

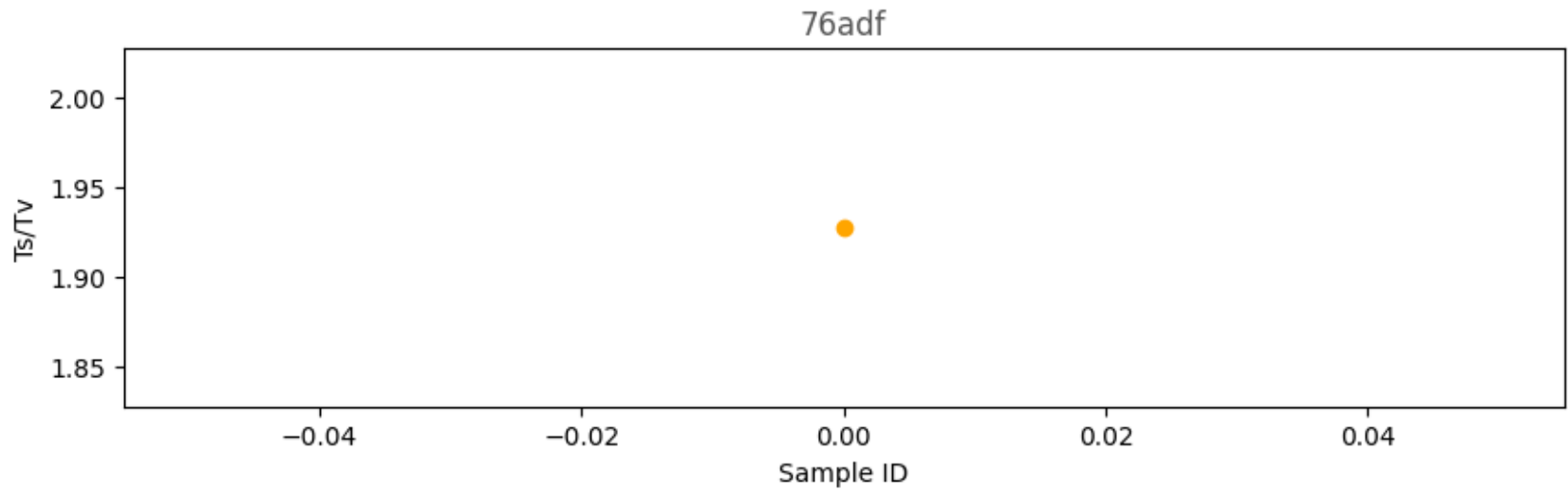
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76adf	4,059,359	1.93	1.93	947,575	–	0	0
* frameshift ratio: out/(out+in)							

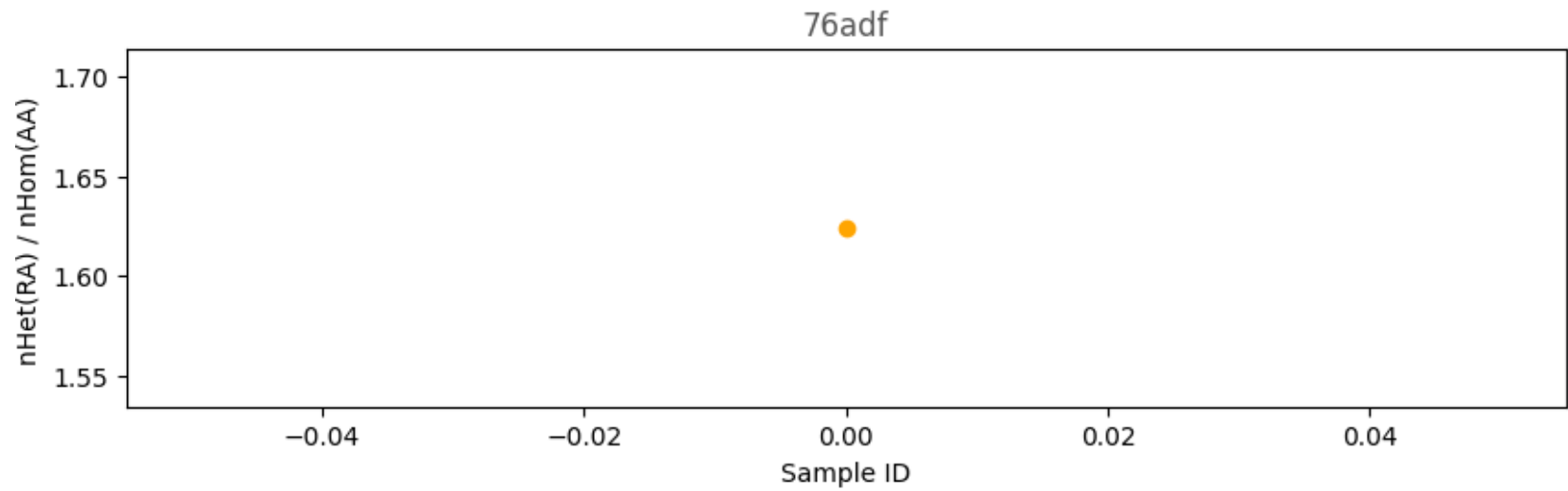
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76adf	61.9%	1.89	66.6%	99,293	1,900

- 76adf .. /ngc/projects2/gm/data/archive/2022/variants/snv/76adf/som-103893773090-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01035-220727_A01176_AHNW3KDSX3-EXT_LAB
KA_NGCWGS-NGCWGS04710_22RKG001443x01_snv_germline_raw.haplotype_caller.vcf.gz

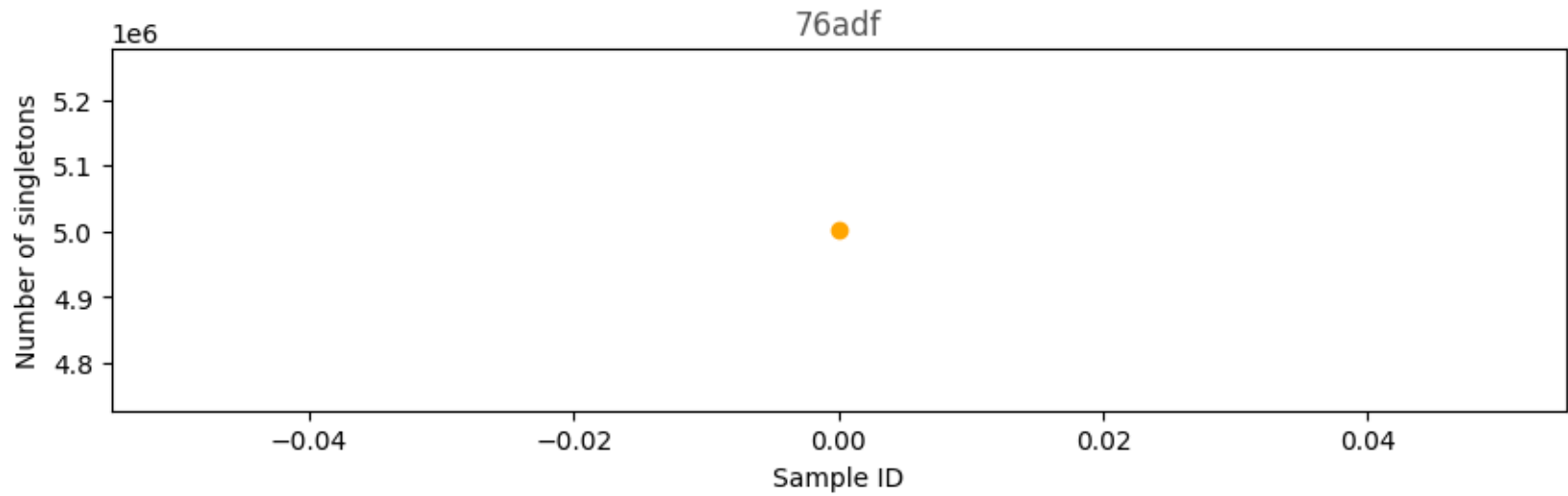
Ts/Tv by sample



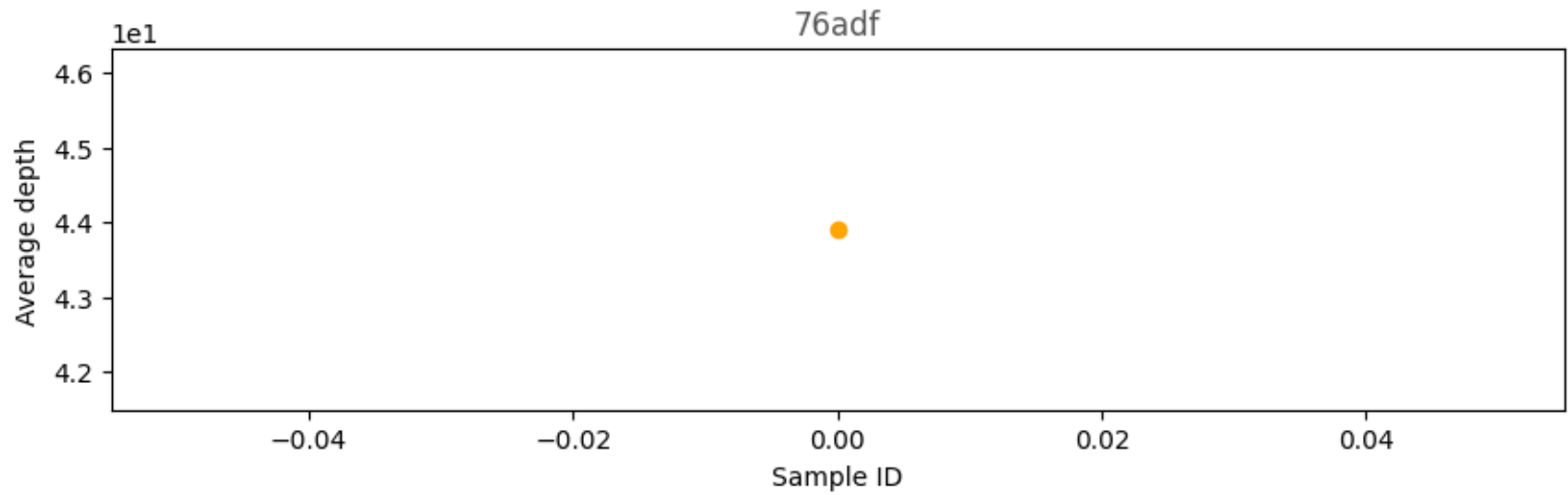
Hets vs non-ref Homs by sample



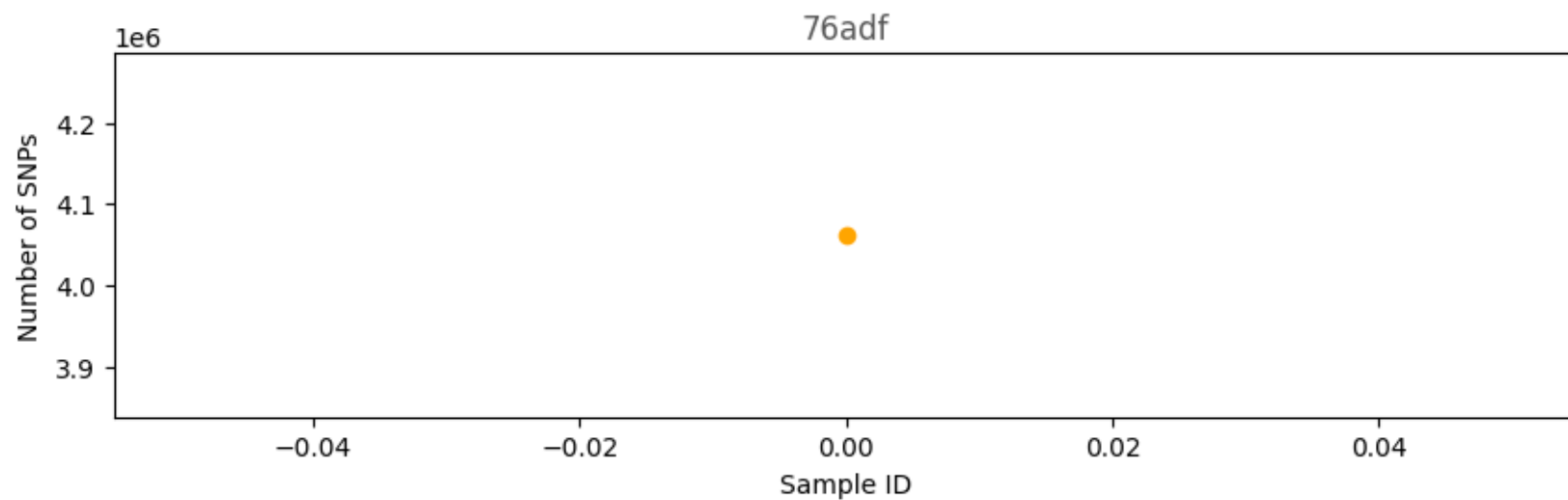
Singletons by sample (hets and homs)



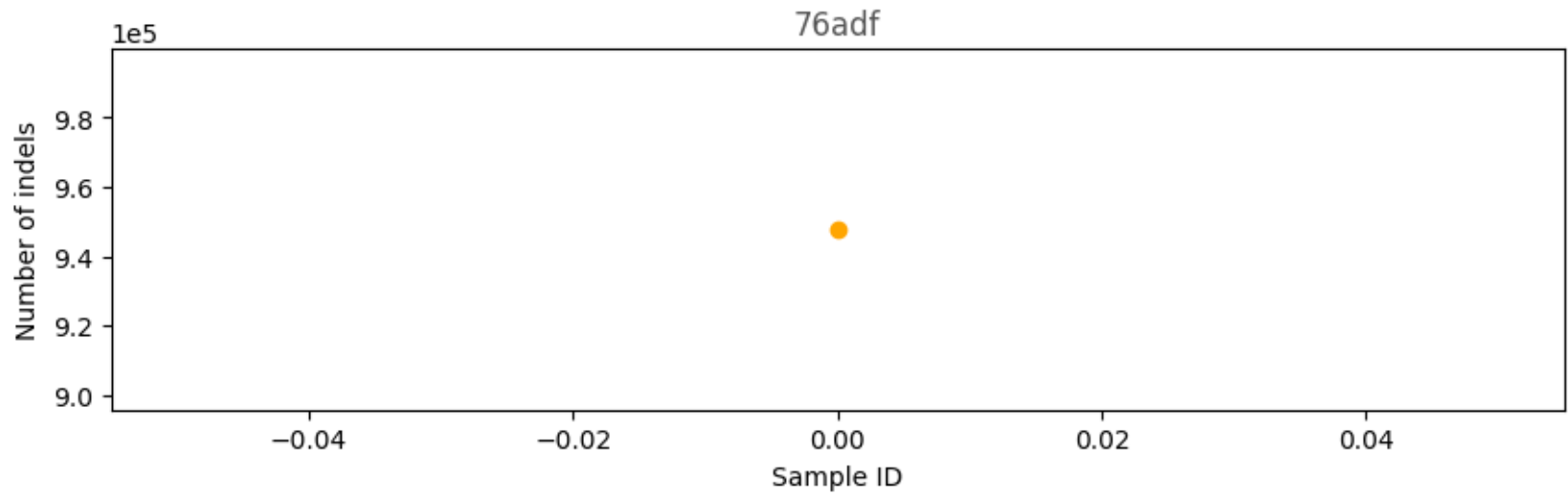
Average depth by sample



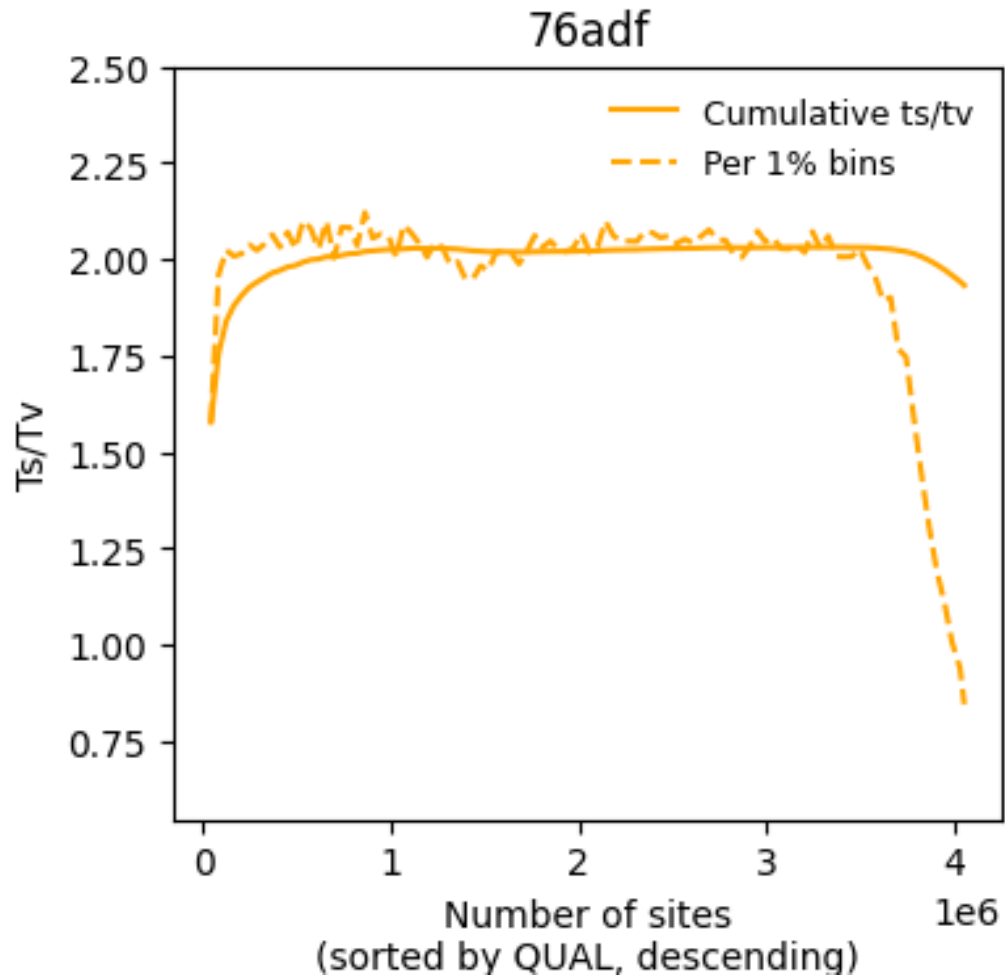
Number of SNPs by sample



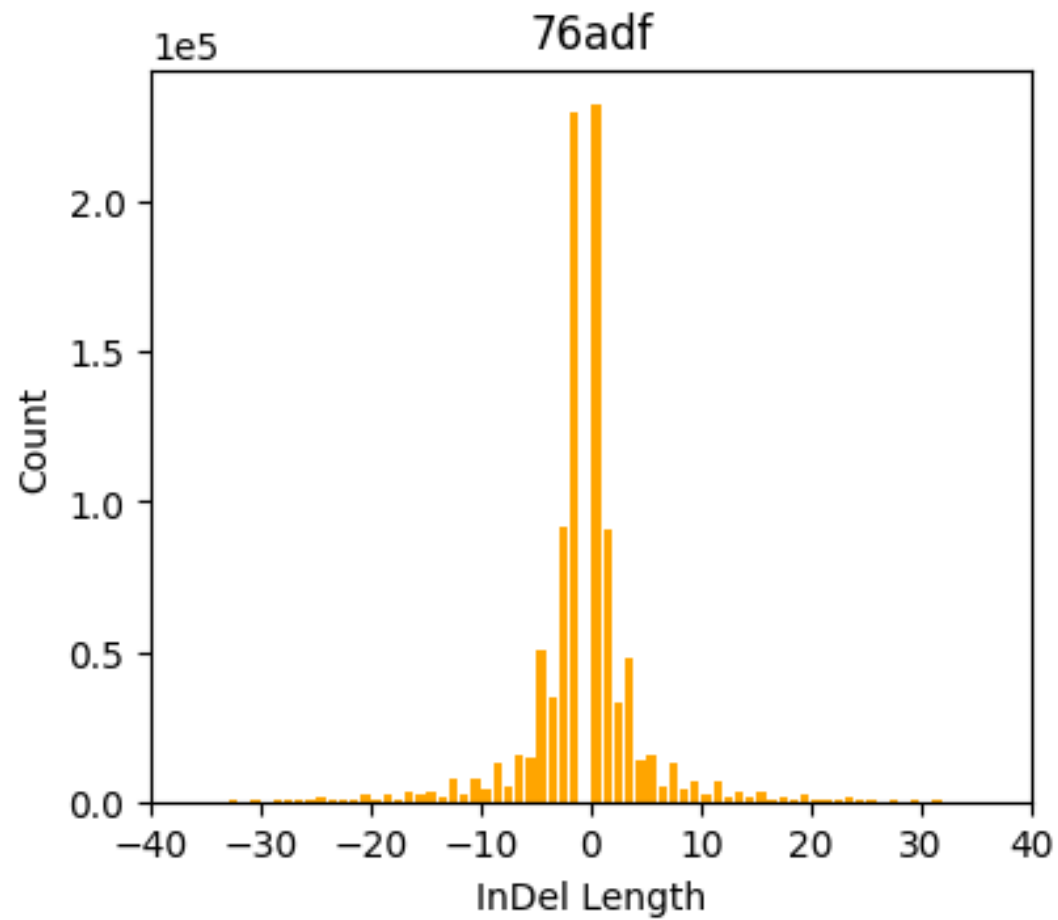
Number of indels by sample



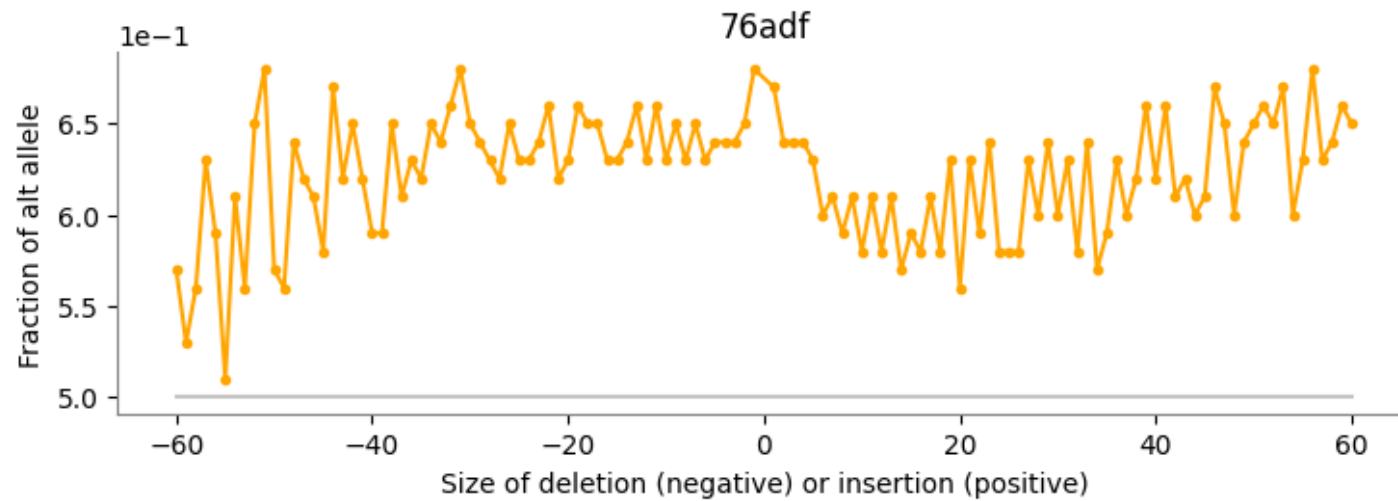
Ts/Tv stratified by QUAL



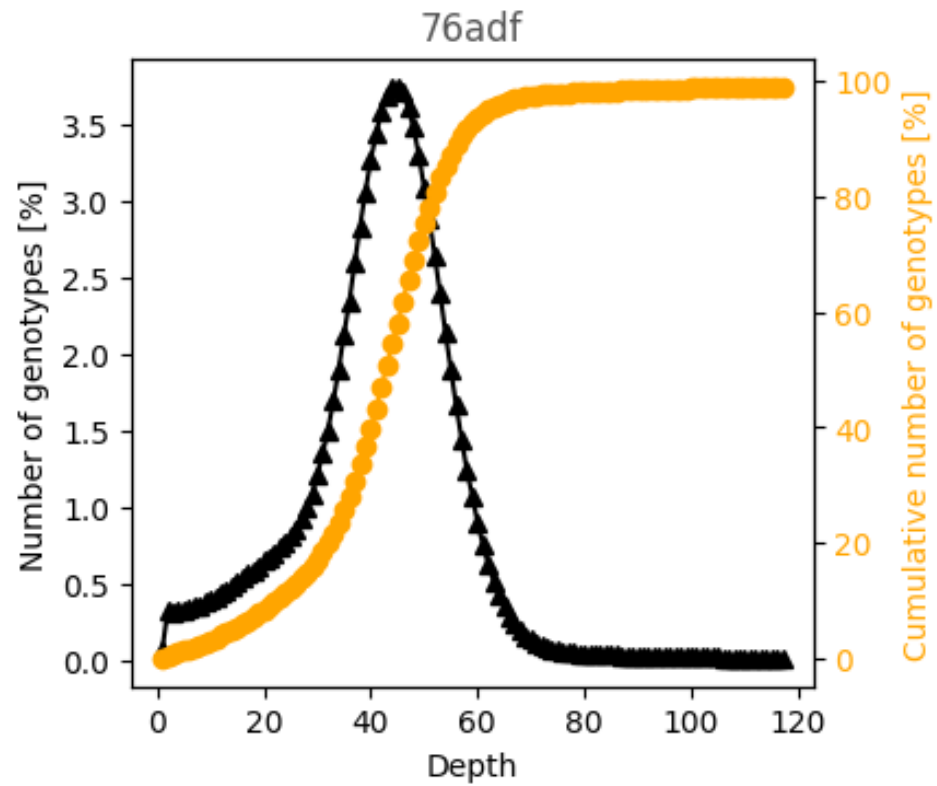
Indel distribution



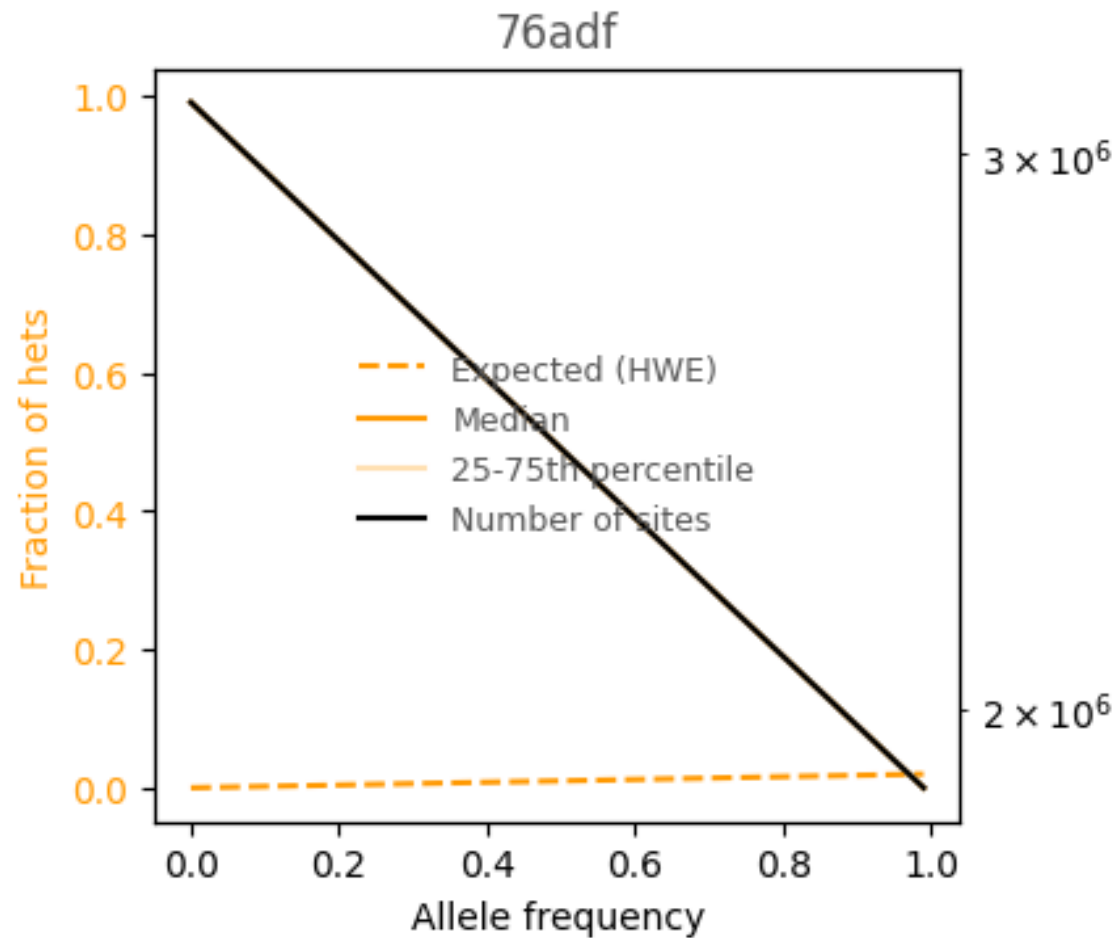
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

