

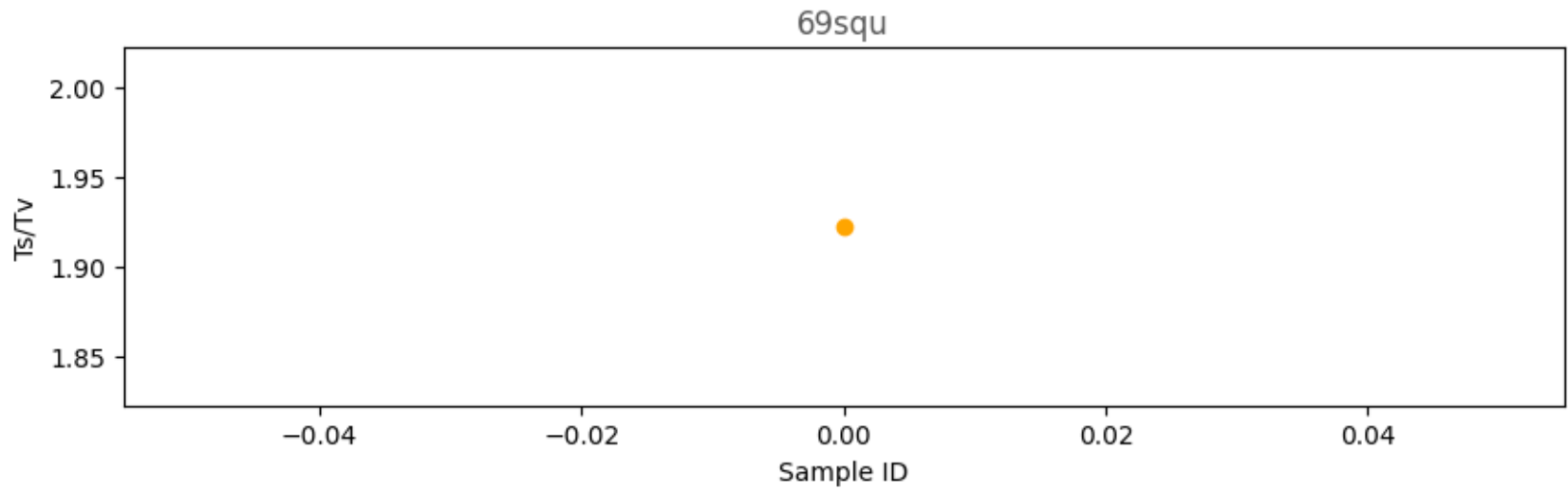
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
69squ	4,072,840	1.92	1.93	950,095	–	0	0
* frameshift ratio: out/(out+in)							

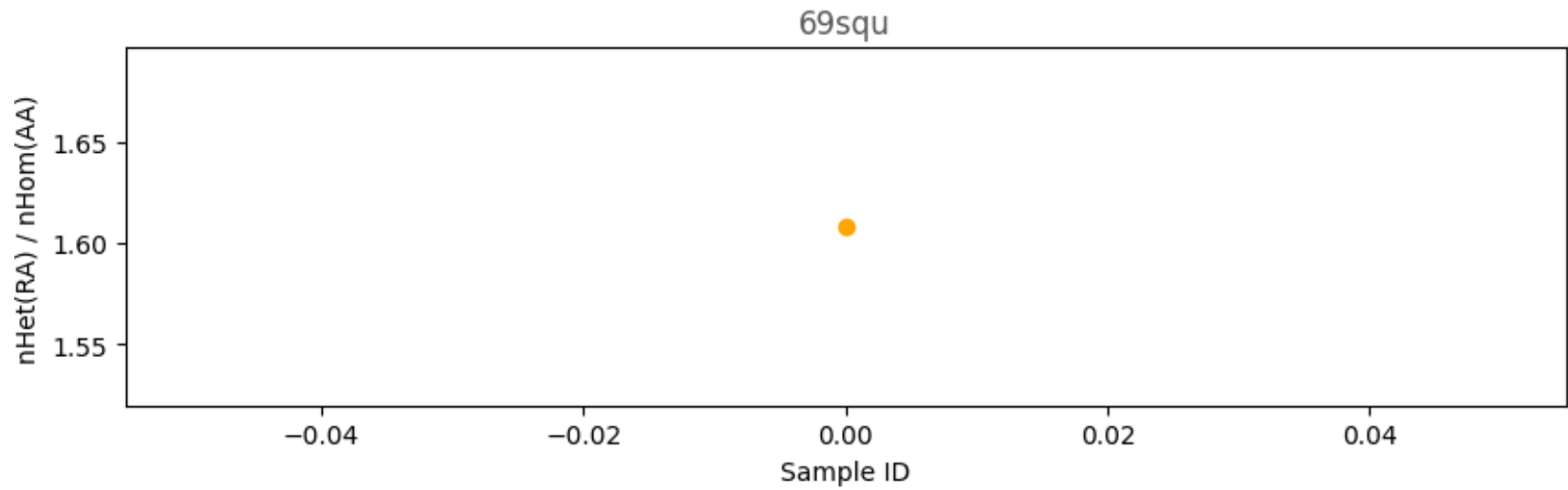
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
69squ	61.7%	1.88	66.8%	102,212	2,014

- 69squ .. /ngc/projects2/gm/data/archive/2022/variants/snv/69squptom-103876114548-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00838-220601_A00559_AHJ5TYDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS04376_snv_germline_raw.haplotype_caller.vcf.gz

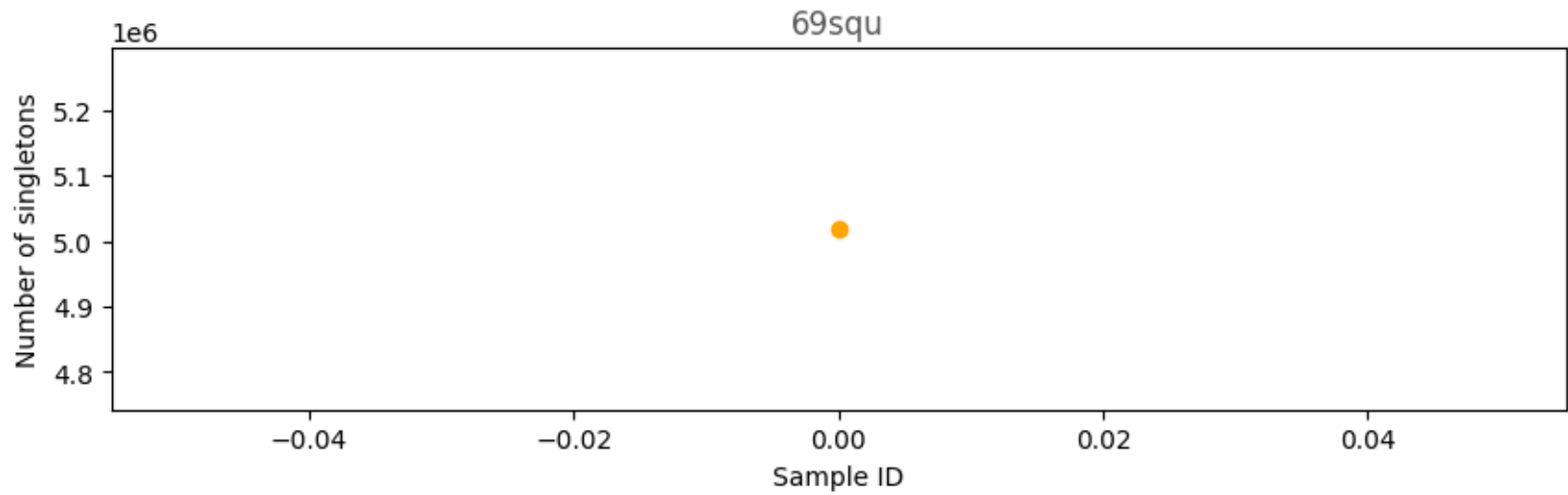
Ts/Tv by sample



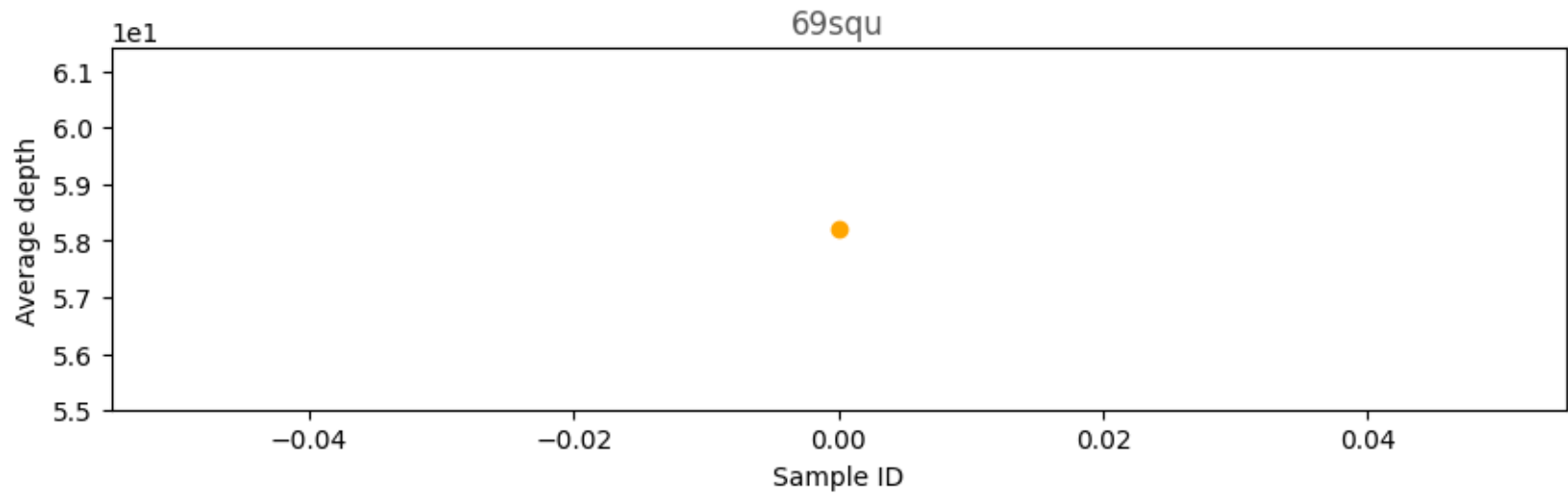
Hets vs non-ref Homs by sample



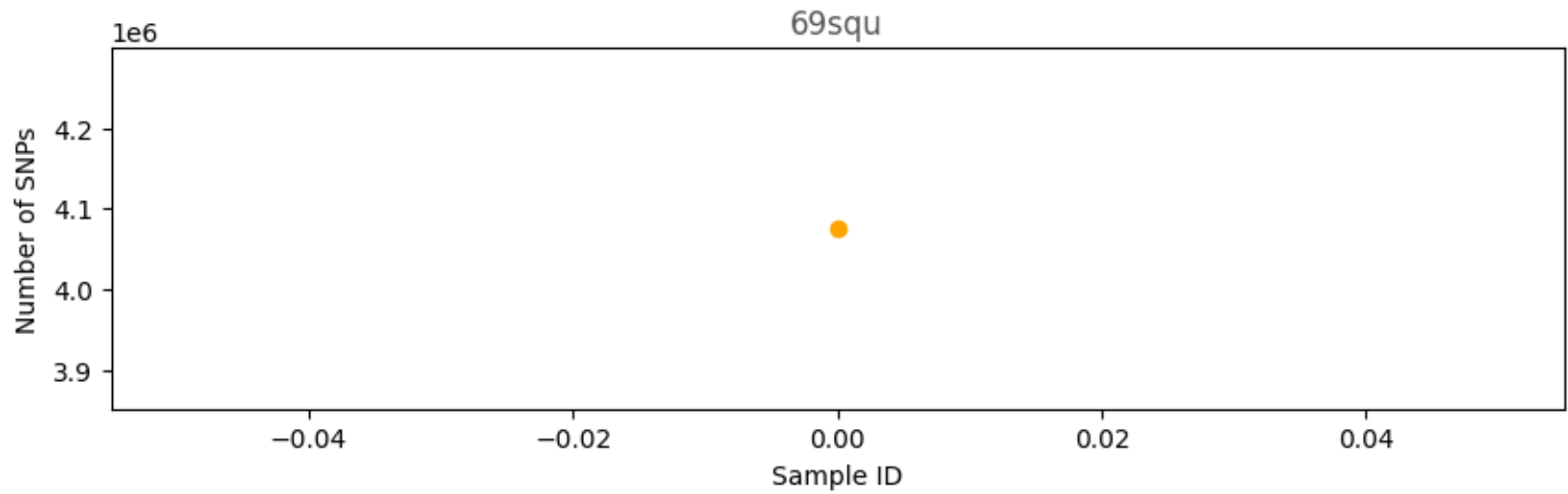
Singletons by sample (hets and homs)



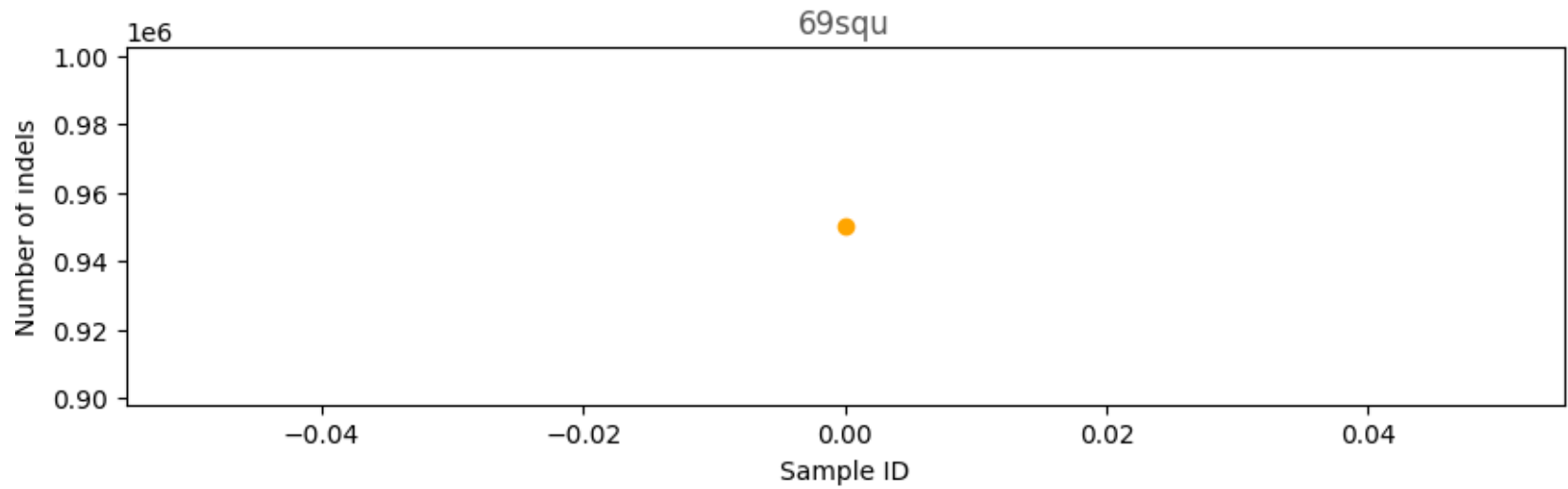
Average depth by sample



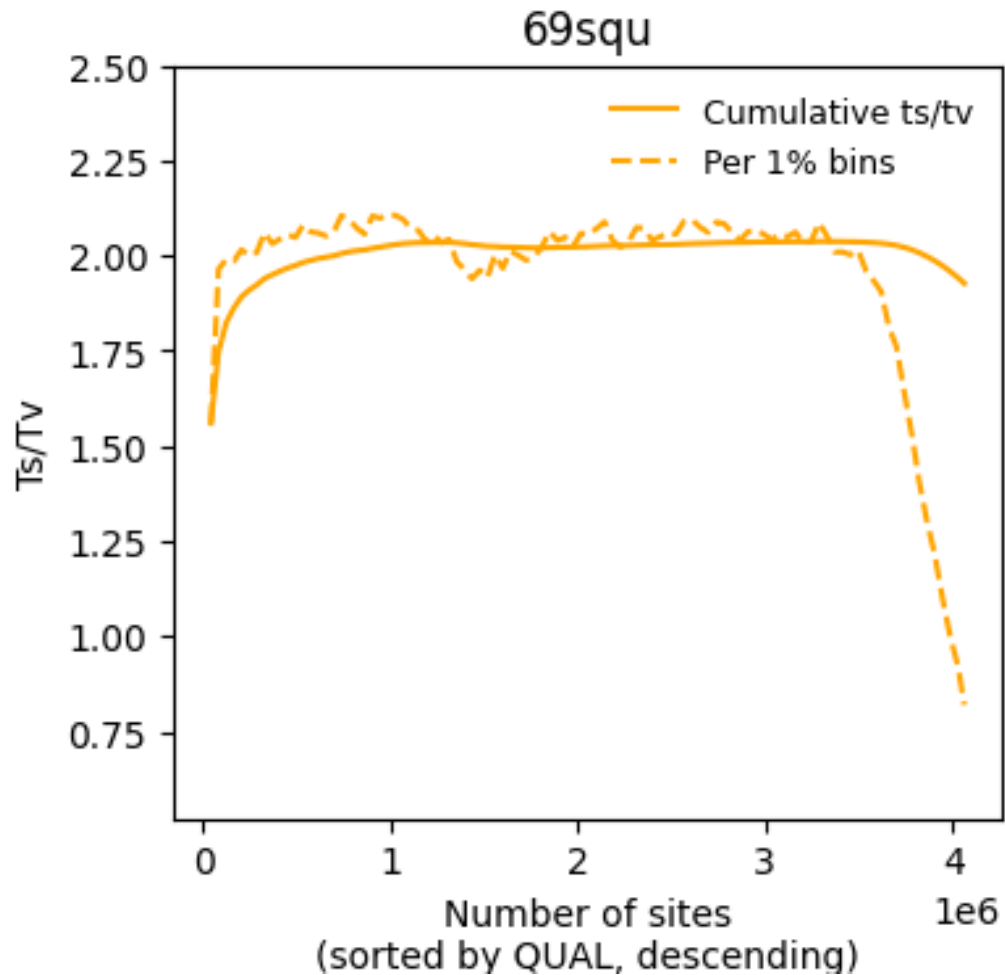
Number of SNPs by sample



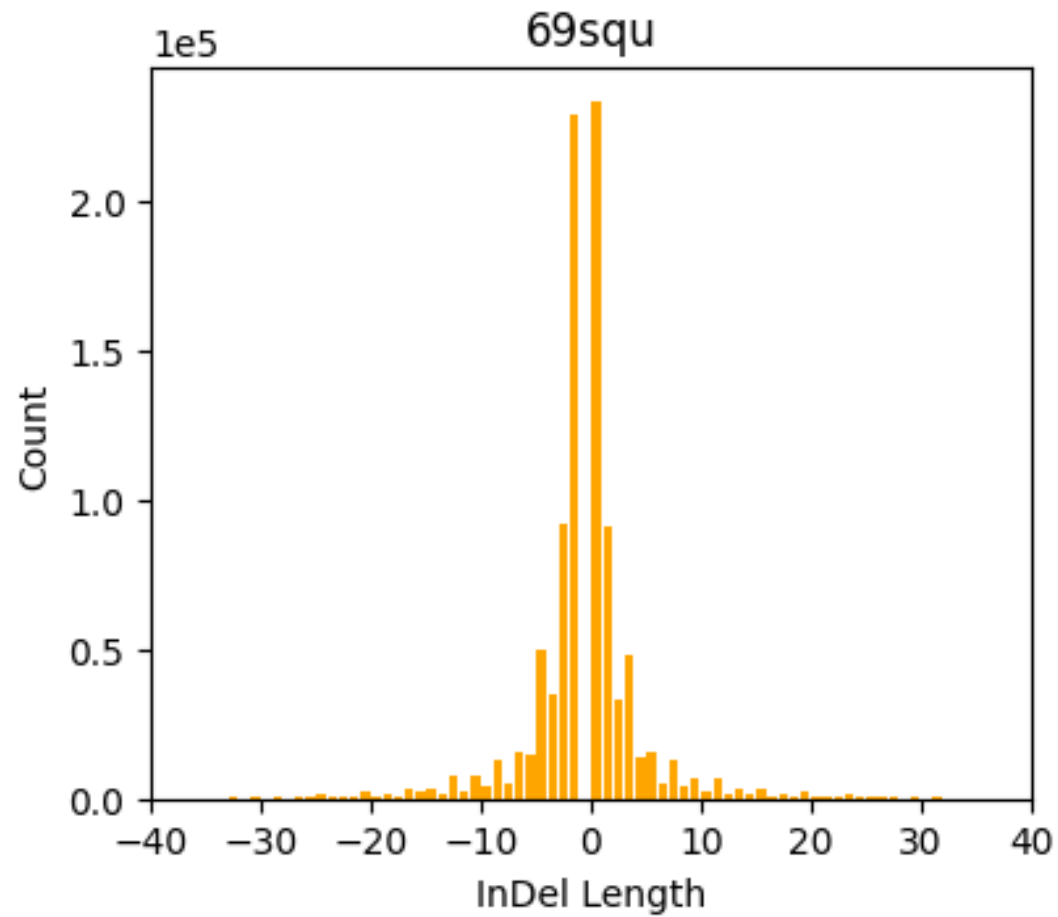
Number of indels by sample



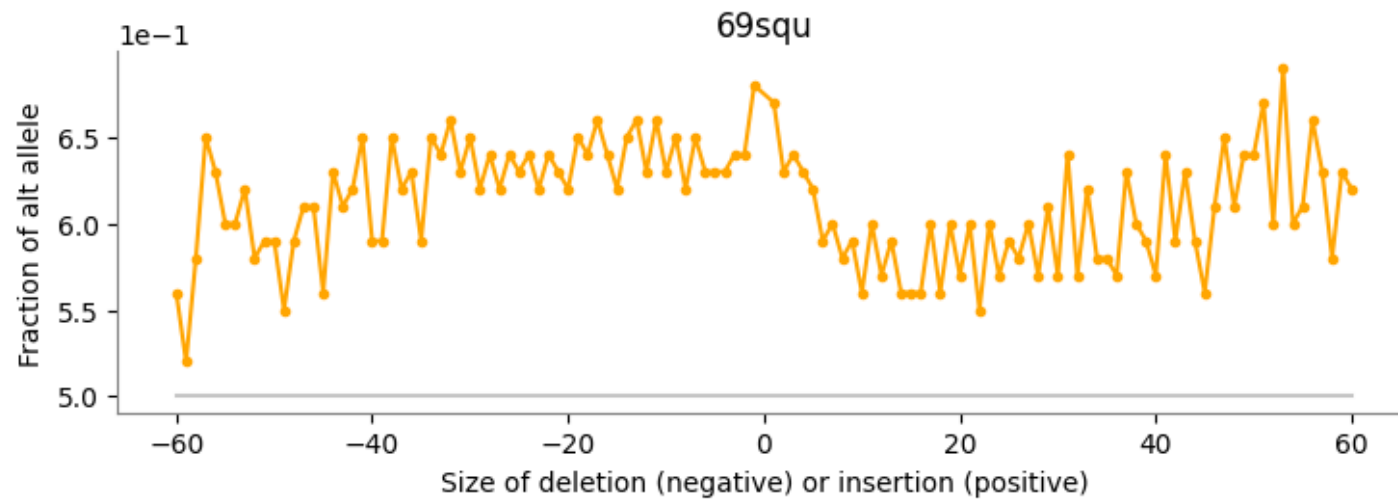
Ts/Tv stratified by QUAL



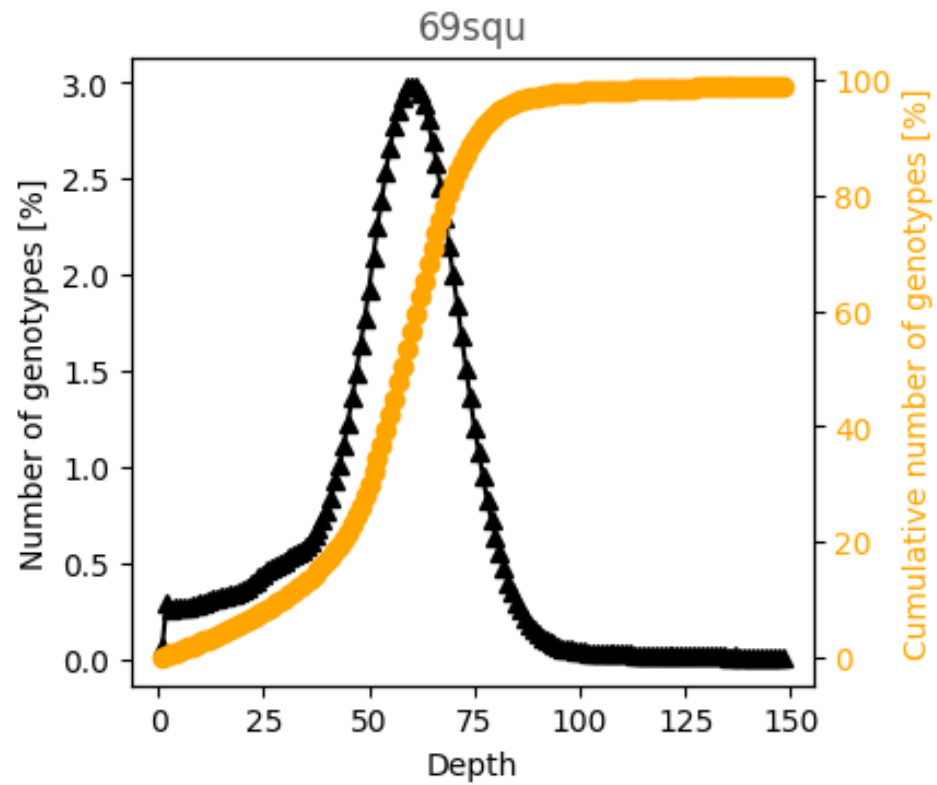
Indel distribution



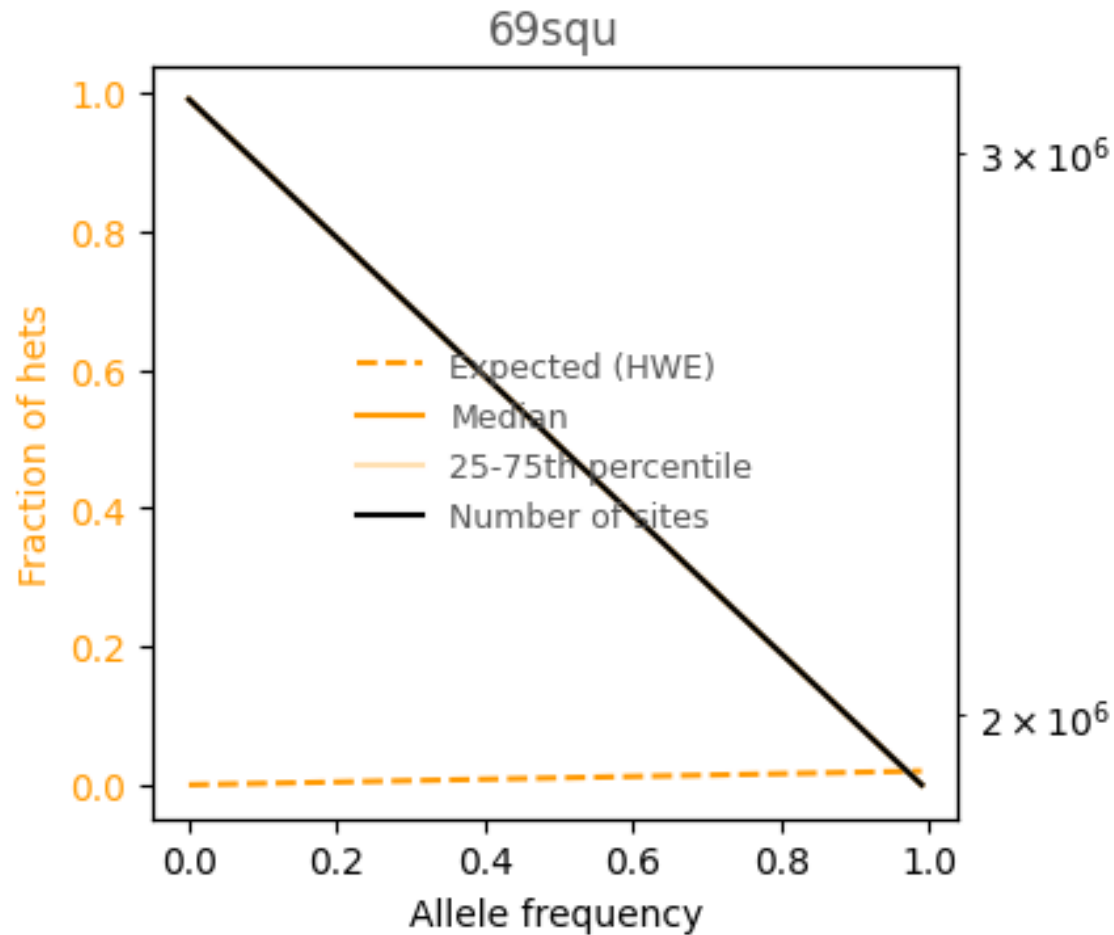
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

