Summary Numbers

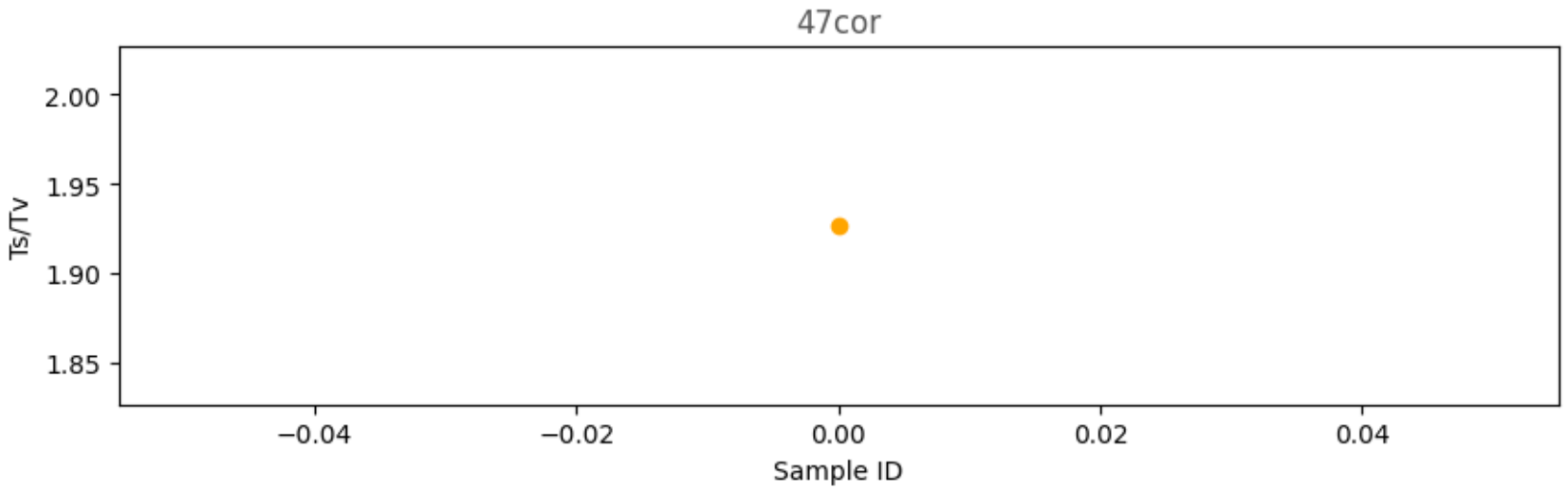
	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
47cor	4,099,238	1.93	1.93	955,336	–	0	0

* frameshift ratio: out/(out+in)

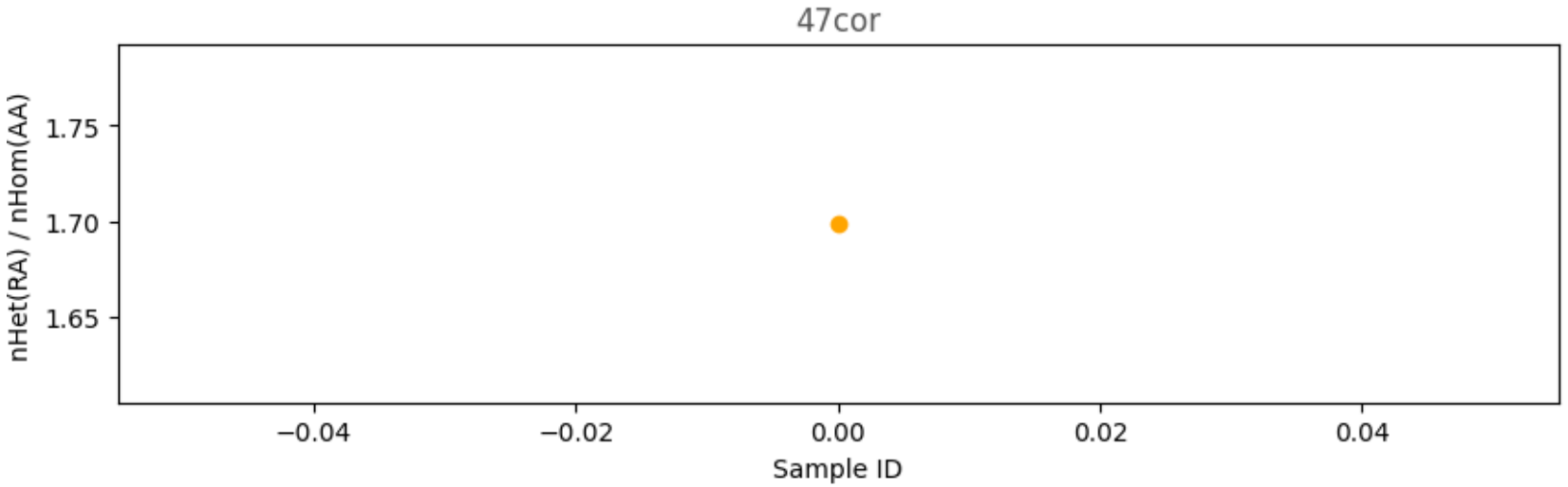
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
47cor	63.0%	1.90	68.2%	102,922	2,386

- 47cor .. `/ngc/projects2/gm/data/archive/2022/variants/snv/47corthaf-103904565053-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01191-220831_A01961_BHTNYFDSX3-EXT_LABKA_NGCWGS-NGCWGS04947_snv_germline_raw.haplotype_caller.vcf.gz`

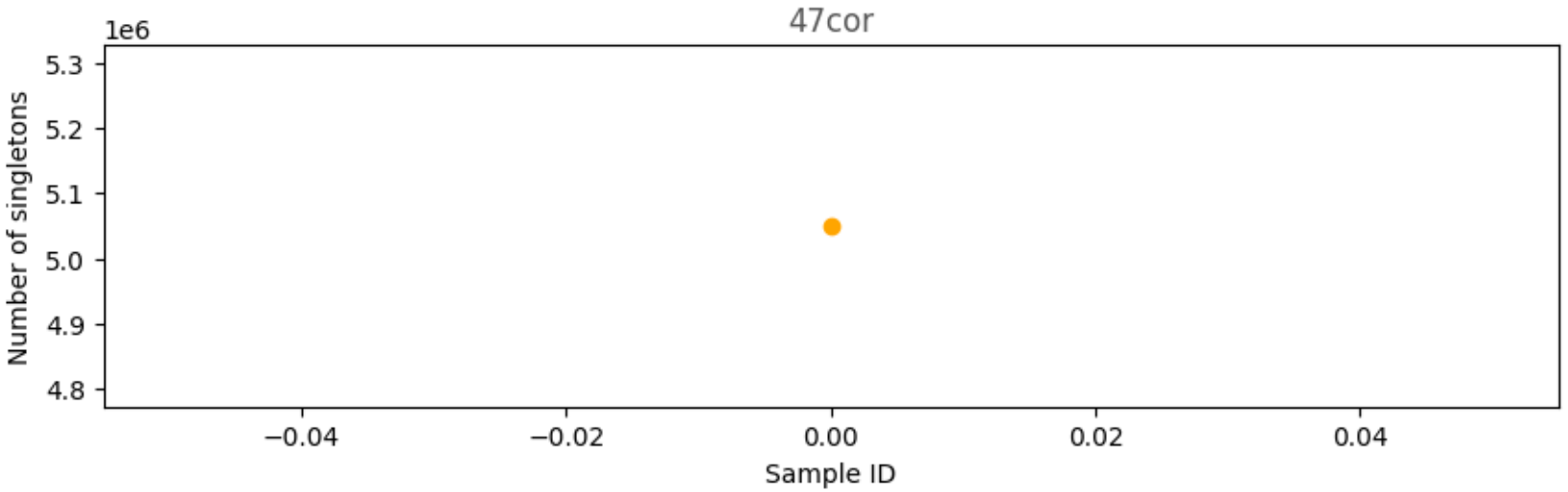
Ts/Tv by sample



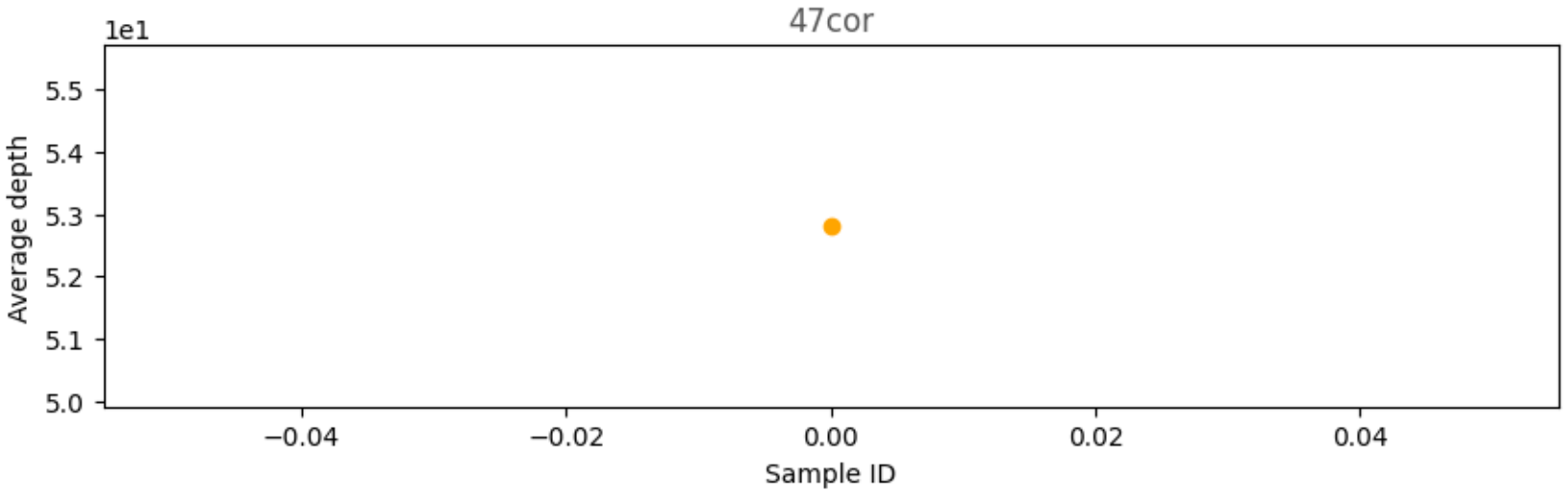
Hets vs non-ref Homs by sample



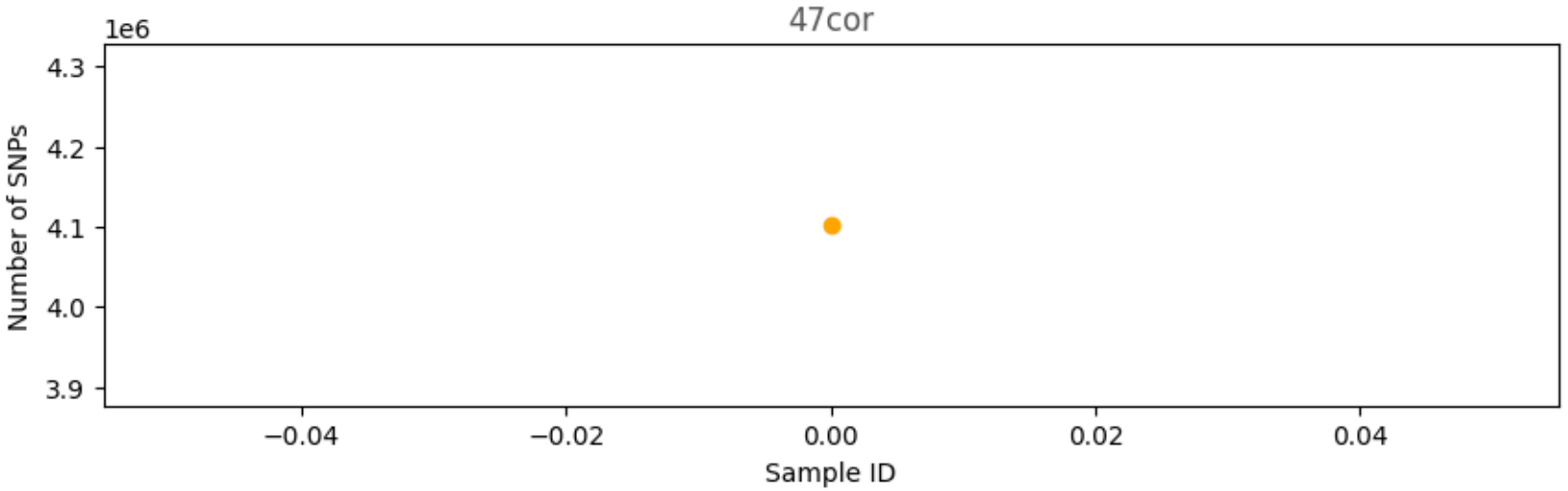
Singletons by sample (hets and homs)



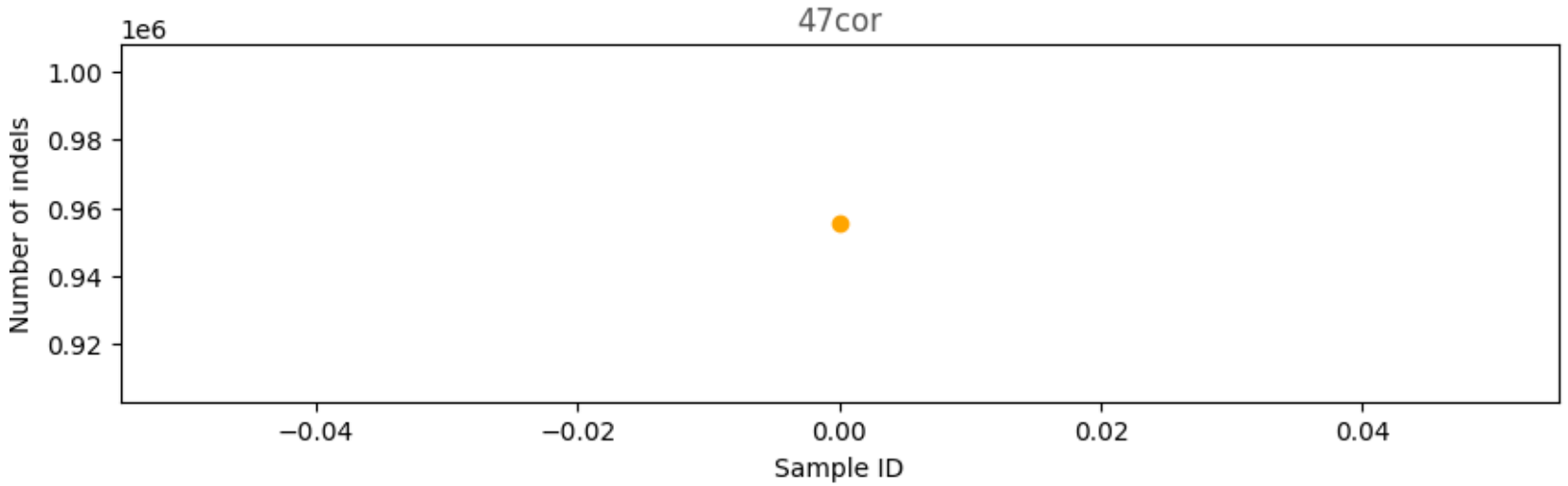
Average depth by sample



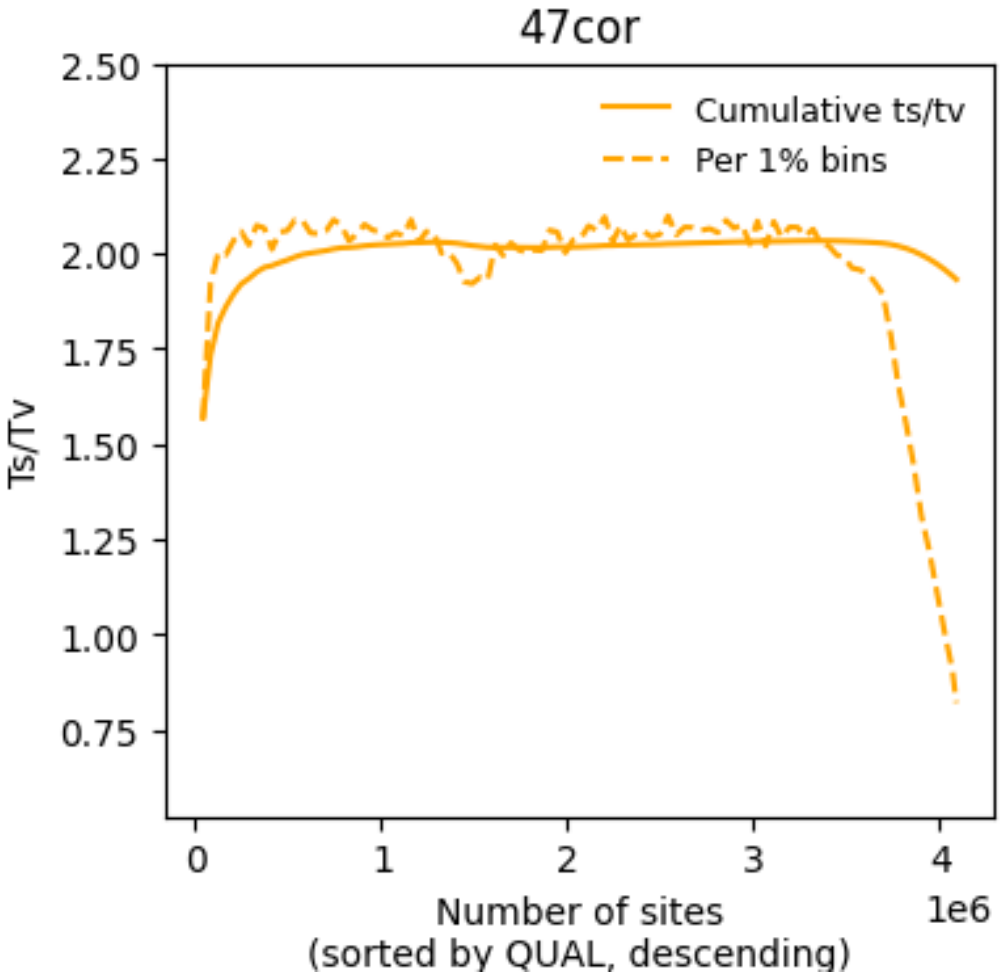
Number of SNPs by sample



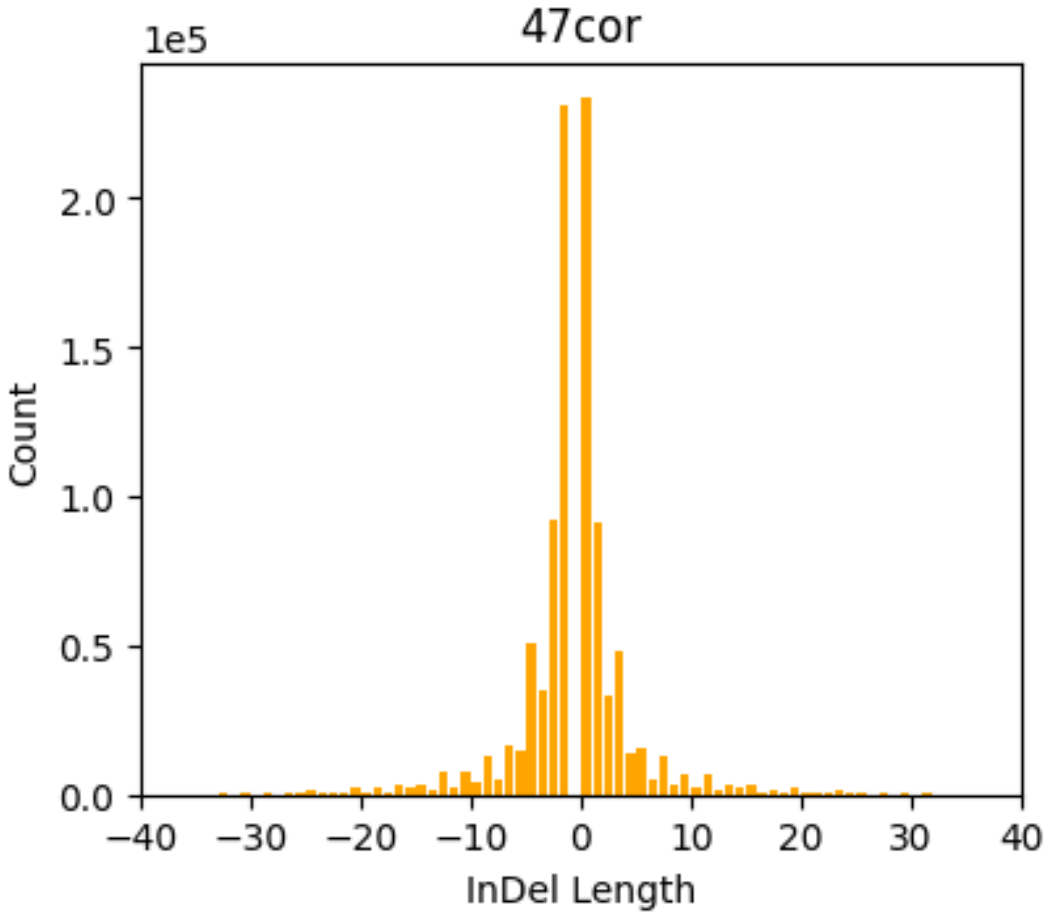
Number of indels by sample



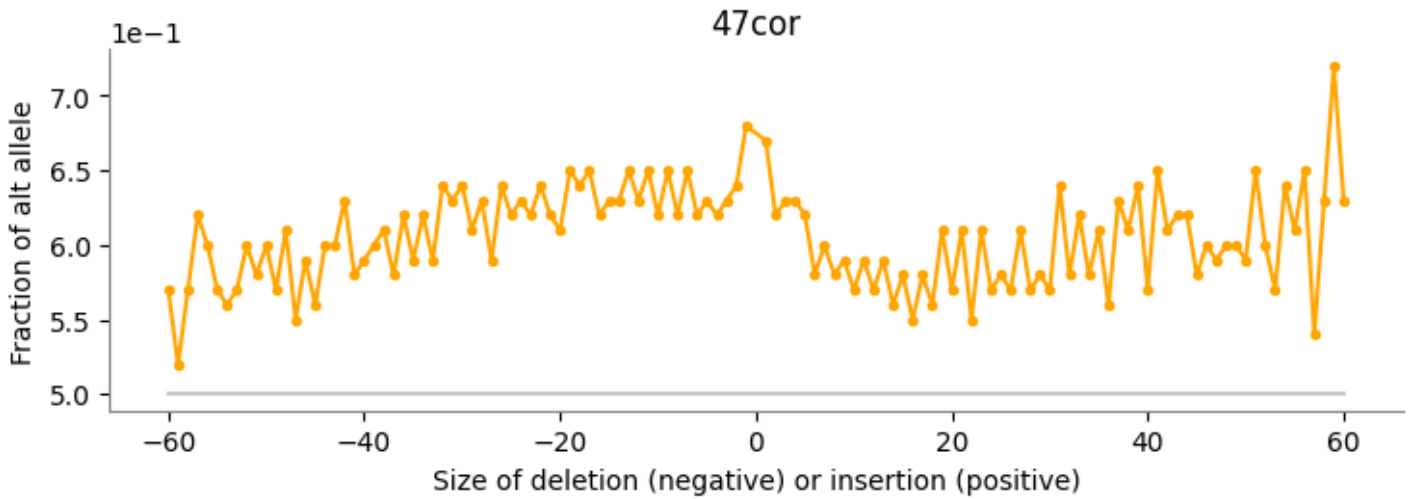
Ts/Tv stratified by QUAL



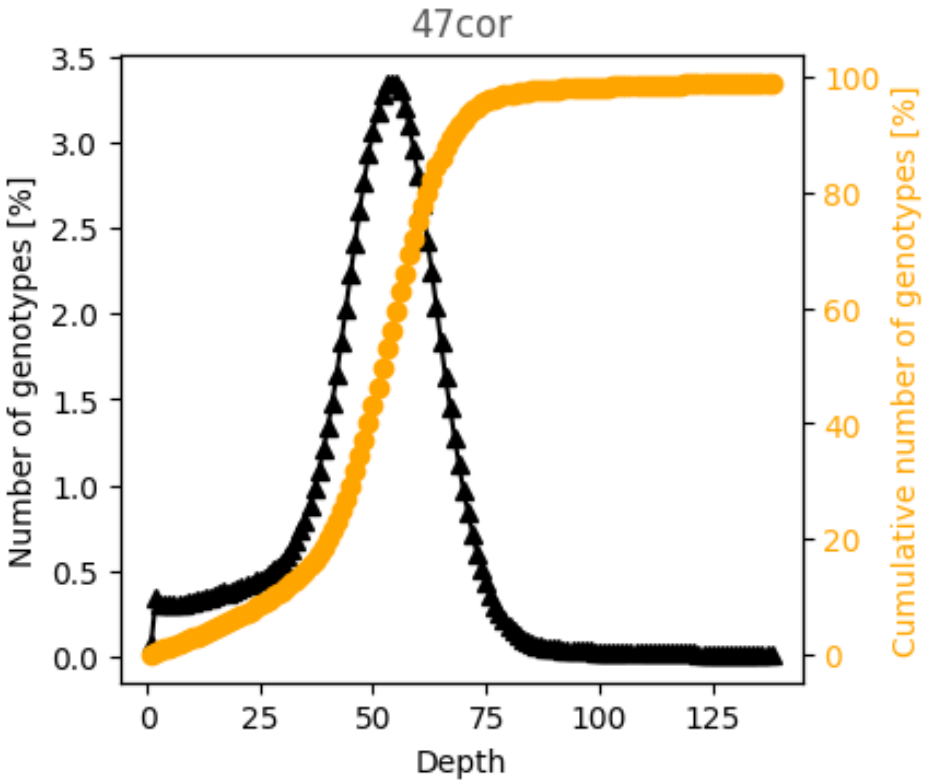
Indel distribution



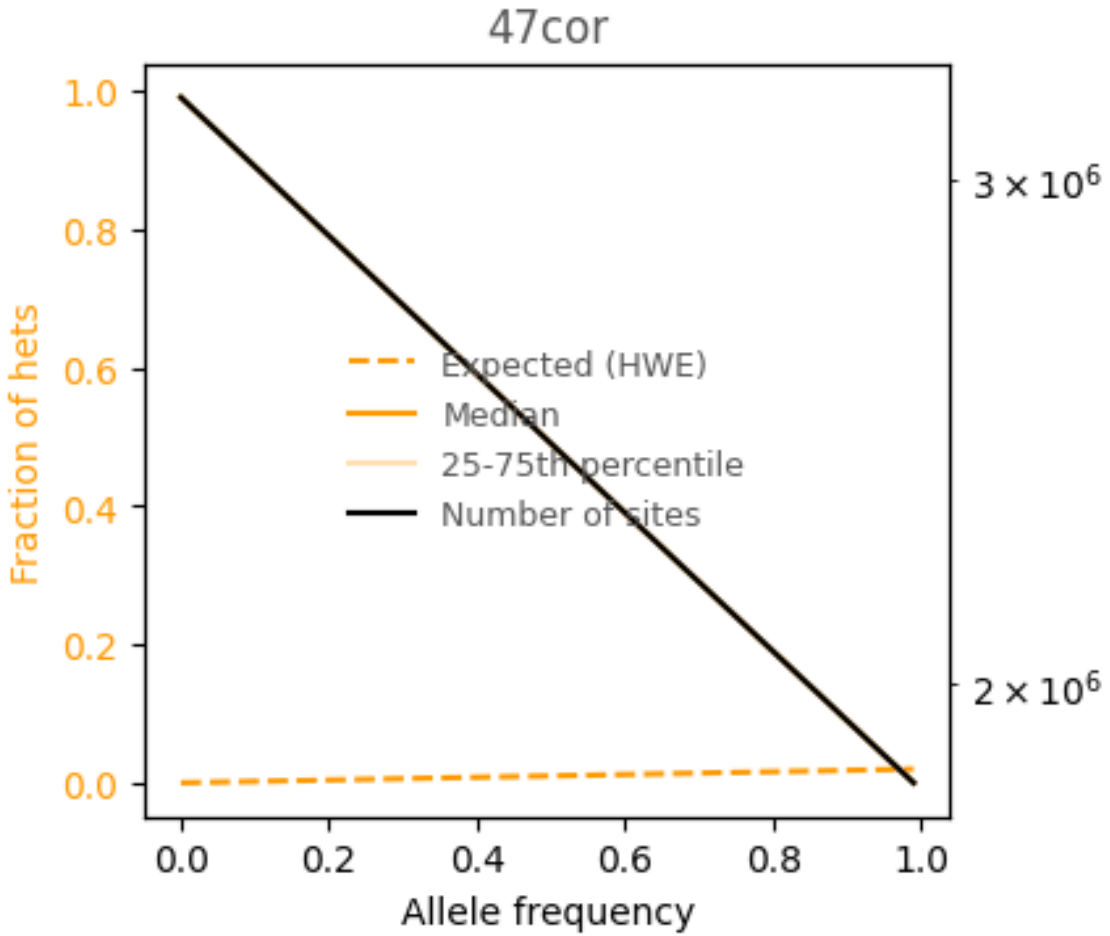
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types