

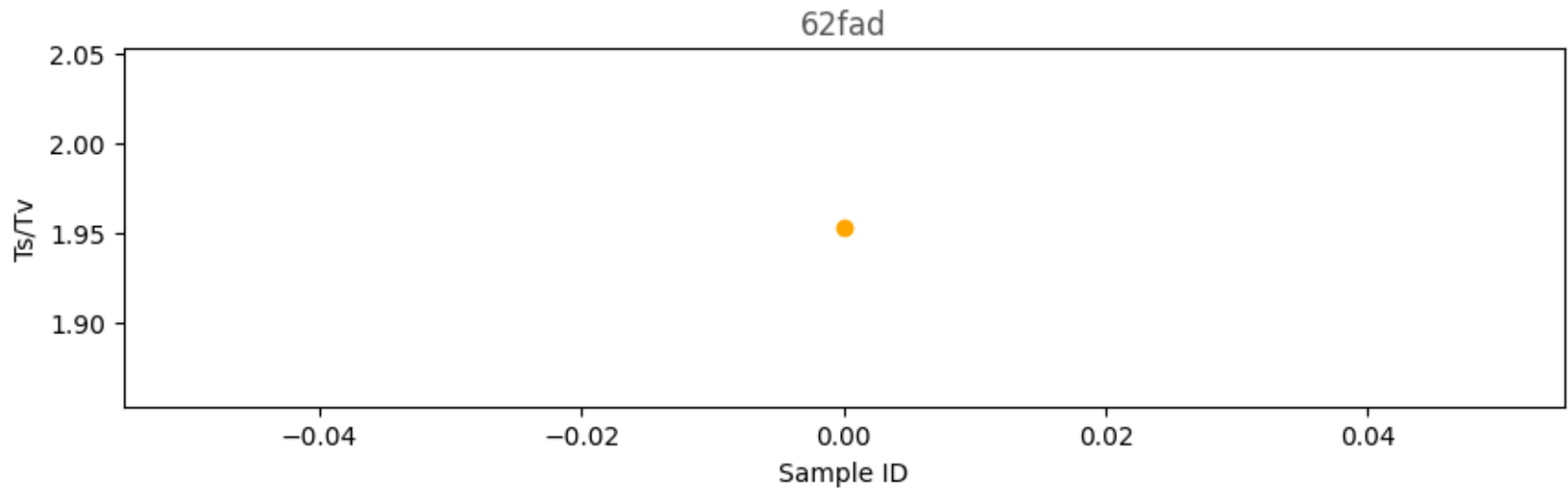
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
62fad	4,054,647	1.95	1.96	951,252	–	0	0
* frameshift ratio: out/(out+in)							

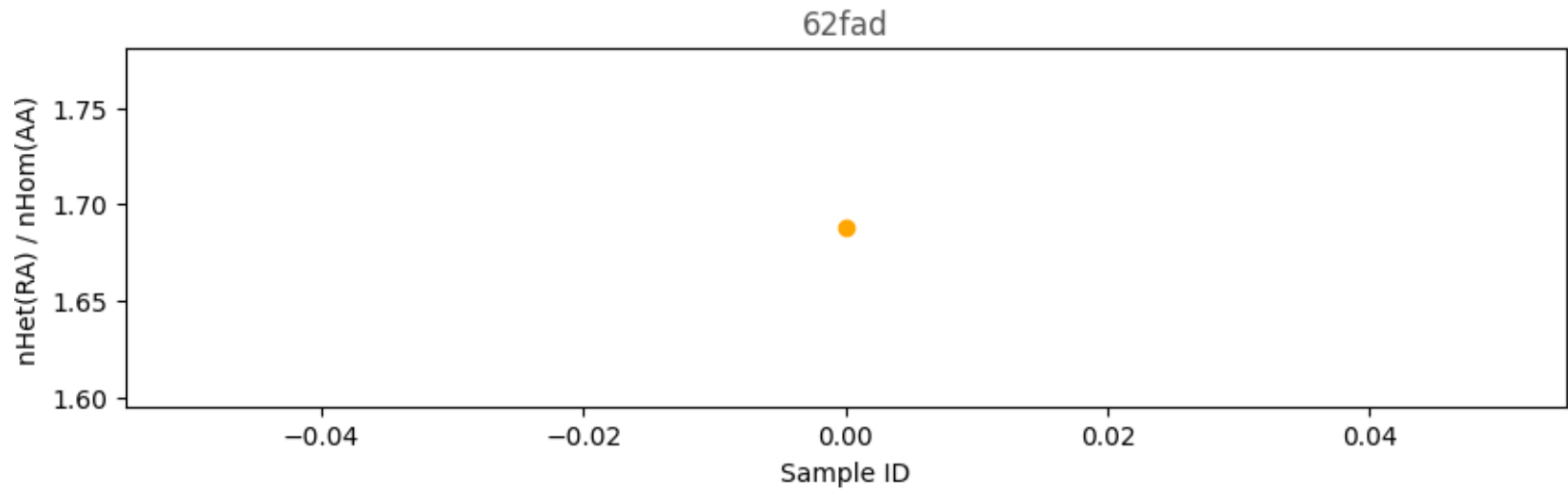
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
62fad	62.8%	1.92	67.8%	97,678	1,819

- 62fad .. /ngc/projects2/gm/data/archive/2022/variants/snv/62fadukef-103806443399-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211124_A01411_BHWHNFDSX2-RHGM_LABKA_WGS-
 WGS03374_snv_germline_raw.haplotype_caller.vcf.gz

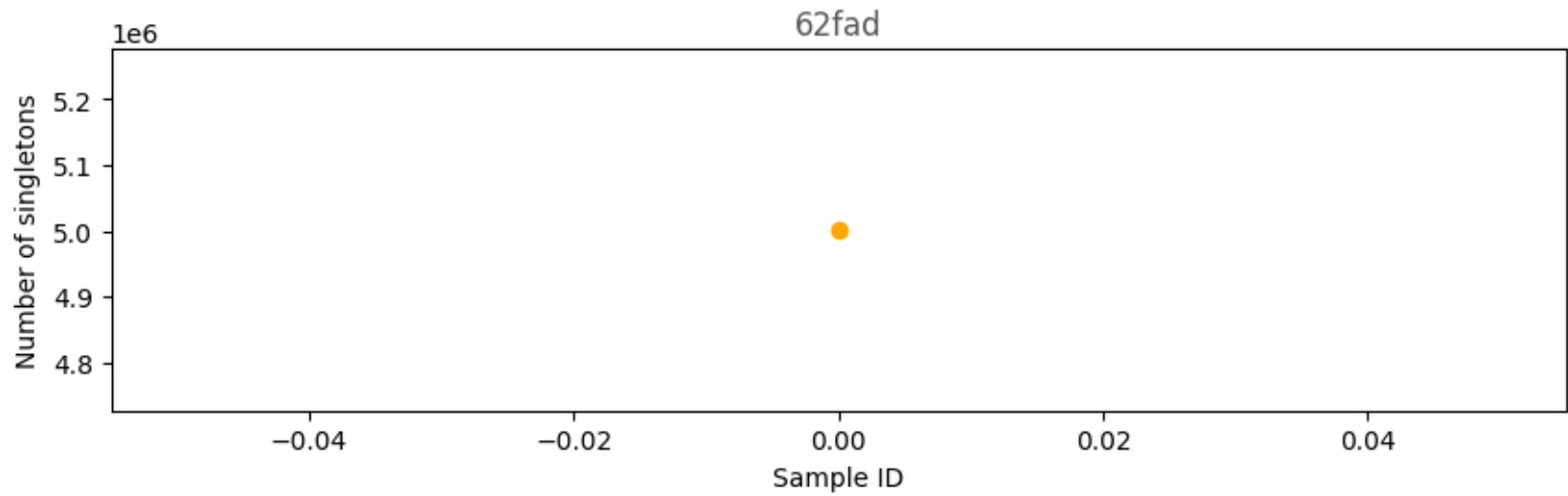
Ts/Tv by sample



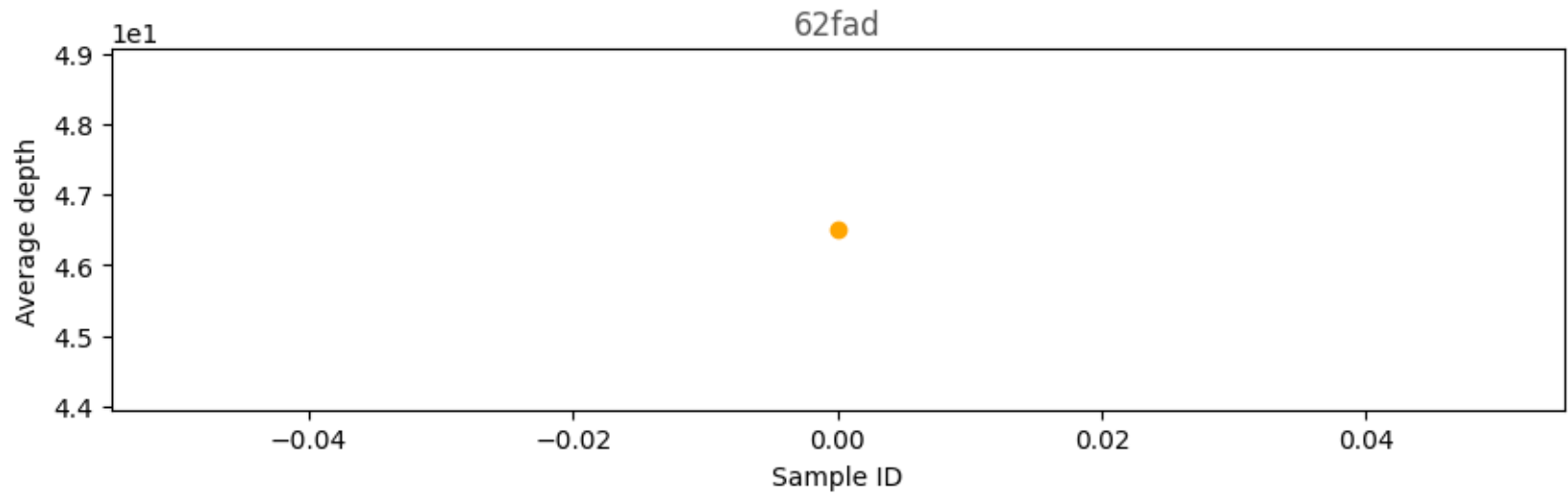
Hets vs non-ref Homs by sample



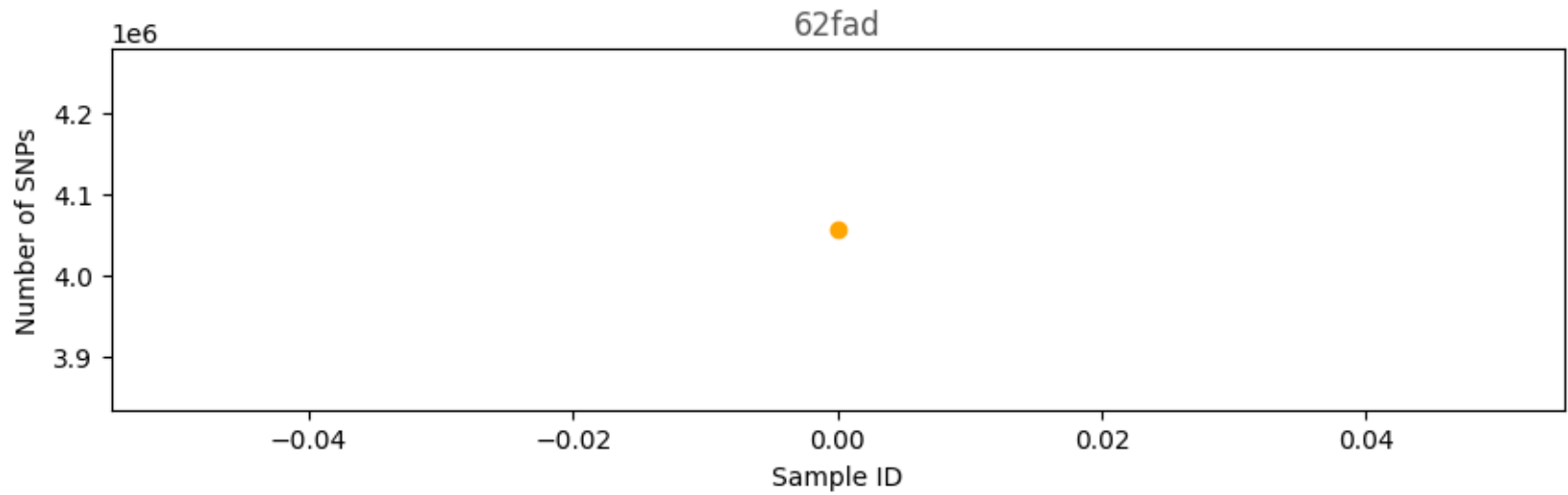
Singletons by sample (hets and homs)



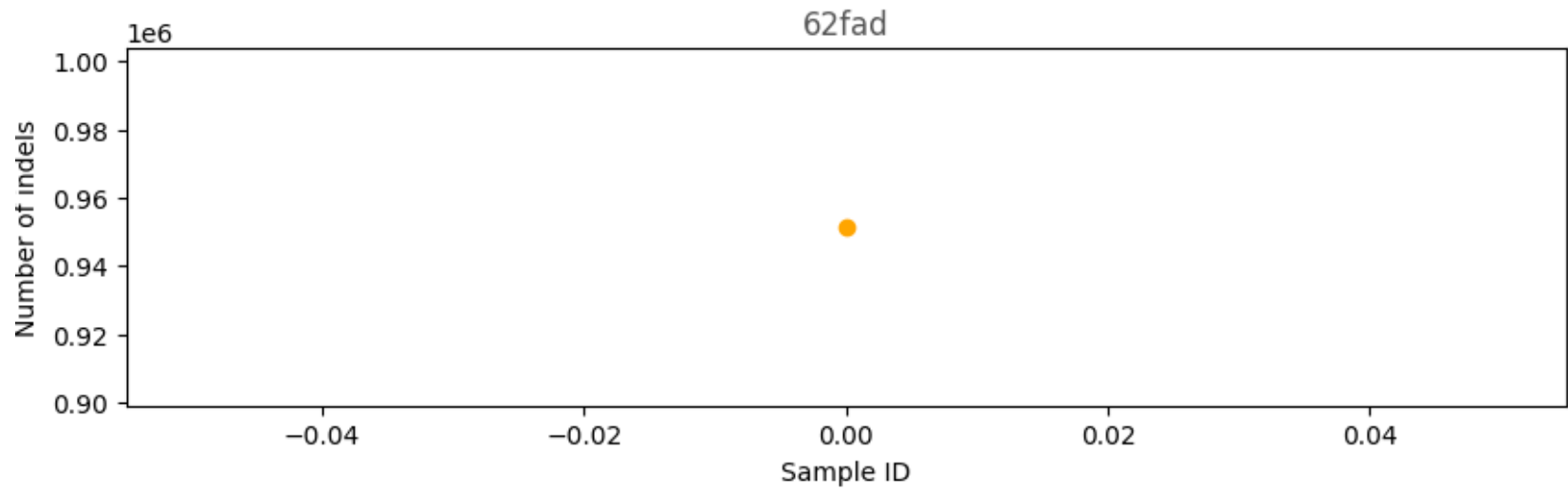
Average depth by sample



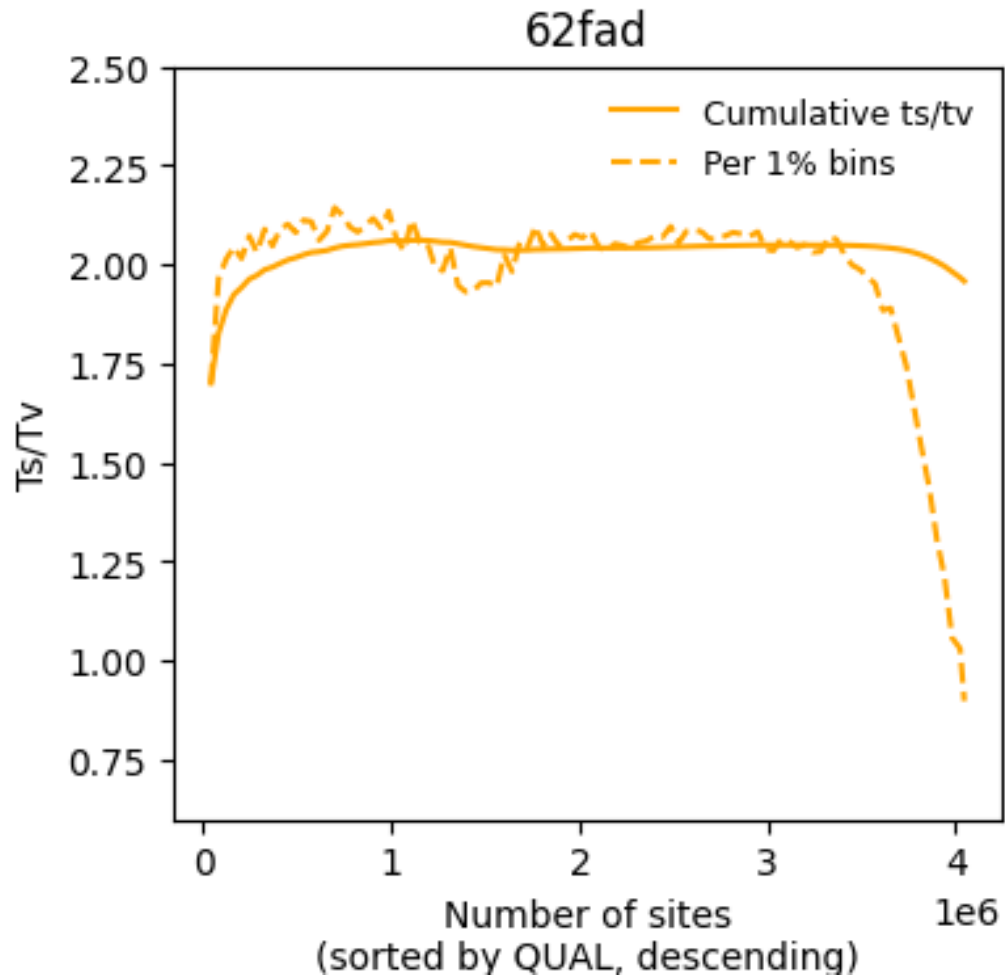
Number of SNPs by sample



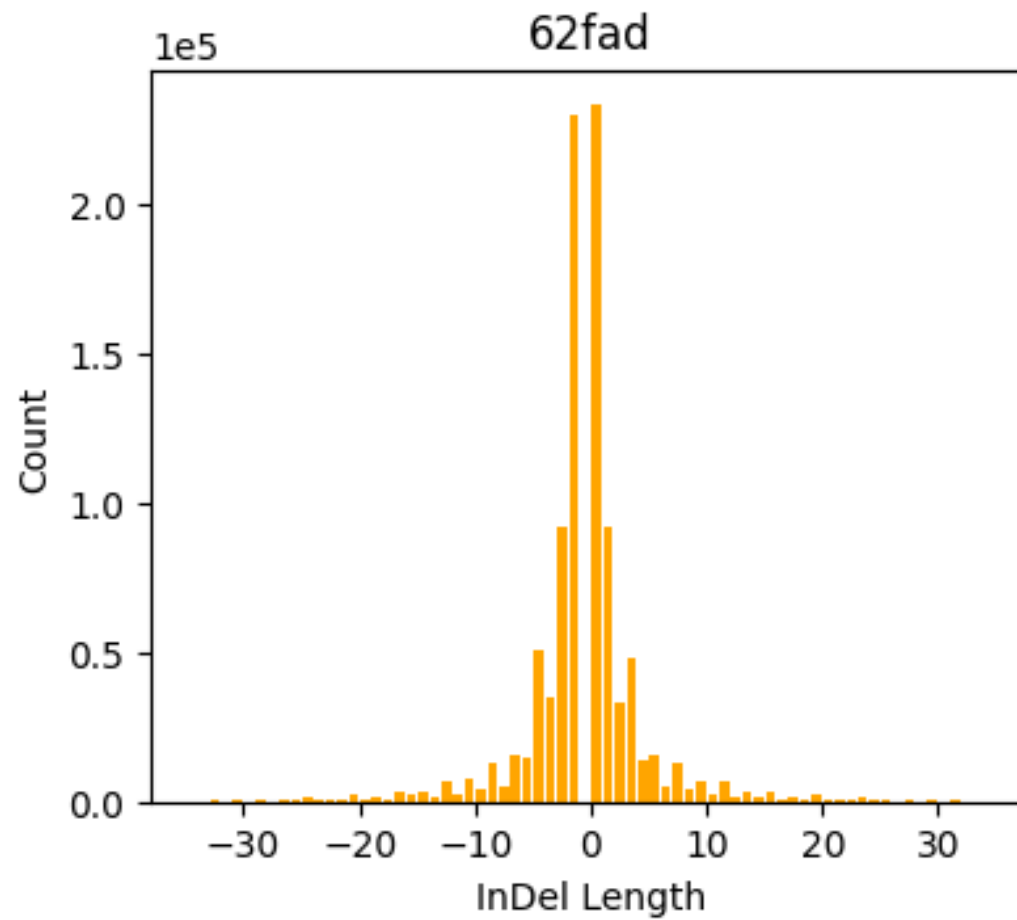
Number of indels by sample



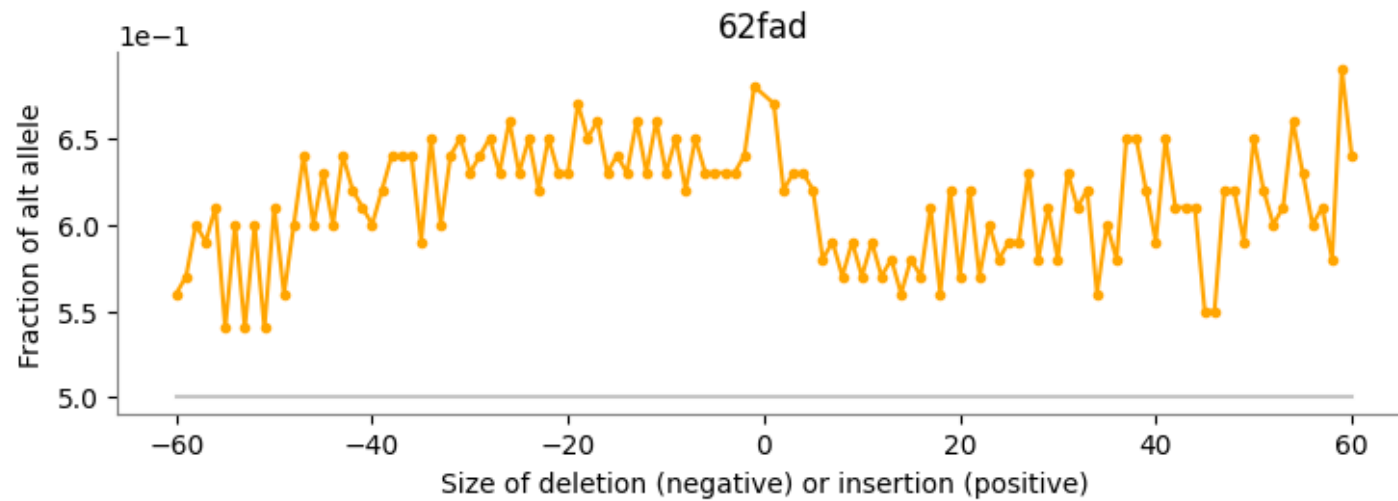
Ts/Tv stratified by QUAL



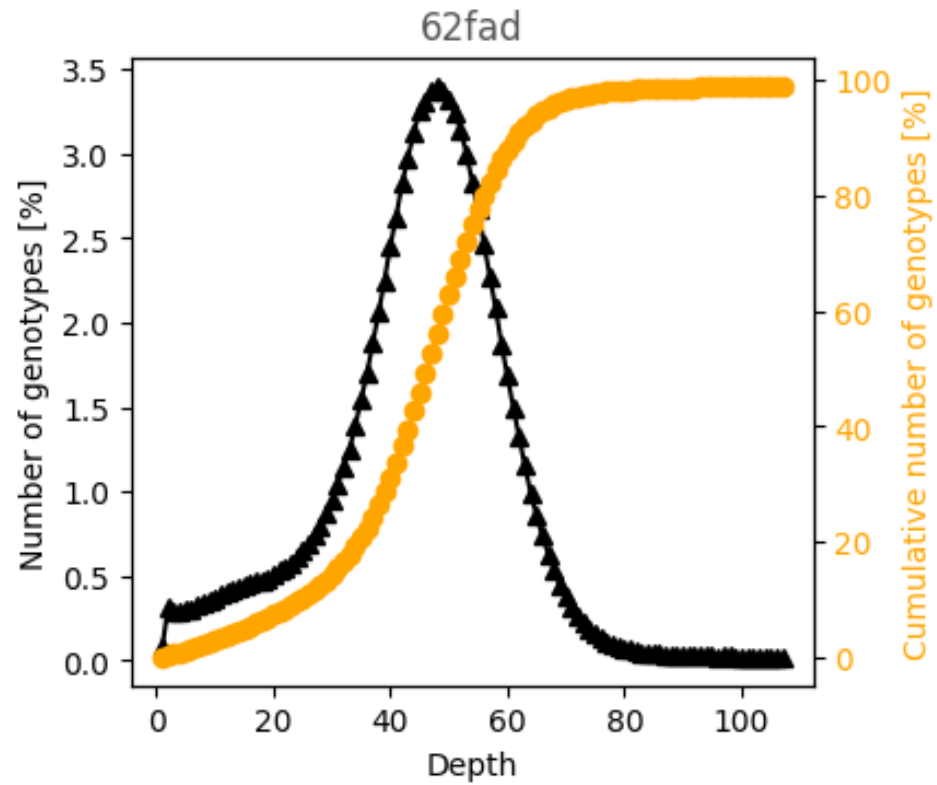
Indel distribution



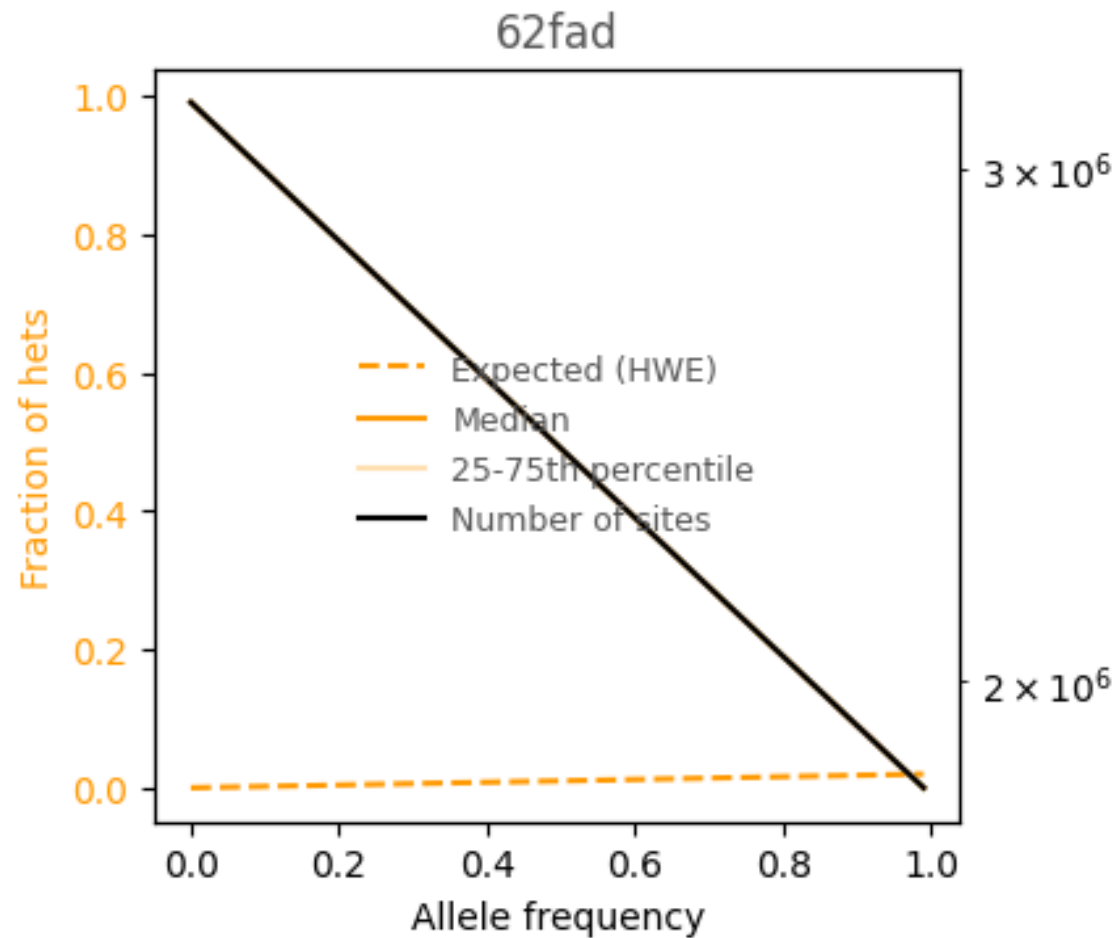
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

