Summary Numbers

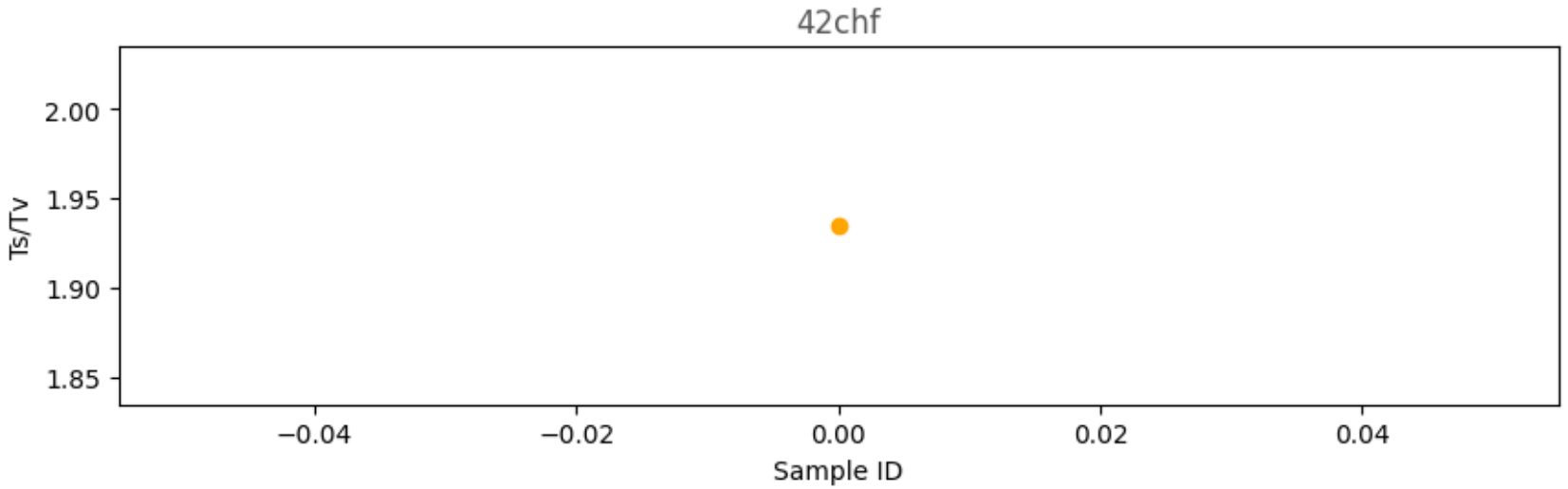
	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
42chf	4,088,882	1.93	1.94	954,325	–	0	0

* frameshift ratio: out/(out+in)

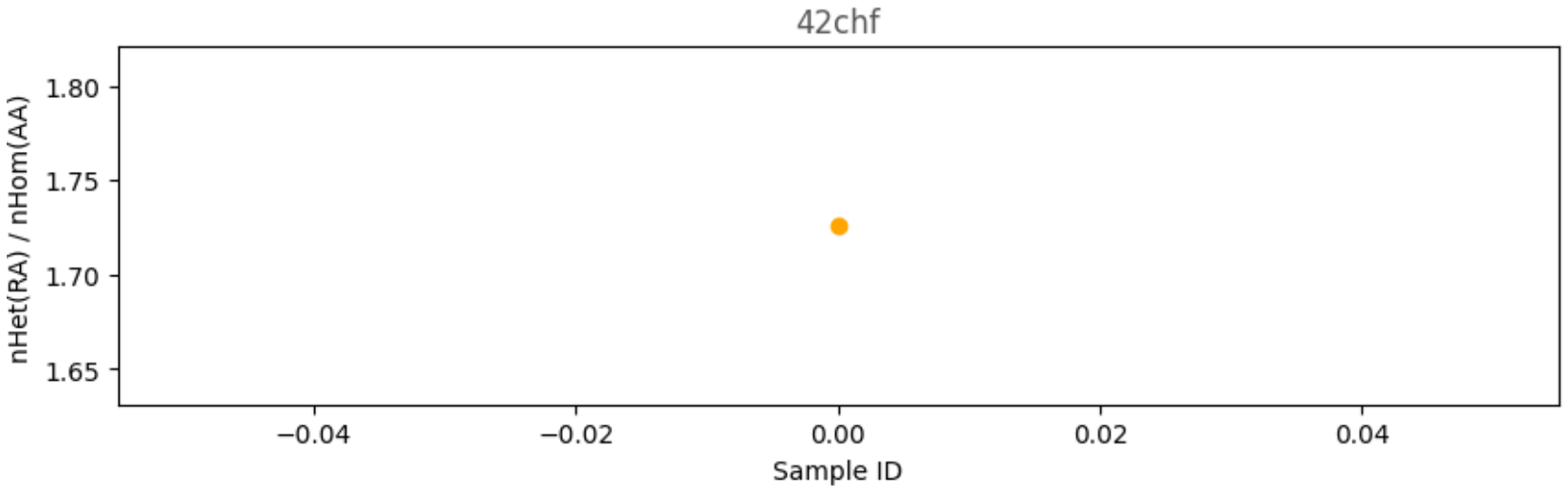
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
42chf	63.3%	1.91	68.5%	101,708	2,141

- 42chf .. `/ngc/projects2/gm/data/archive/2022/variants/snv/42chfuelf-110296344862-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02028-221019_A01176_BH3FN7DSX5-EXT_LABKA_NGCWGS-NGCWGS05902_snv_germline_raw.haplotype_caller.vcf.gz`

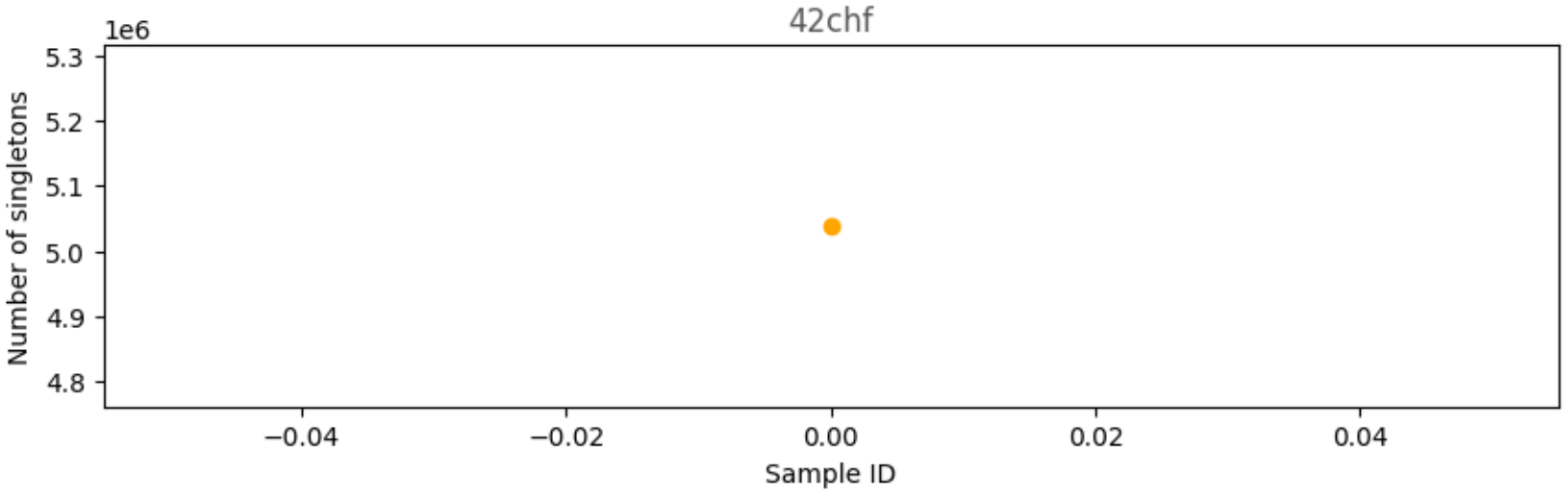
Ts/Tv by sample



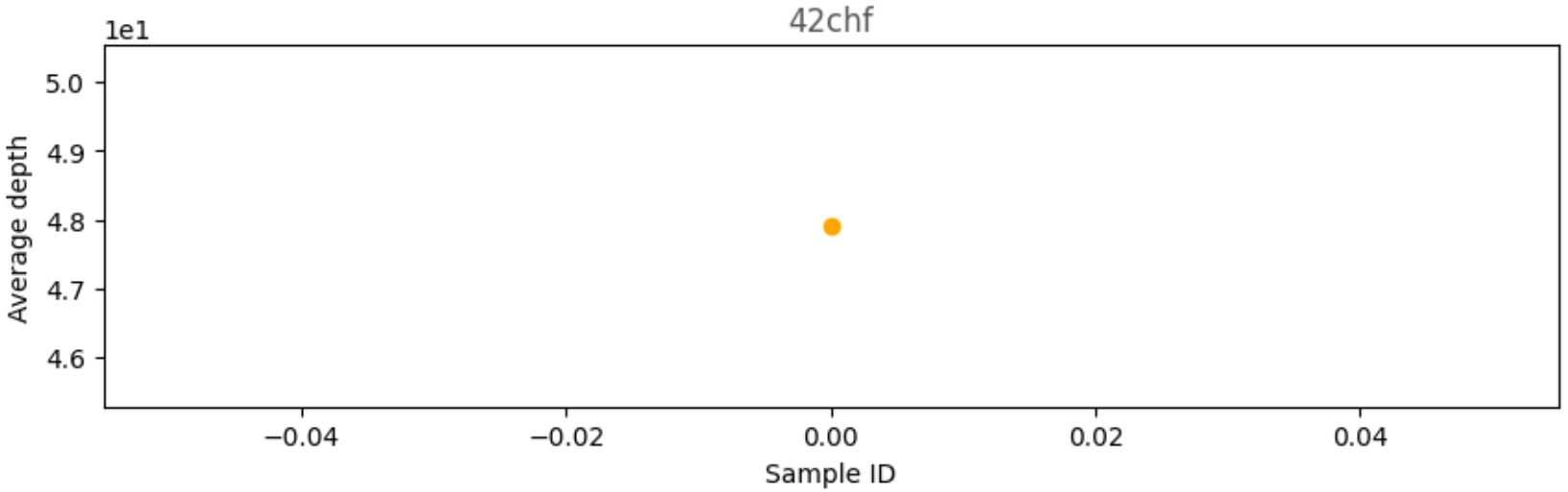
Hets vs non-ref Homs by sample



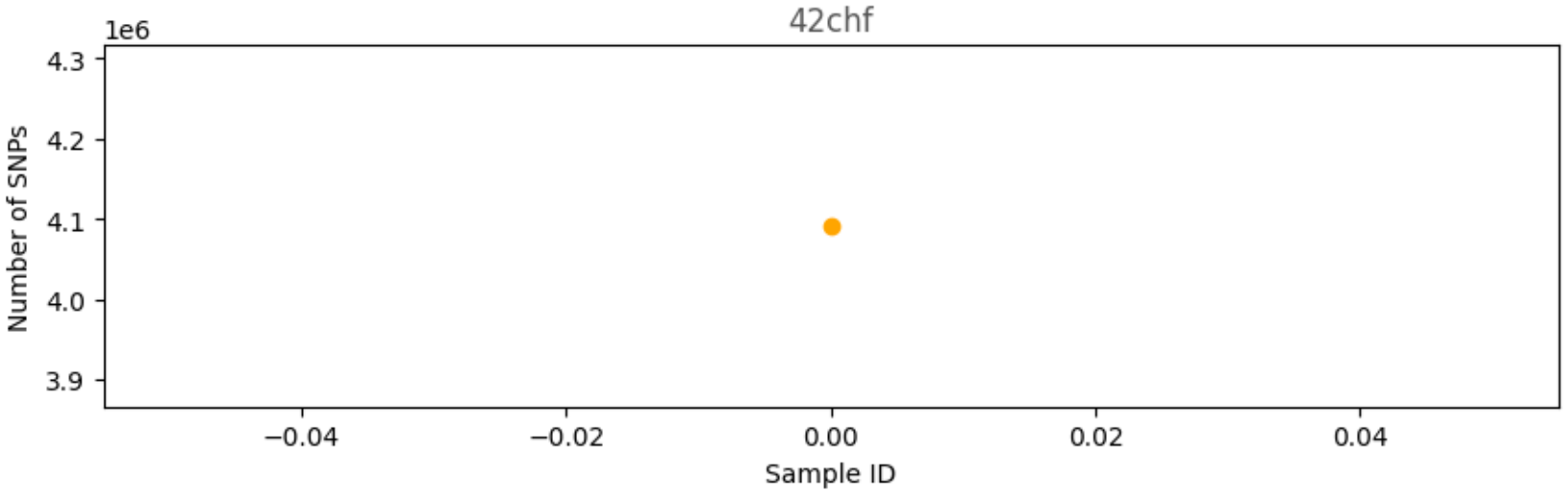
Singletons by sample (hets and homs)



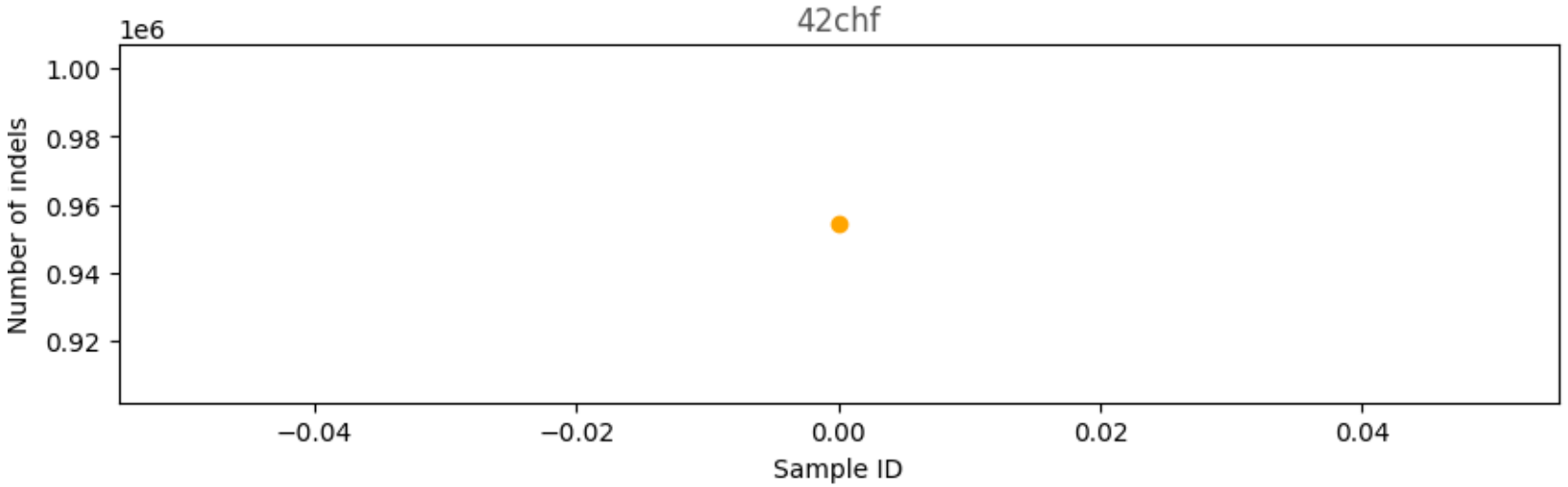
Average depth by sample



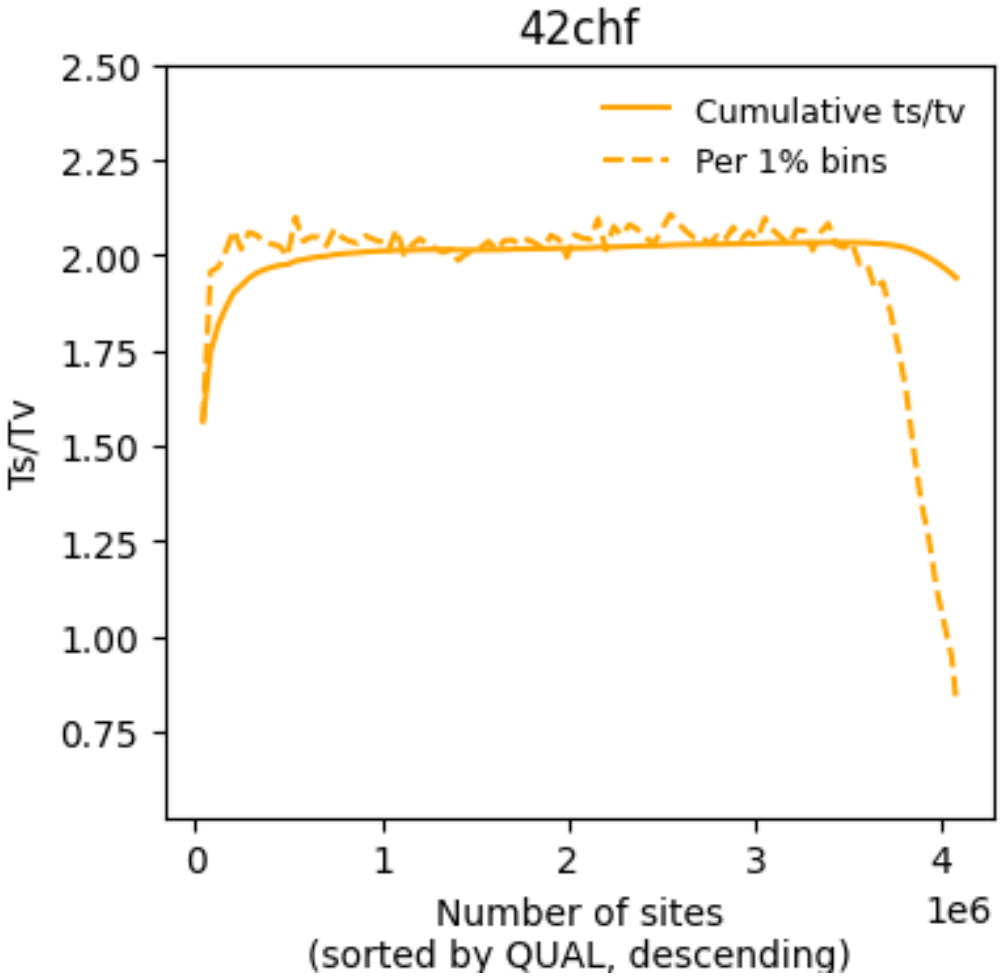
Number of SNPs by sample



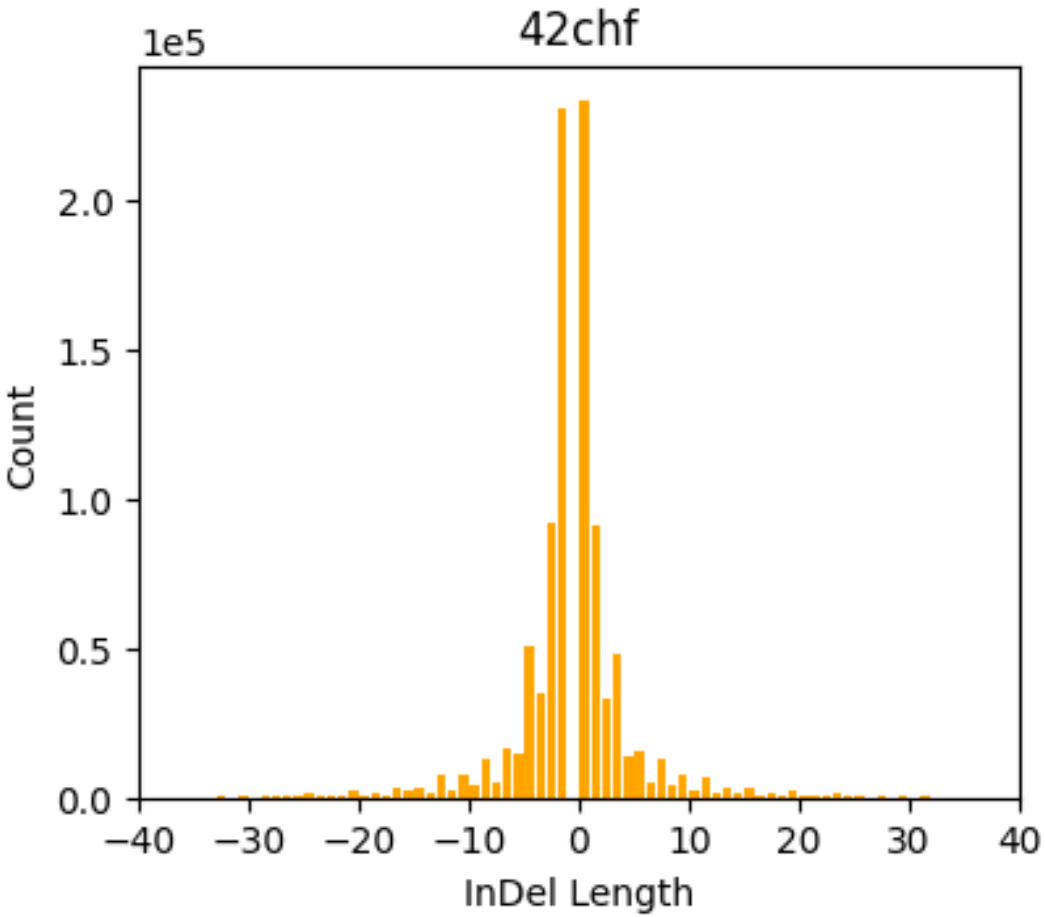
Number of indels by sample



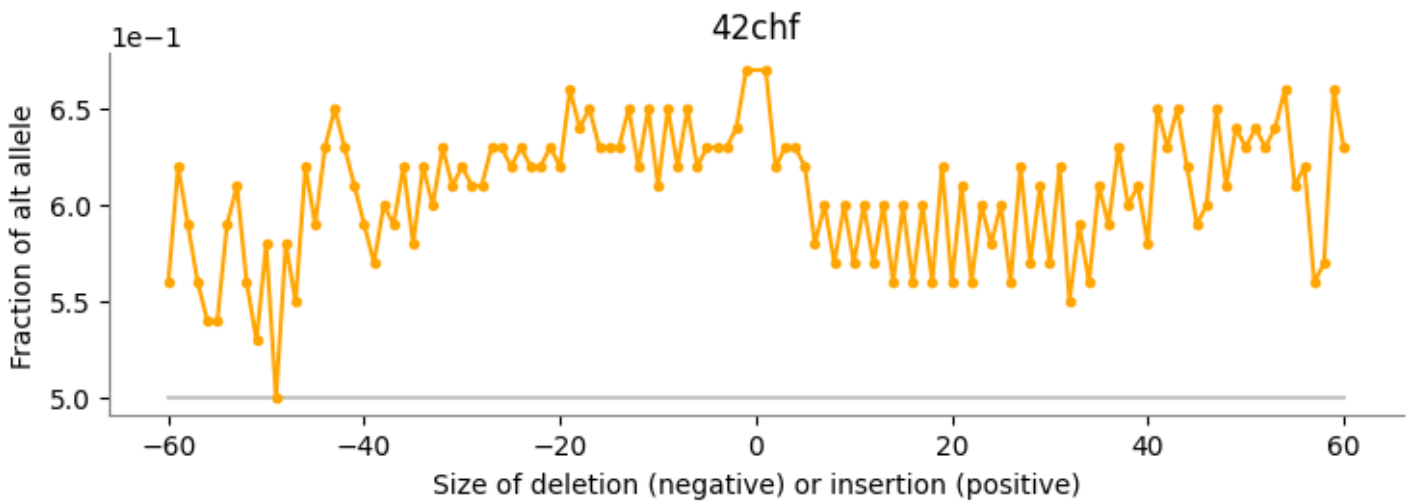
Ts/Tv stratified by QUAL



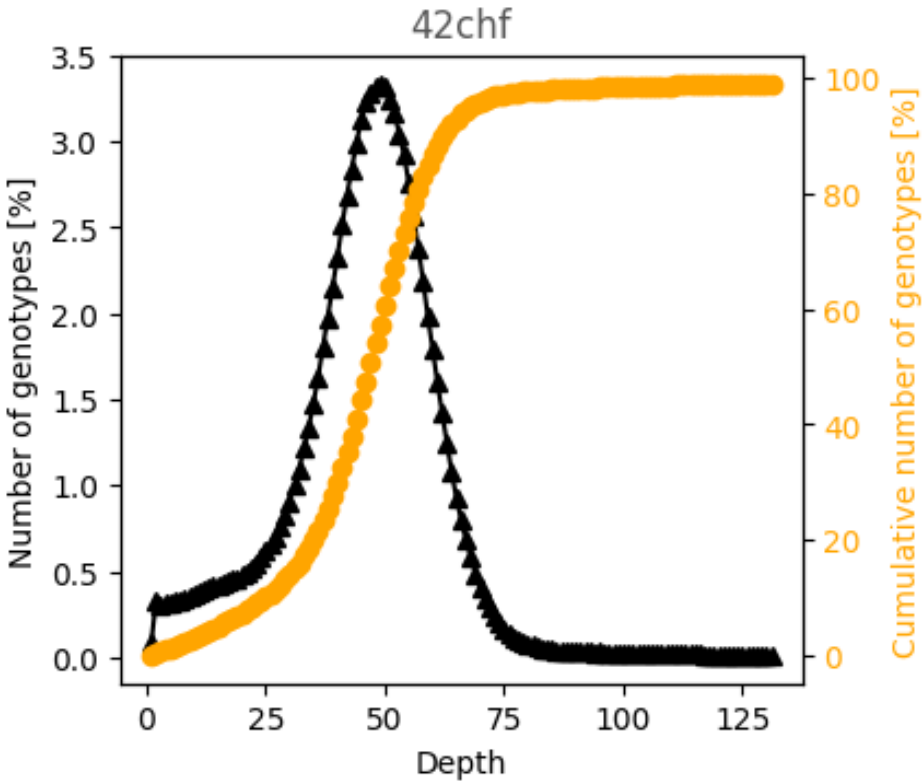
Indel distribution



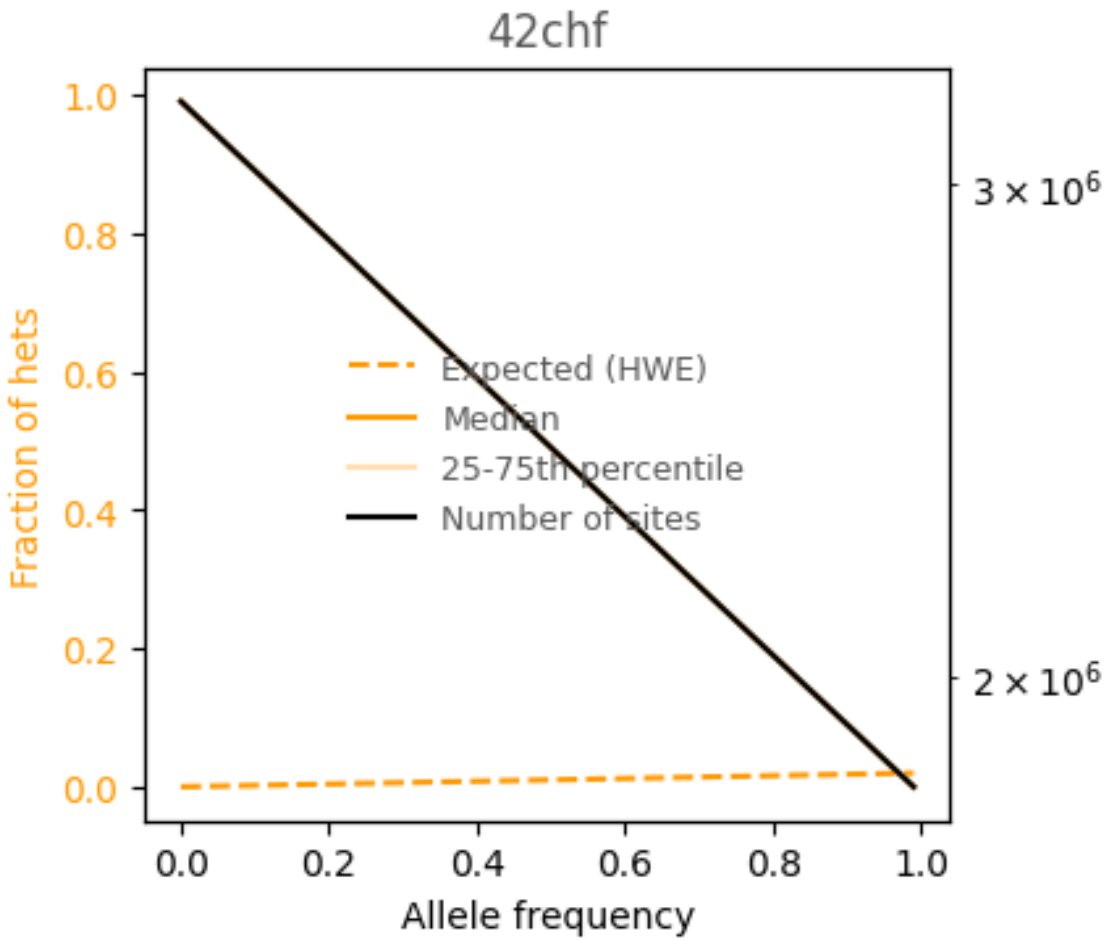
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types