

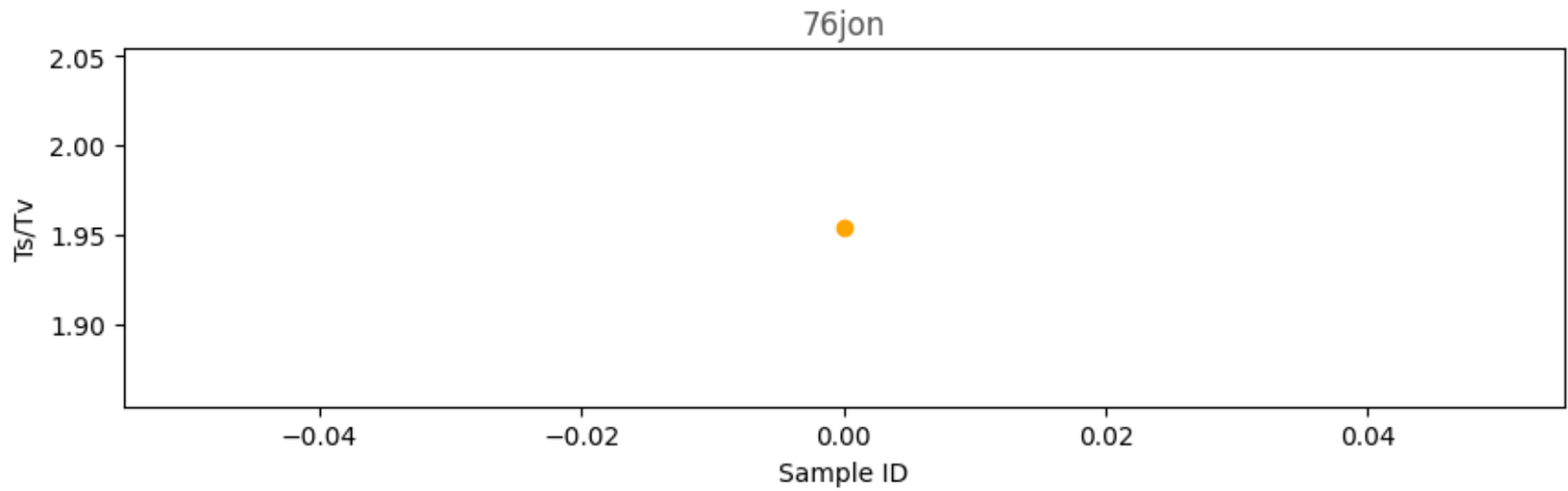
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76jon	4,035,788	1.95	1.96	946,783	–	0	0
* frameshift ratio: out/(out+in)							

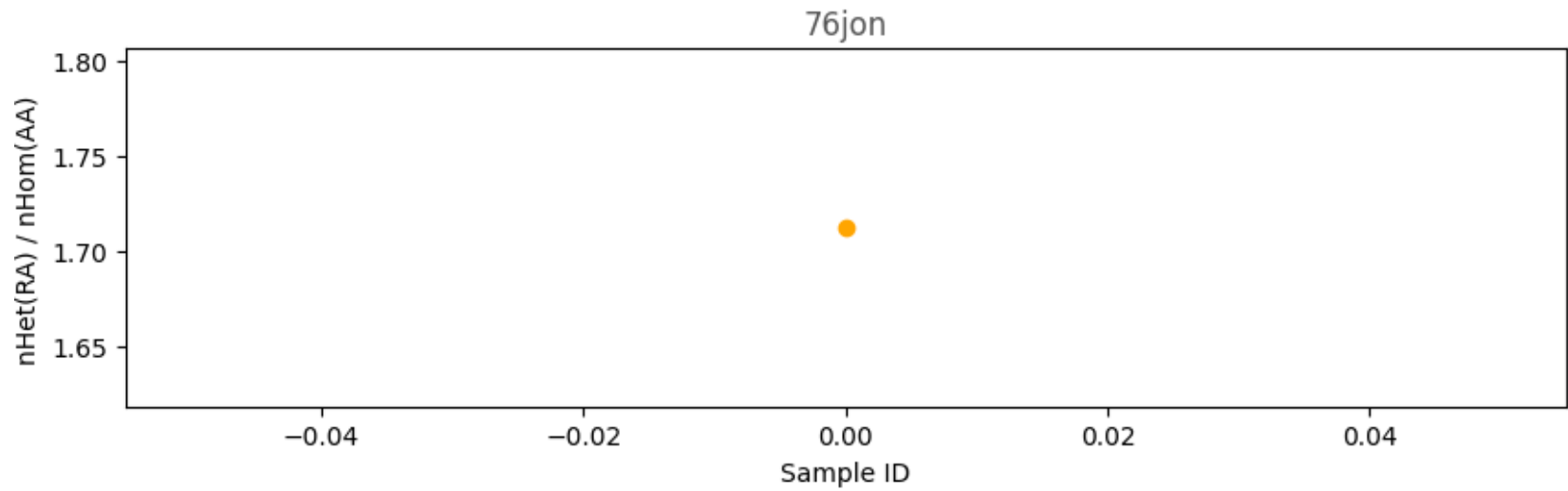
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76jon	63.1%	1.93	68.1%	98,772	1,821

- 76jon .. /ngc/projects2/gm/data/archive/2022/variants/snv/76jonthif-103817923885-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00476-211222_A01176_AHT7KCDSX2-EXT_LAB
 KA_NGCWGS-NGCWGS03573_21RKG030130x01_snv_germline_raw.haplotype_caller.vcf.gz

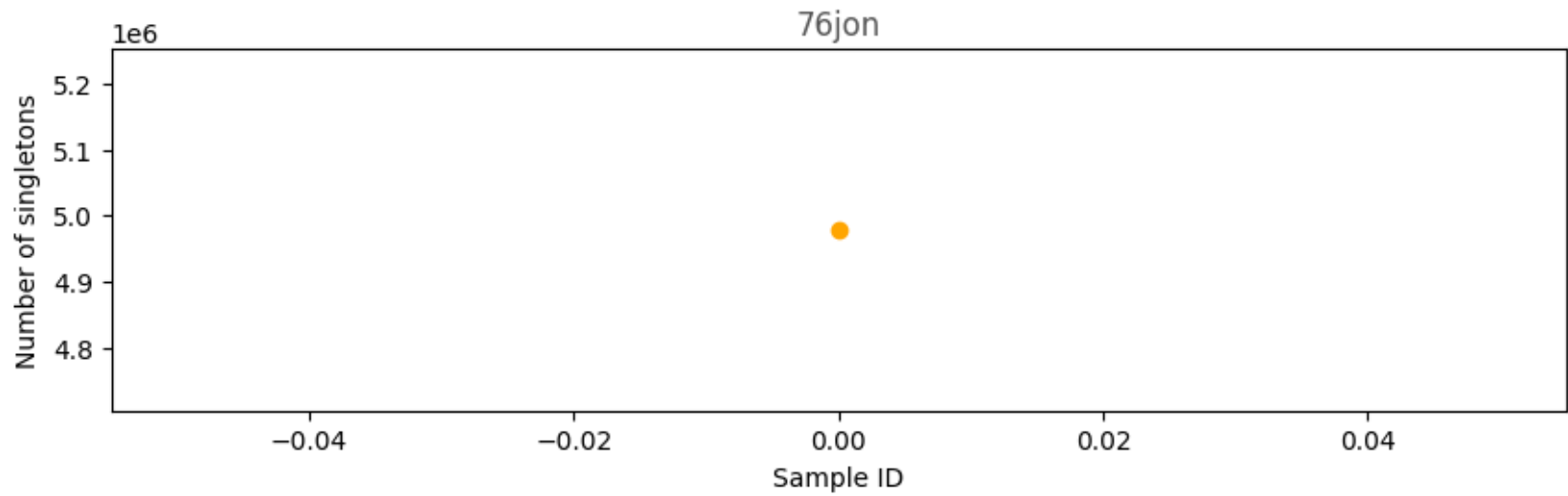
Ts/Tv by sample



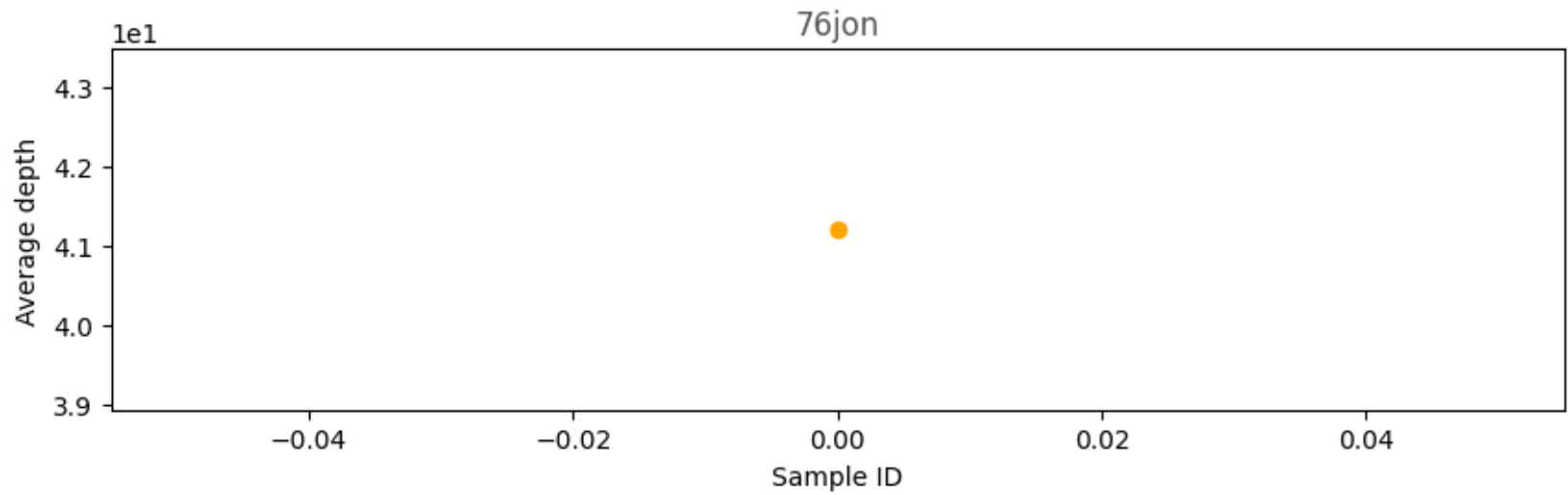
Hets vs non-ref Homs by sample



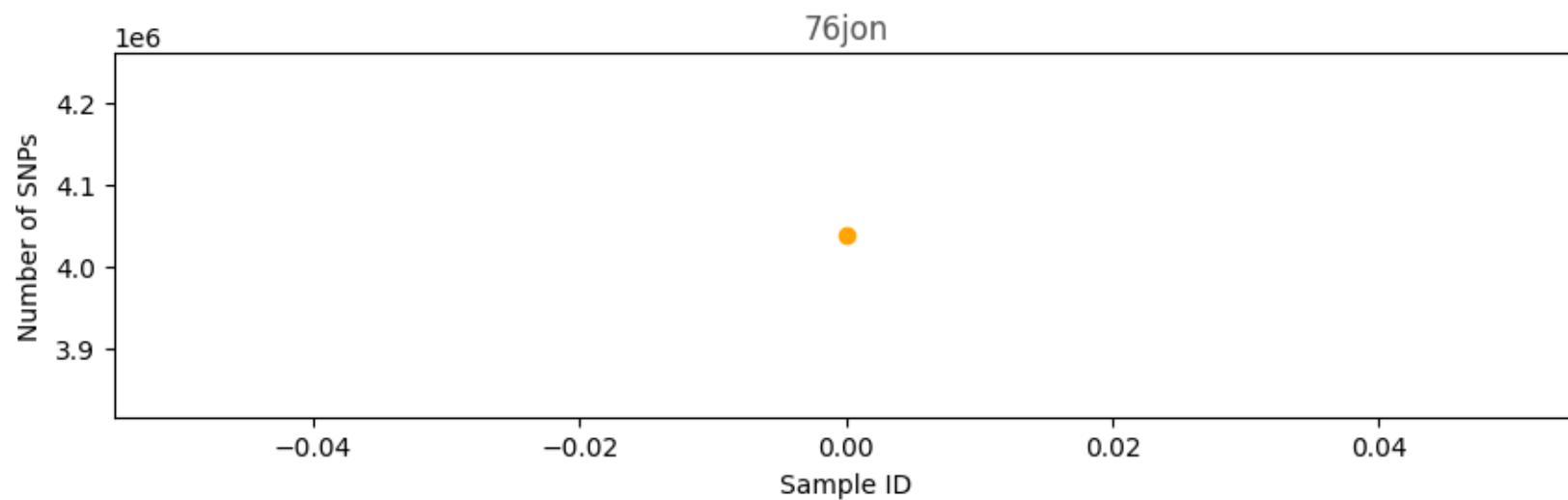
Singletons by sample (hets and homs)



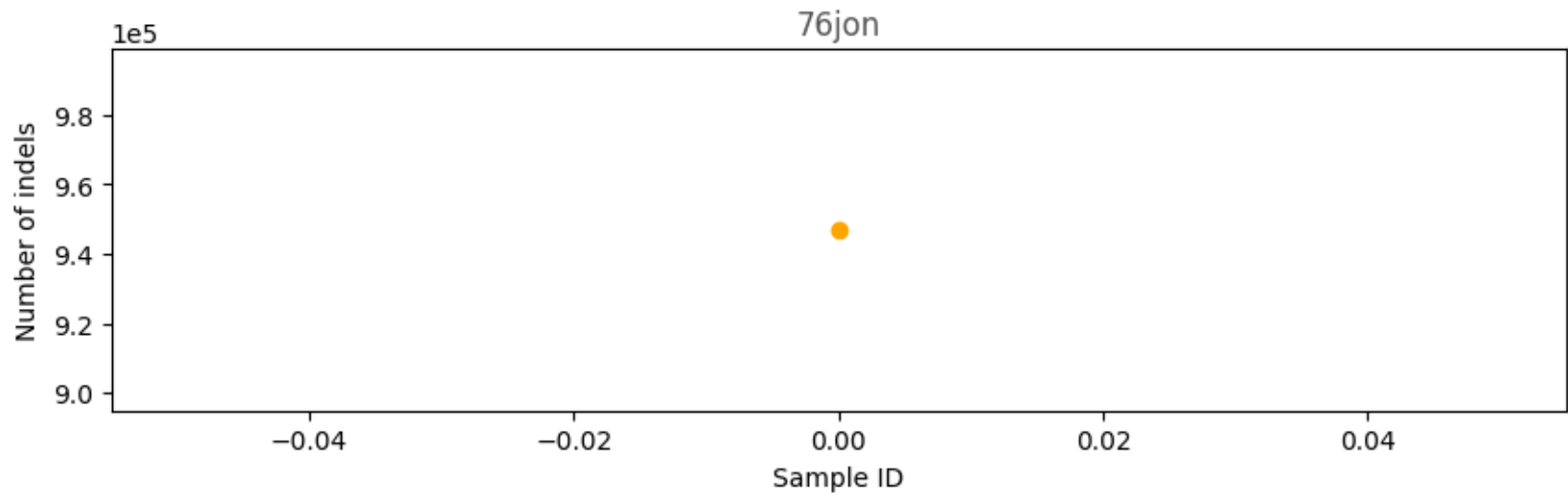
Average depth by sample



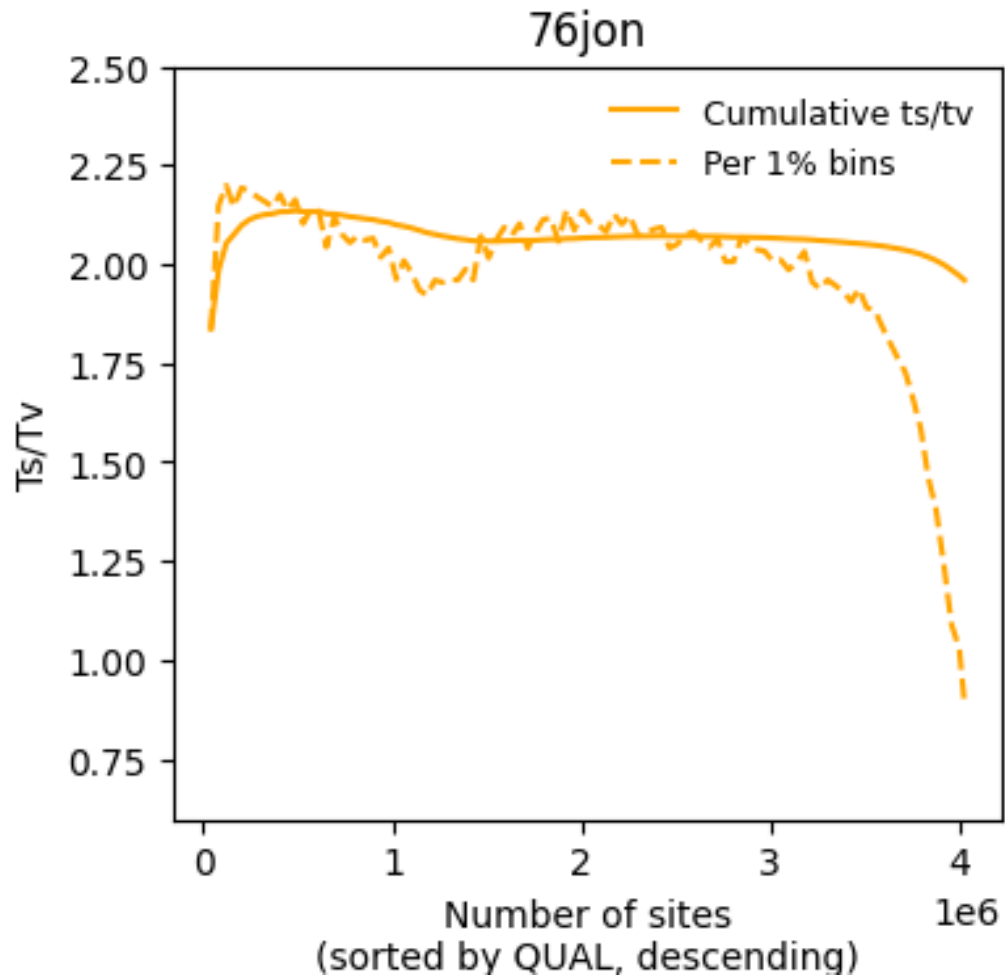
Number of SNPs by sample



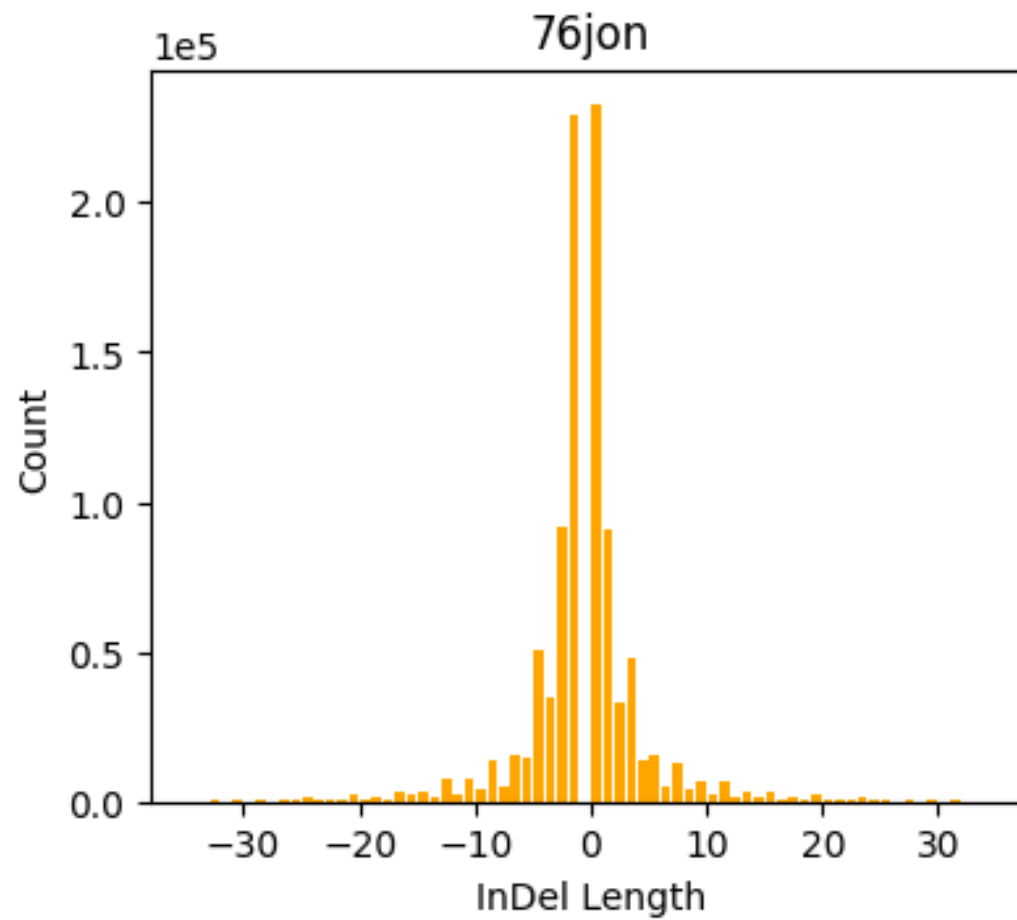
Number of indels by sample



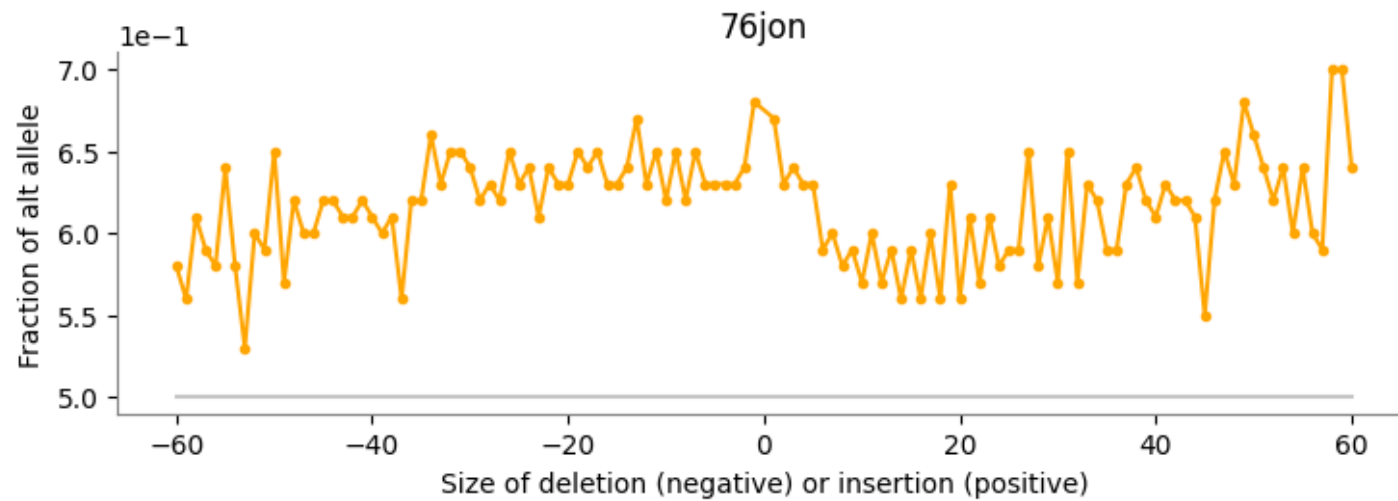
Ts/Tv stratified by QUAL



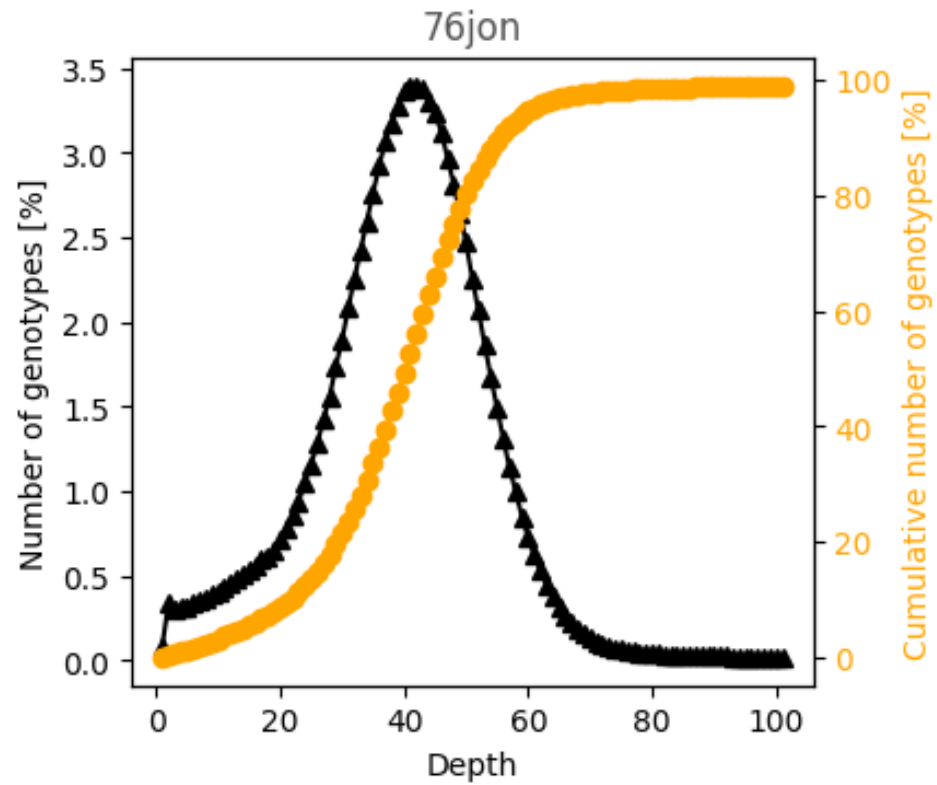
Indel distribution



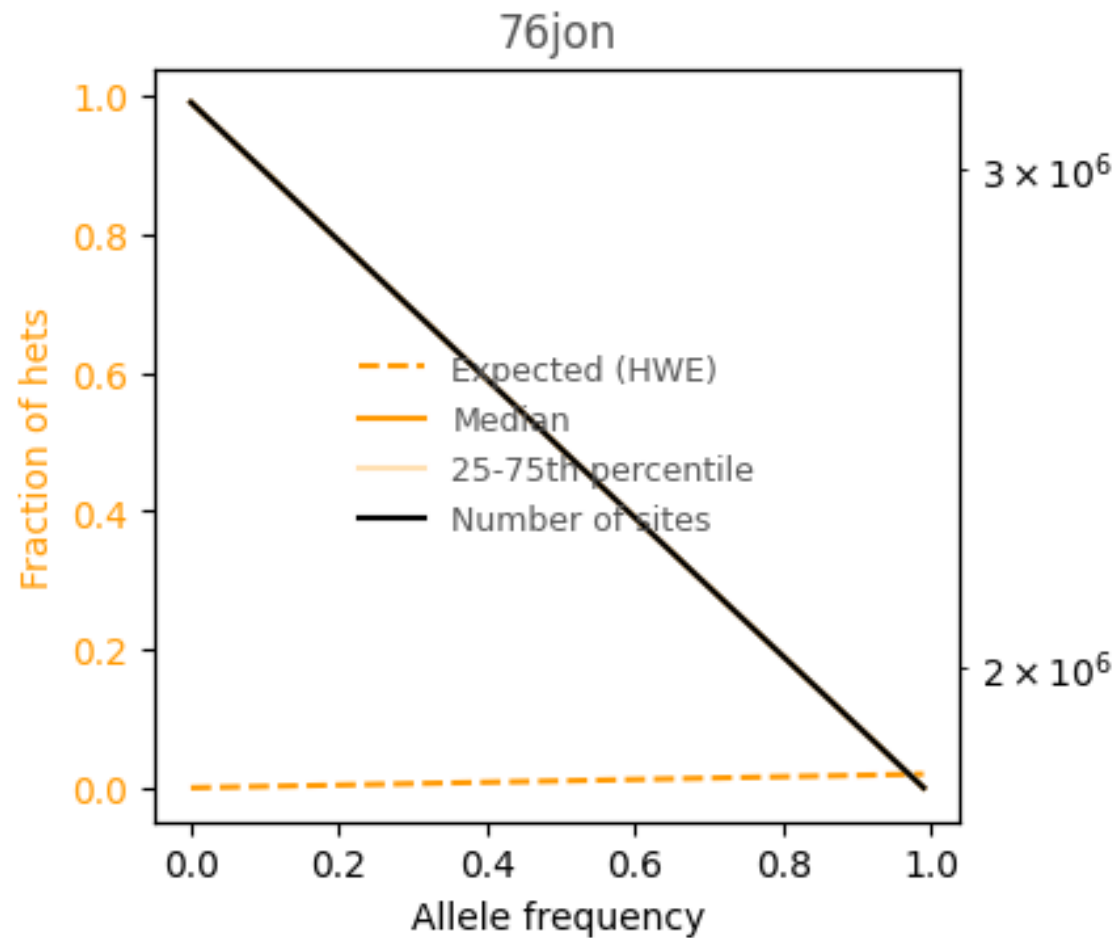
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

