

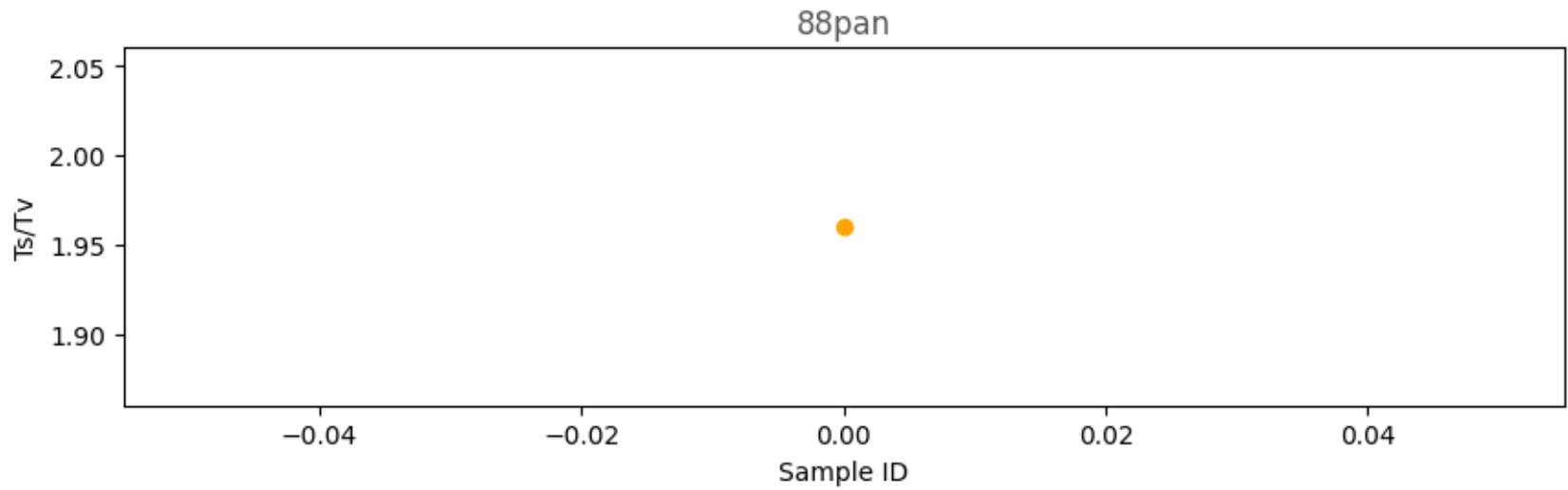
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
88pan	4,040,851	1.96	1.97	944,719	–	0	0
* frameshift ratio: out/(out+in)							

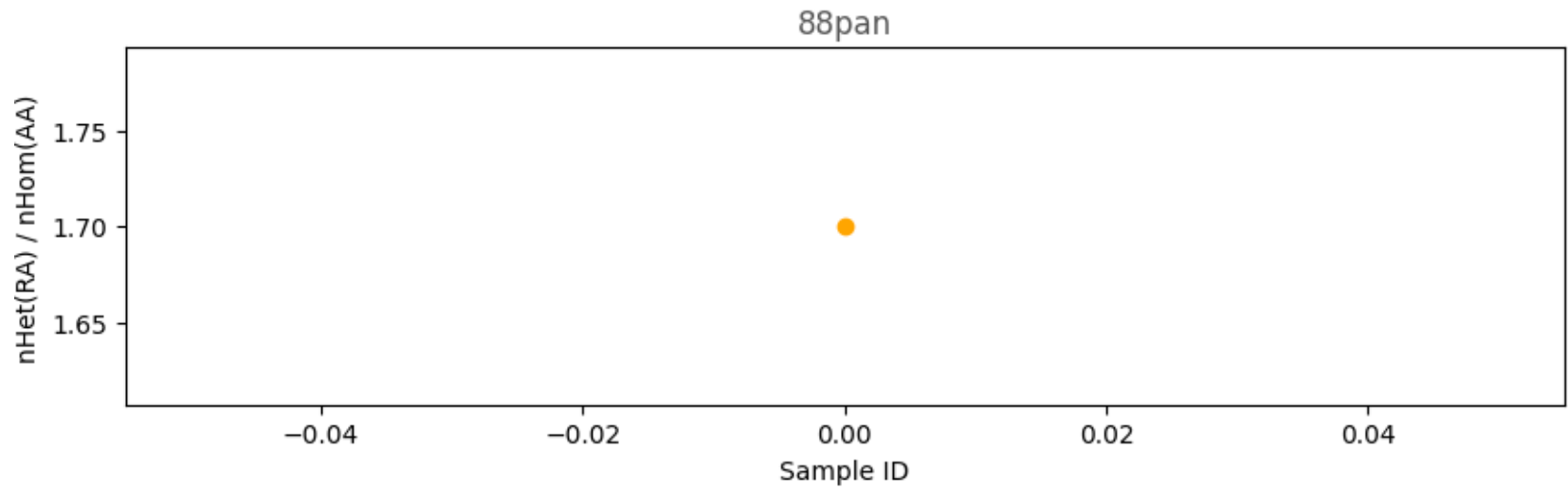
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
88pan	63.0%	1.94	67.8%	96,826	1,902

- 88pan .. /ngc/projects2/gm/data/archive/2022/variants/snv/88pandtcf-103776924290-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00290-210907_A01176_AHJVNLDSX2-EXT_LAB
 KA_NGCWGS-NGCWGS00184_21RKG020840x01_snv_germline_raw.haplotype_caller.vcf.gz

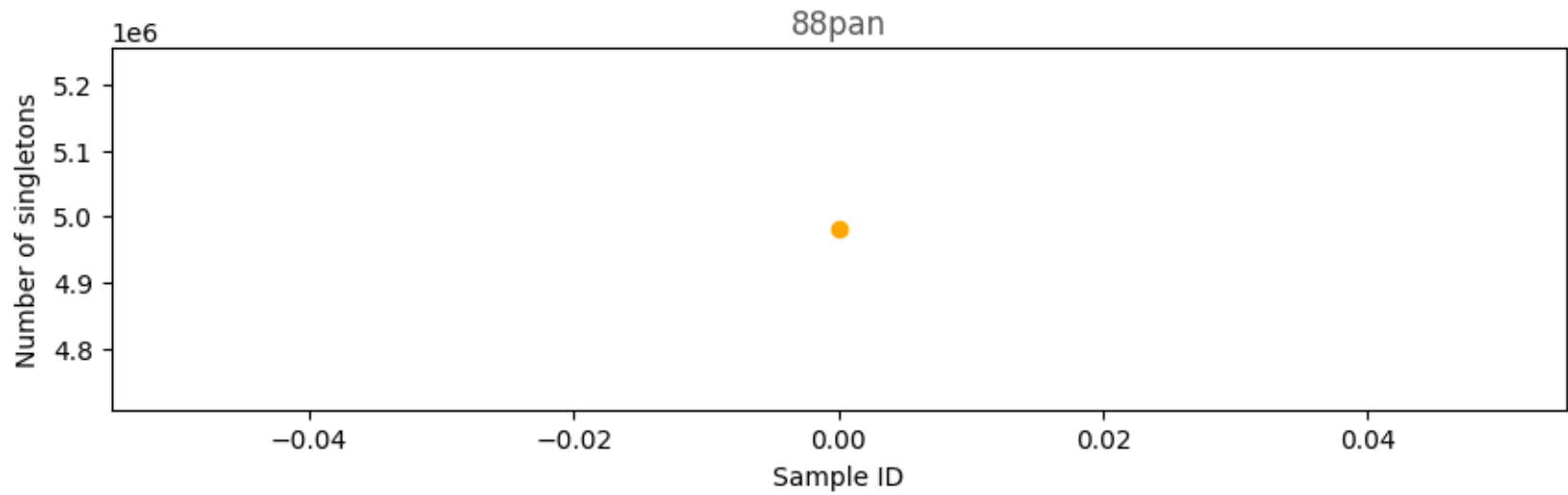
Ts/Tv by sample



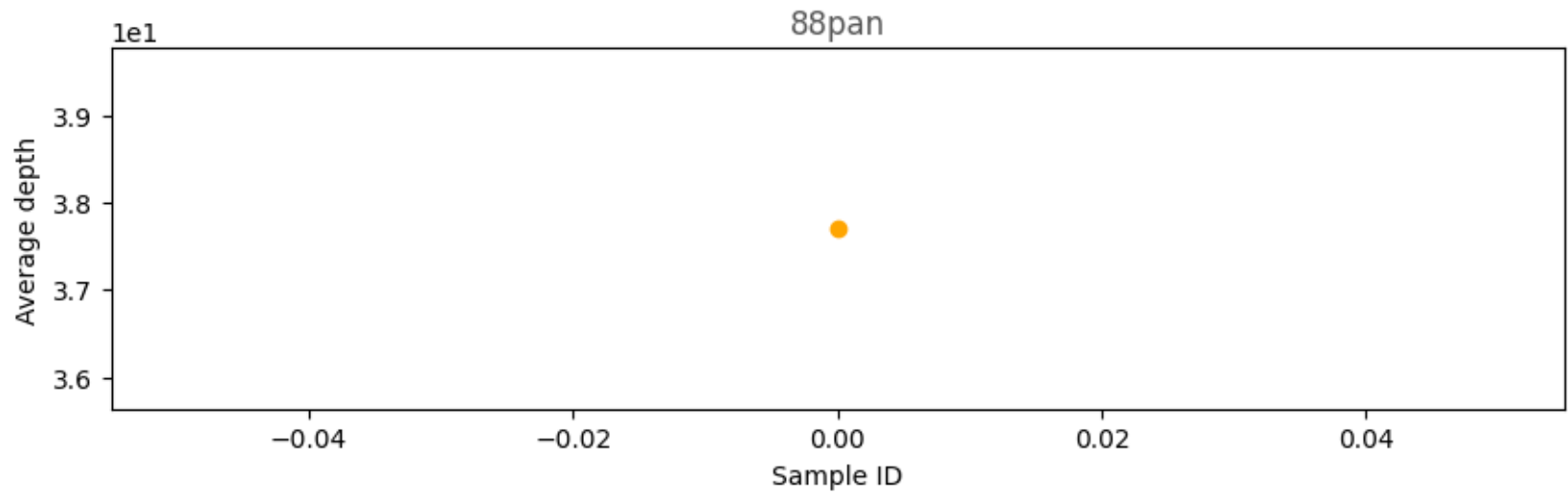
Hets vs non-ref Homs by sample



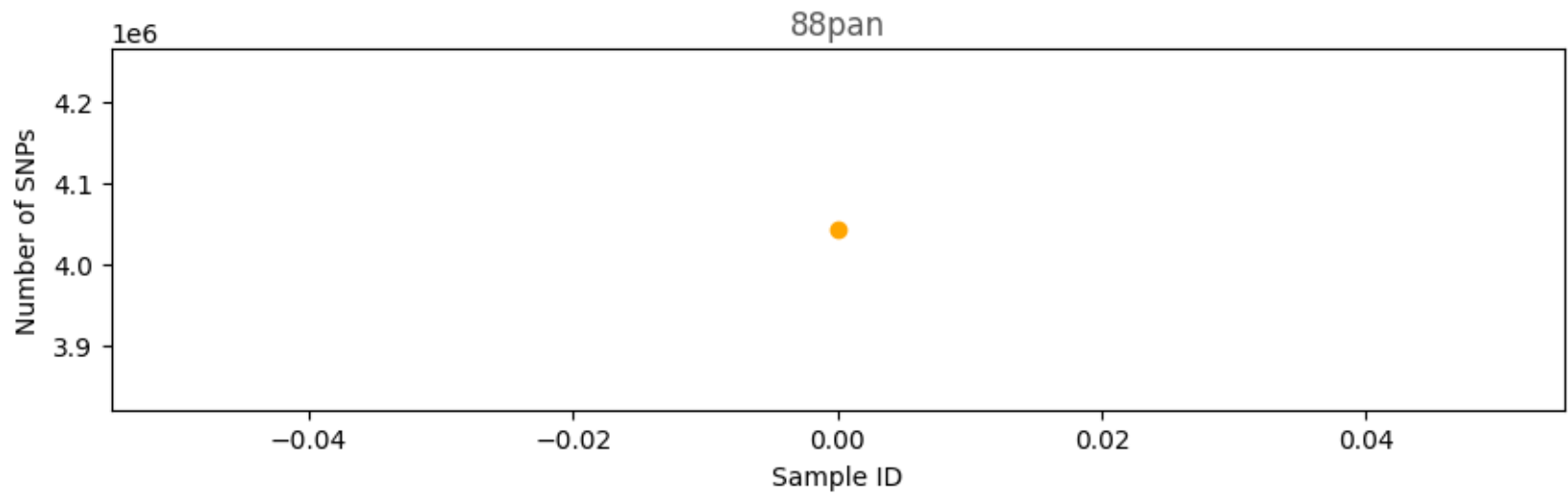
Singletons by sample (hets and homs)



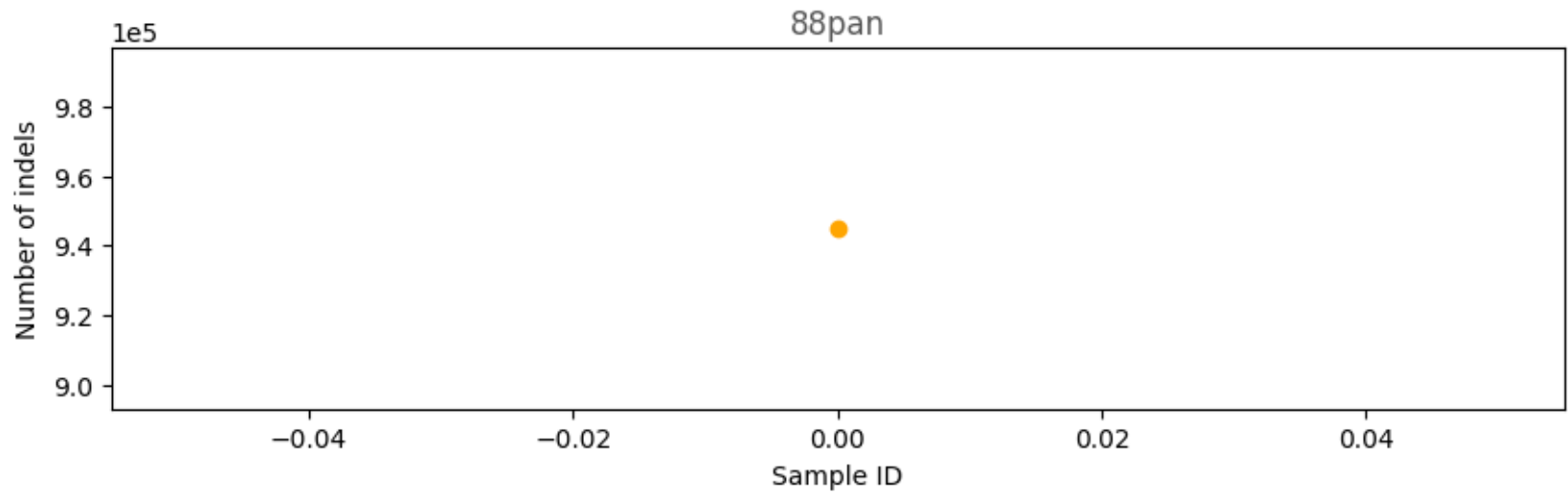
Average depth by sample



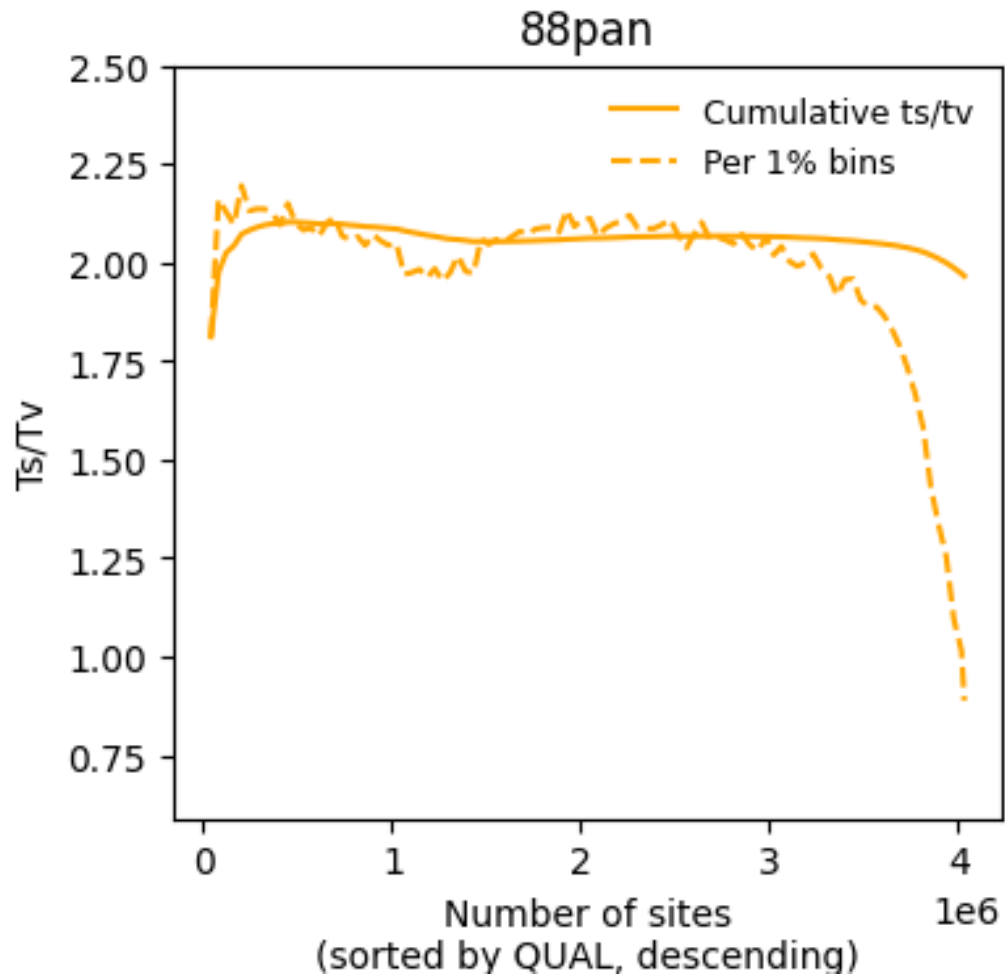
Number of SNPs by sample



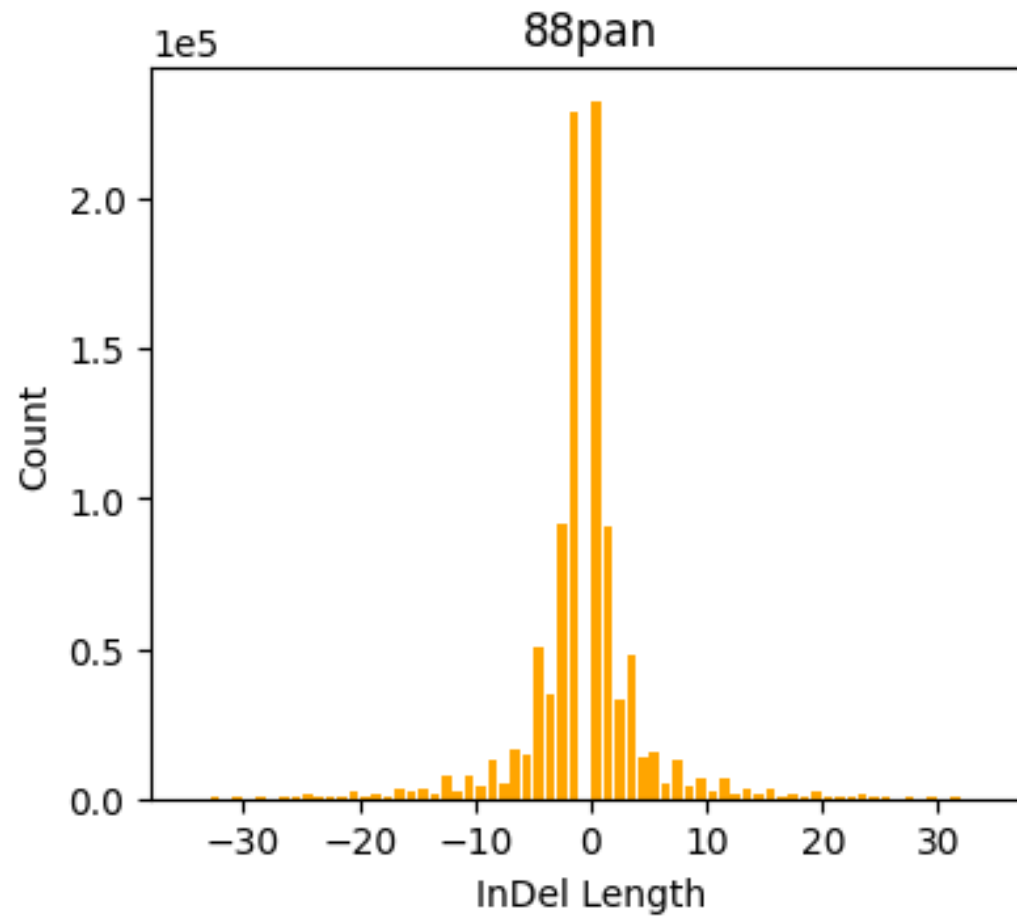
Number of indels by sample



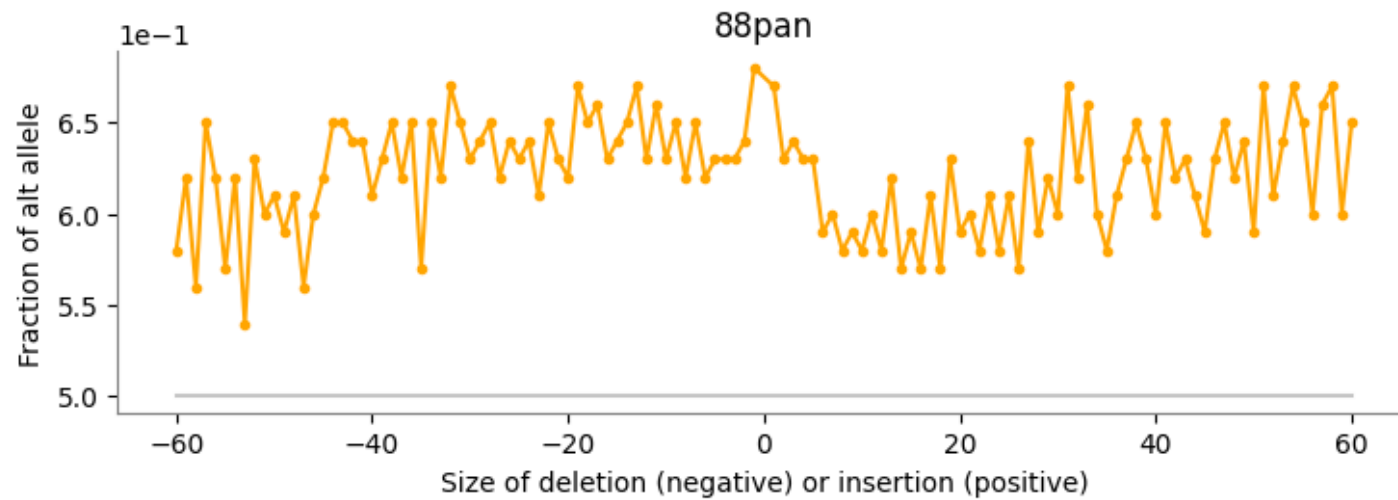
Ts/Tv stratified by QUAL



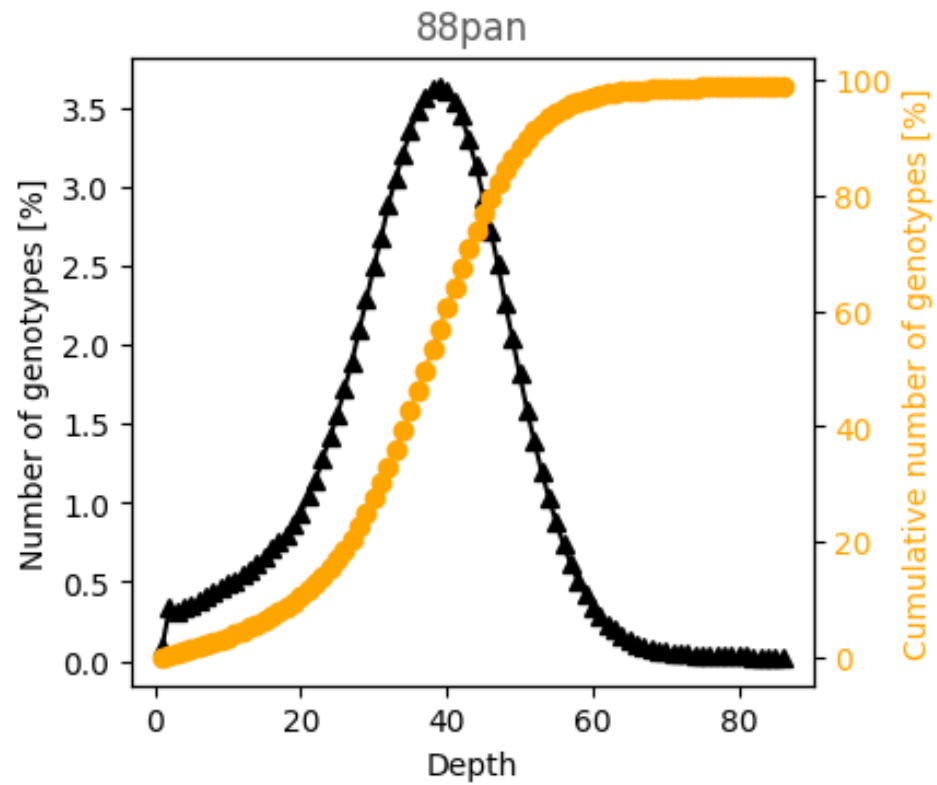
Indel distribution



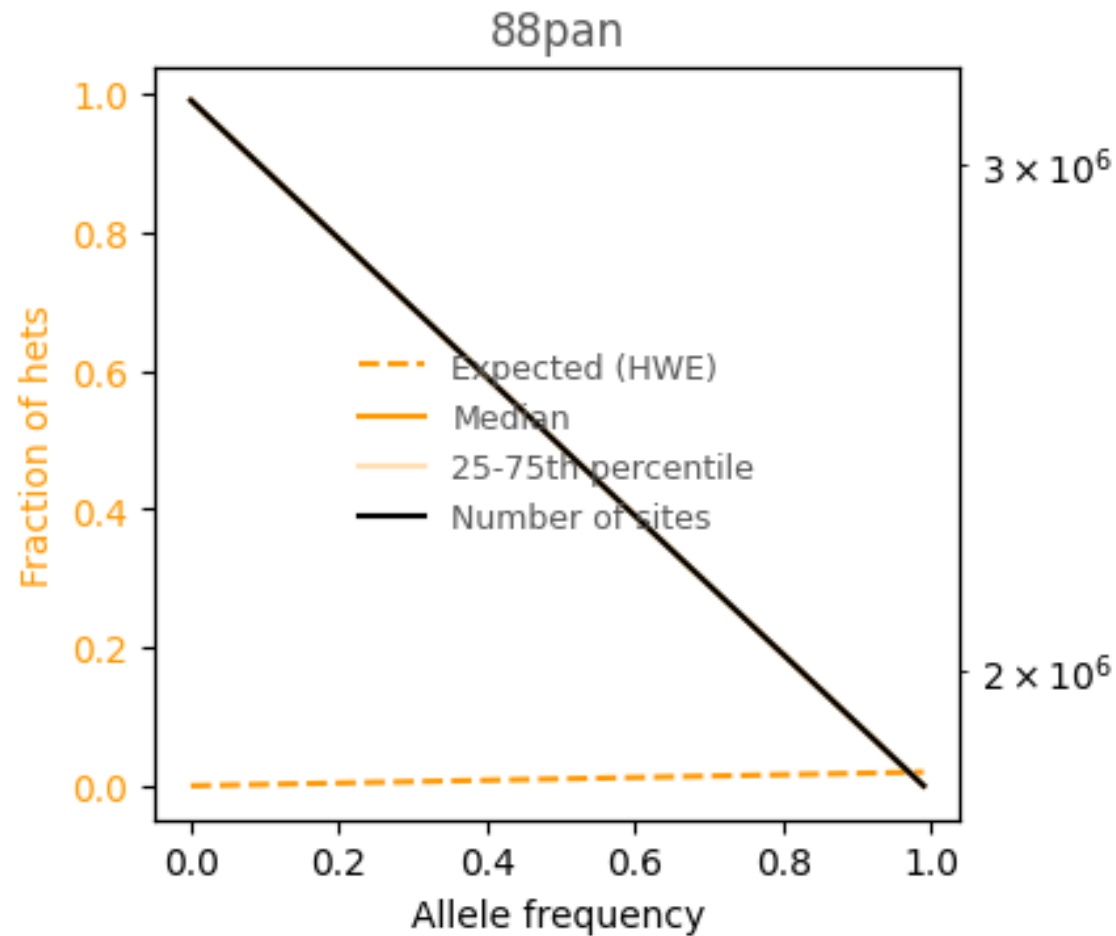
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

