

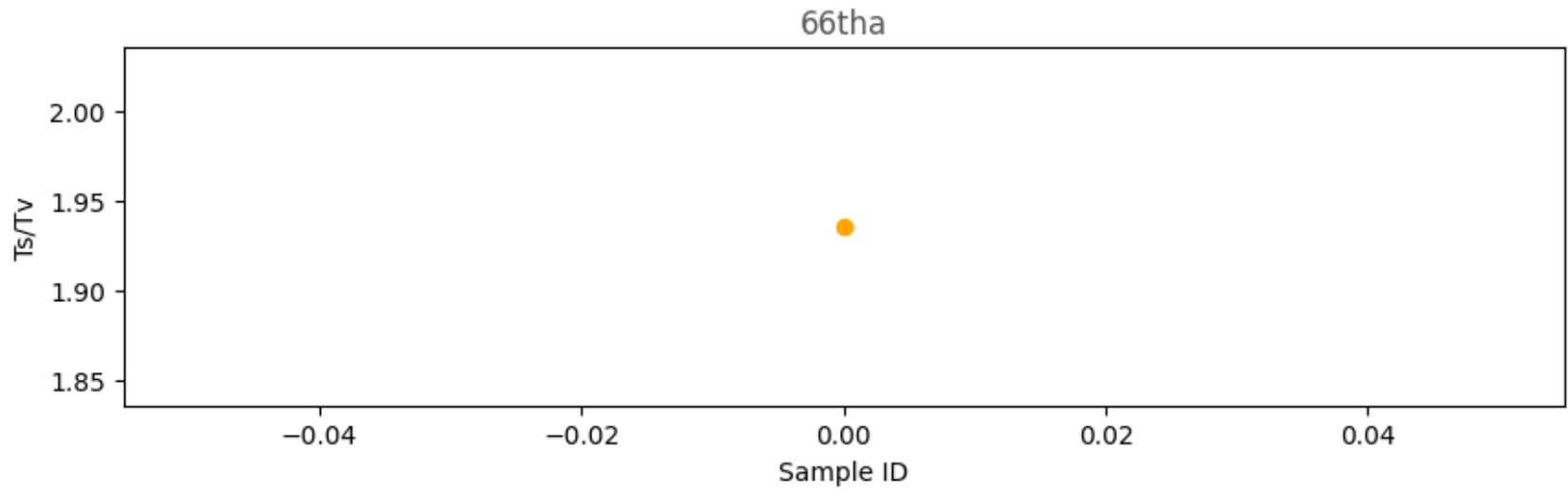
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
66tha	4,039,433	1.94	1.94	942,883	–	0	0
* frameshift ratio: out/(out+in)							

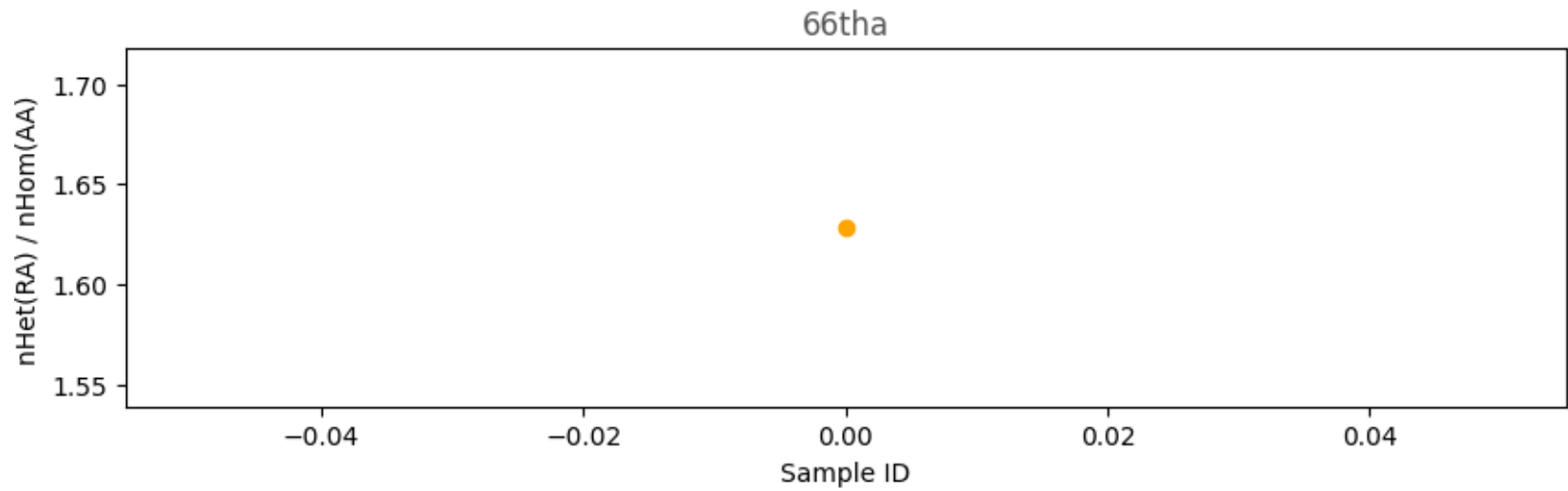
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
66tha	62.0%	1.90	66.4%	96,985	1,984

- 66tha .. /ngc/projects2/gm/data/archive/2022/variants/snv/66thasrem-103908482799-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01406-220912_A01961_BHYNTFDSX3-EXT_LAB
KA_NGCWGS-NGCWGS05182-snv_germline_raw.haplotype_caller.vcf.gz

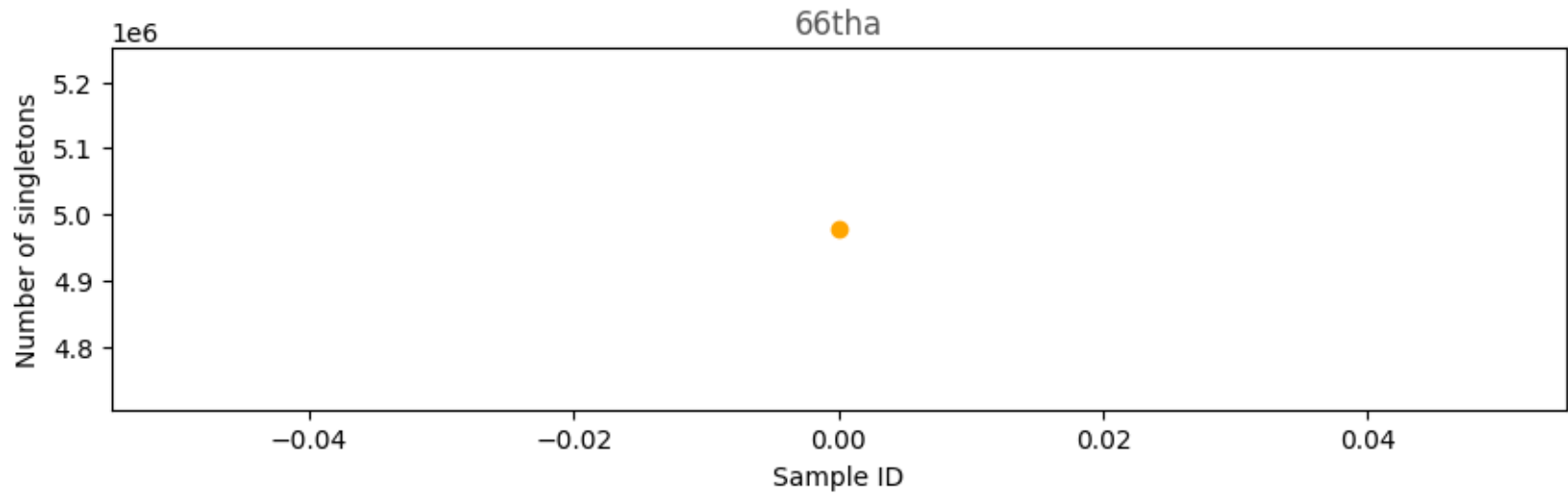
Ts/Tv by sample



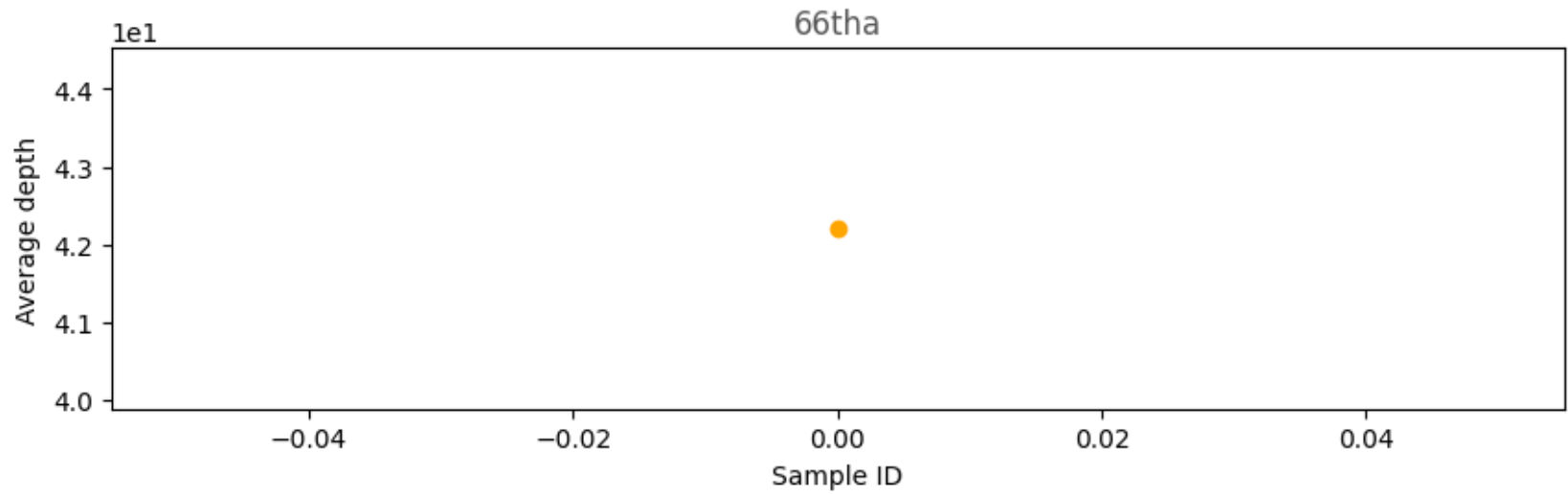
Hets vs non-ref Homs by sample



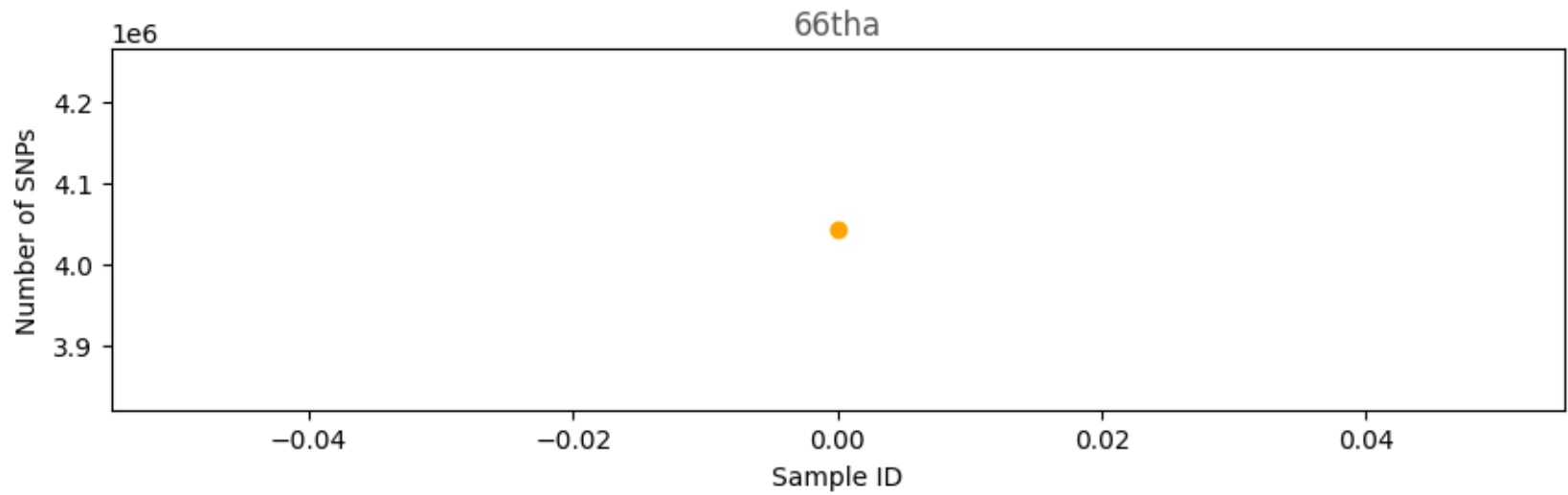
Singletons by sample (hets and homs)



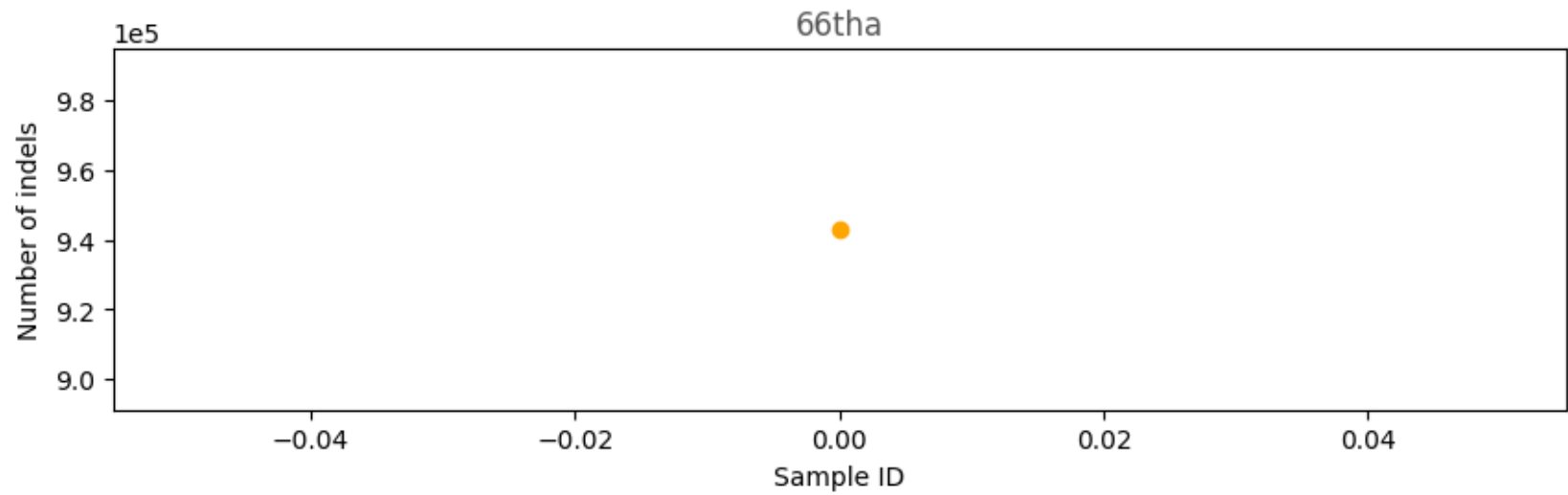
Average depth by sample



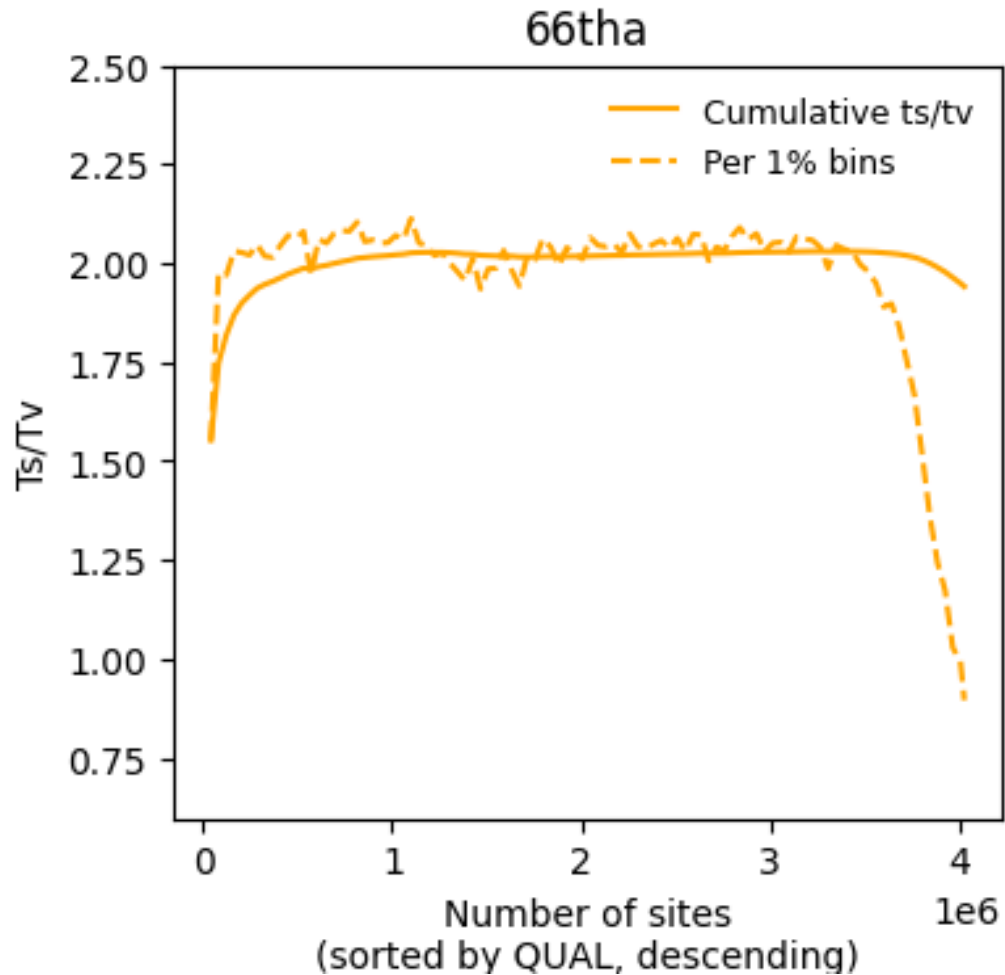
Number of SNPs by sample



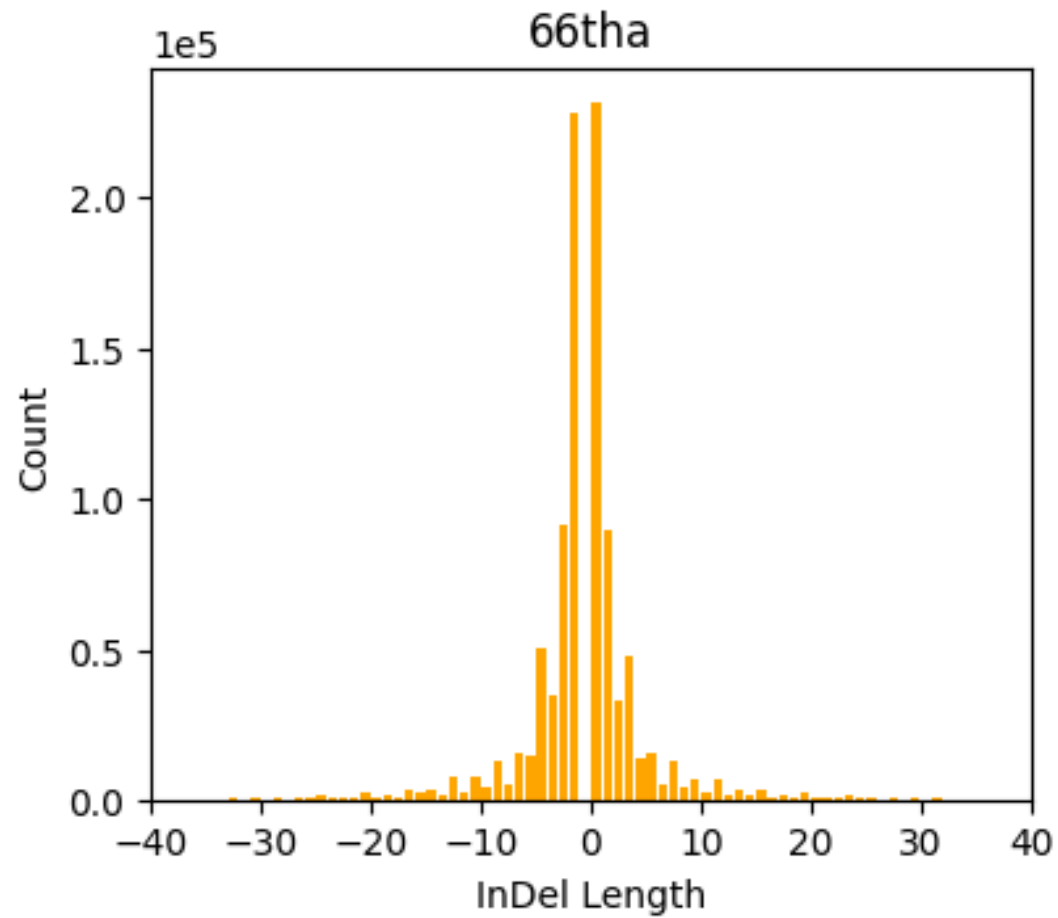
Number of indels by sample



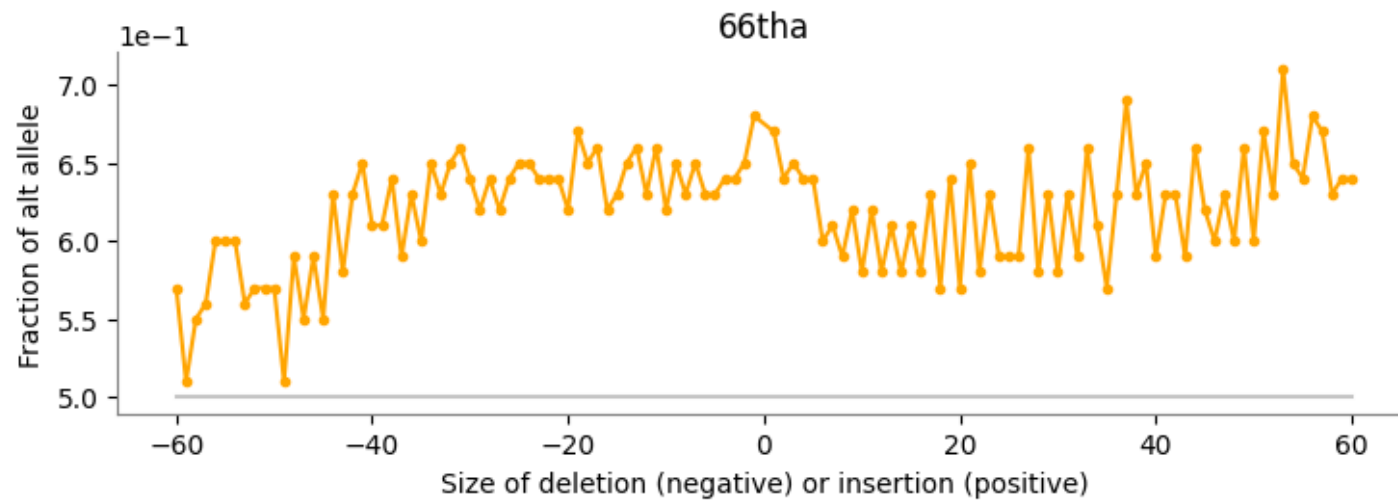
Ts/Tv stratified by QUAL



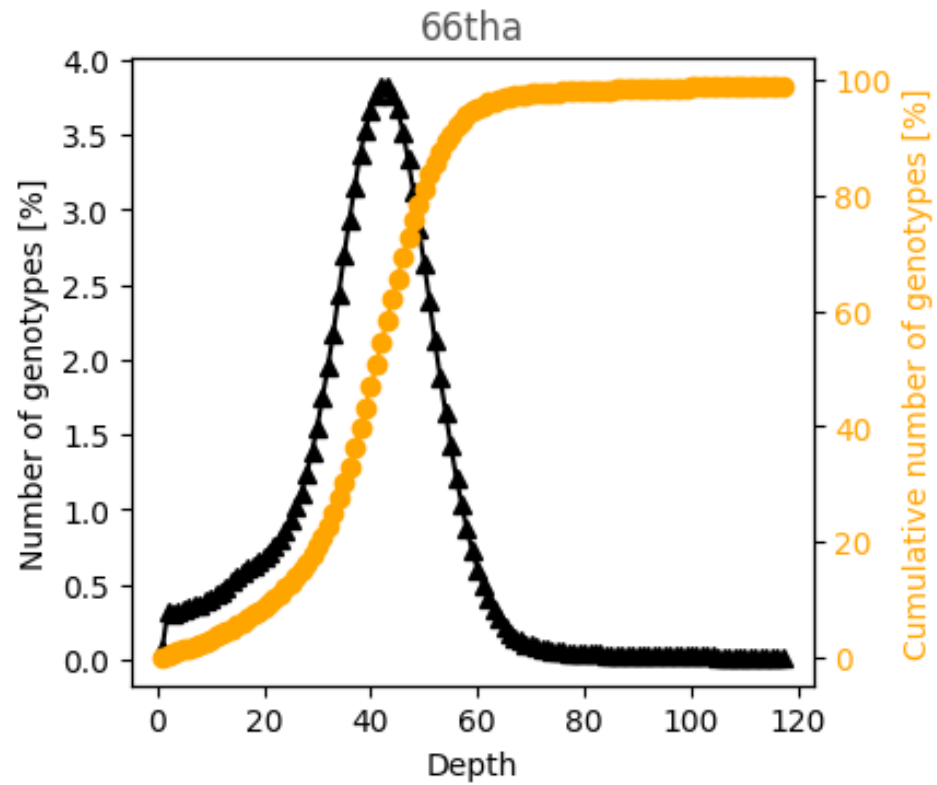
Indel distribution



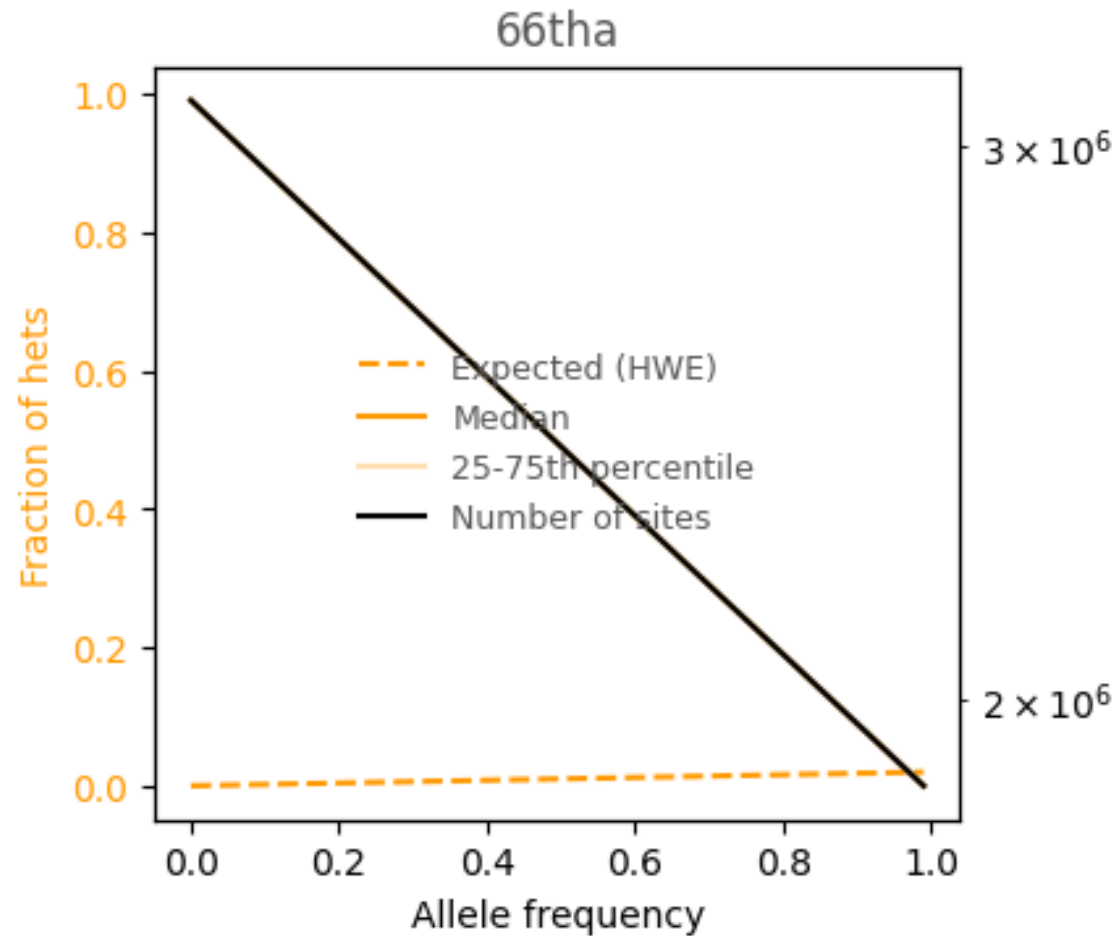
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

