

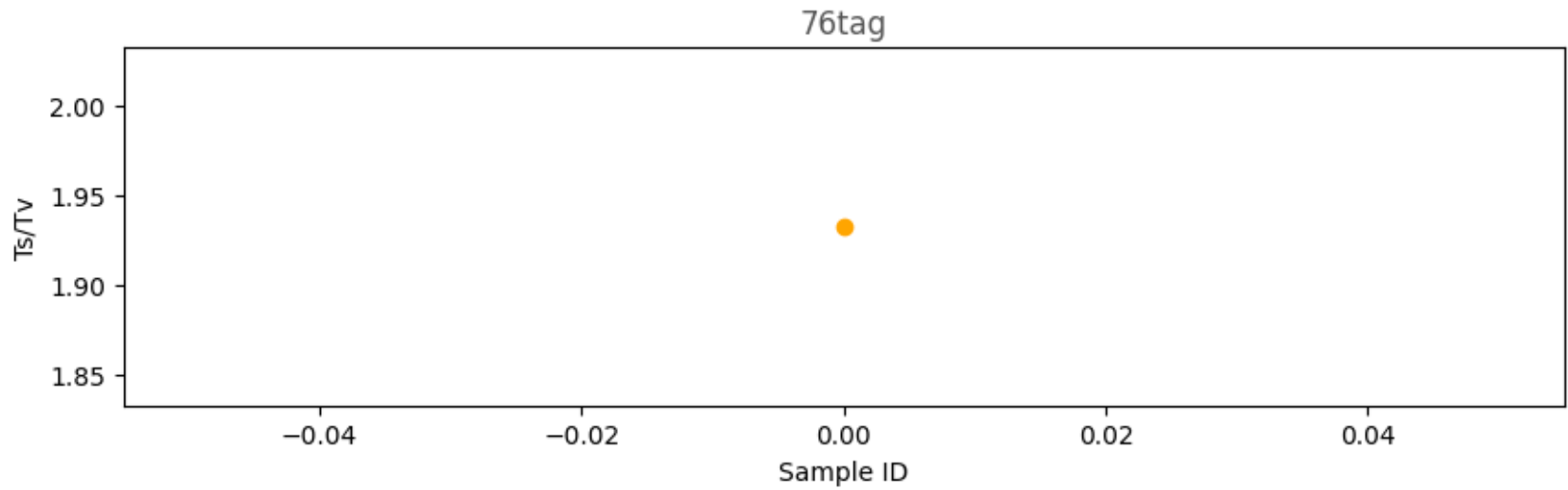
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76tag	4,113,392	1.93	1.94	955,817	–	0	0
* frameshift ratio: out/(out+in)							

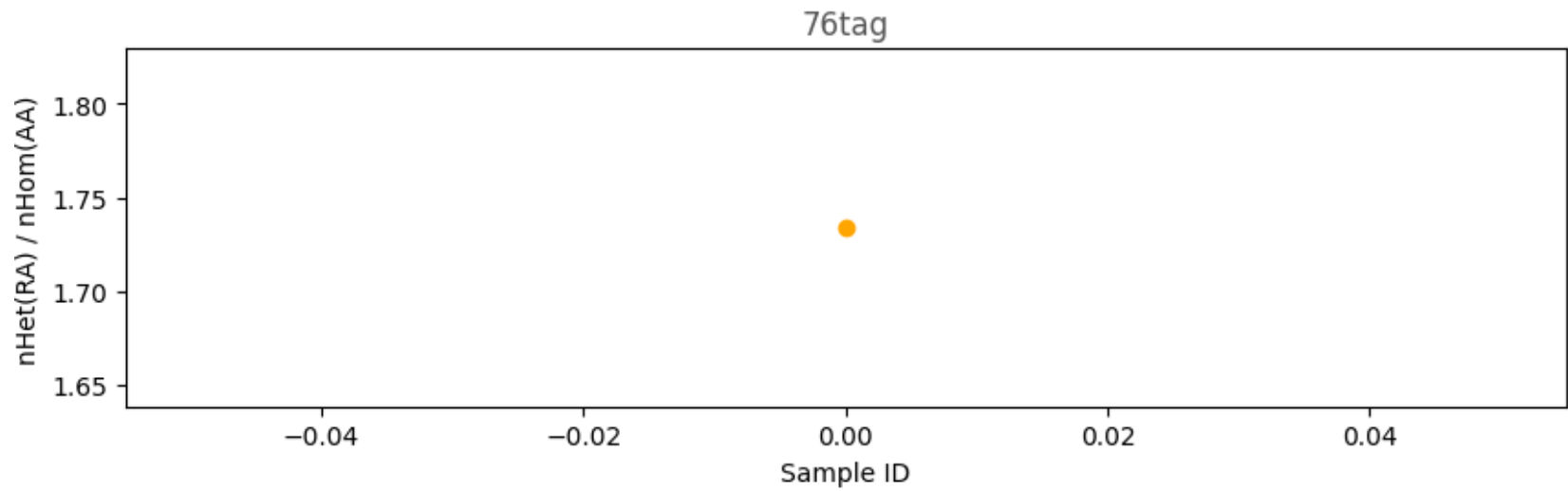
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76tag	63.4%	1.91	68.3%	100,529	2,154

- 76tag .. /ngc/projects2/gm/data/archive/2022/variants/snv/76tagstaf-103909079459-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01499-220914_A01176_BHTNVLDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS05273_22RKG021102_snv_germline_raw.haplotype_caller.vcf.gz

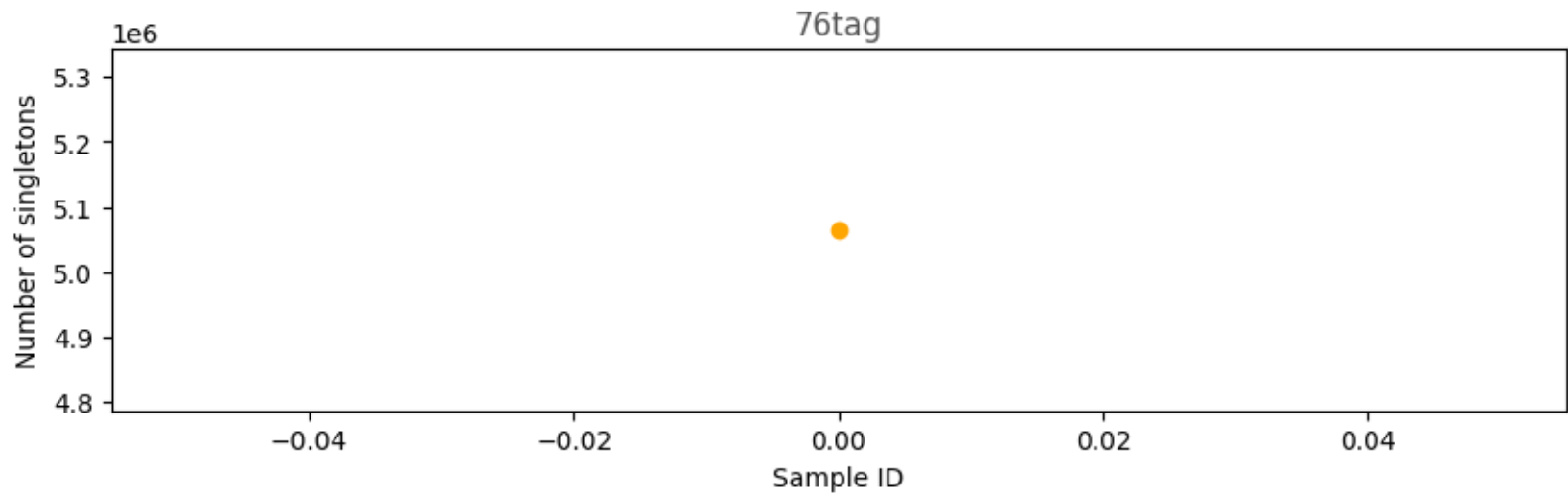
Ts/Tv by sample



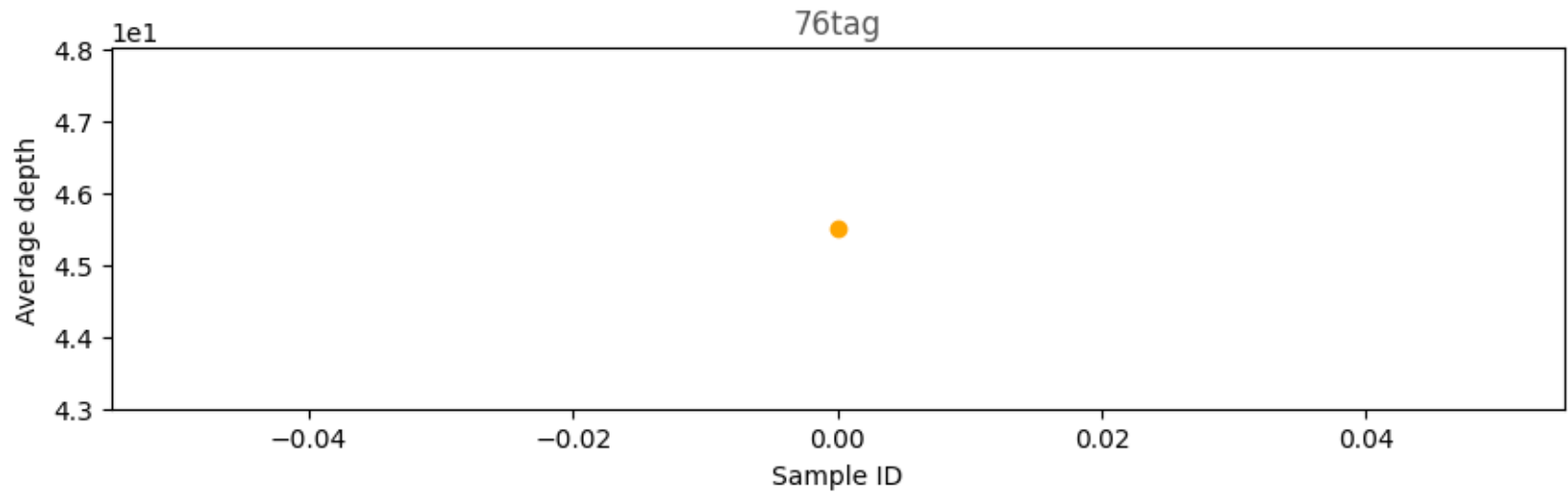
Hets vs non-ref Homs by sample



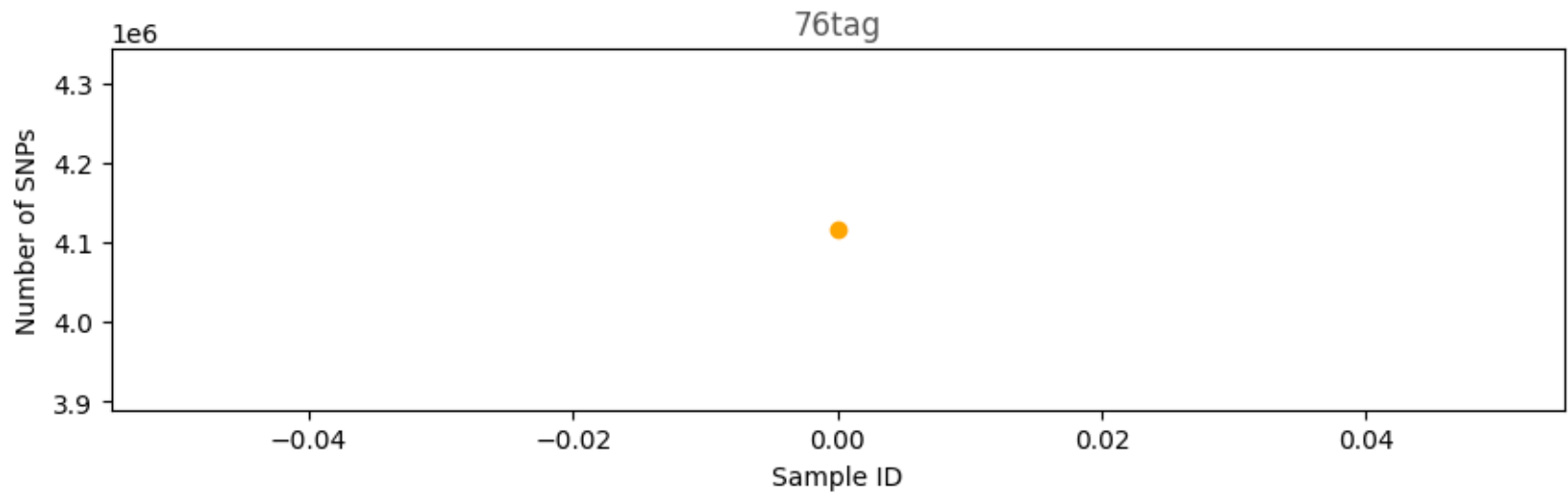
Singletons by sample (hets and homs)



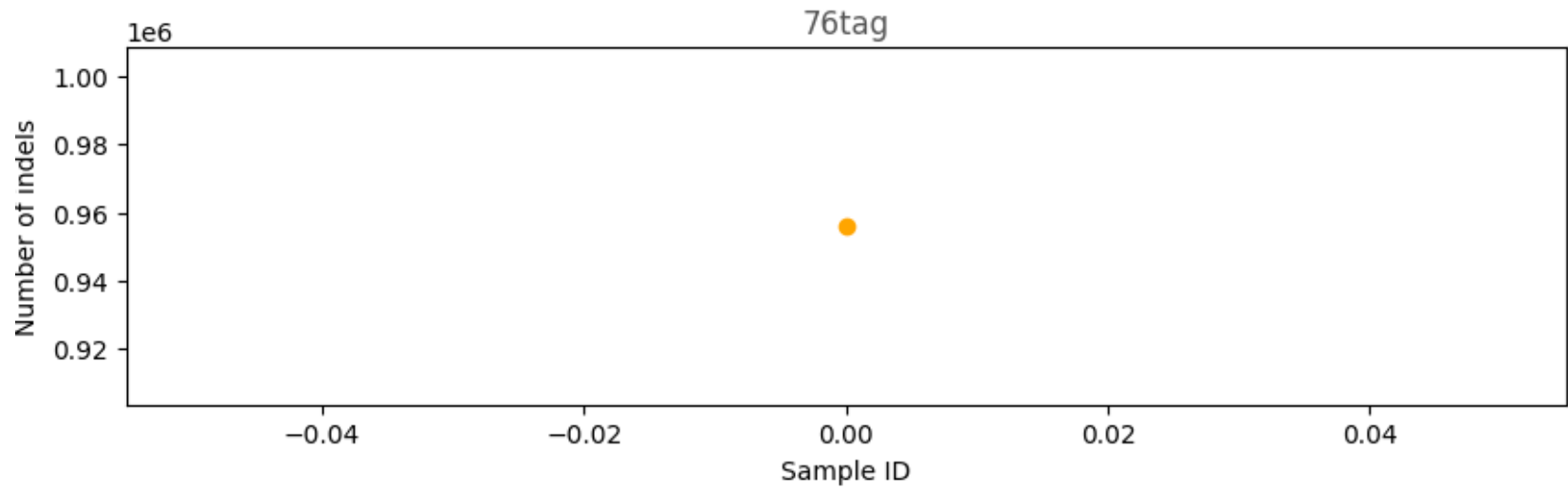
Average depth by sample



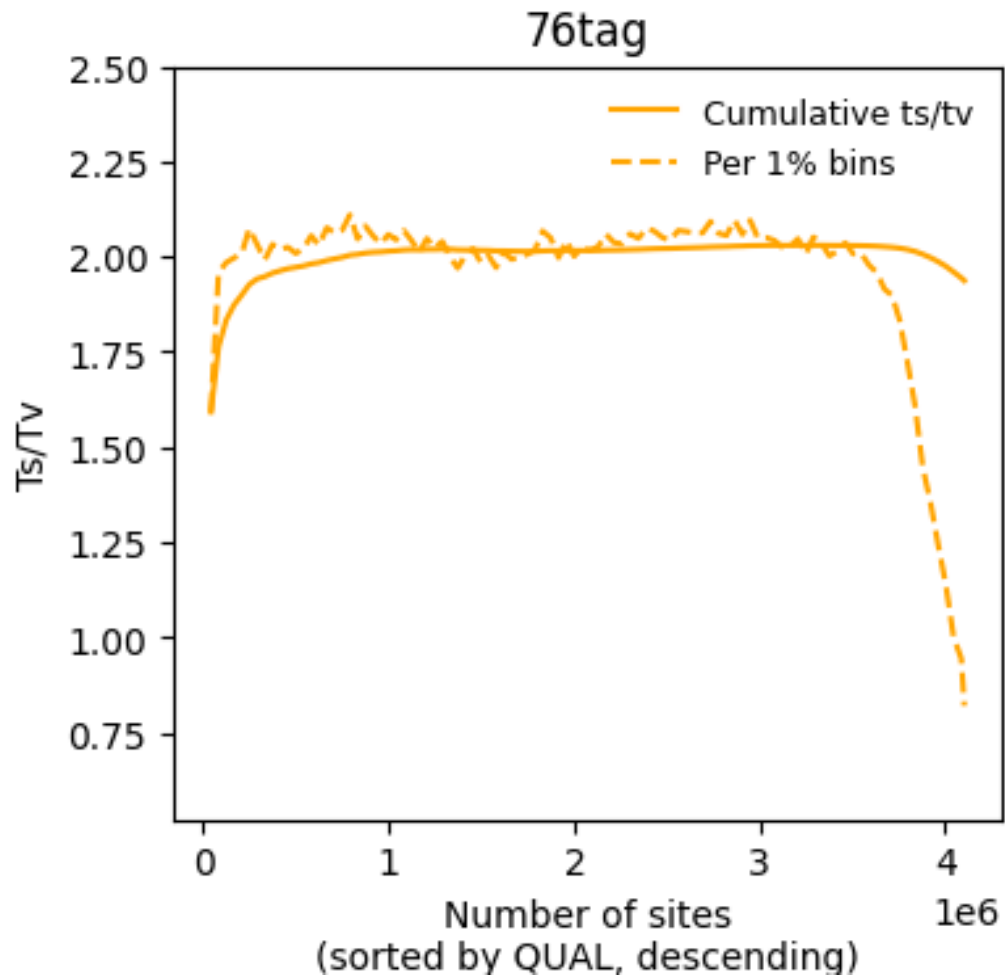
Number of SNPs by sample



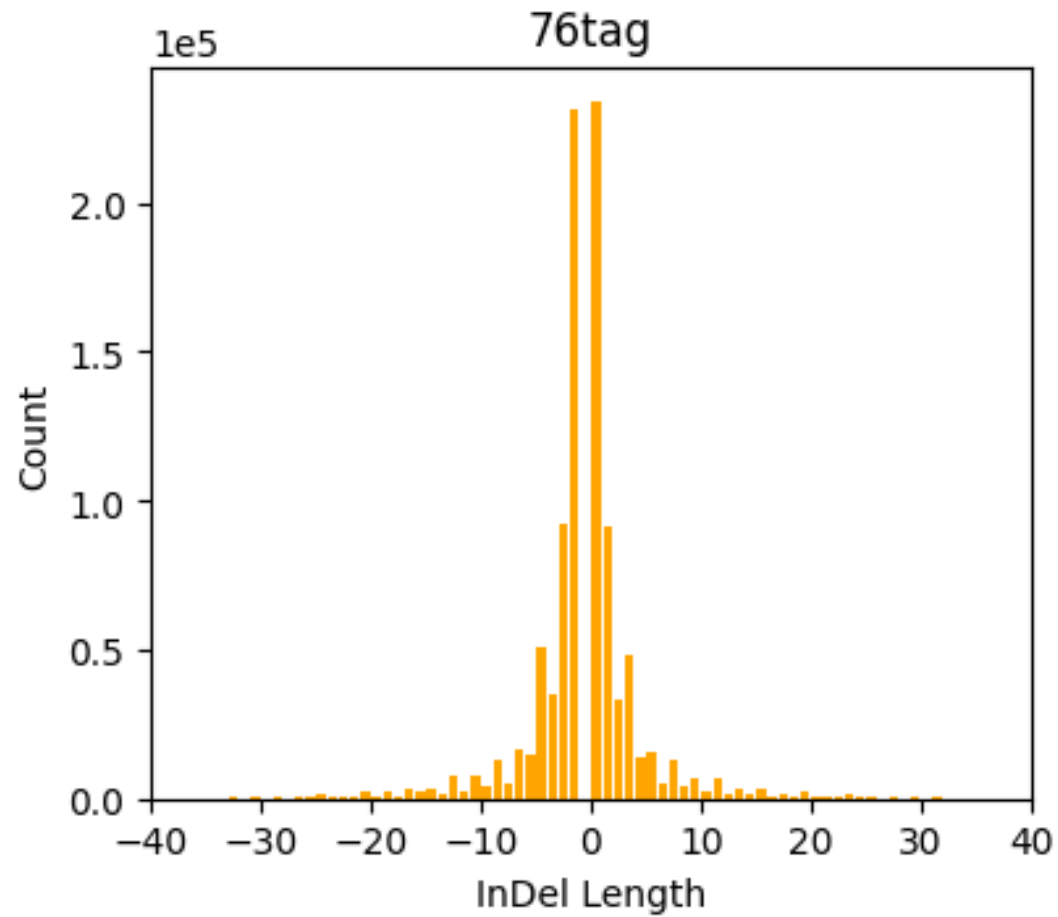
Number of indels by sample



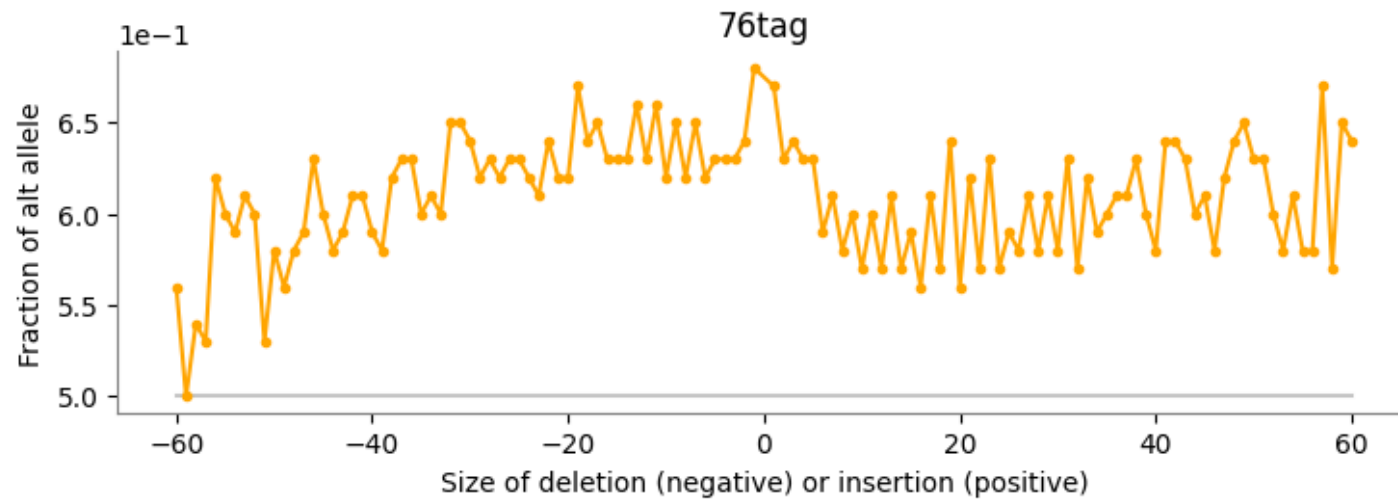
Ts/Tv stratified by QUAL



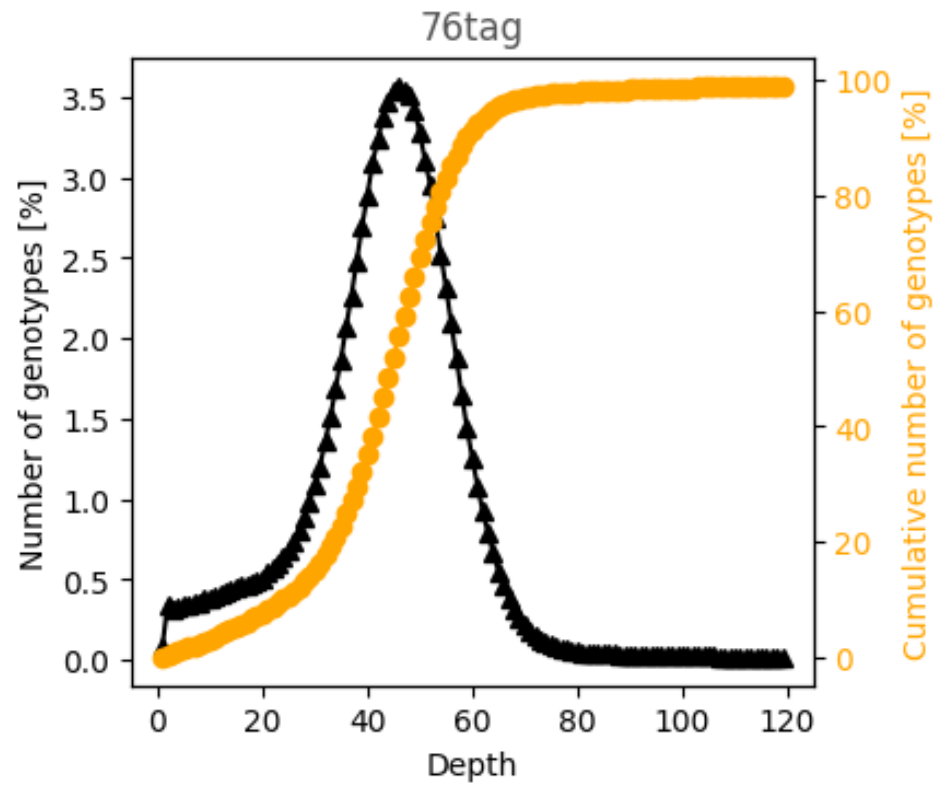
Indel distribution



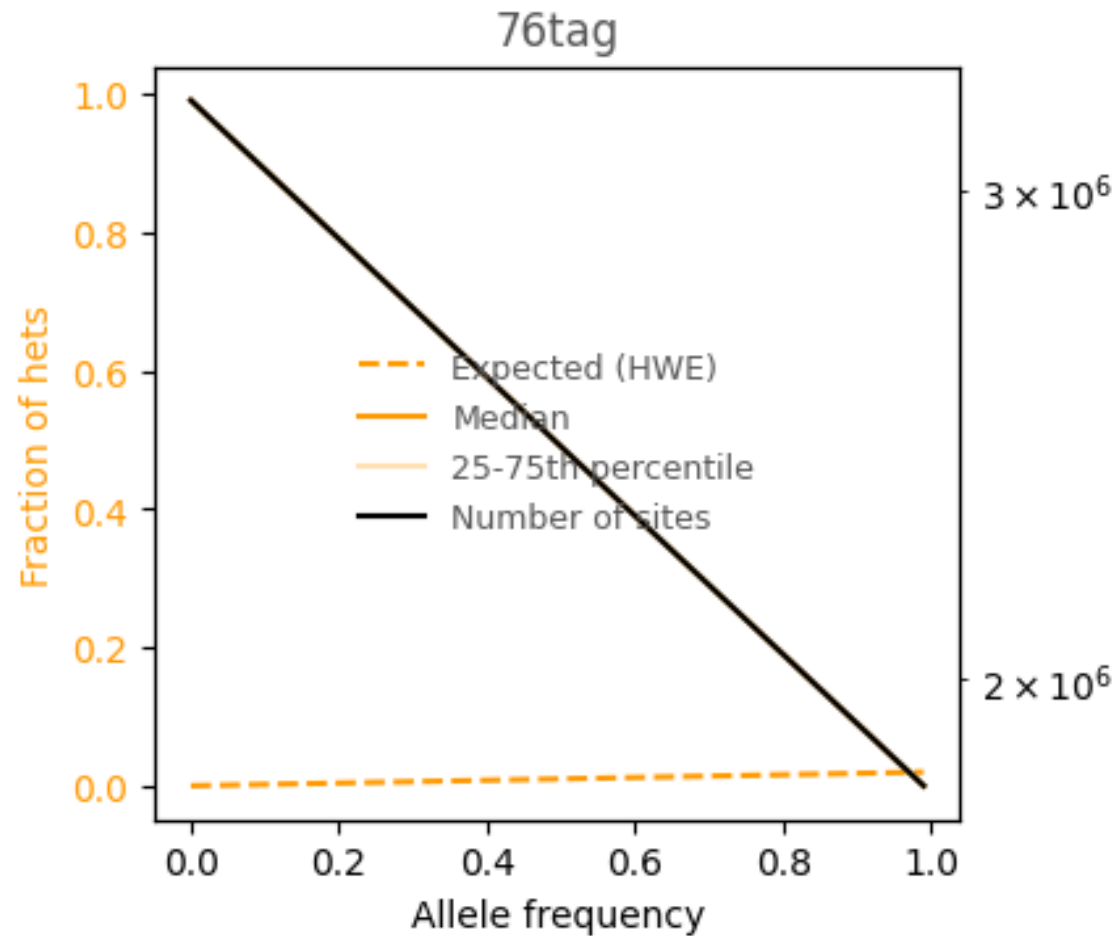
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

