

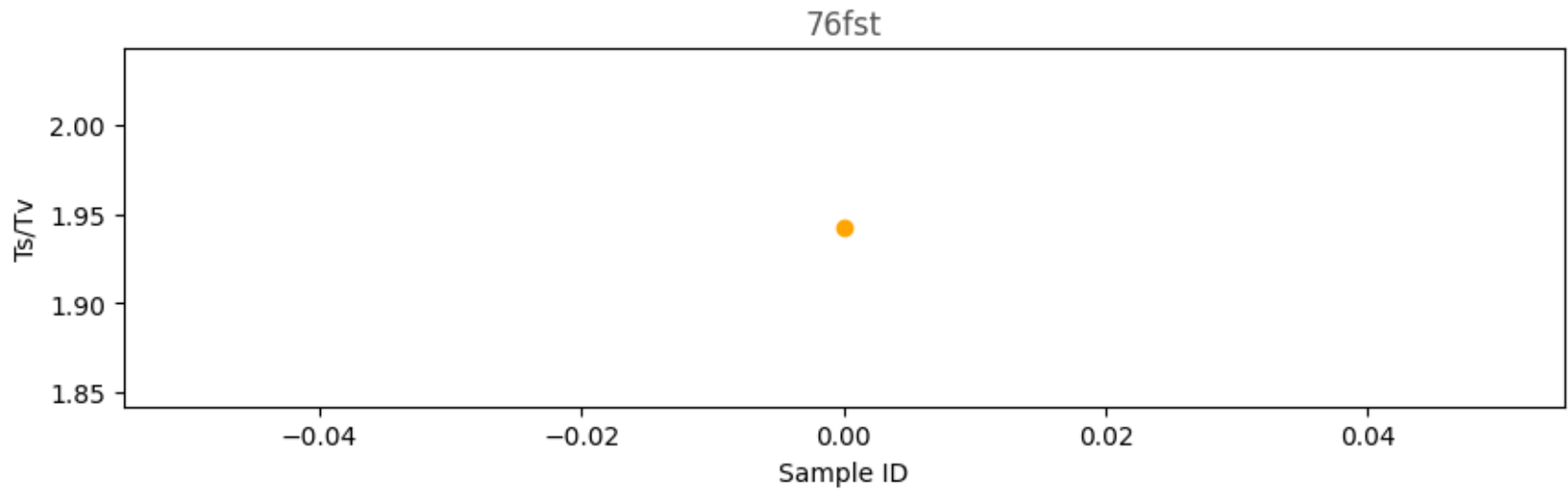
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76fst	4,057,834	1.94	1.95	952,717	–	0	0
* frameshift ratio: out/(out+in)							

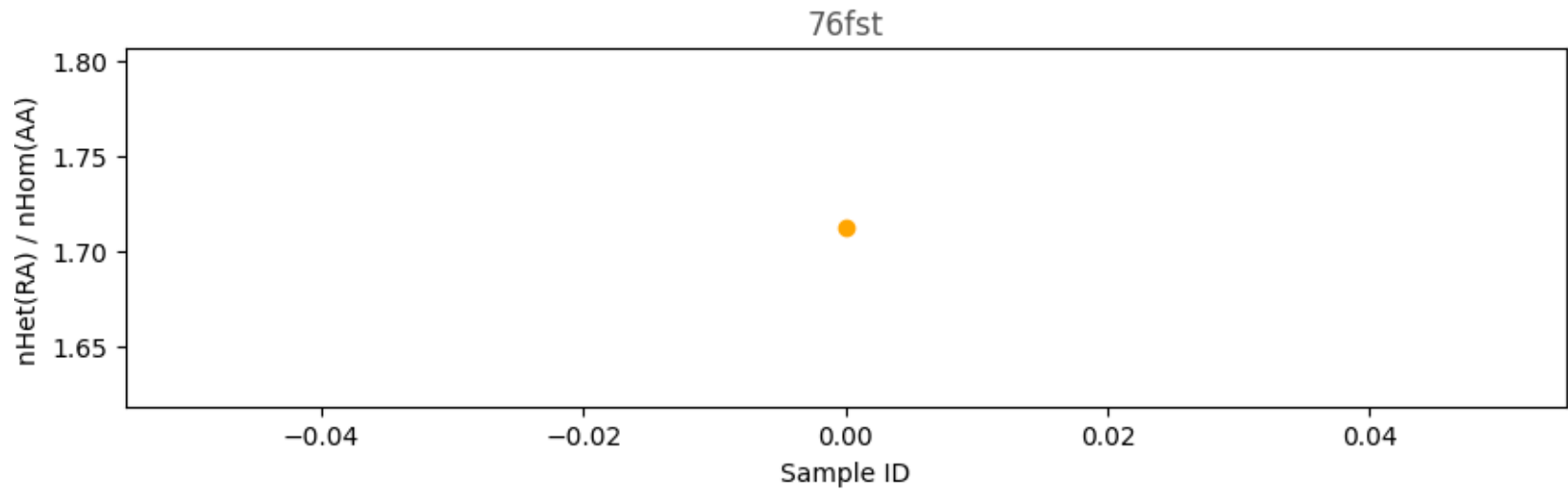
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76fst	63.1%	1.92	68.3%	100,079	1,857

- 76fst .. /ngc/projects2/gm/data/archive/2022/variants/snv/76fstasnf-103917770254-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01879-221005_A01961_AH35TTDSX5-EXT_LAB_KA_NGCWGS-NGCWGS05692-19RKG027308_snv_germline_raw.haplotype_caller.vcf.gz

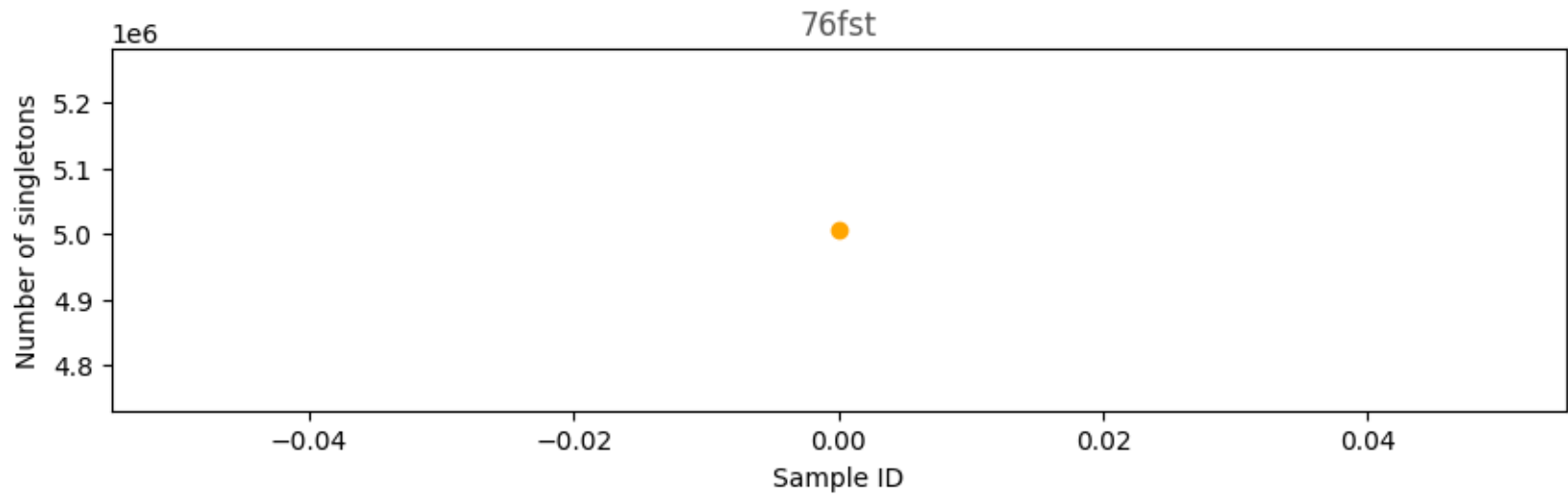
Ts/Tv by sample



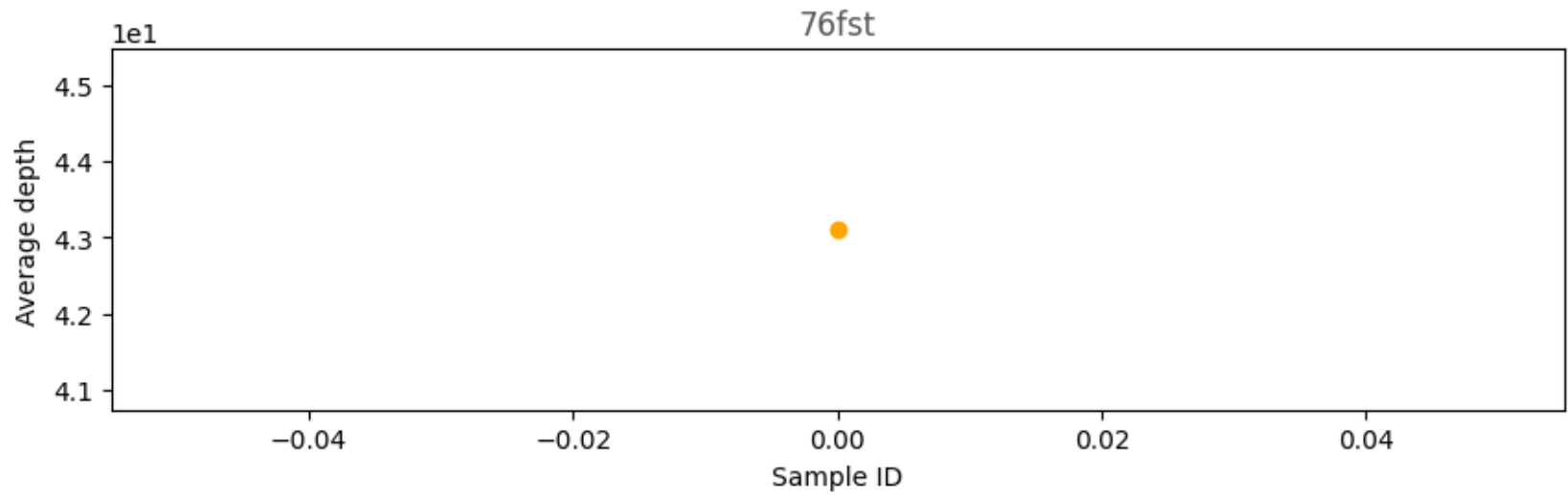
Hets vs non-ref Homs by sample



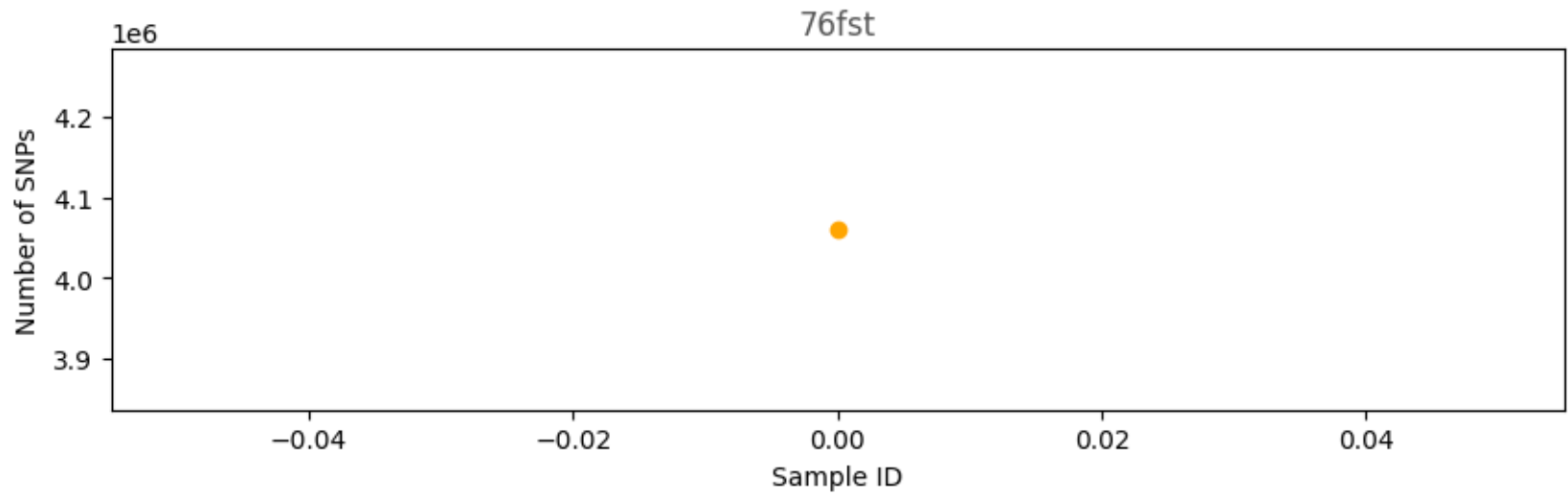
Singletons by sample (hets and homs)



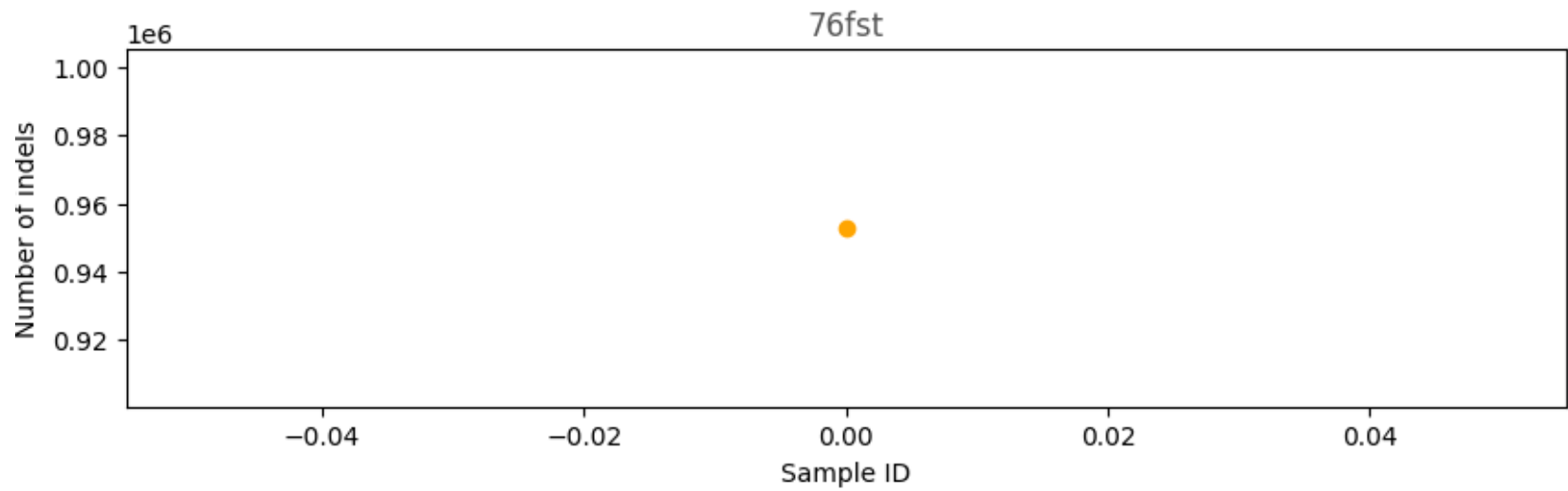
Average depth by sample



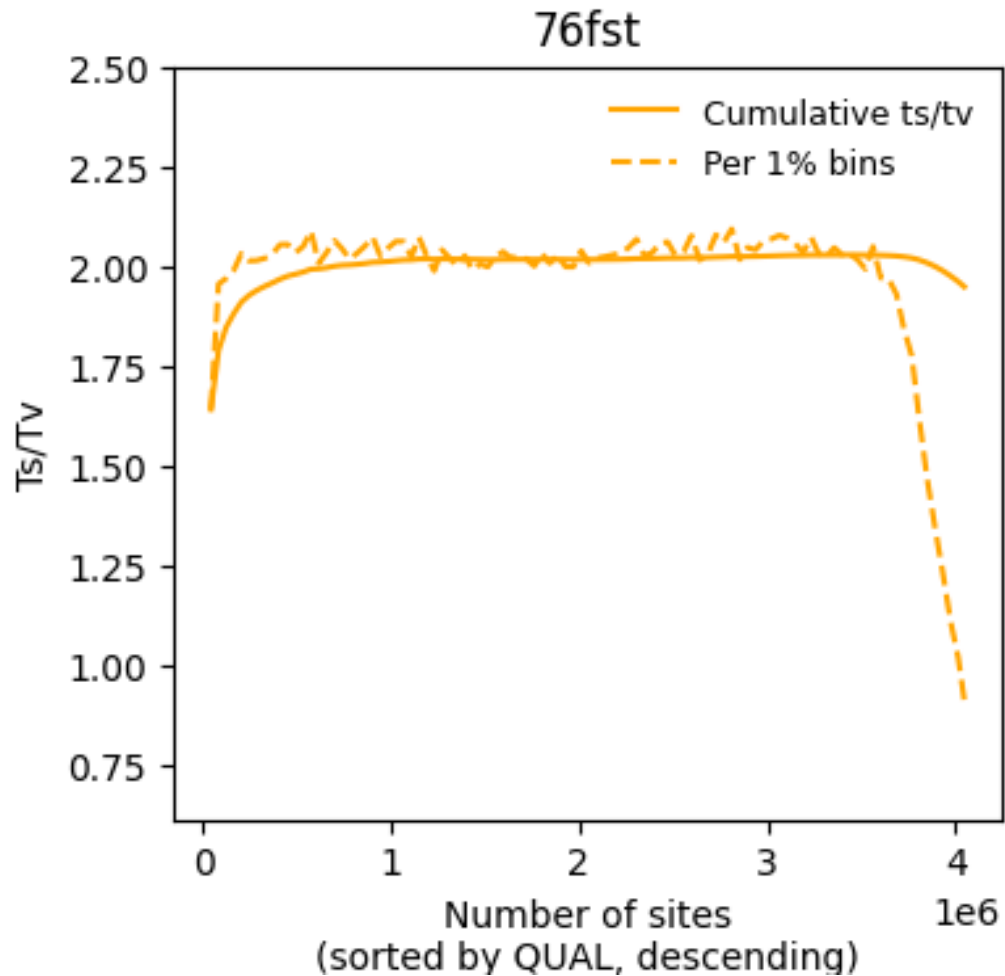
Number of SNPs by sample



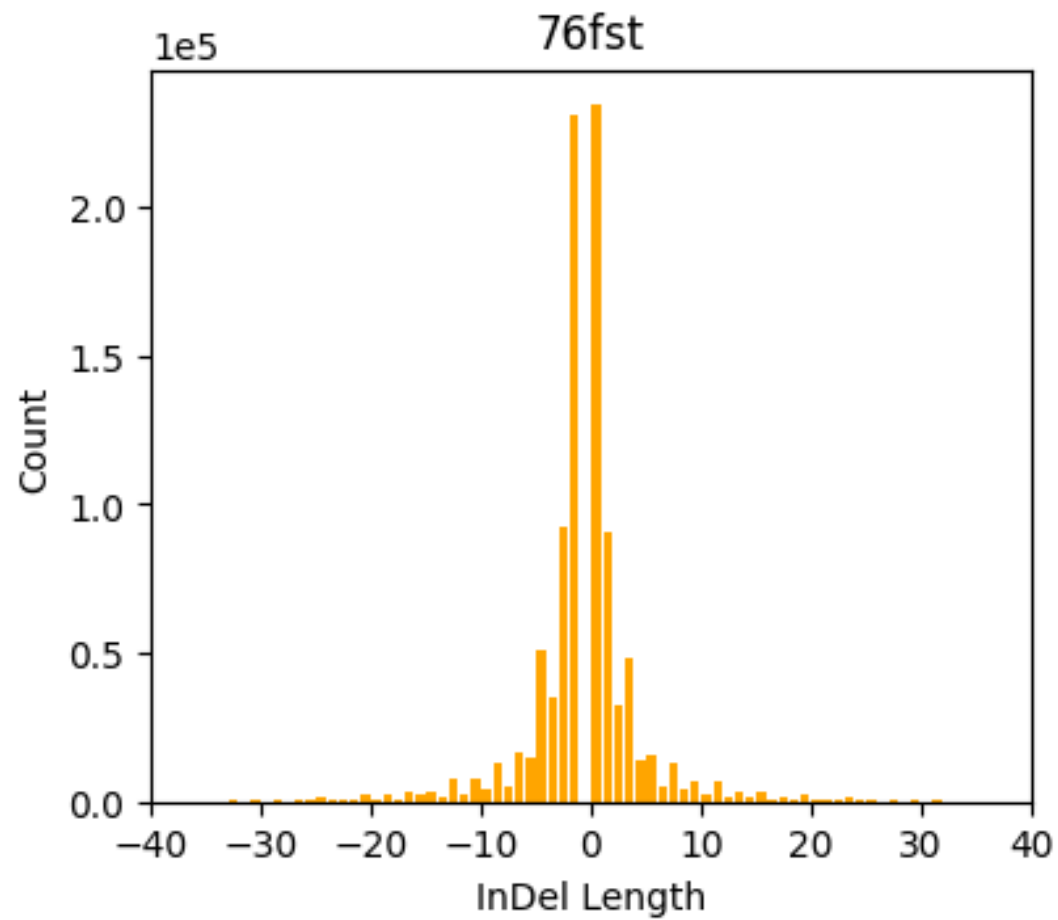
Number of indels by sample



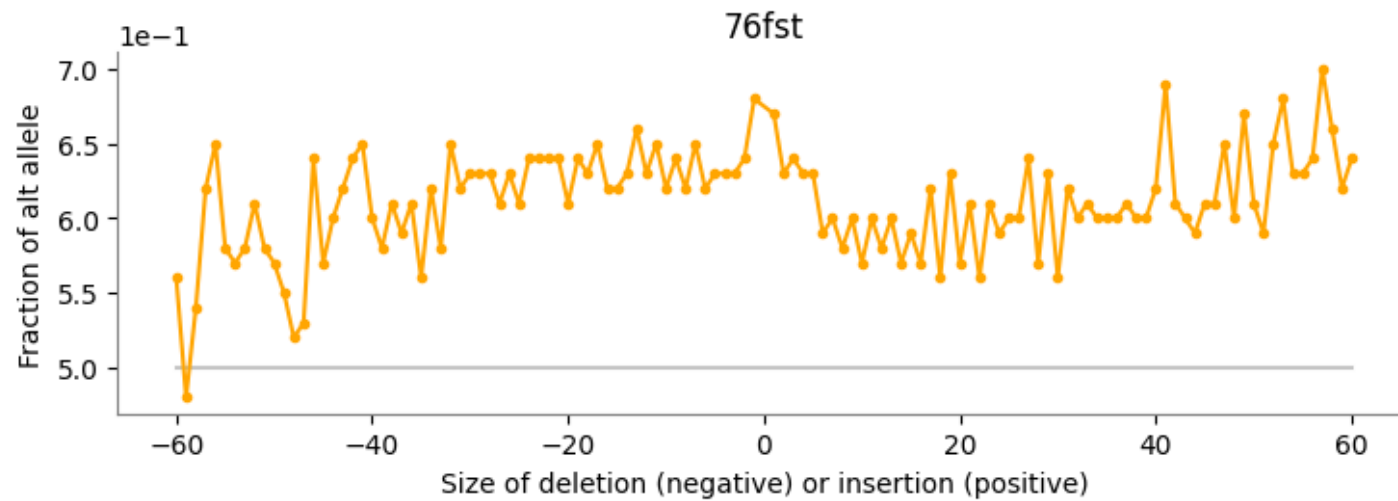
Ts/Tv stratified by QUAL



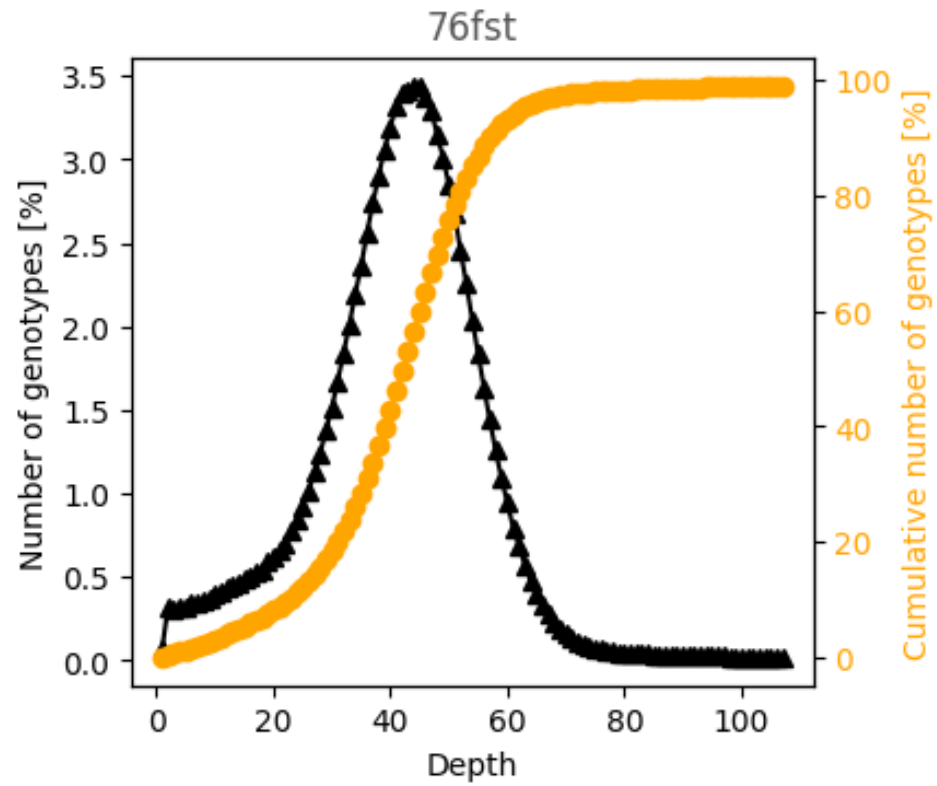
Indel distribution



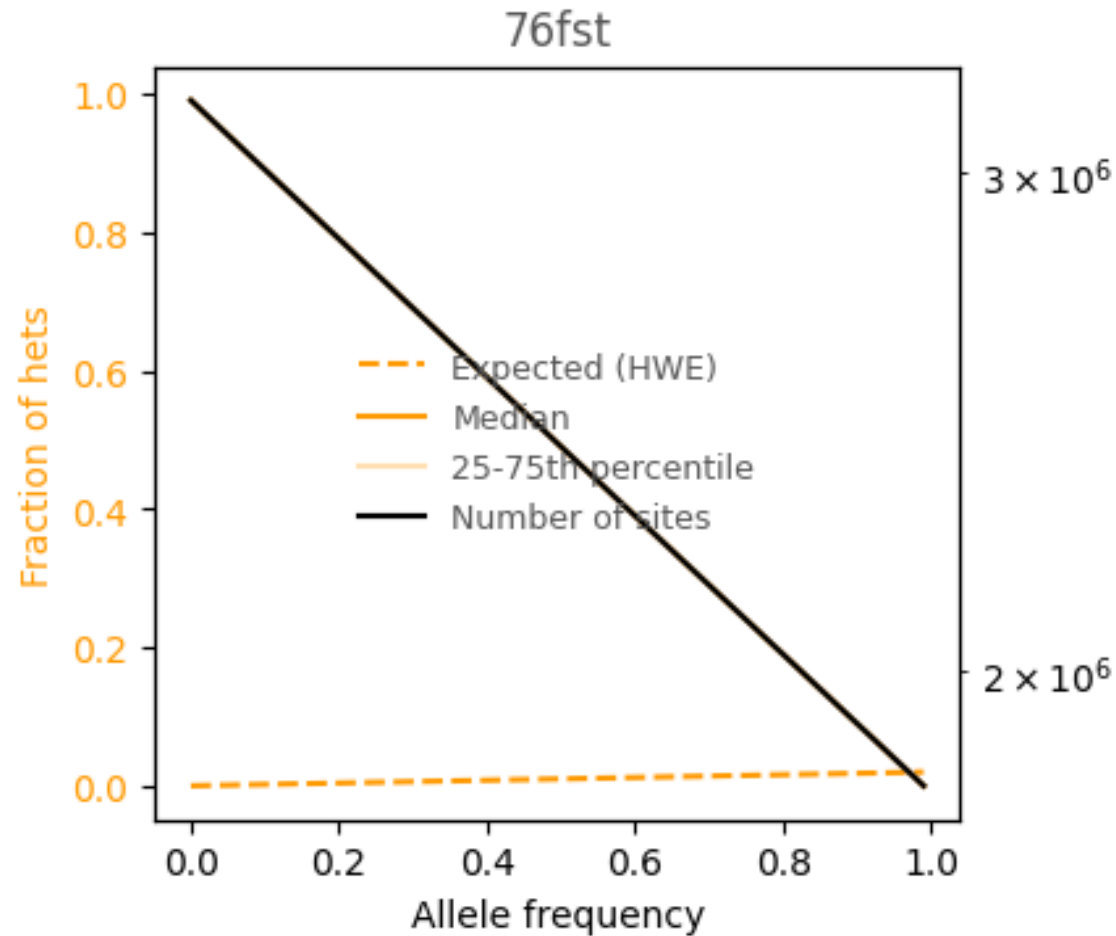
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

