

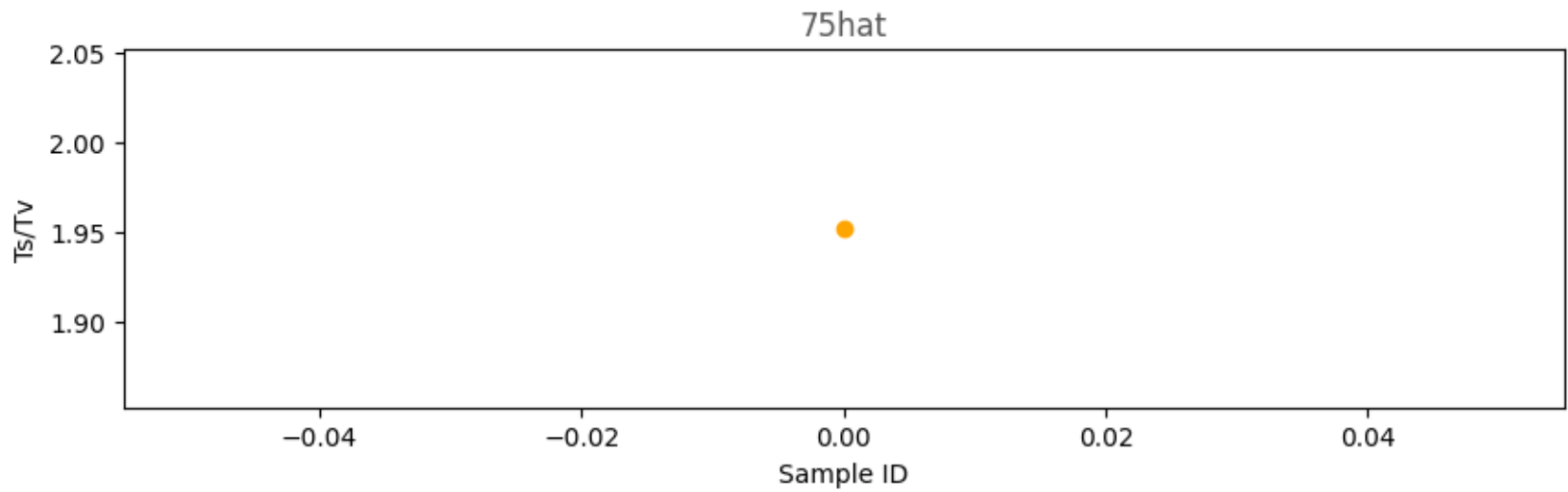
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
75hat	4,050,225	1.95	1.96	961,780	–	0	0
* frameshift ratio: out/(out+in)							

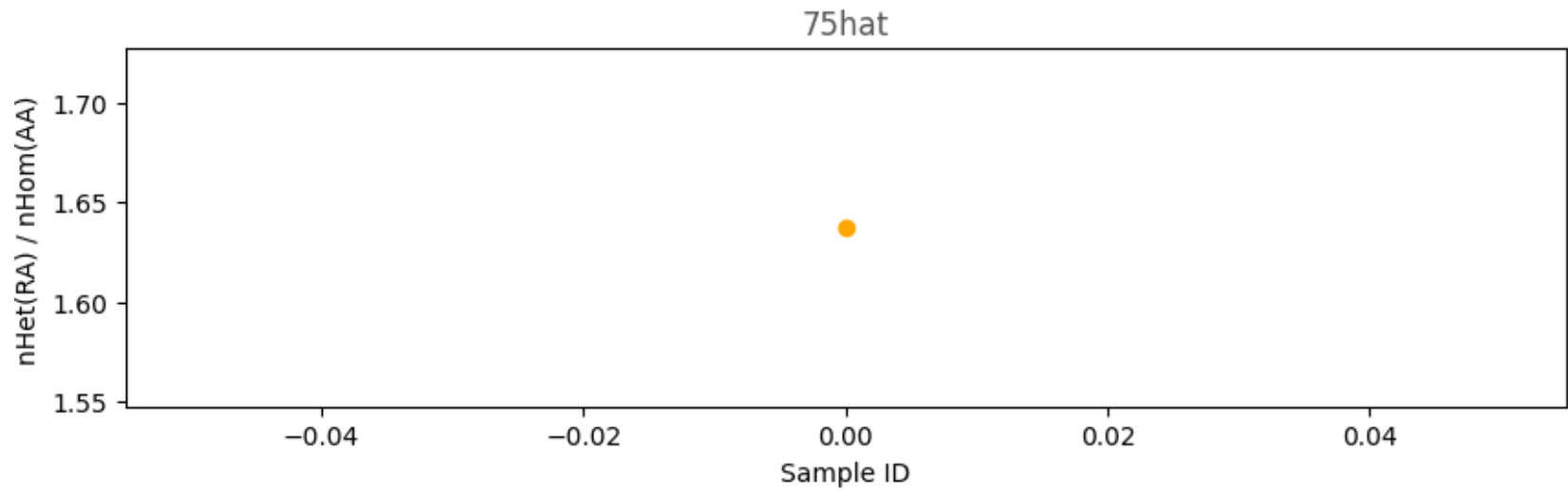
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
75hat	62.1%	1.93	68.1%	105,879	2,078

- 75hat .. /ngc/projects2/gm/data/archive/2022/variants/snv/75hatupaf-103904416805-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01185-220831_A01961_AHTNYGDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS04941_22RKG012090x01_snv_germline_raw.haplotype_caller.vcf.gz

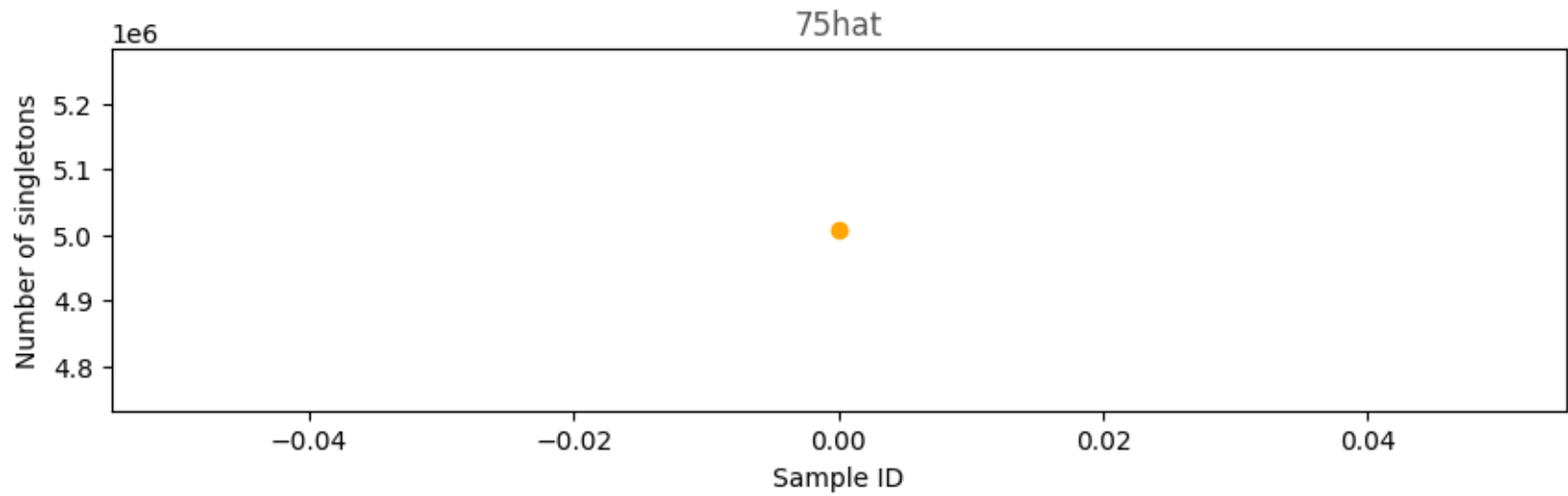
Ts/Tv by sample



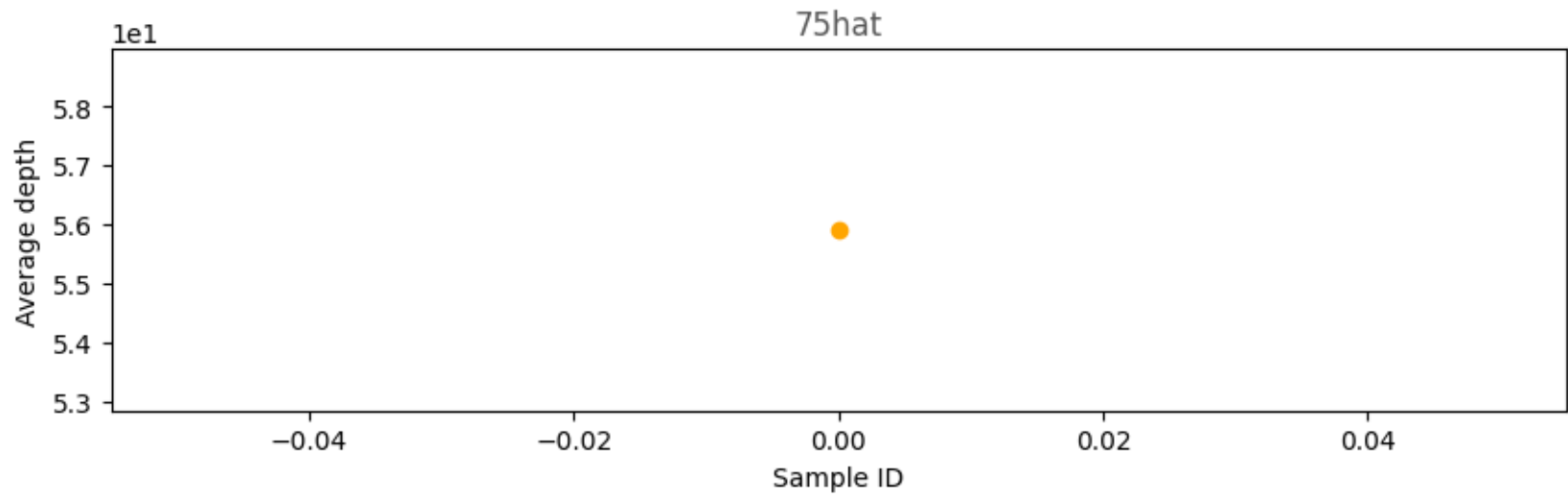
Hets vs non-ref Homs by sample



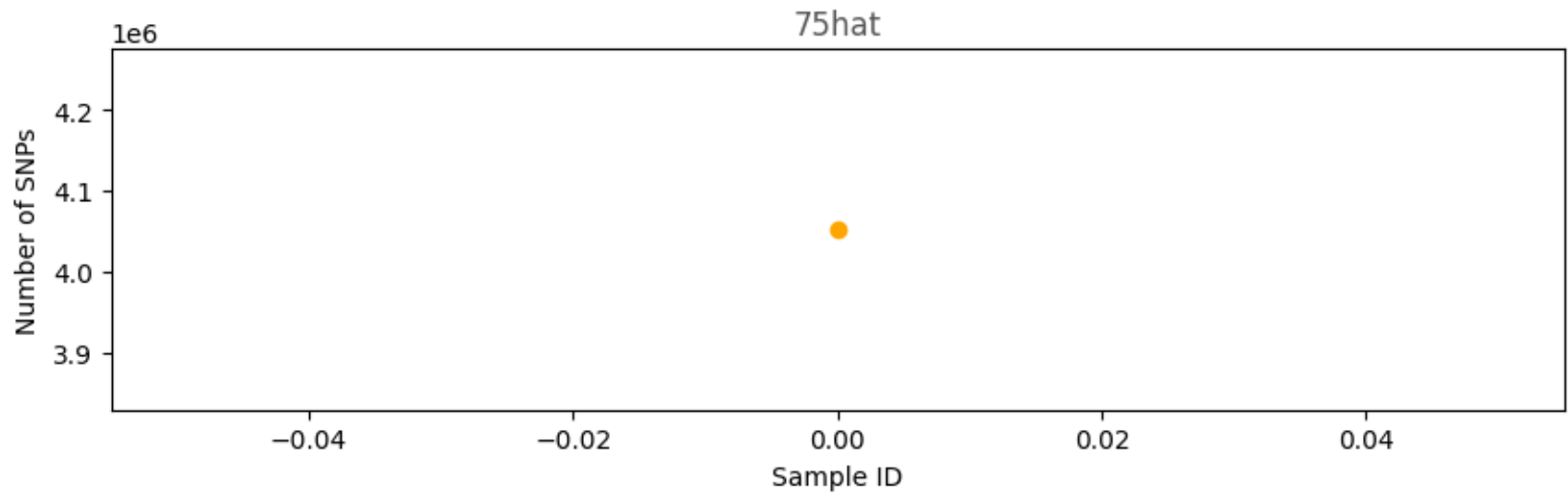
Singletons by sample (hets and homs)



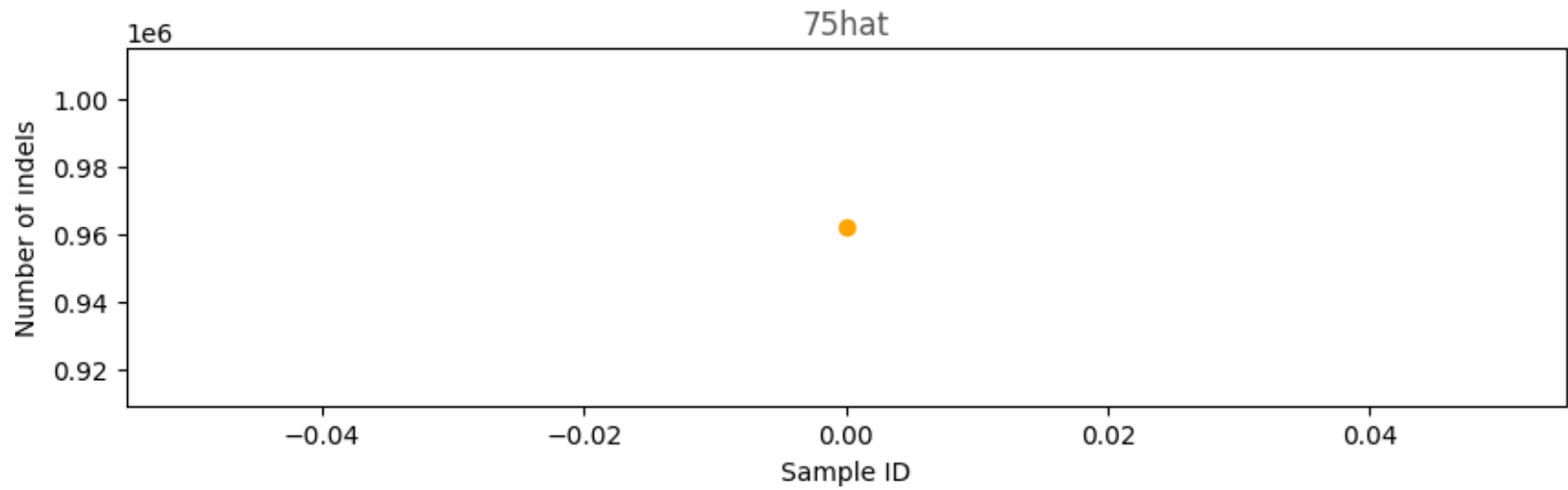
Average depth by sample



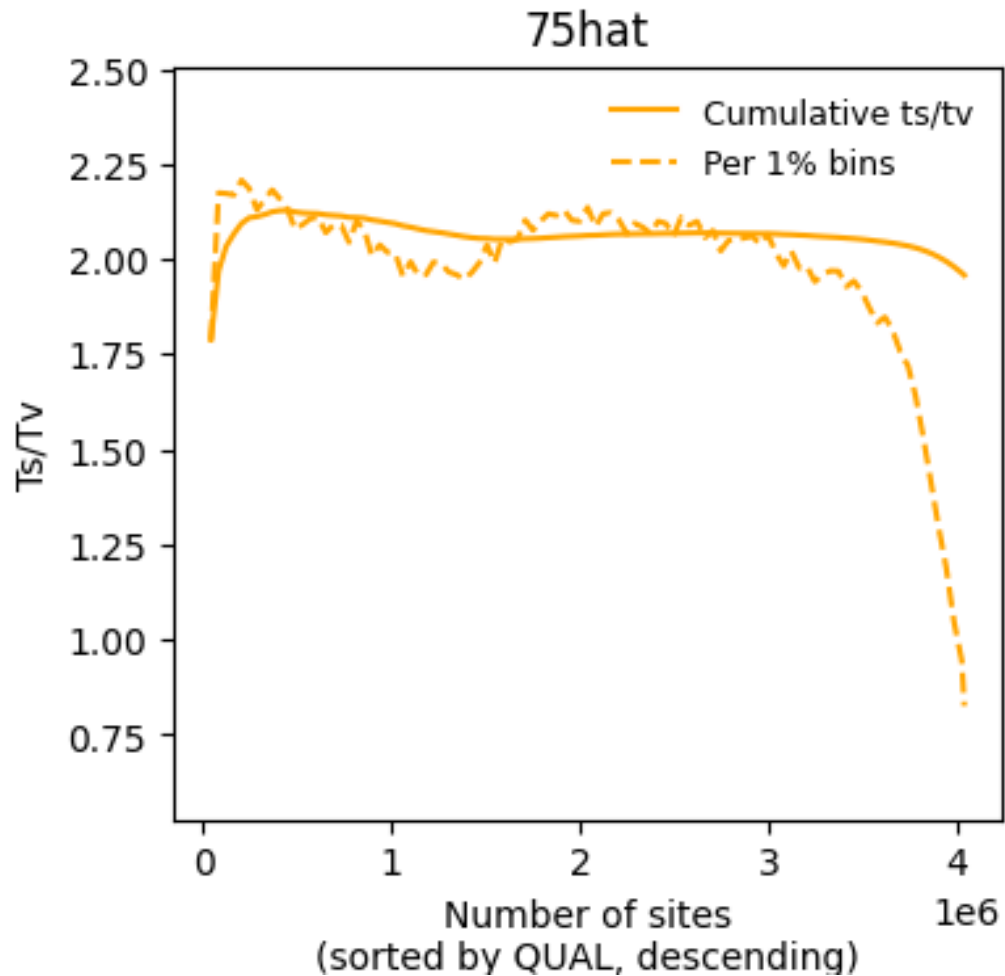
Number of SNPs by sample



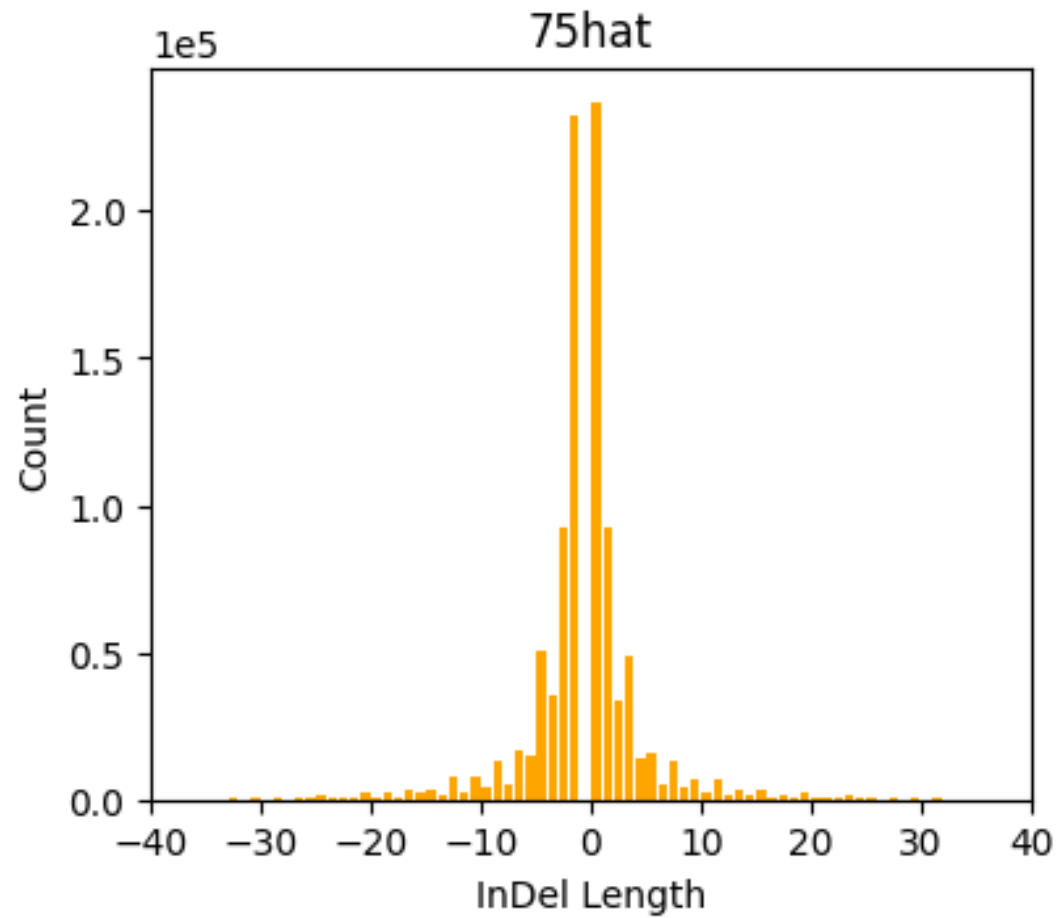
Number of indels by sample



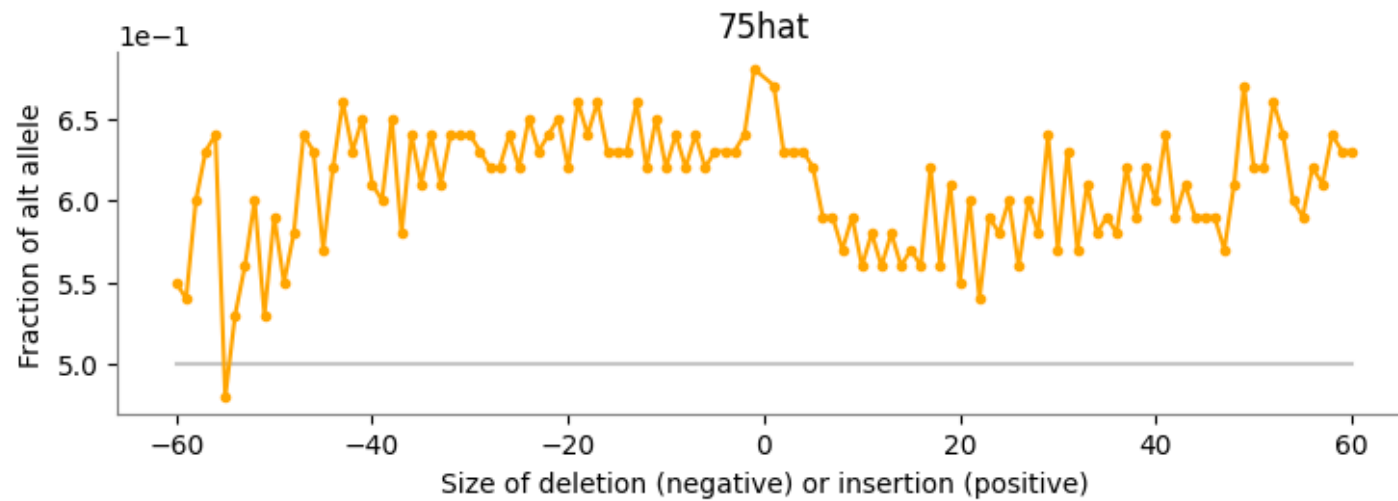
Ts/Tv stratified by QUAL



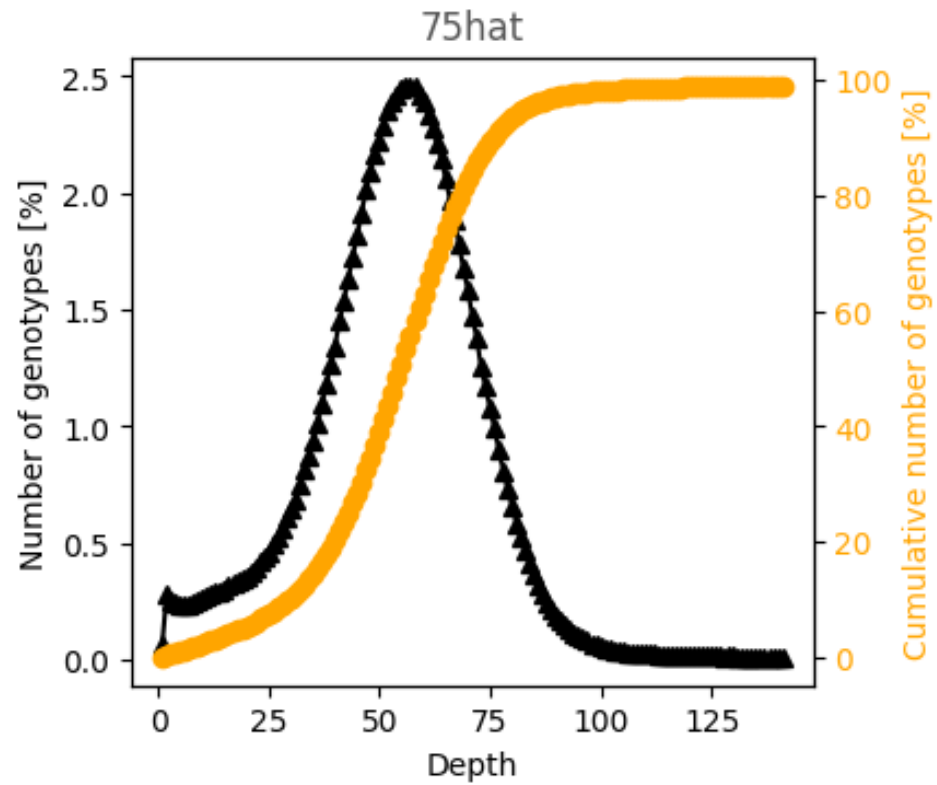
Indel distribution



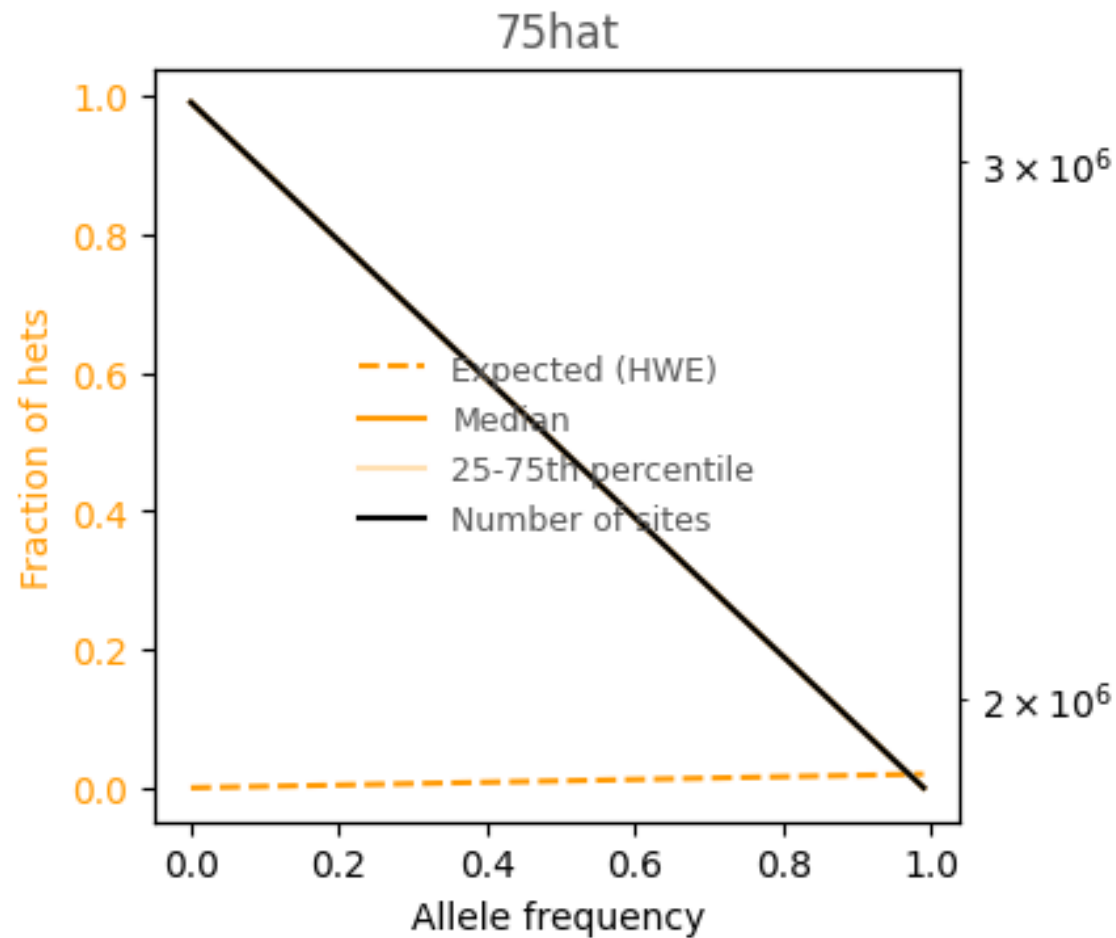
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

