

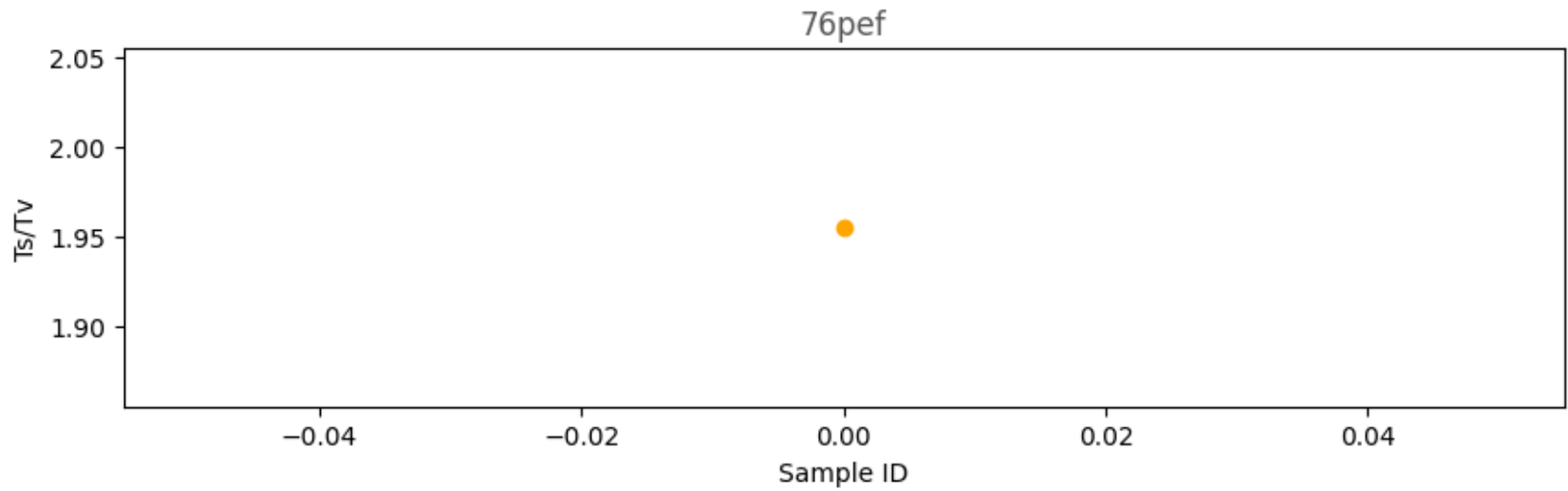
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76pef	4,079,135	1.96	1.96	959,139	–	0	0
* frameshift ratio: out/(out+in)							

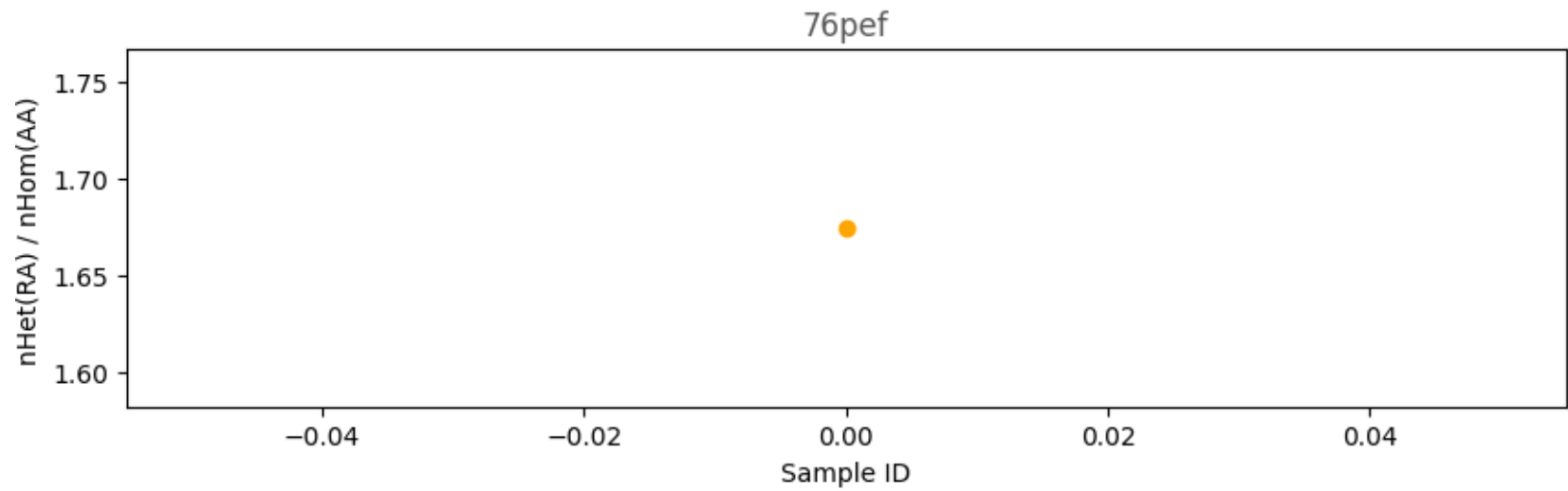
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76pef	62.6%	1.93	67.4%	99,896	1,967

- 76pef .. /ngc/projects2/gm/data/archive/2022/variants/snv/76pefedcm-103794007067-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00387-211026_A01411_AHN23JDSX2-EXT_LAB
 KA_NGCWGS-NGCWGS00286_21RKG024579x01_snv_germline_raw.haplotype_caller.vcf.gz

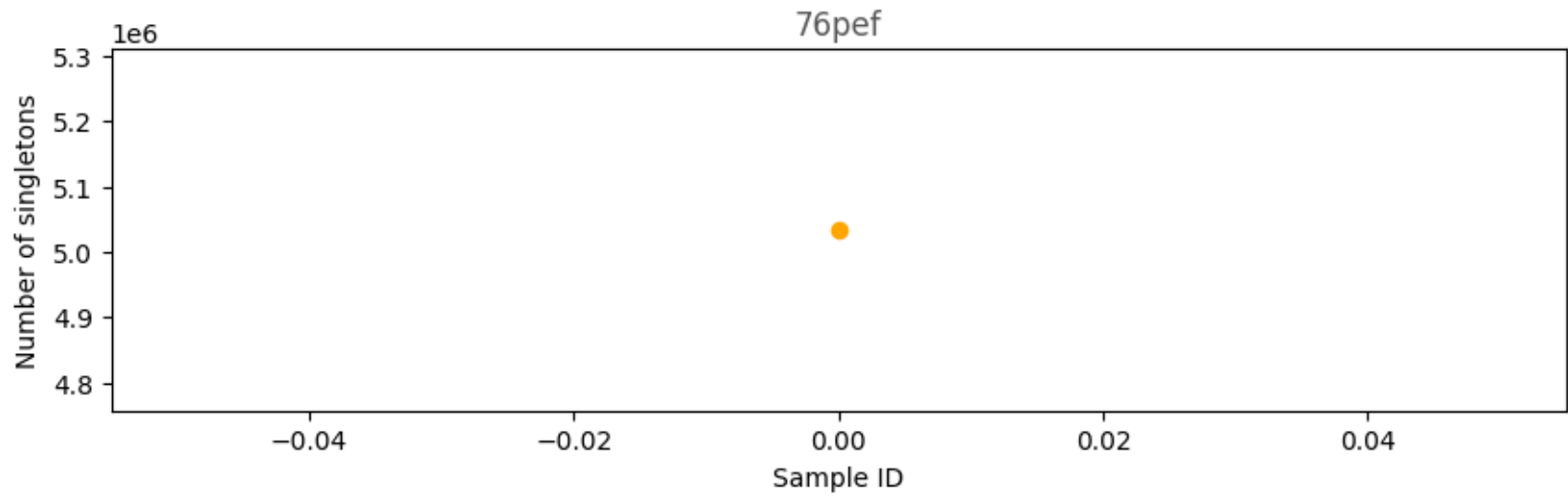
Ts/Tv by sample



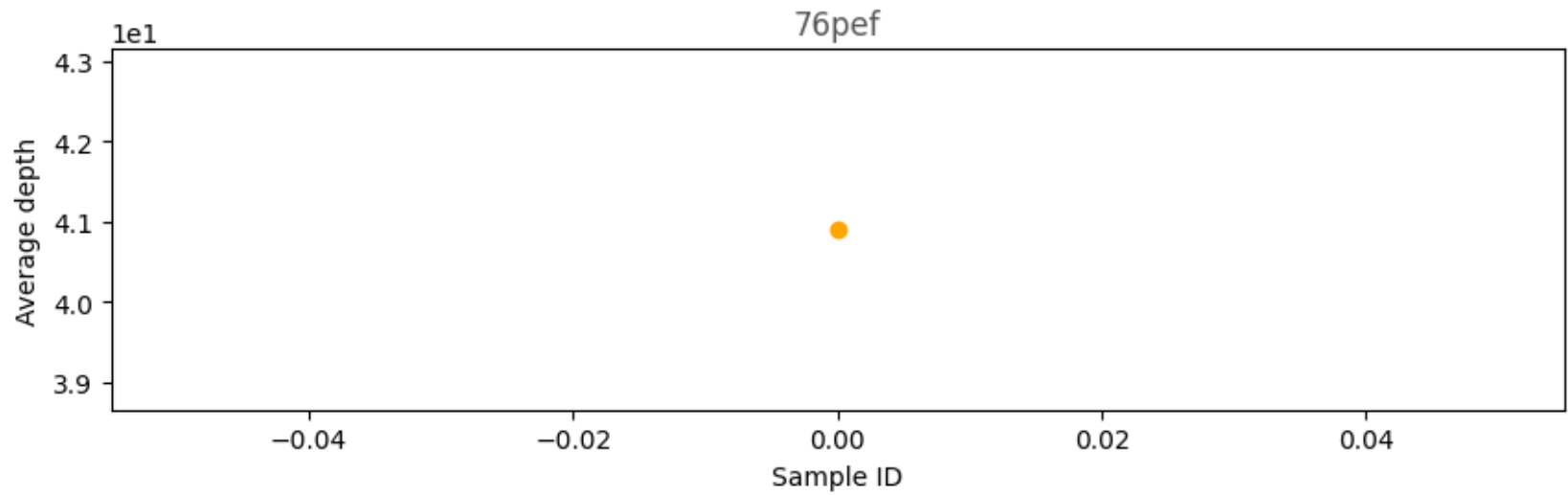
Hets vs non-ref Homs by sample



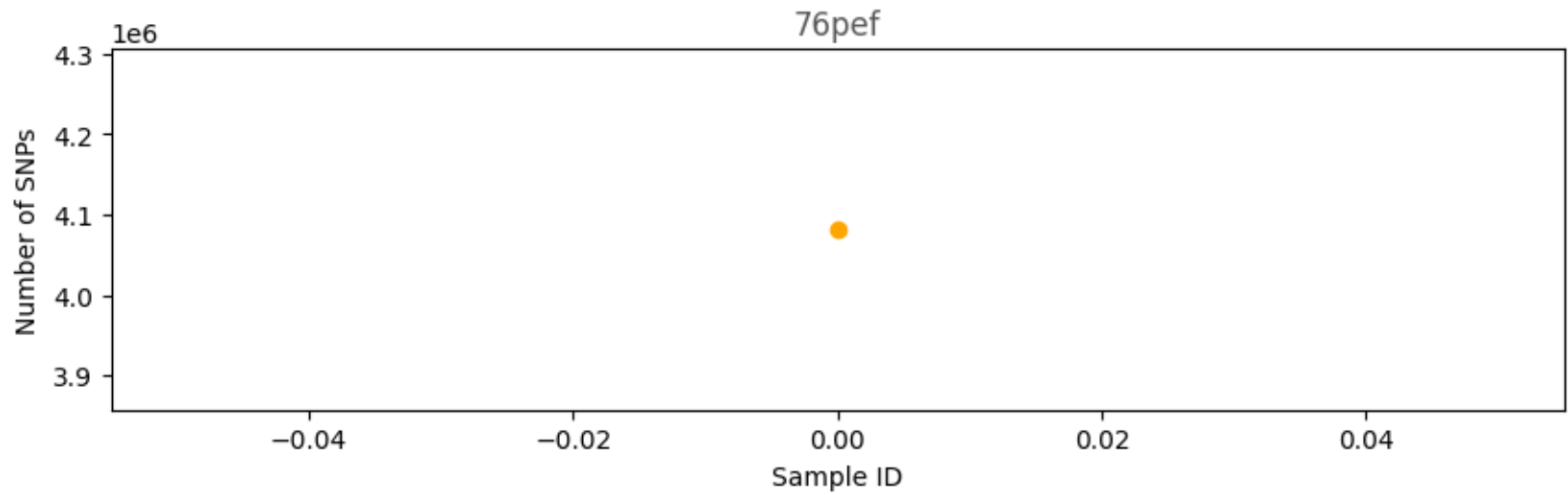
Singletons by sample (hets and homs)



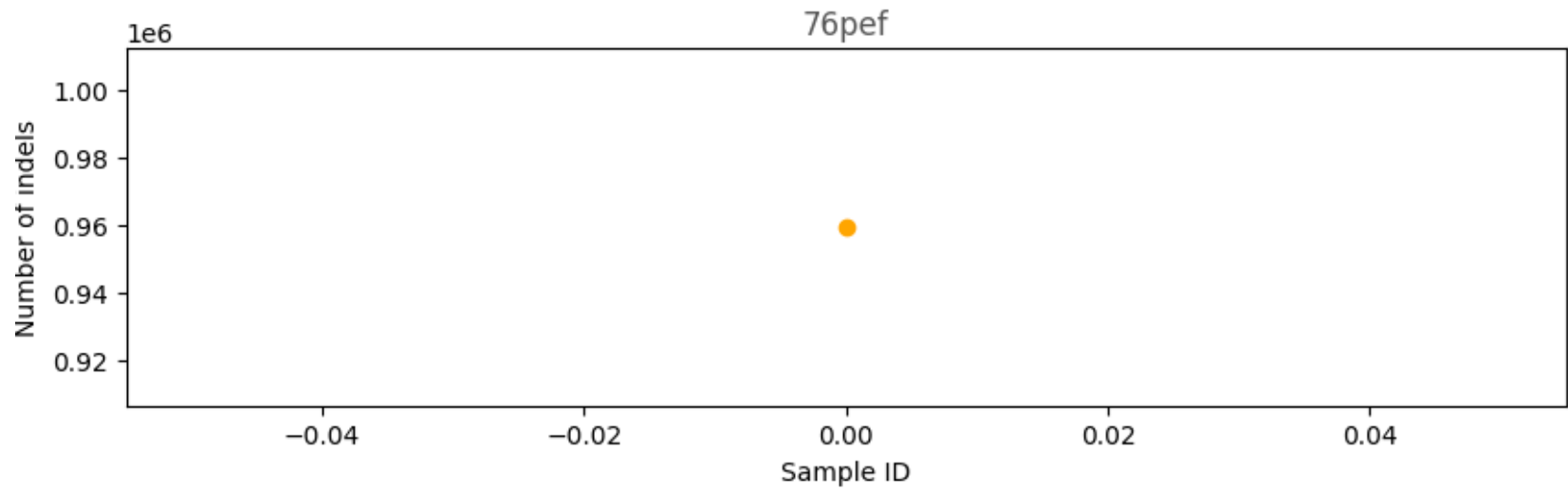
Average depth by sample



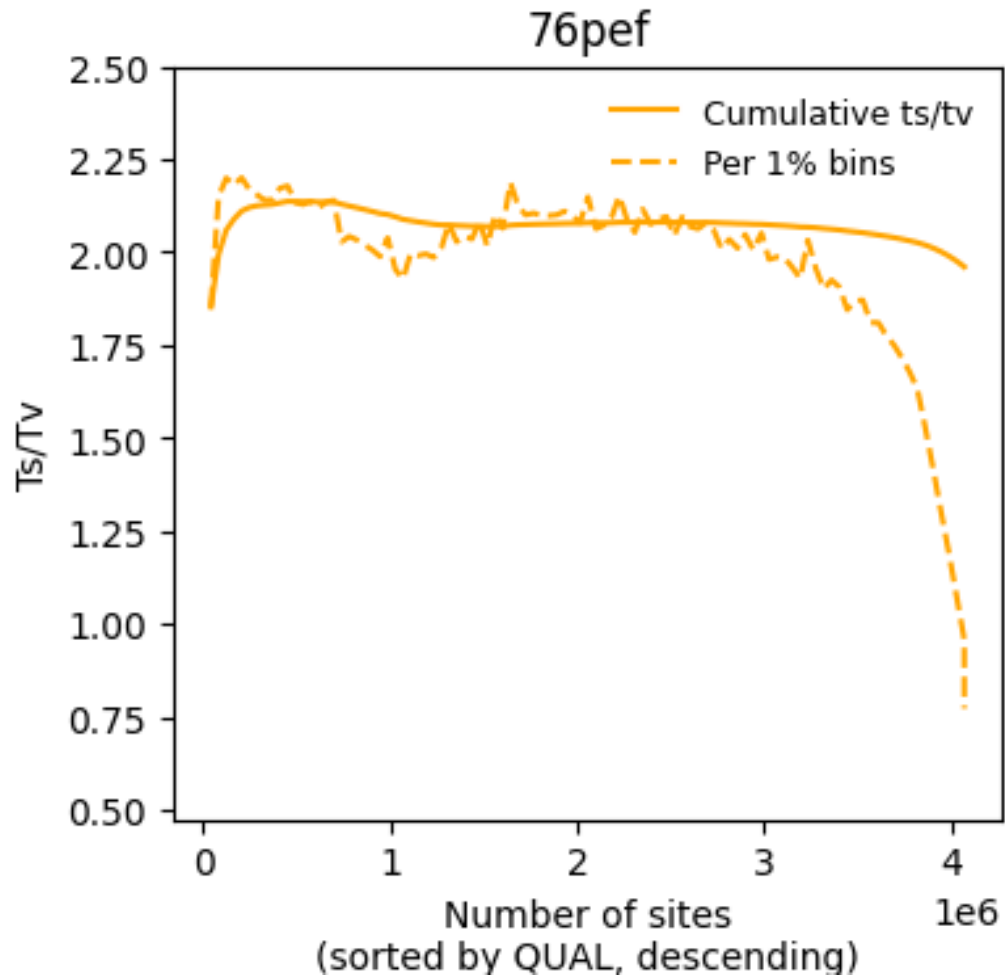
Number of SNPs by sample



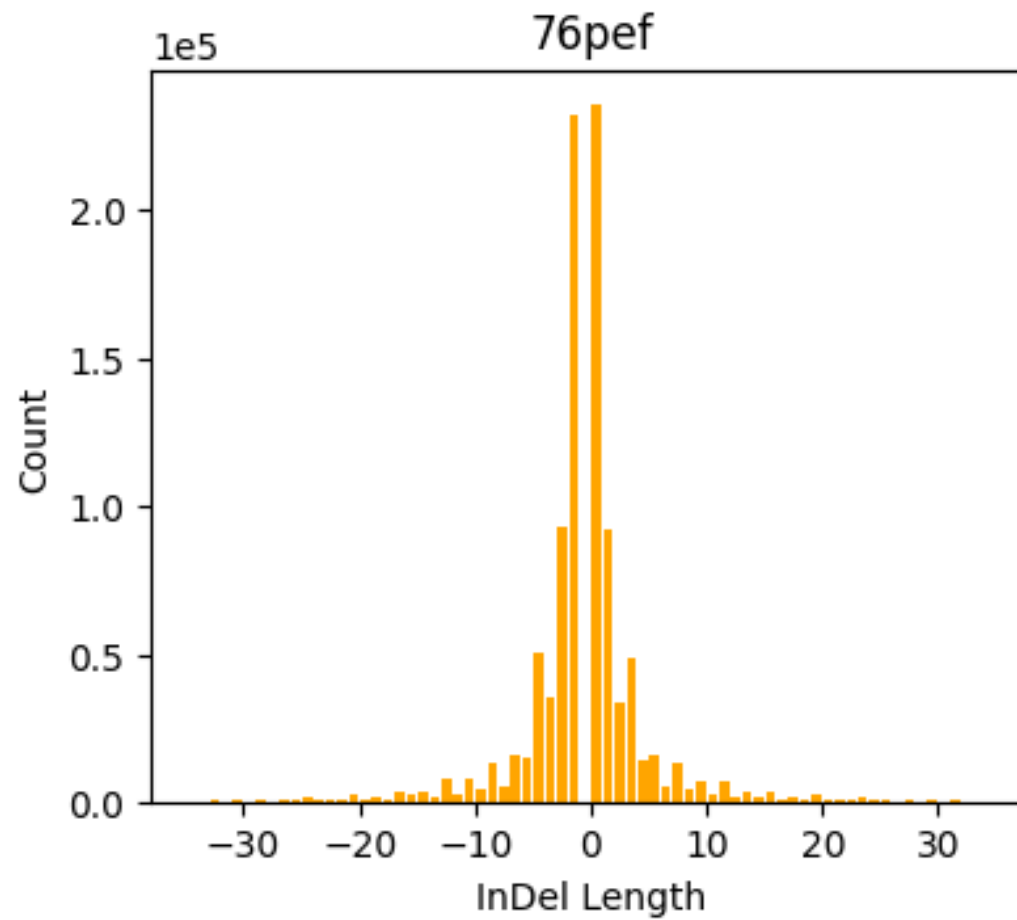
Number of indels by sample



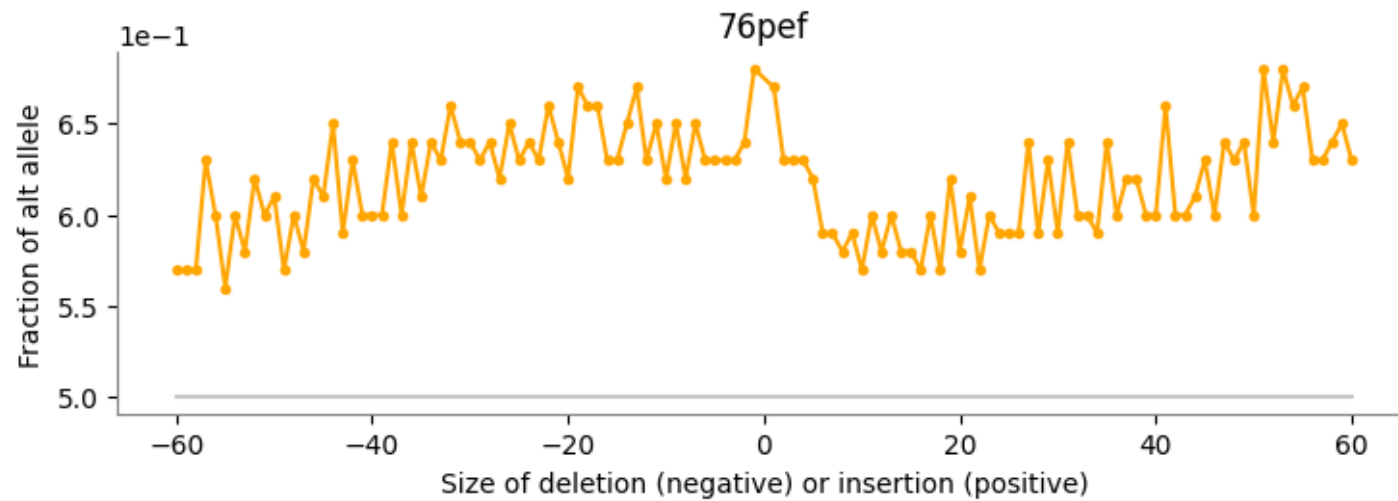
Ts/Tv stratified by QUAL



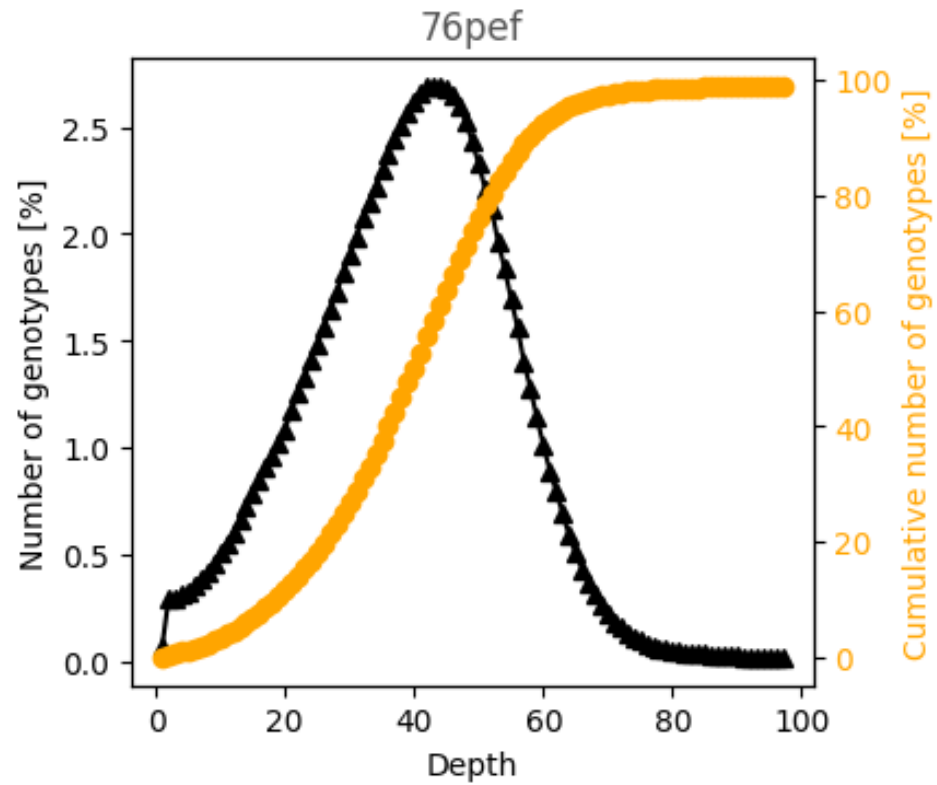
Indel distribution



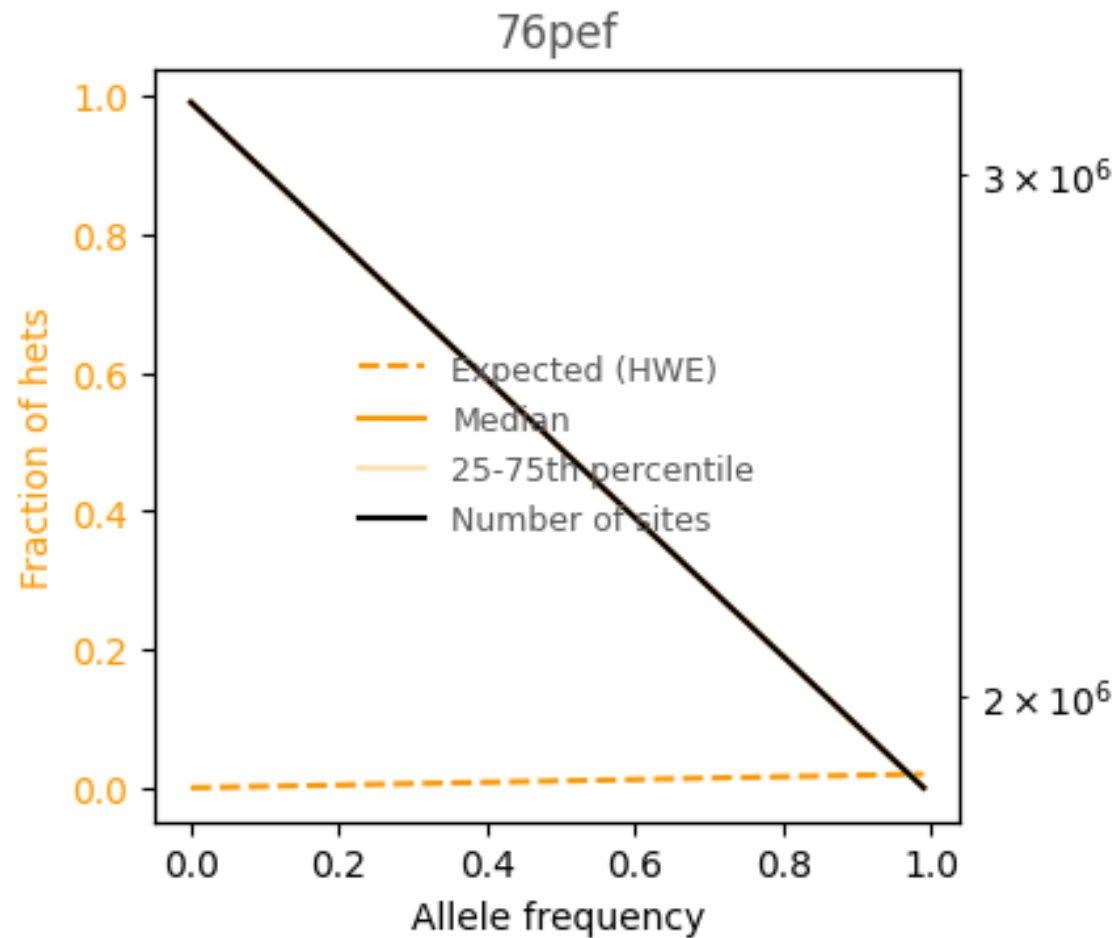
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

