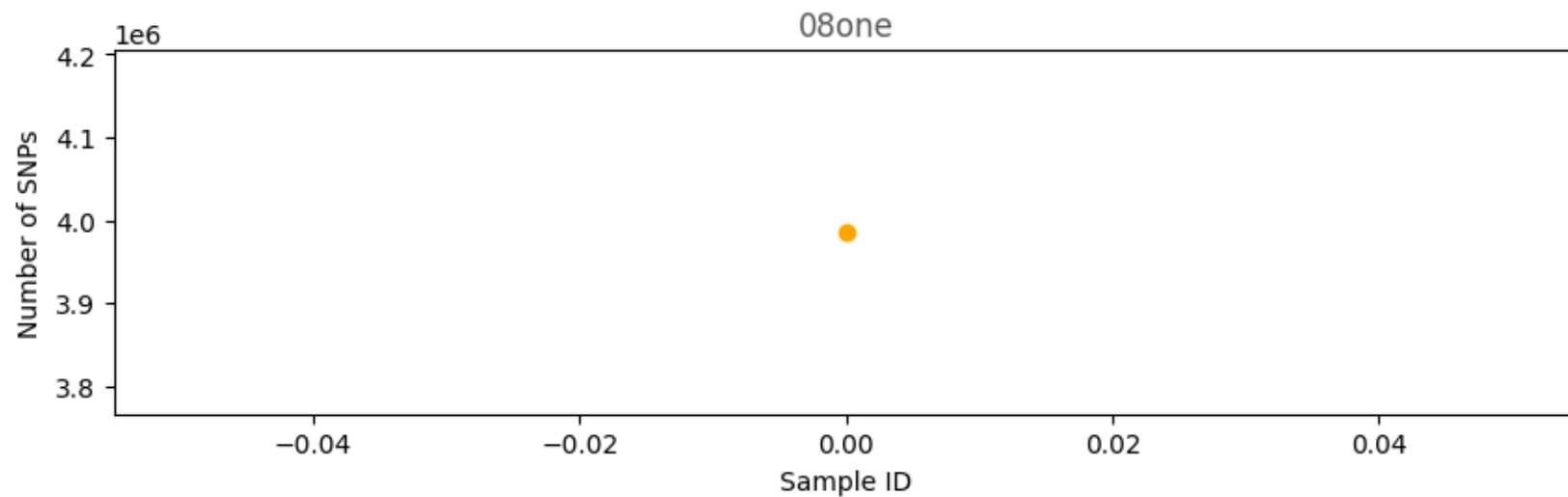
Singletons by sample (hets and homs)



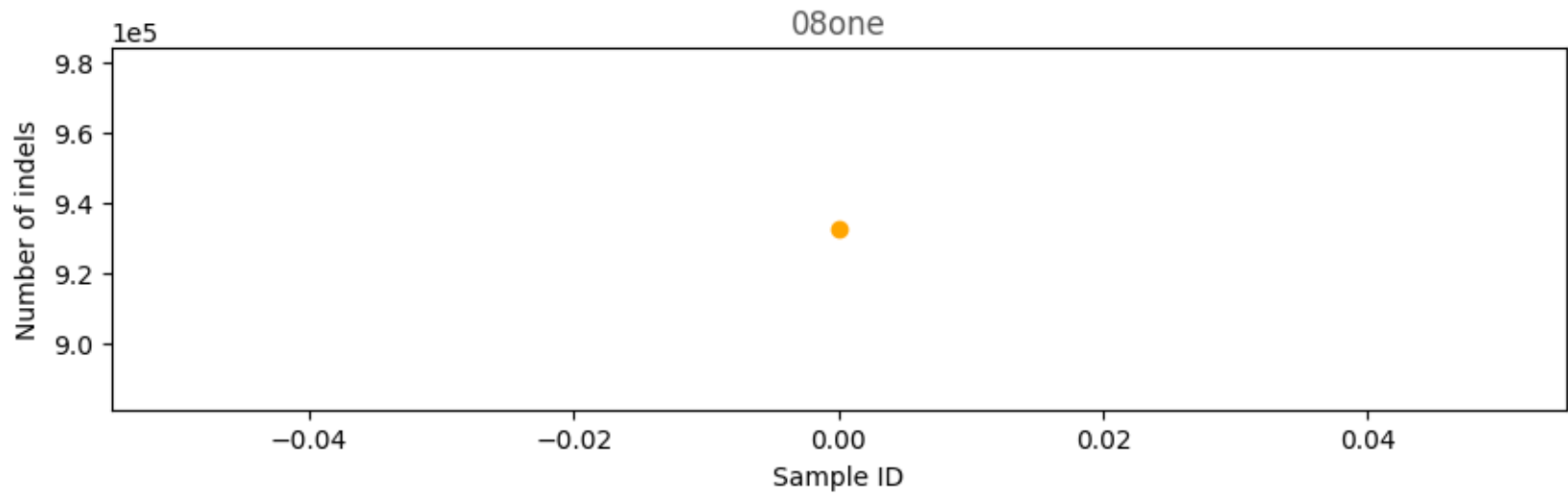
Average depth by sample



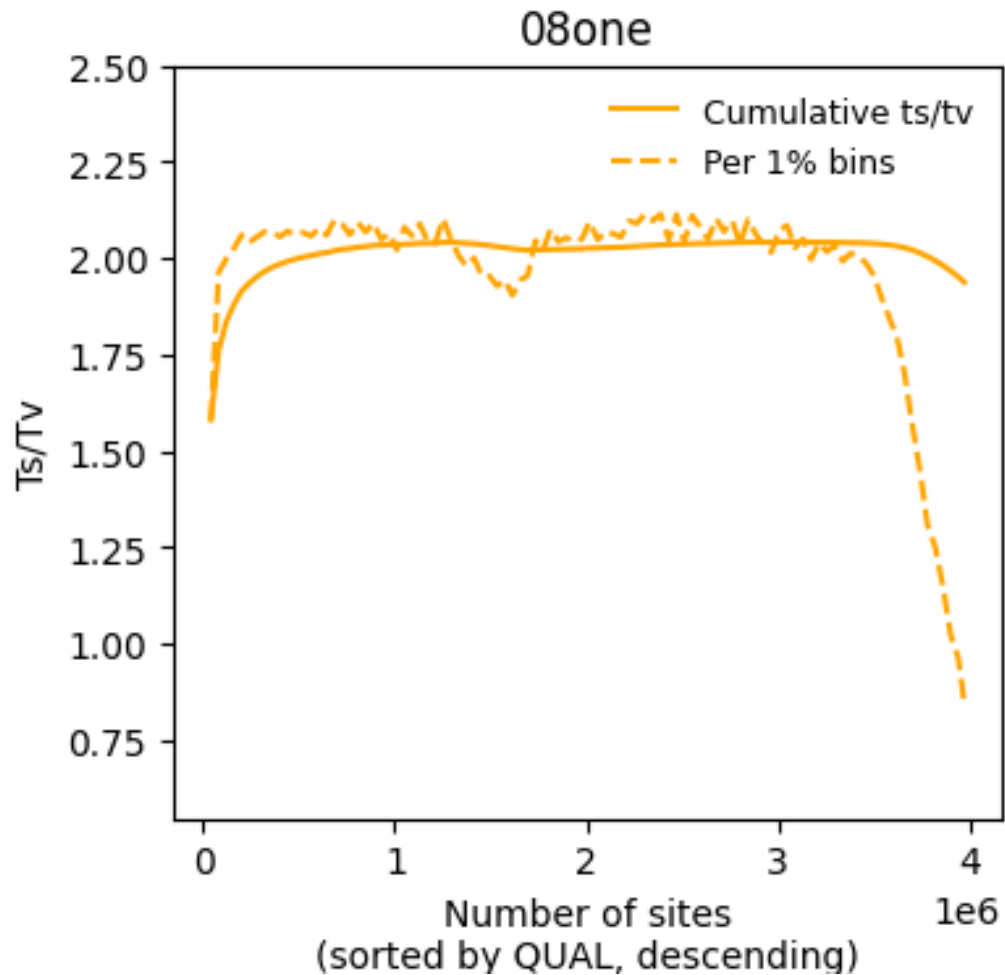
Number of SNPs by sample



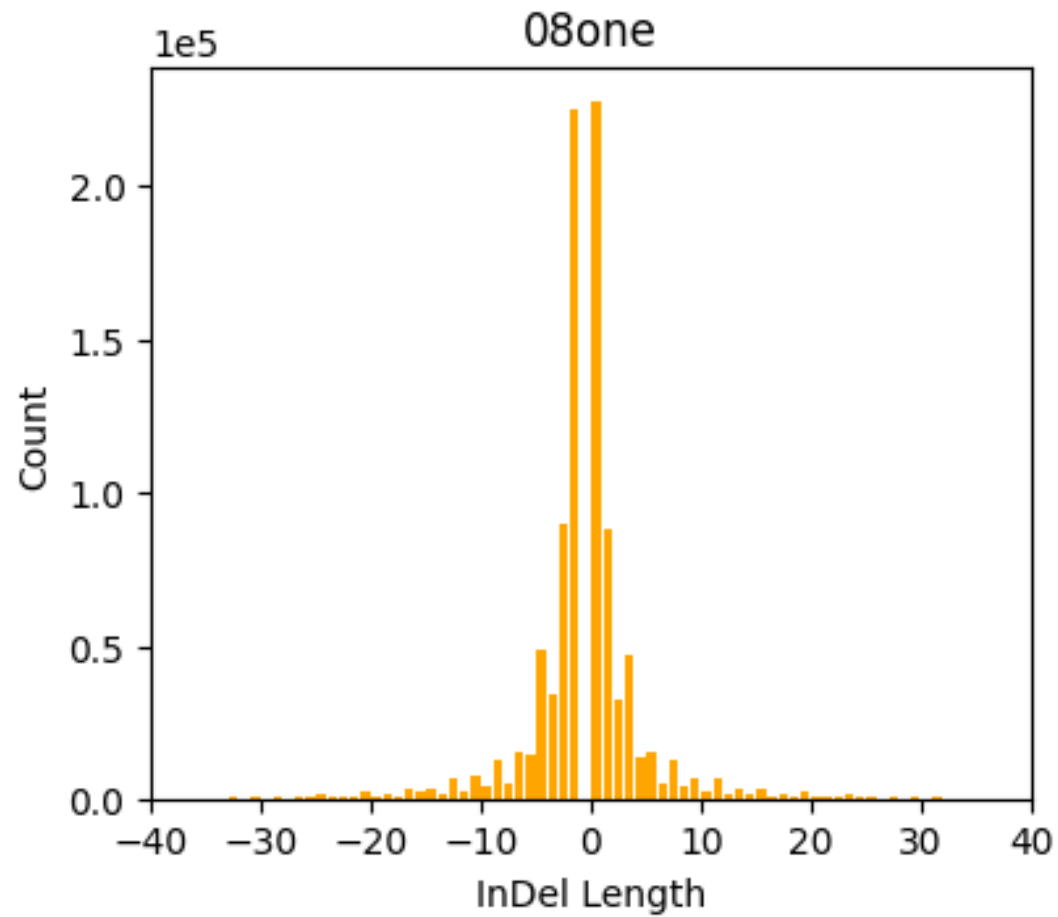
Number of indels by sample



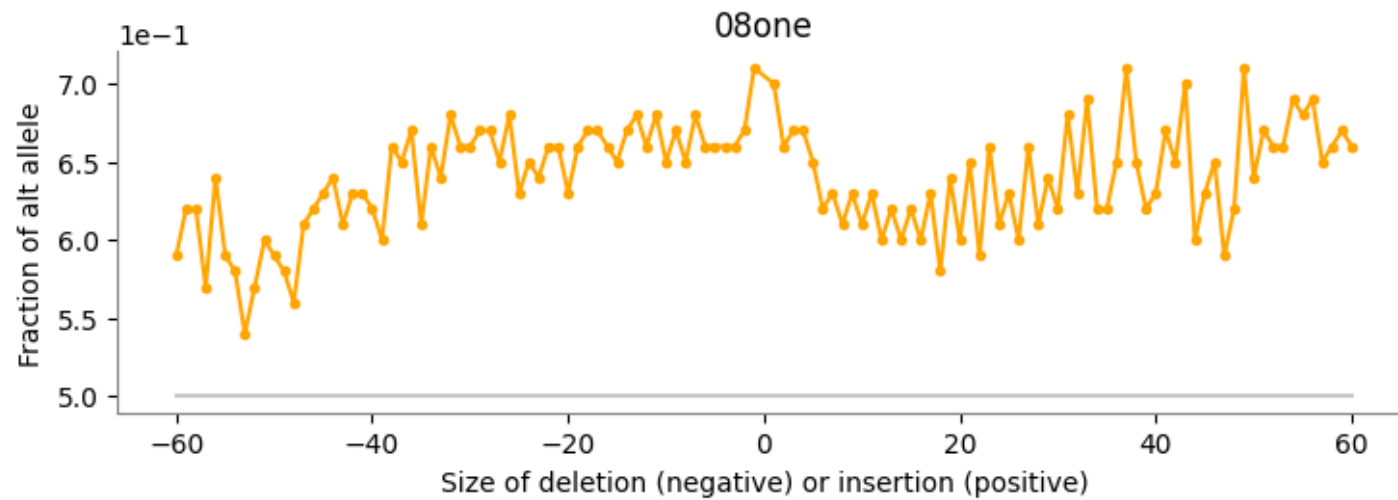
Ts/Tv stratified by QUAL



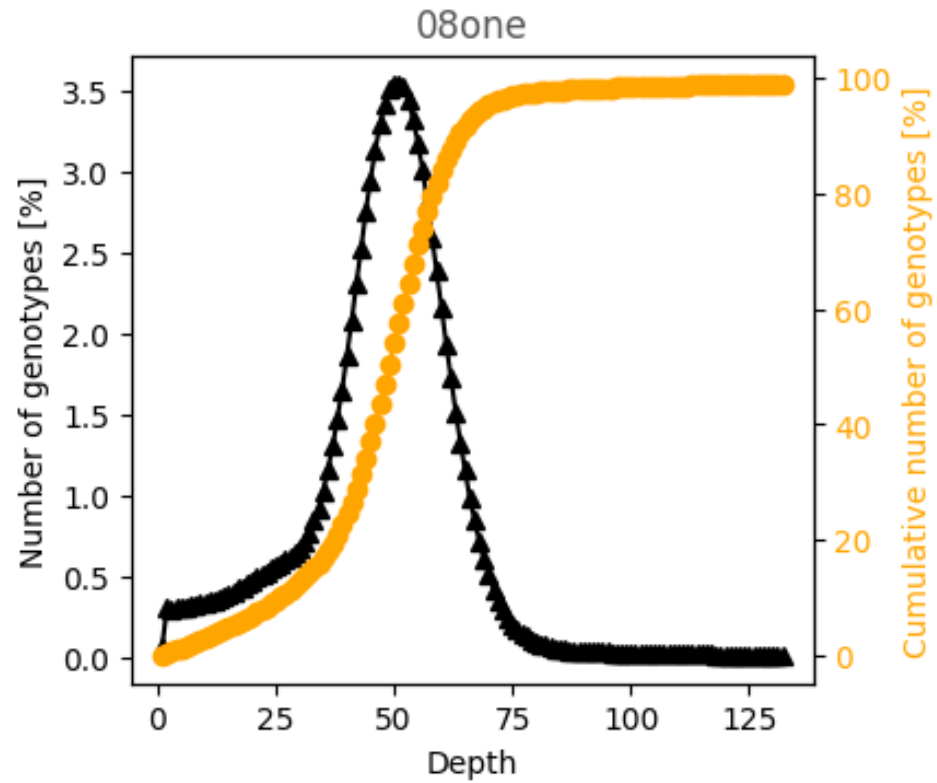
Indel distribution



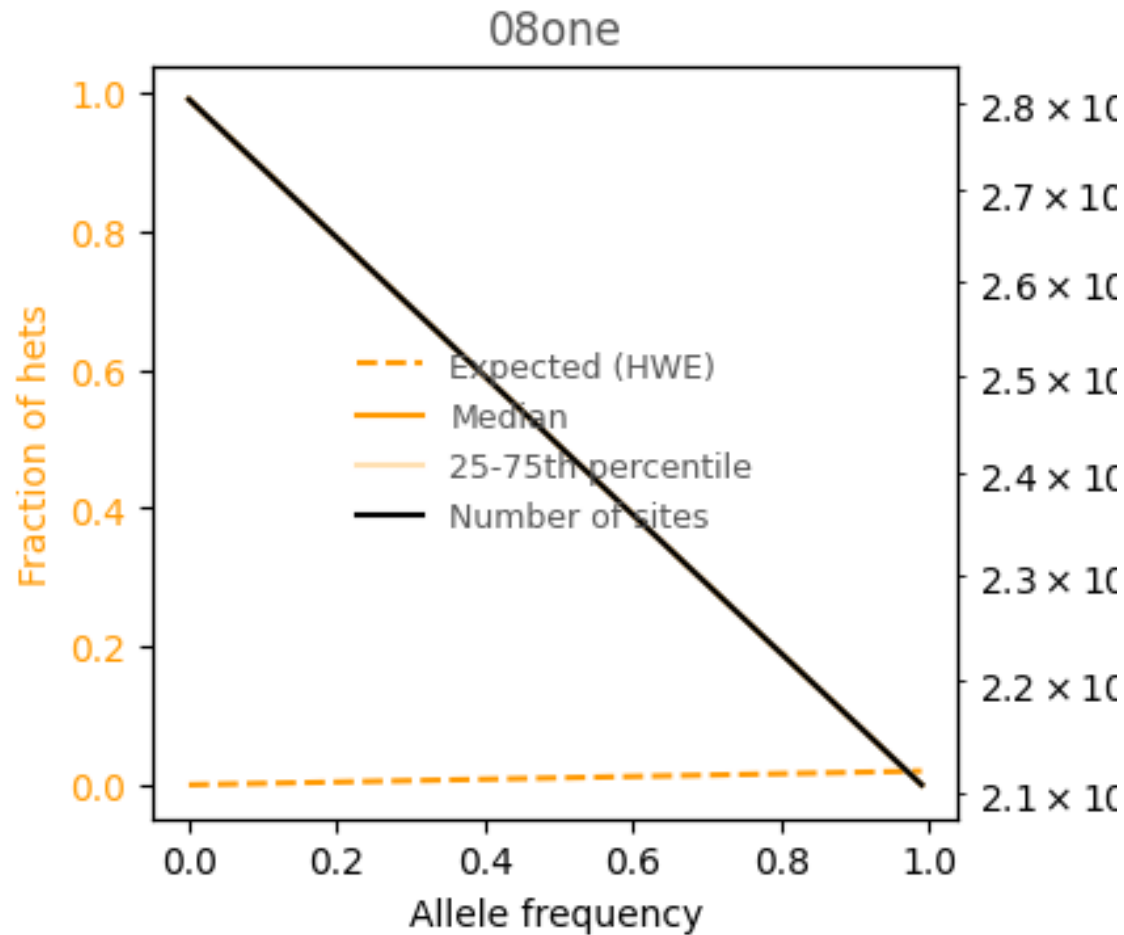
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

