

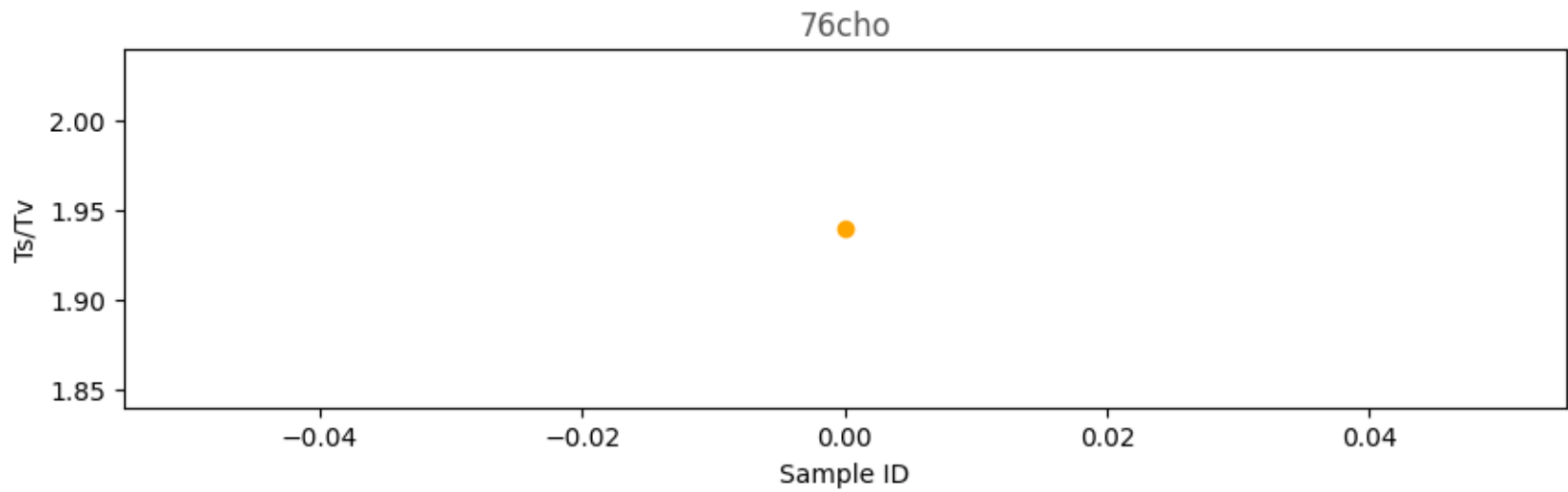
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76cho	4,058,343	1.94	1.94	943,325	–	0	0
* frameshift ratio: out/(out+in)							

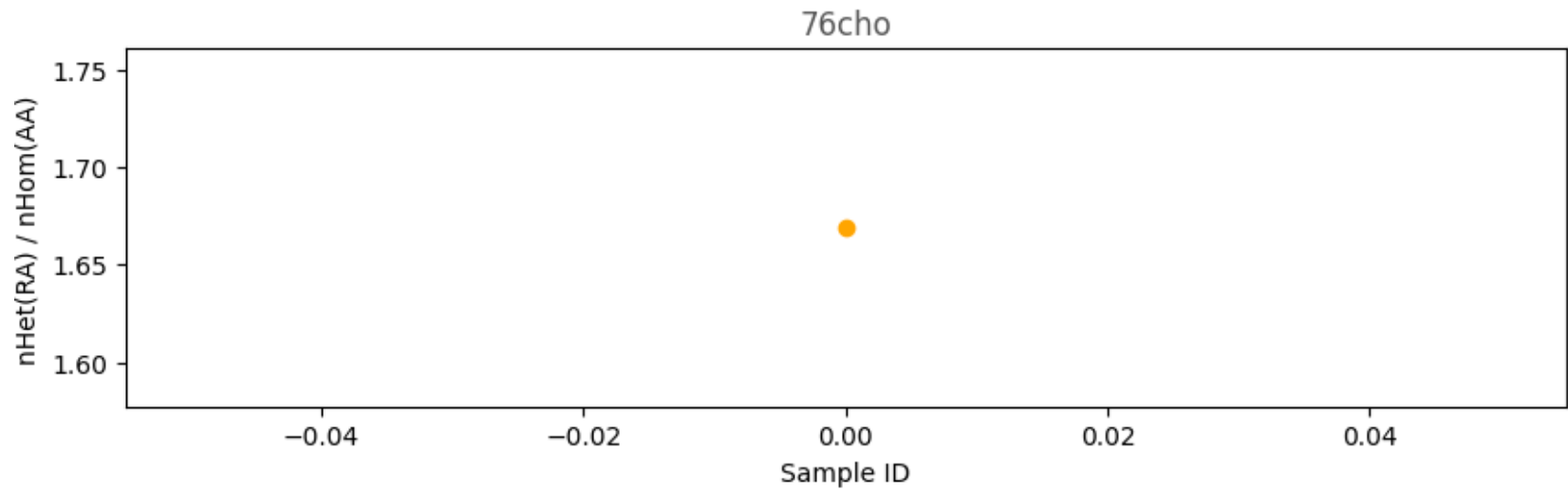
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76cho	62.5%	1.91	67.4%	96,870	2,135

- 76cho .. /ngc/projects2/gm/data/archive/2022/variants/snv/76chonyhf-103794070125-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00357-211019_A01411_BHHHY3DSX2-EXT_LAB
 KA_NGCWGS-NGCWGS00256_21RKG022879x01_snv_germline_raw.haplotype_caller.vcf.gz

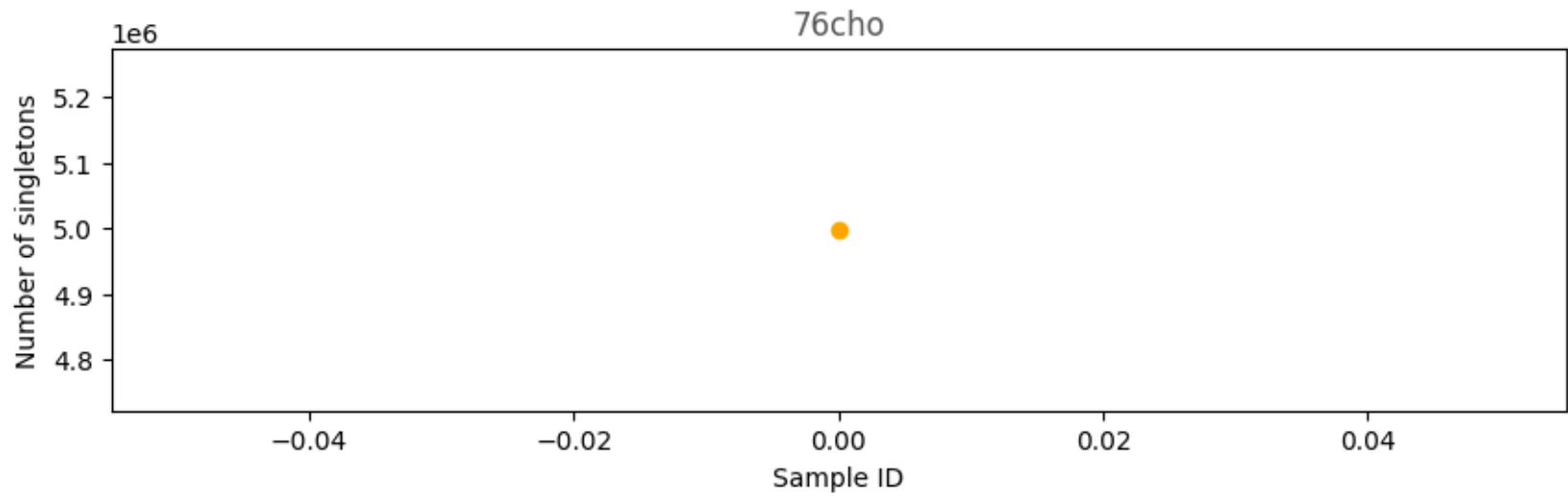
Ts/Tv by sample



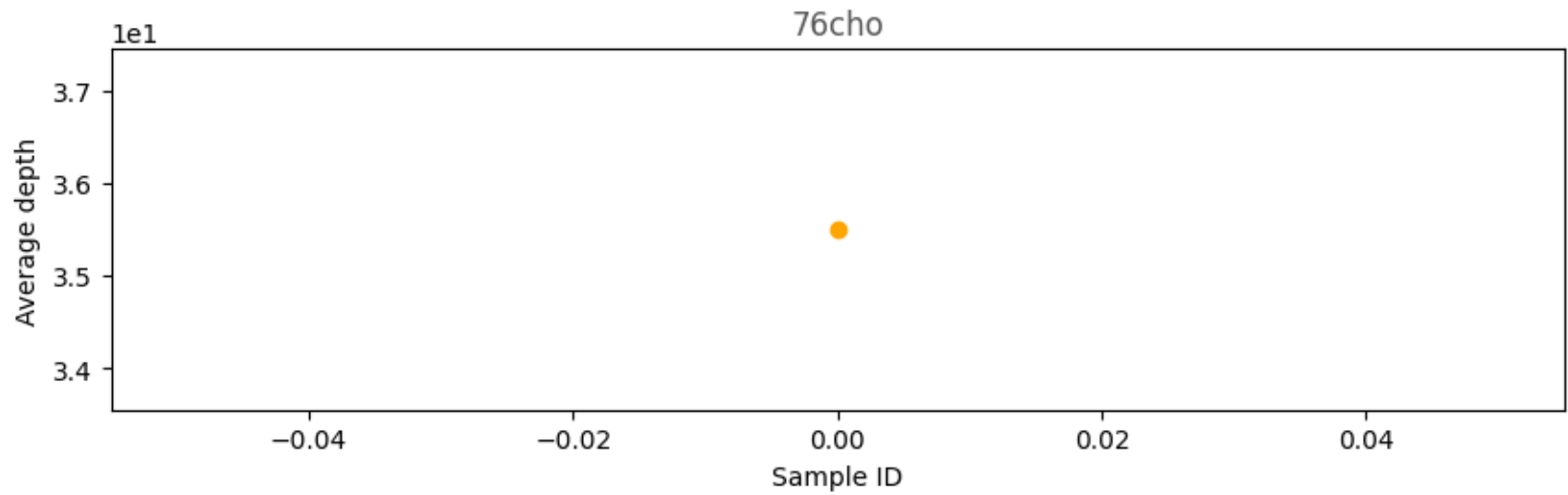
Hets vs non-ref Homs by sample



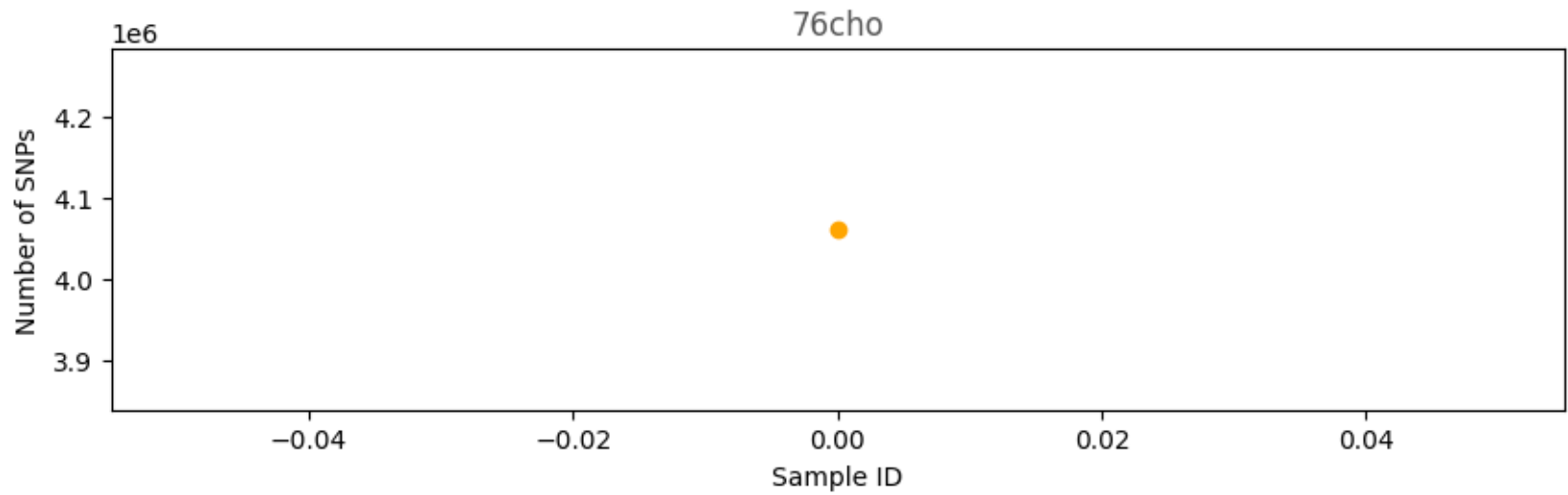
Singletons by sample (hets and homs)



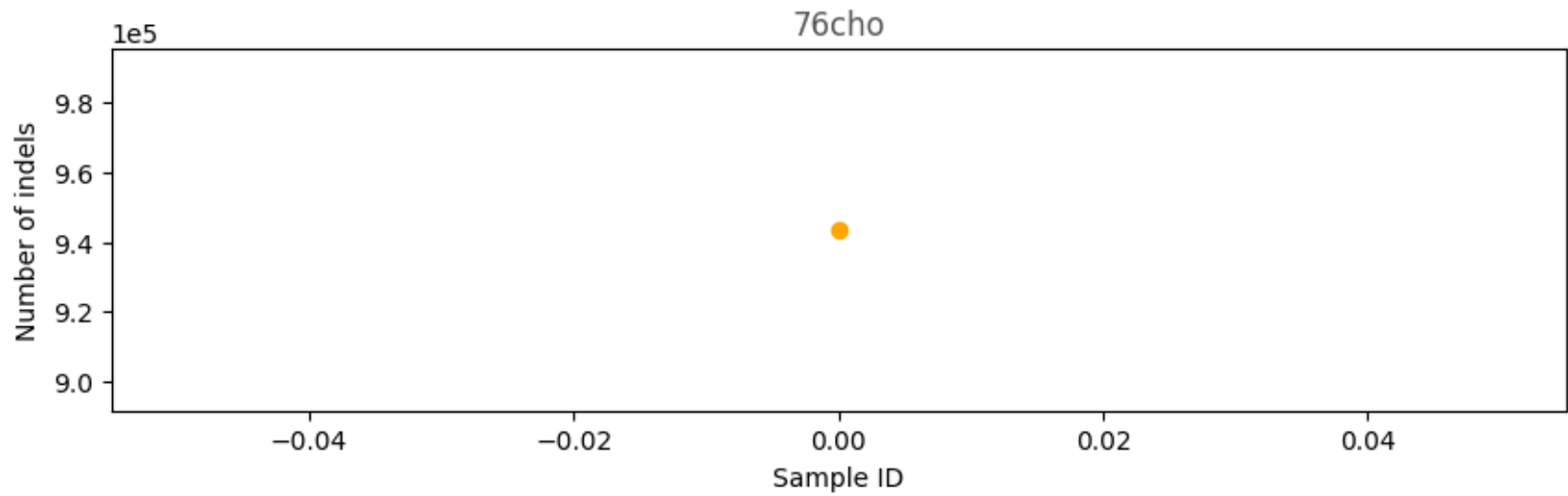
Average depth by sample



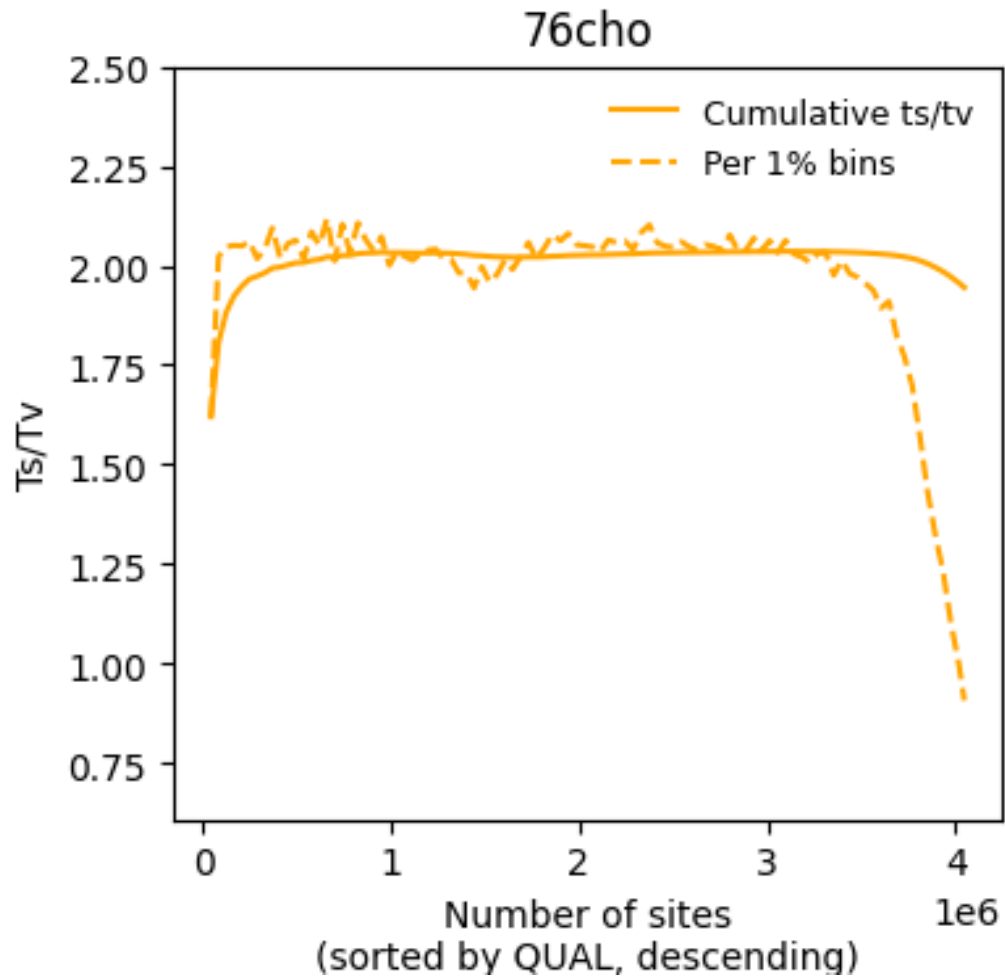
Number of SNPs by sample



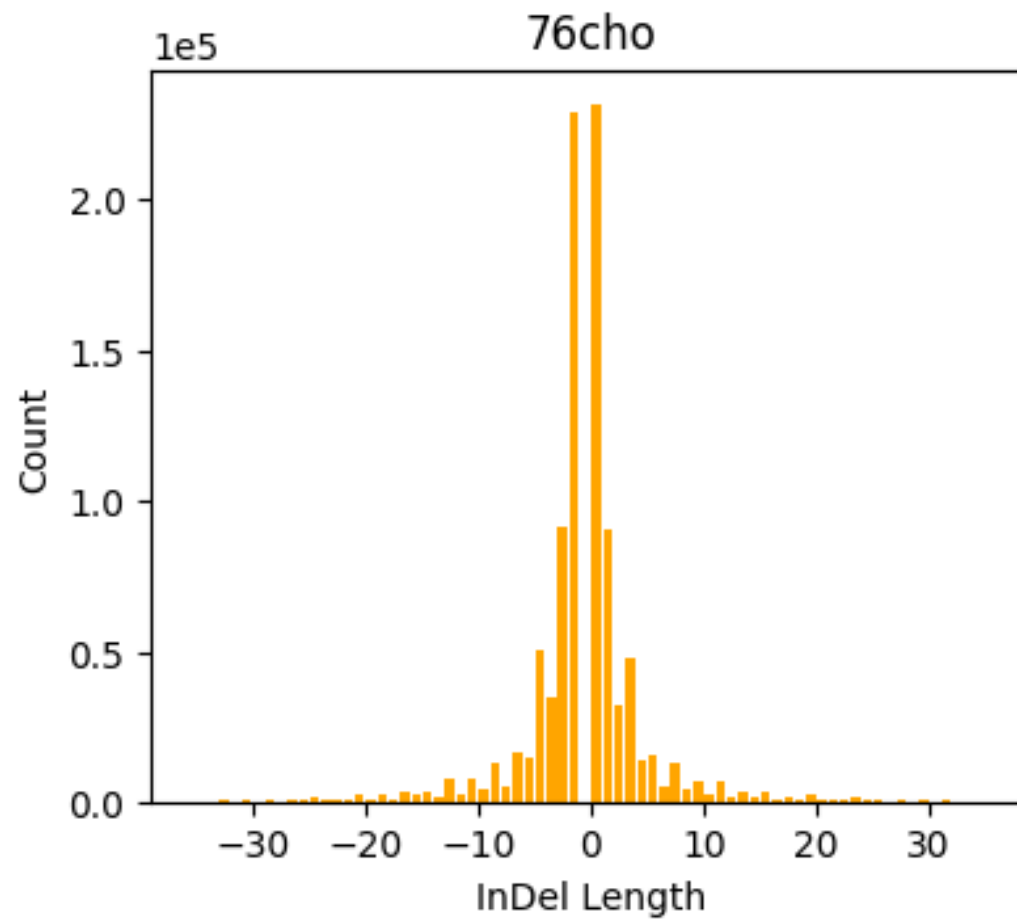
Number of indels by sample



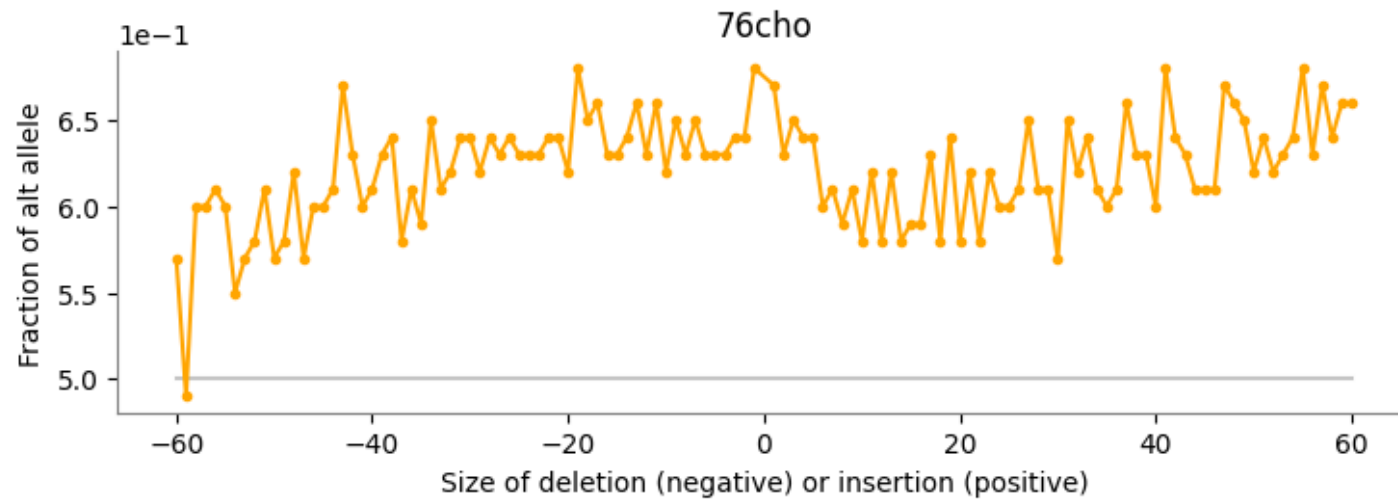
Ts/Tv stratified by QUAL



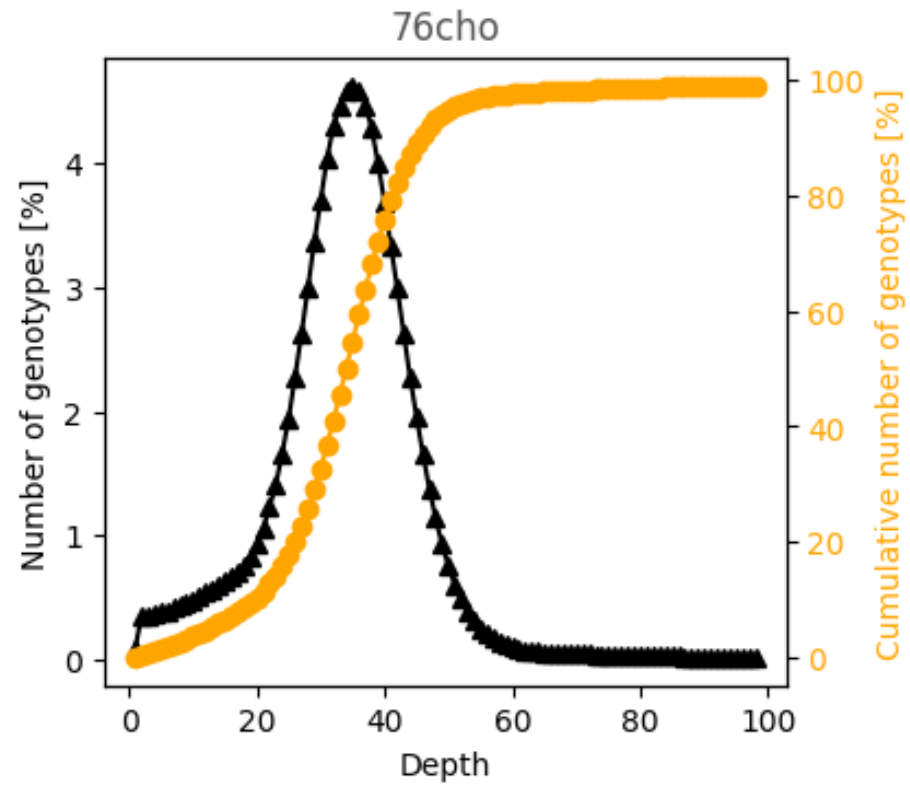
Indel distribution



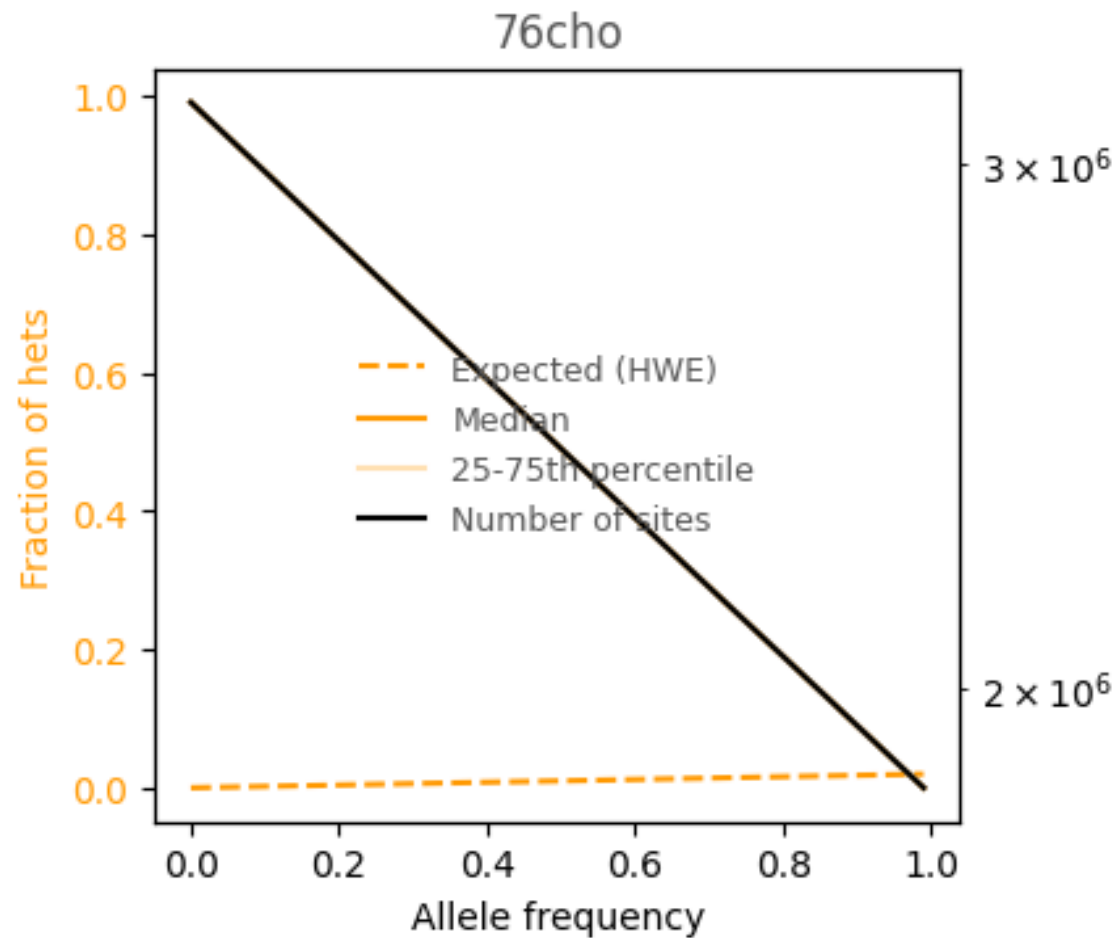
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

