

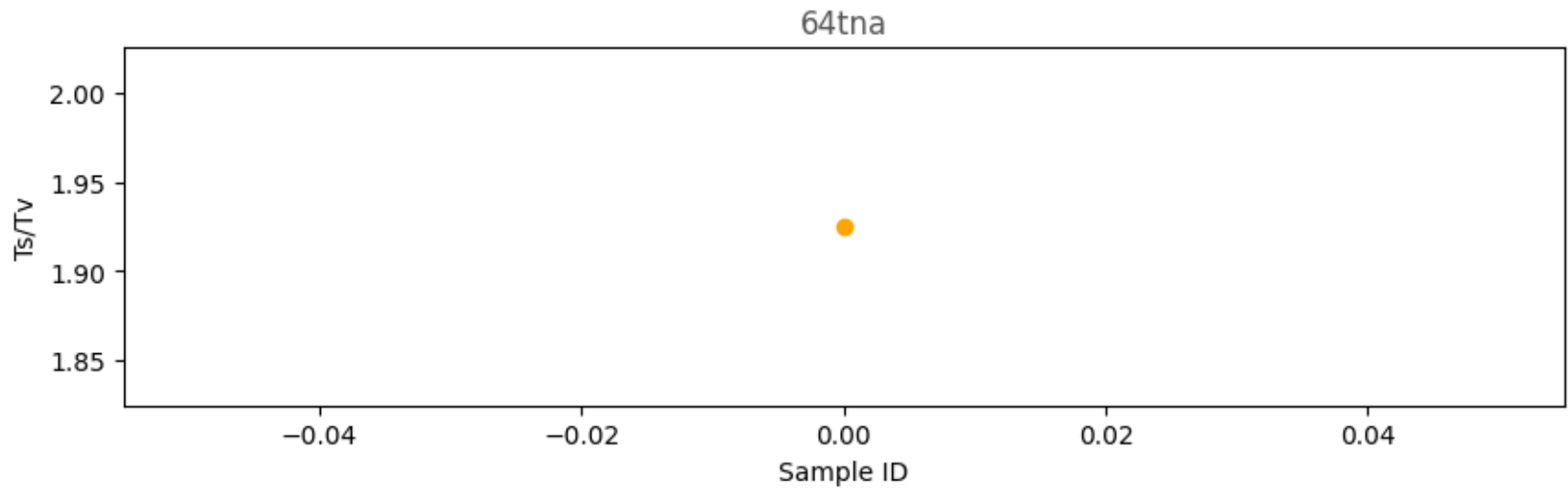
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
64tna	4,120,780	1.92	1.93	964,269	–	0	0
* frameshift ratio: out/(out+in)							

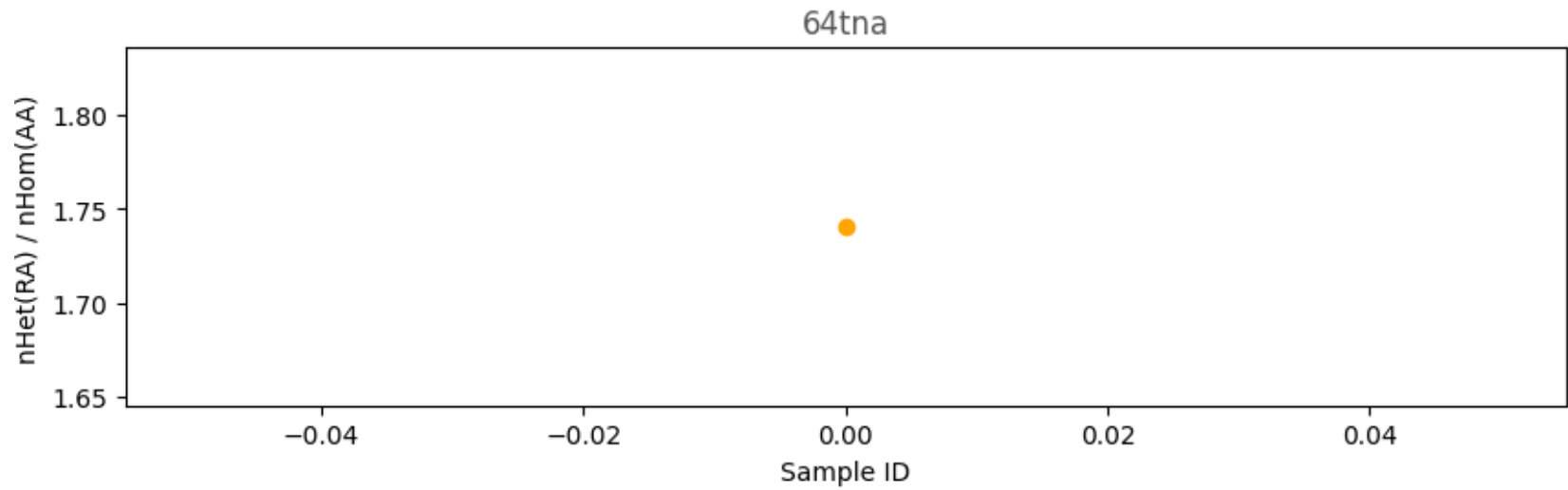
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
64tna	63.5%	1.89	68.8%	104,355	2,069

- 64tna .. /ngc/projects2/gm/data/archive/2022/variants/snv/64tnatusf-4138942654-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220518_A01411_AHJ5W2DSX3-RHGM_LABKA_WGS-WG_S04283_snv_germline_raw.haplotype_caller.vcf.gz

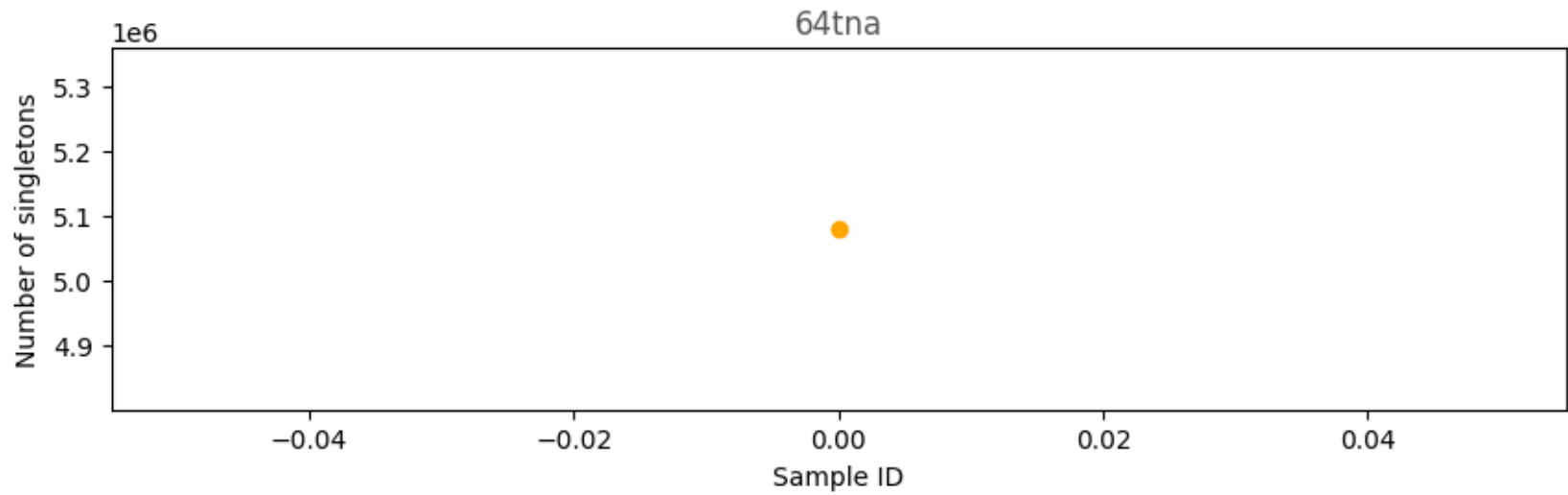
Ts/Tv by sample



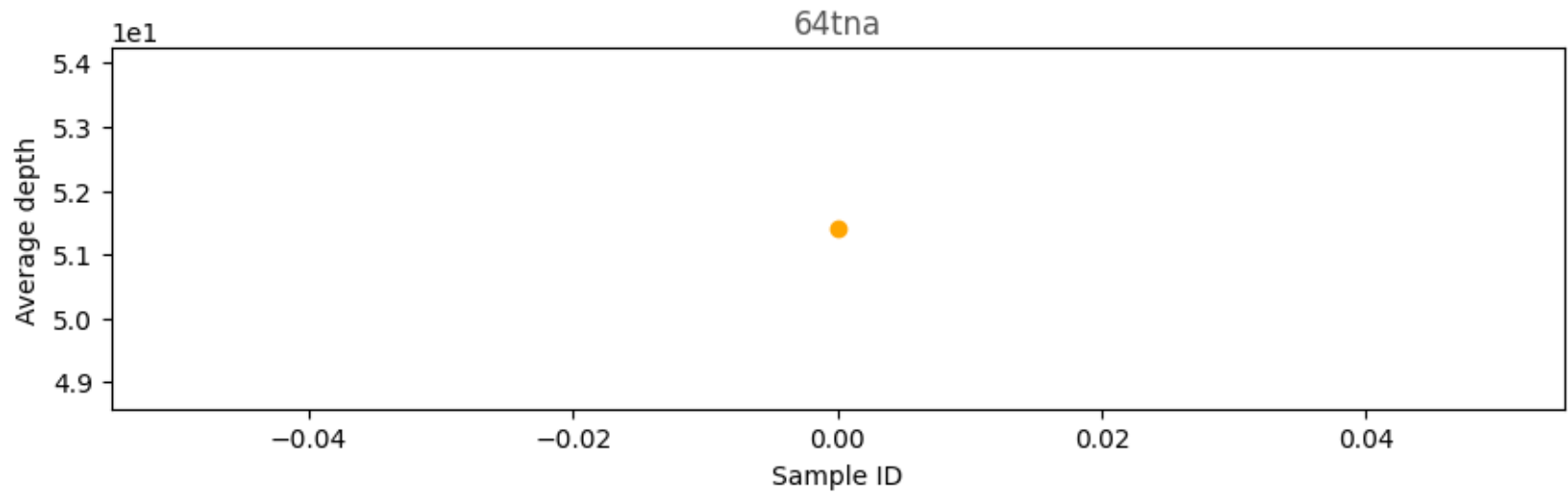
Hets vs non-ref Homs by sample



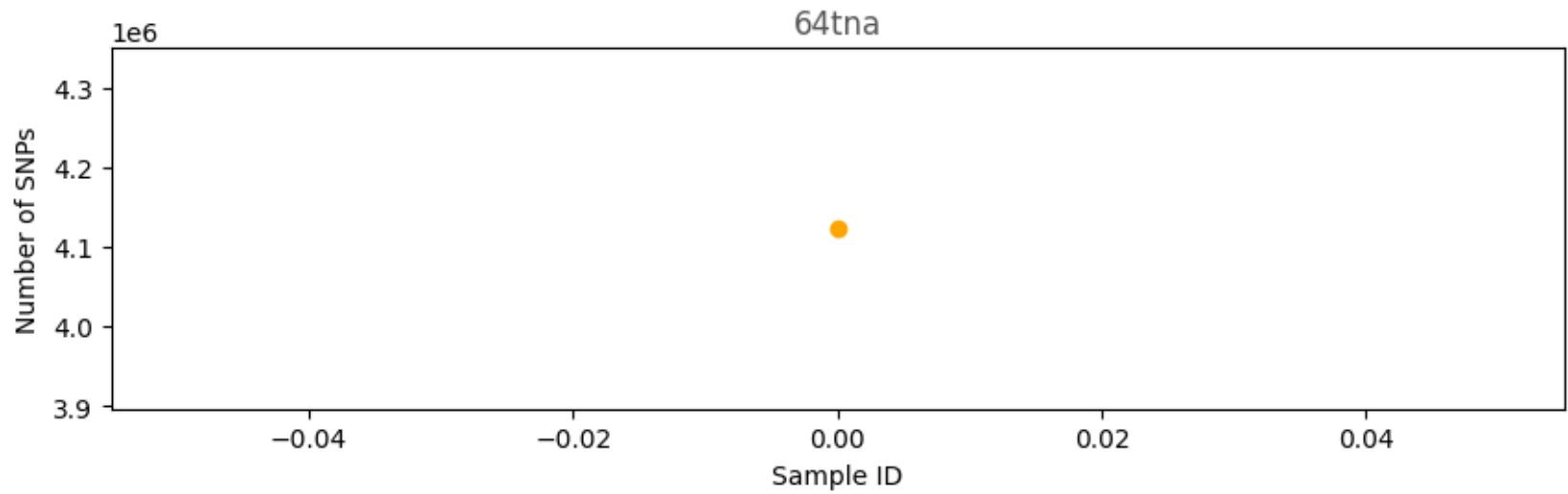
Singletons by sample (hets and homs)



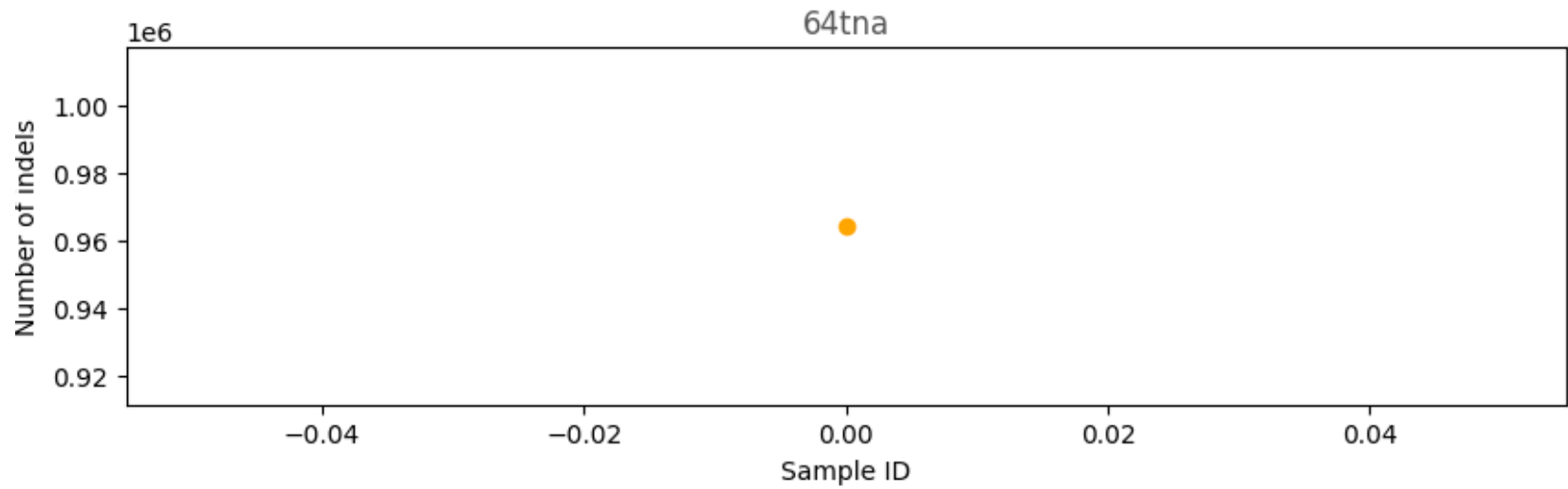
Average depth by sample



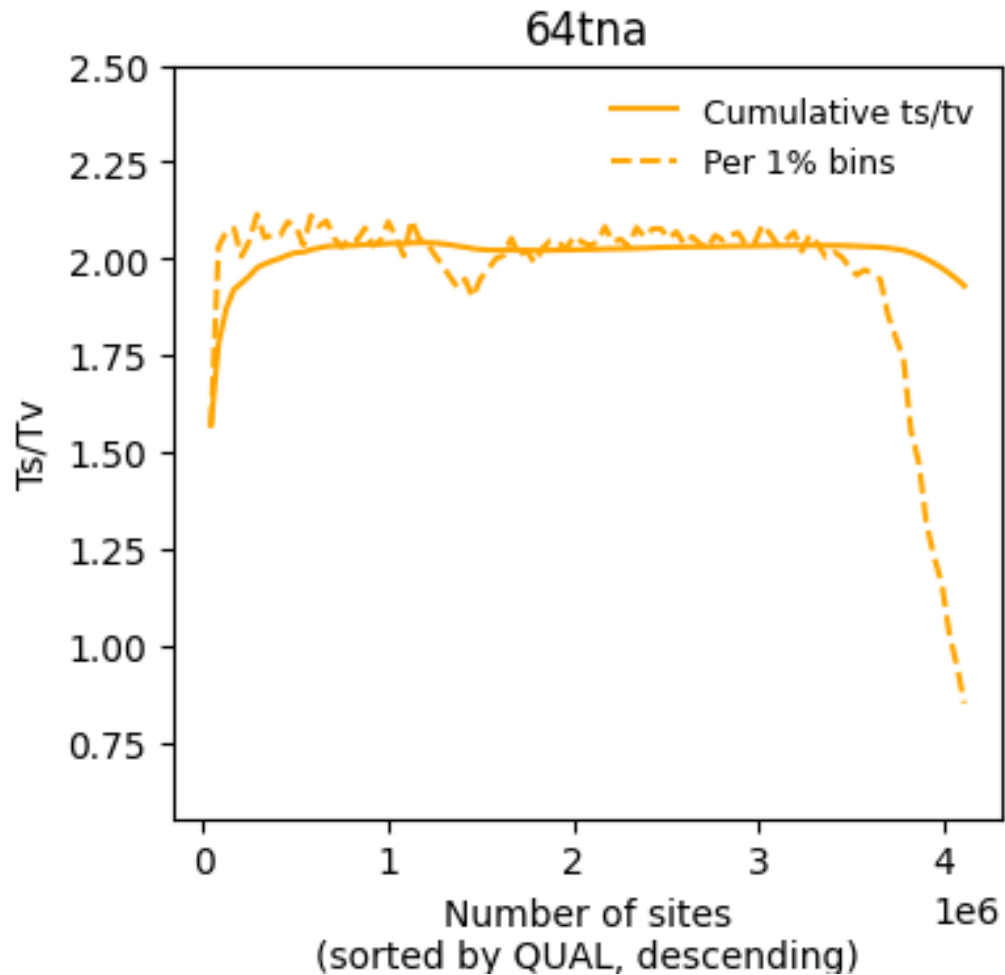
Number of SNPs by sample



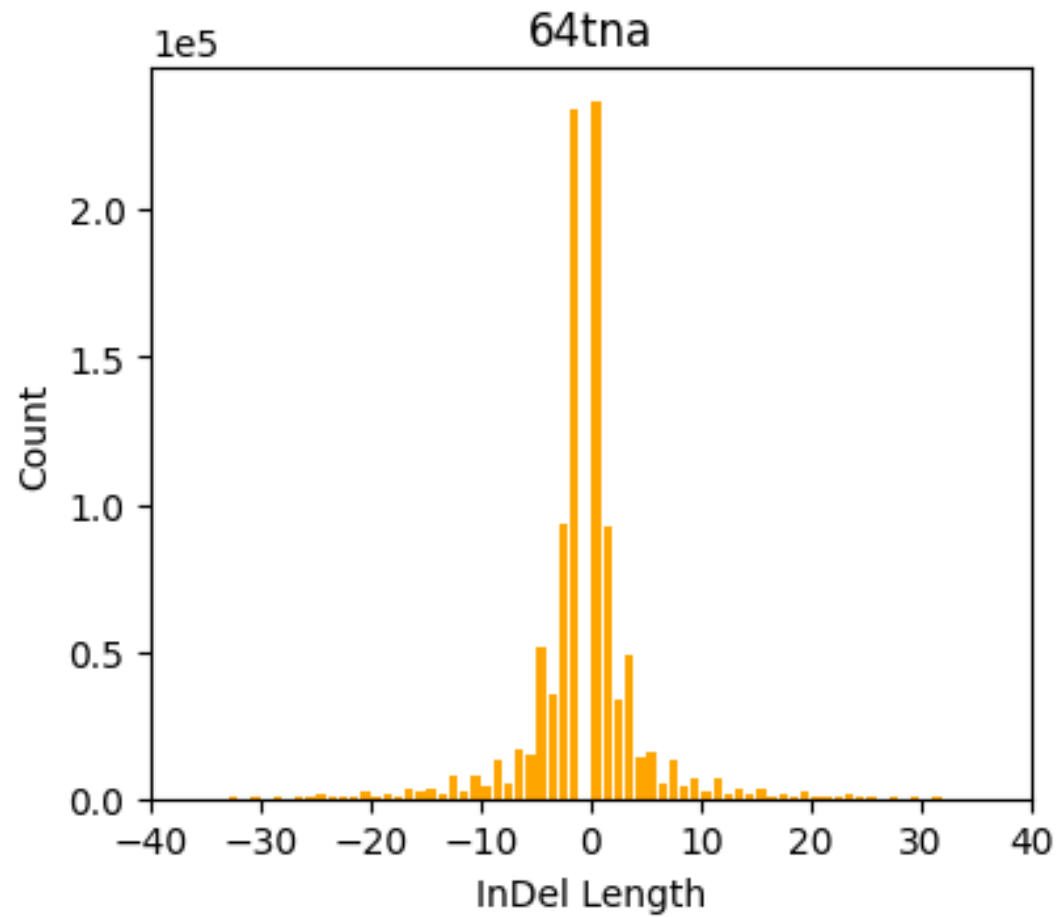
Number of indels by sample



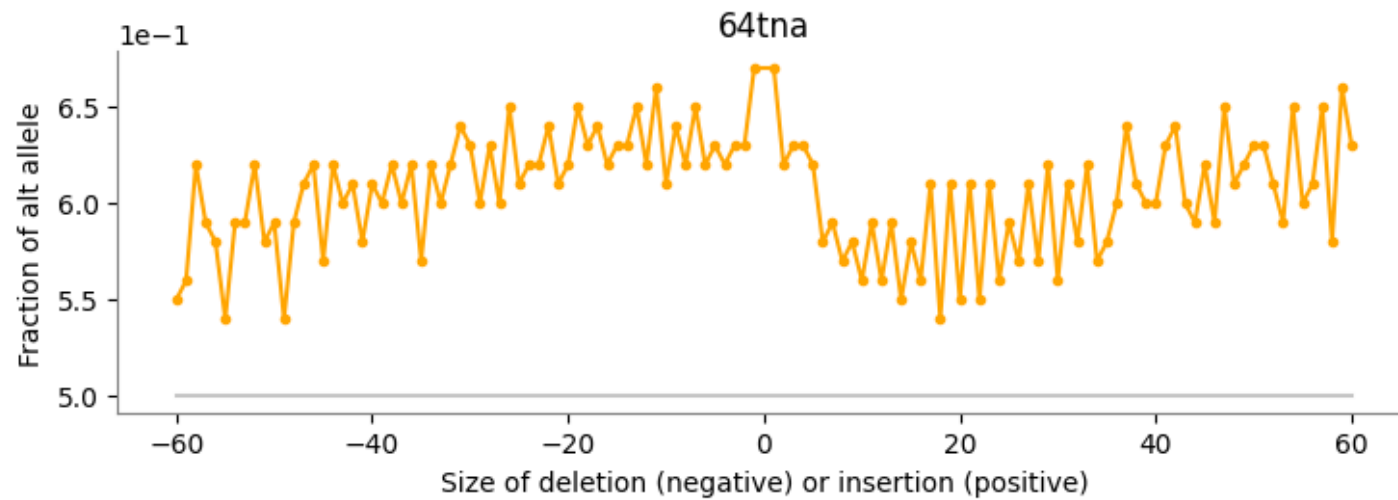
Ts/Tv stratified by QUAL



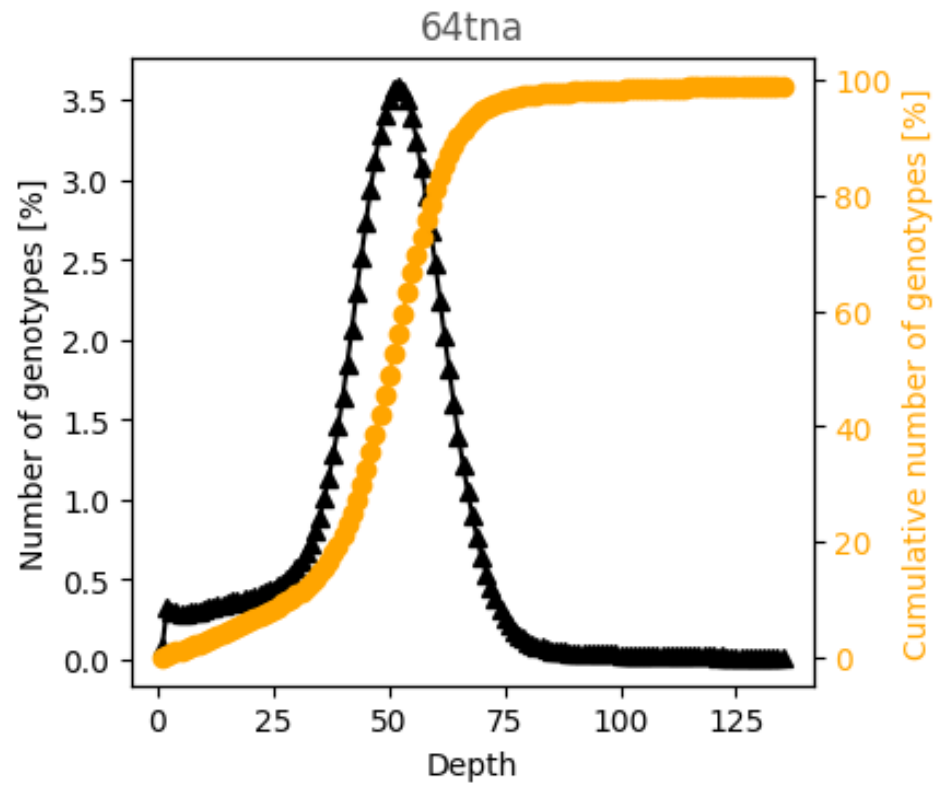
Indel distribution



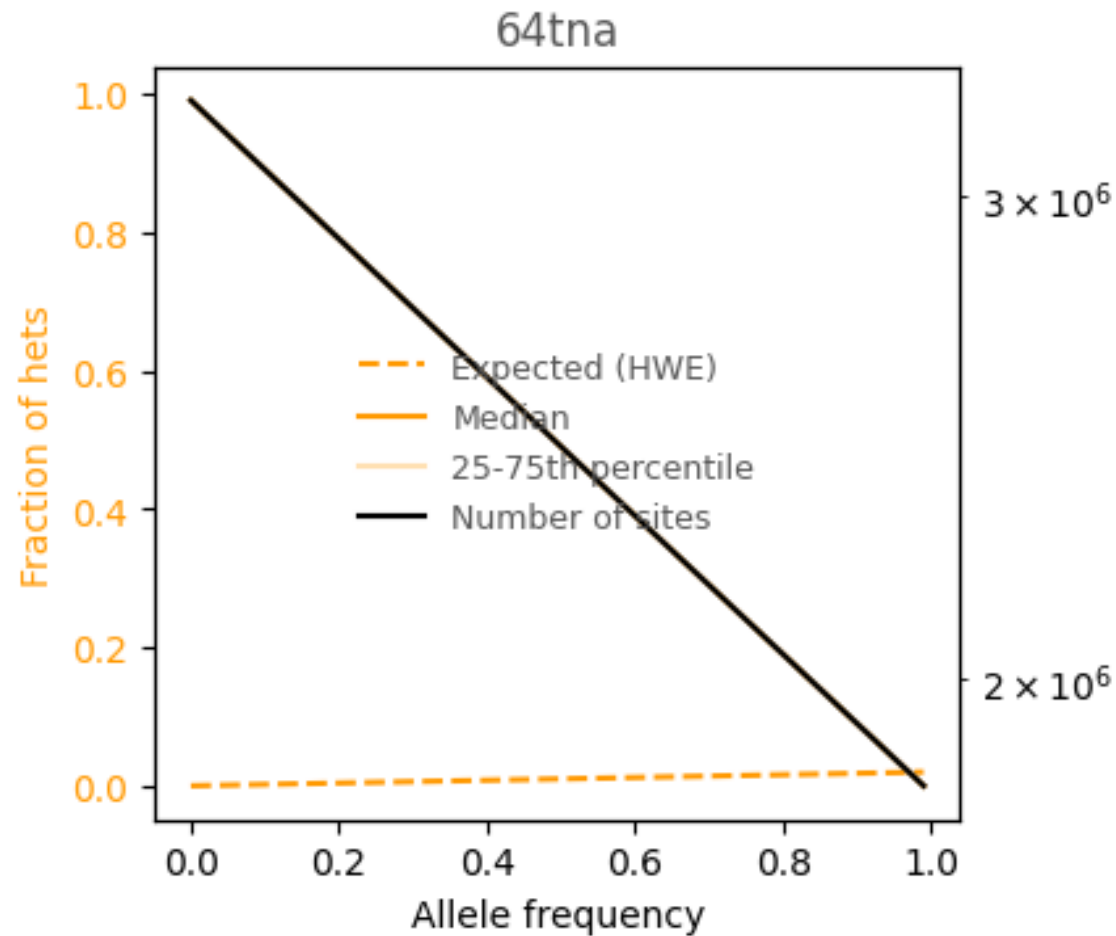
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

