

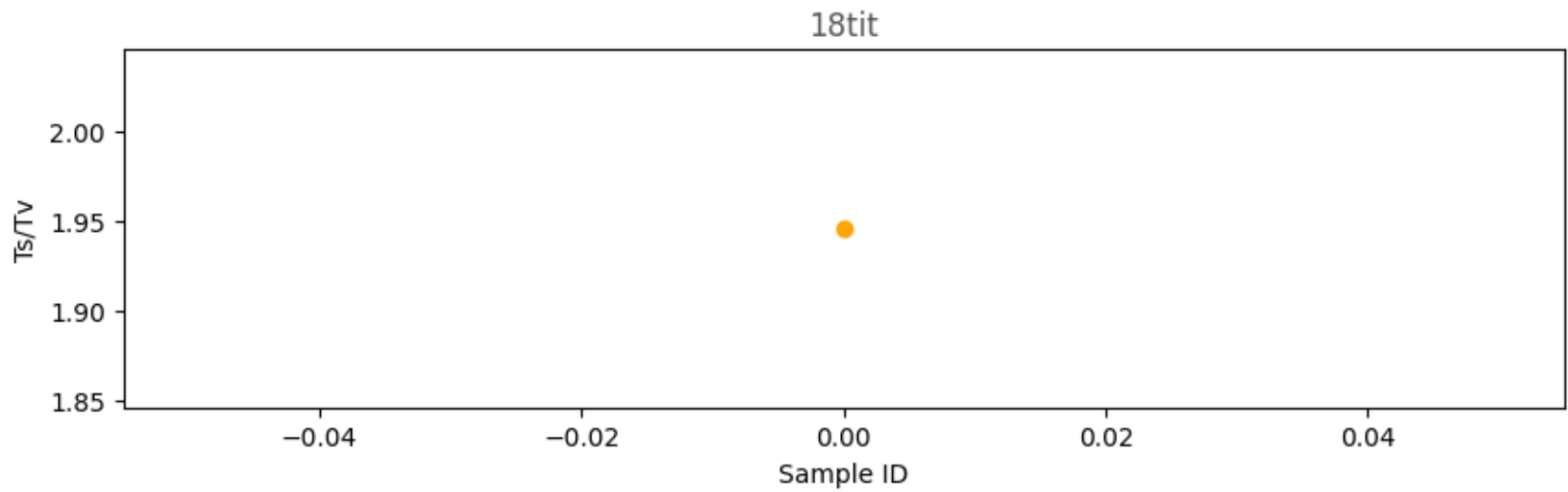
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
18tit	4,094,877	1.95	1.95	953,387	–	0	0
* frameshift ratio: out/(out+in)							

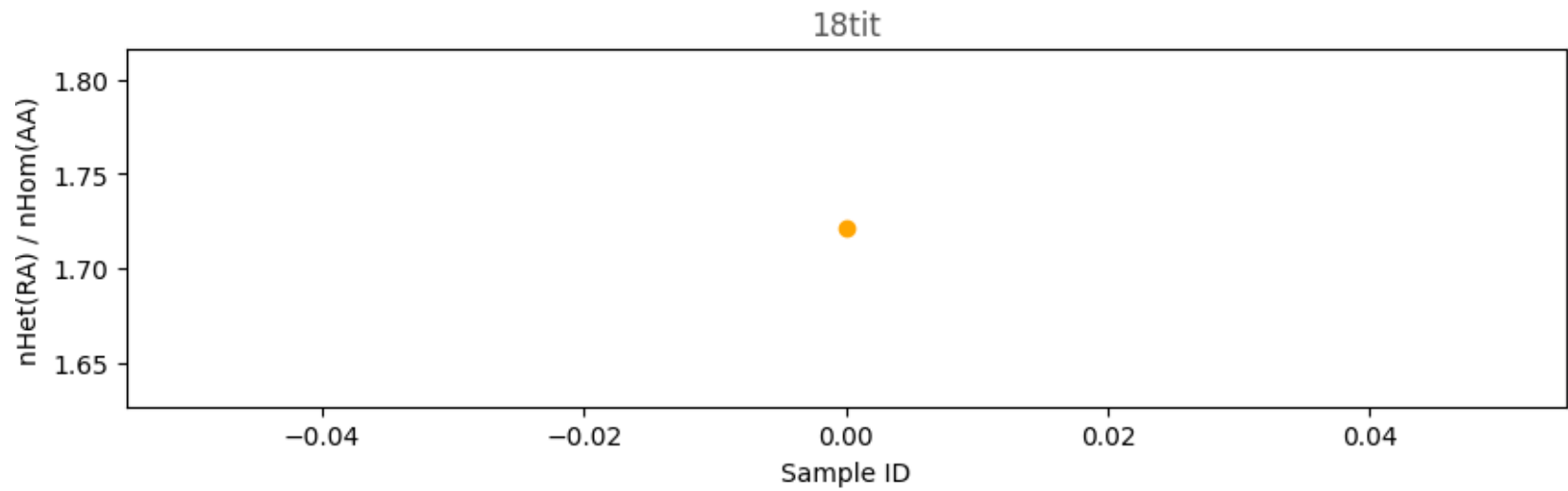
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
18tit	63.3%	1.92	68.2%	100,157	2,054

- 18tit .. /ngc/projects2/gm/data/archive/2022/variants/snv/18tithfof-103889936169-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01624-220921_A00559_AHYM7WDSX3-EXT_LAB_KA_NGCWGS-NGCWGS05414_22RKG016939_snv_germline_raw.haplotype_caller.vcf.gz

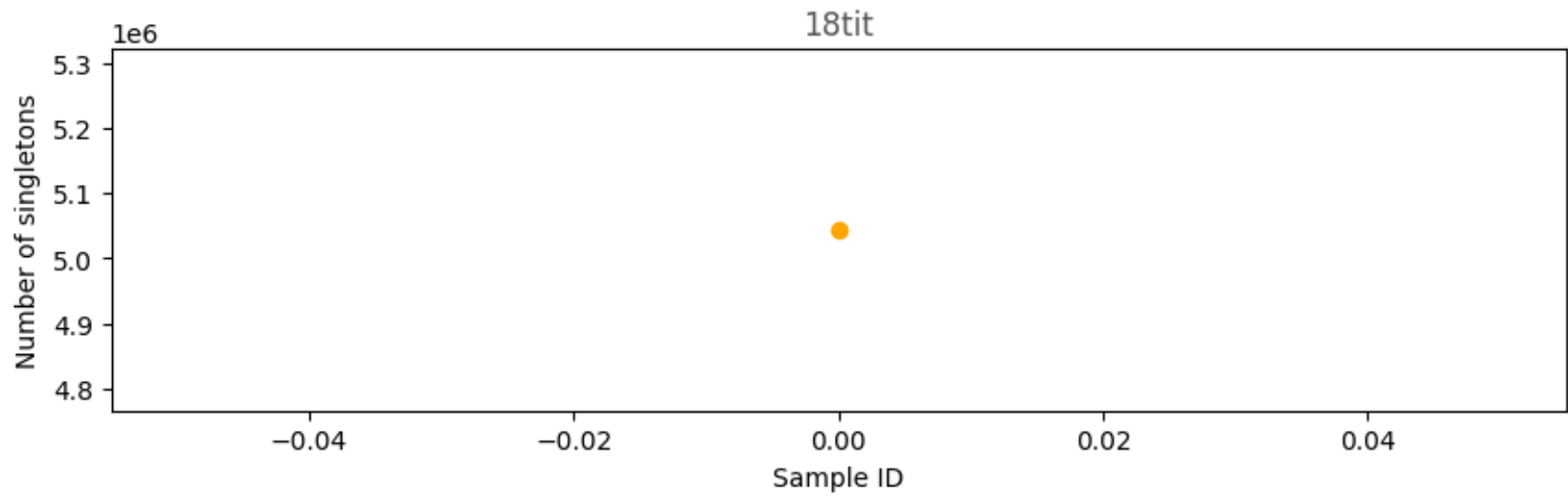
Ts/Tv by sample



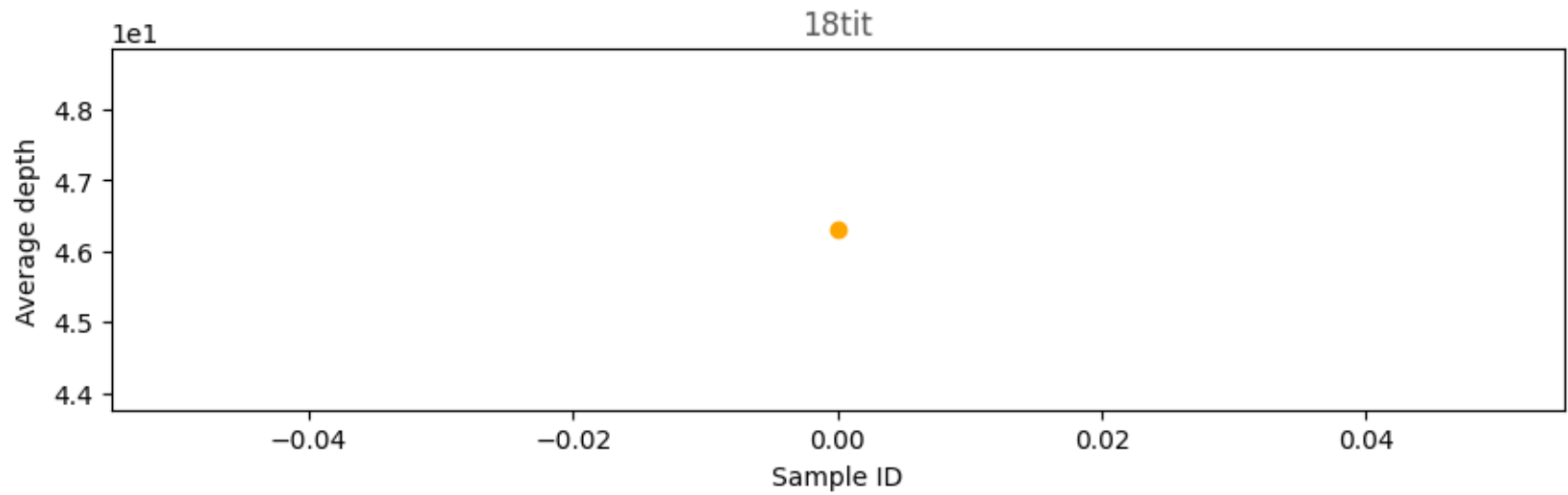
Hets vs non-ref Homs by sample



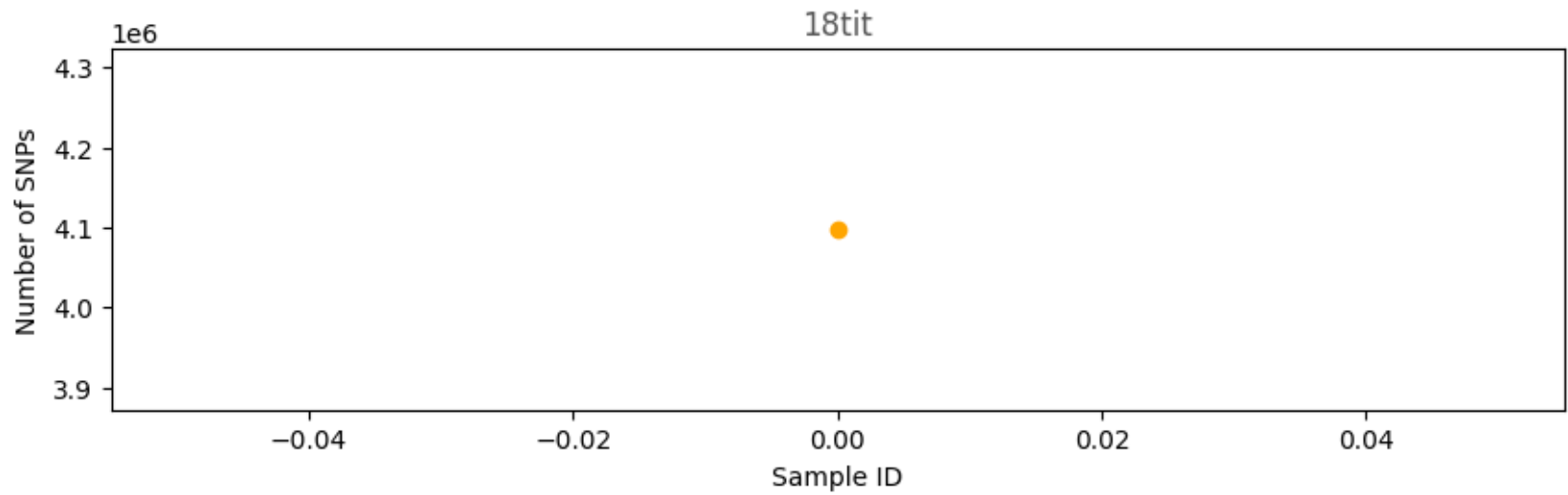
Singletons by sample (hets and homs)



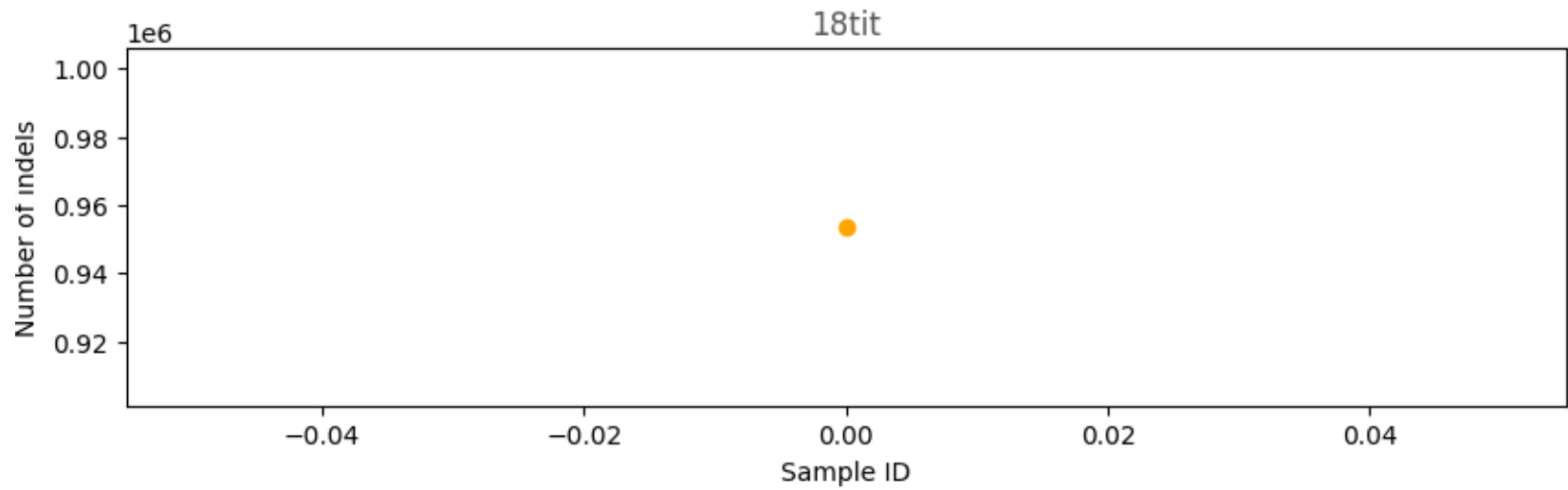
Average depth by sample



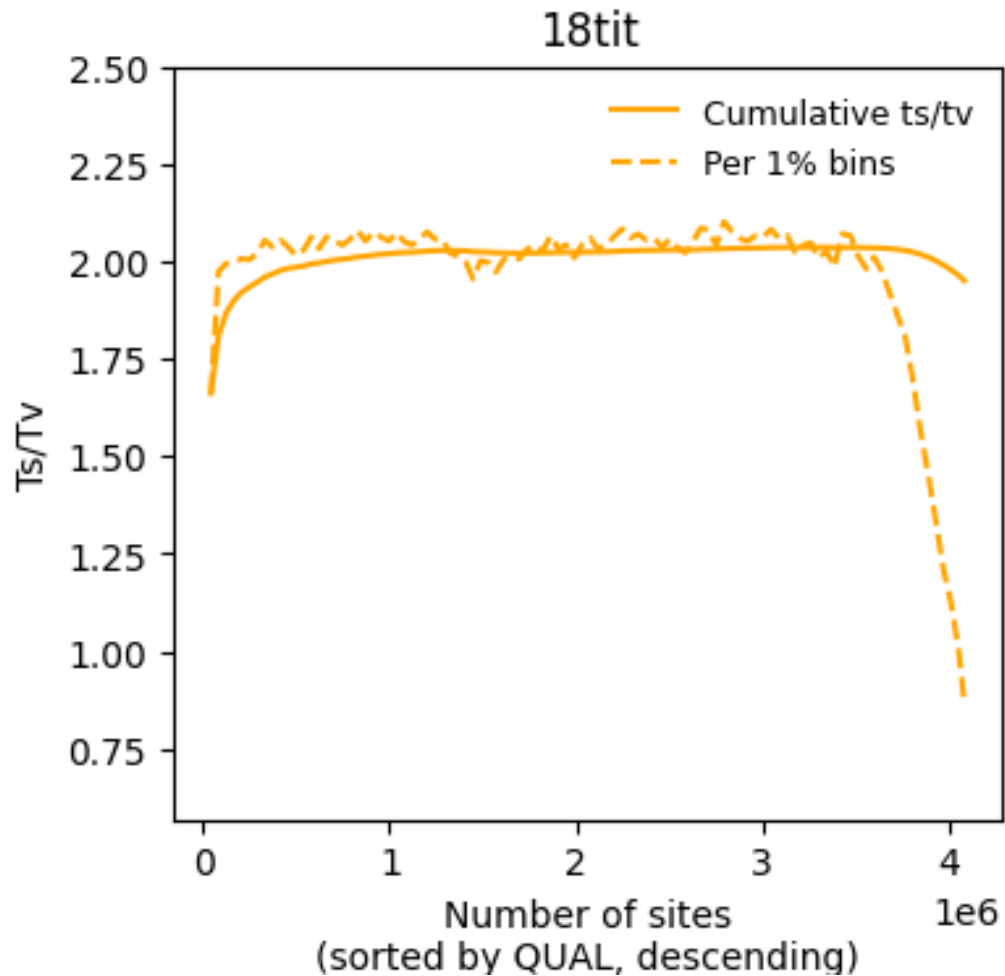
Number of SNPs by sample



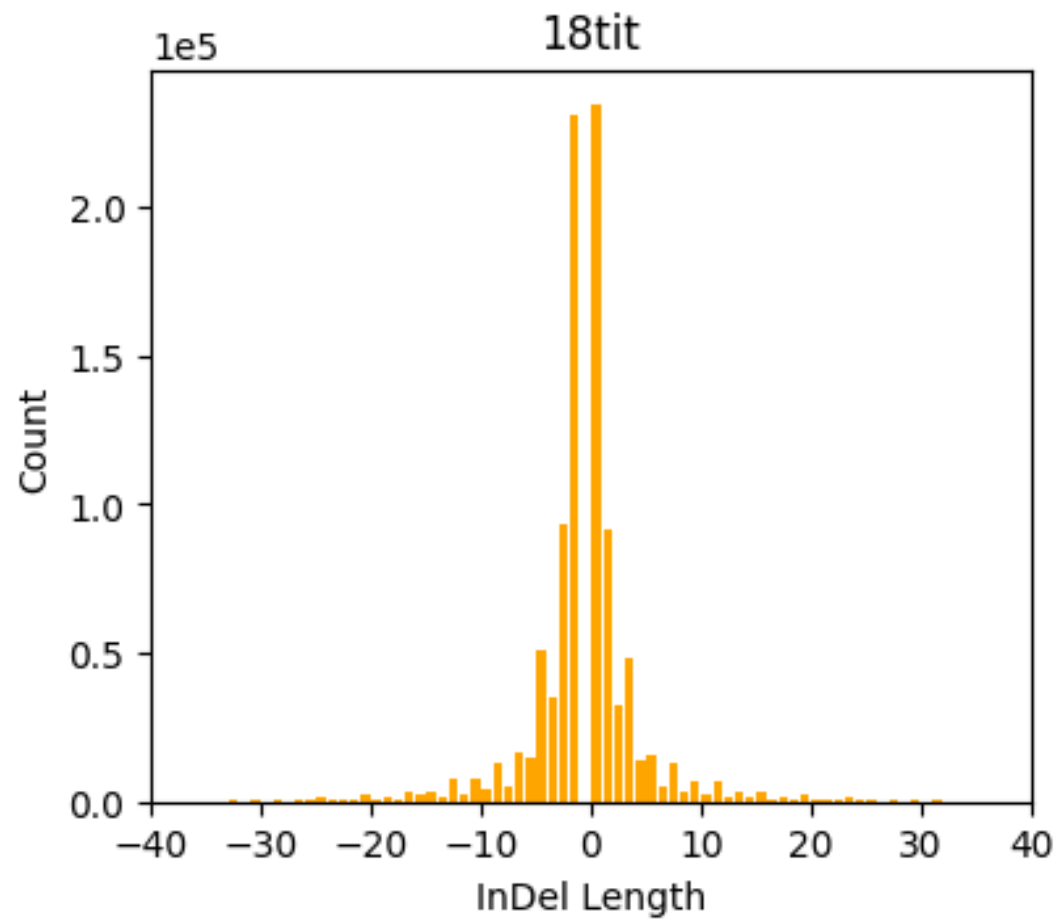
Number of indels by sample



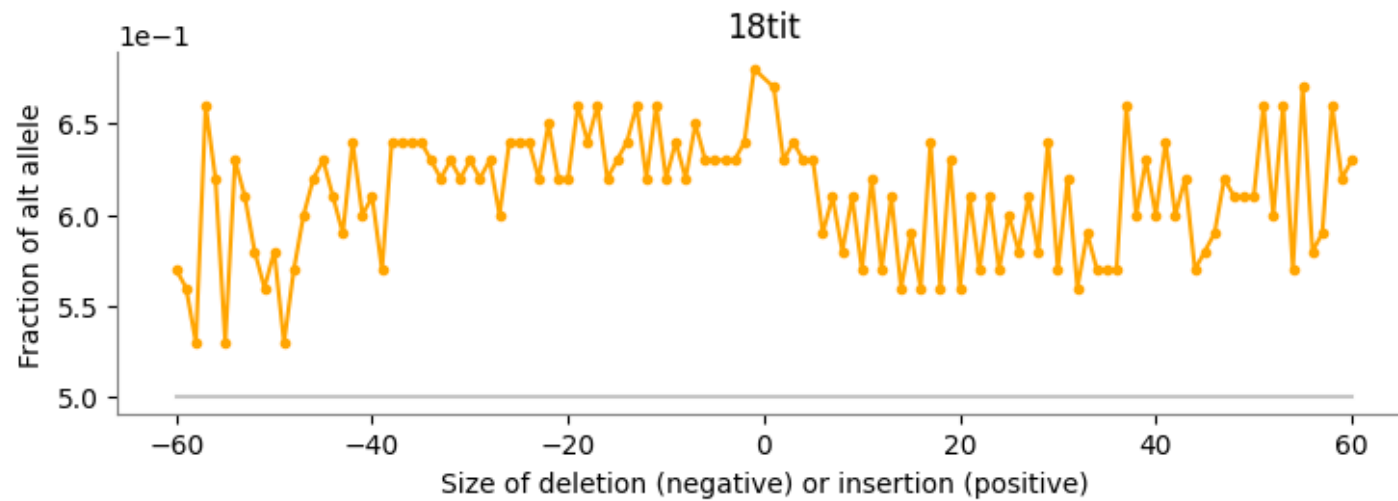
Ts/Tv stratified by QUAL



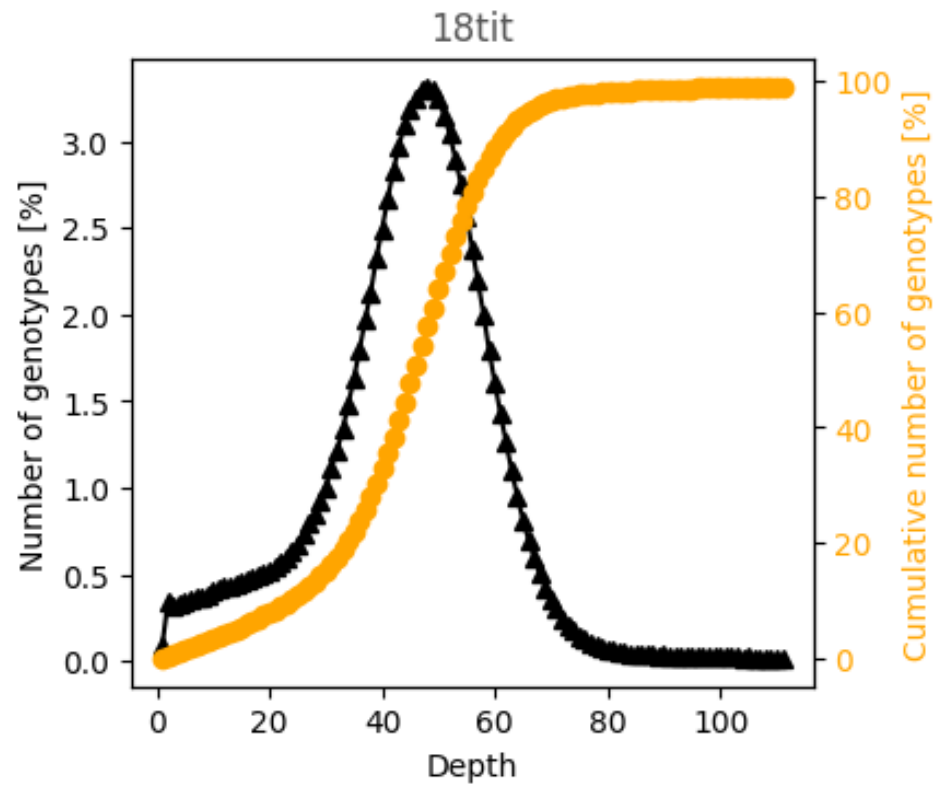
Indel distribution



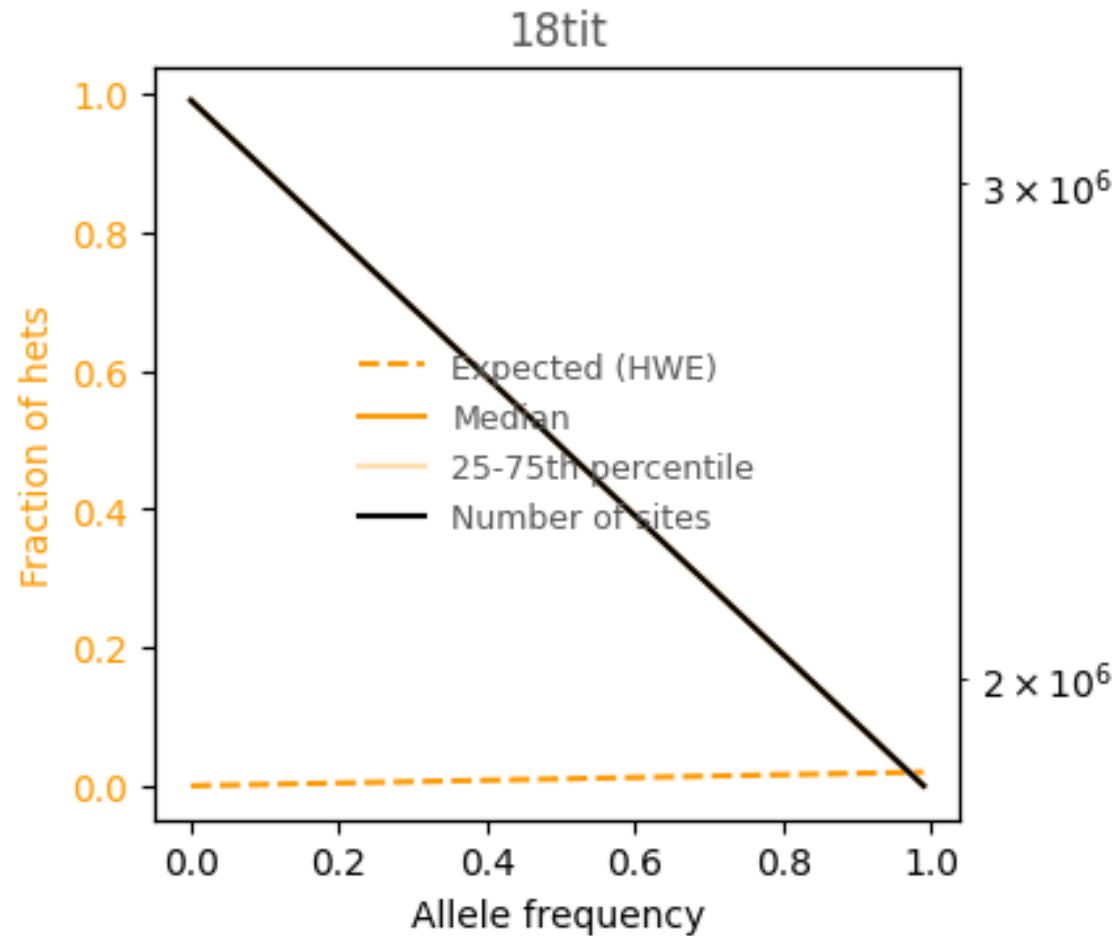
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

