

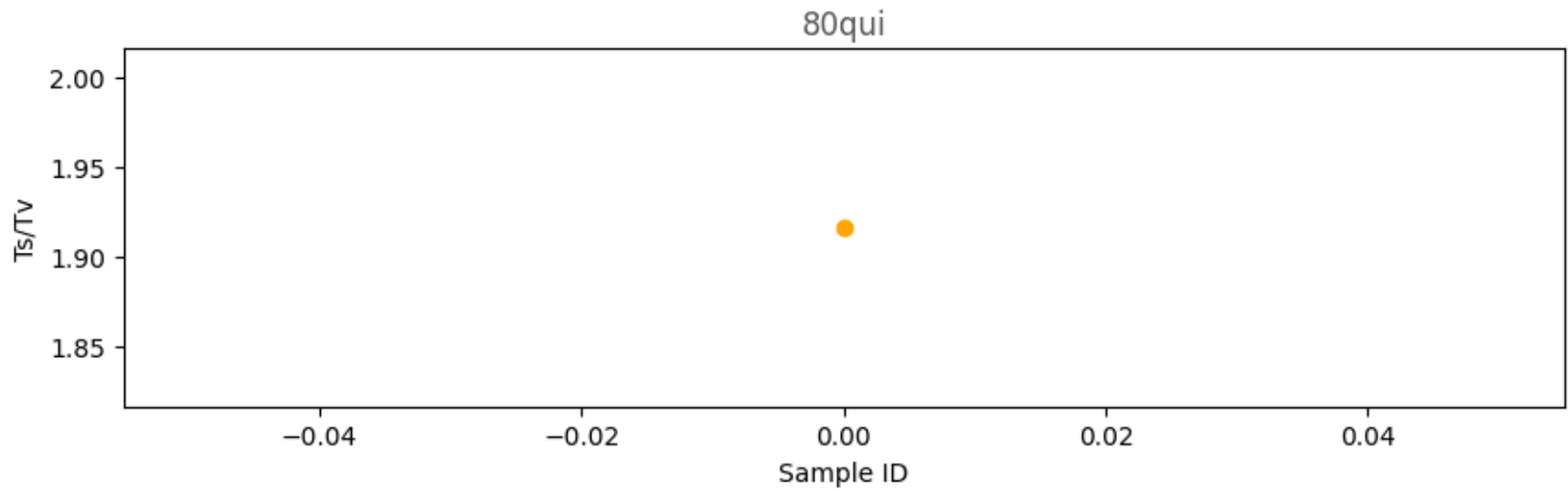
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
80qui	4,133,702	1.92	1.92	966,648	–	0	0
* frameshift ratio: out/(out+in)							

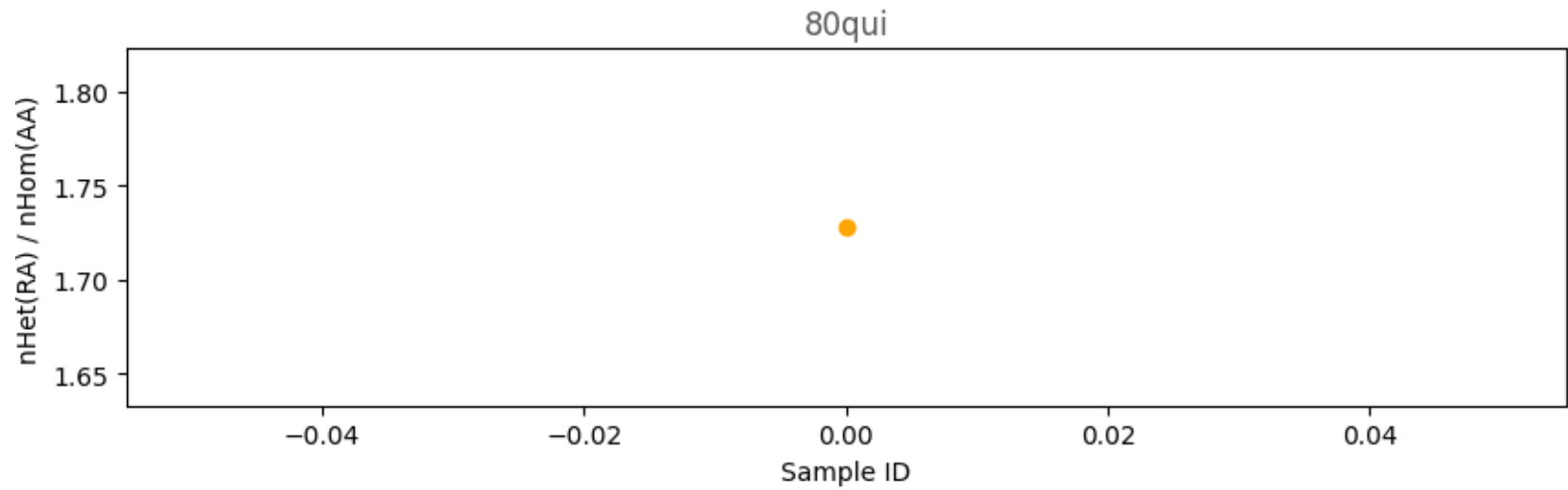
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
80qui	63.4%	1.88	69.0%	108,797	2,423

- 80qui .. /ngc/projects2/gm/data/archive/2022/variants/snv/80quindnf-103745695745-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00167-210608_A01176_AHCYWGDSX2-EXT_LAB
 KA_NGCWGS-NGCWGS00082-808vdpv56_snv_germline_raw.haplotype_caller.vcf.gz

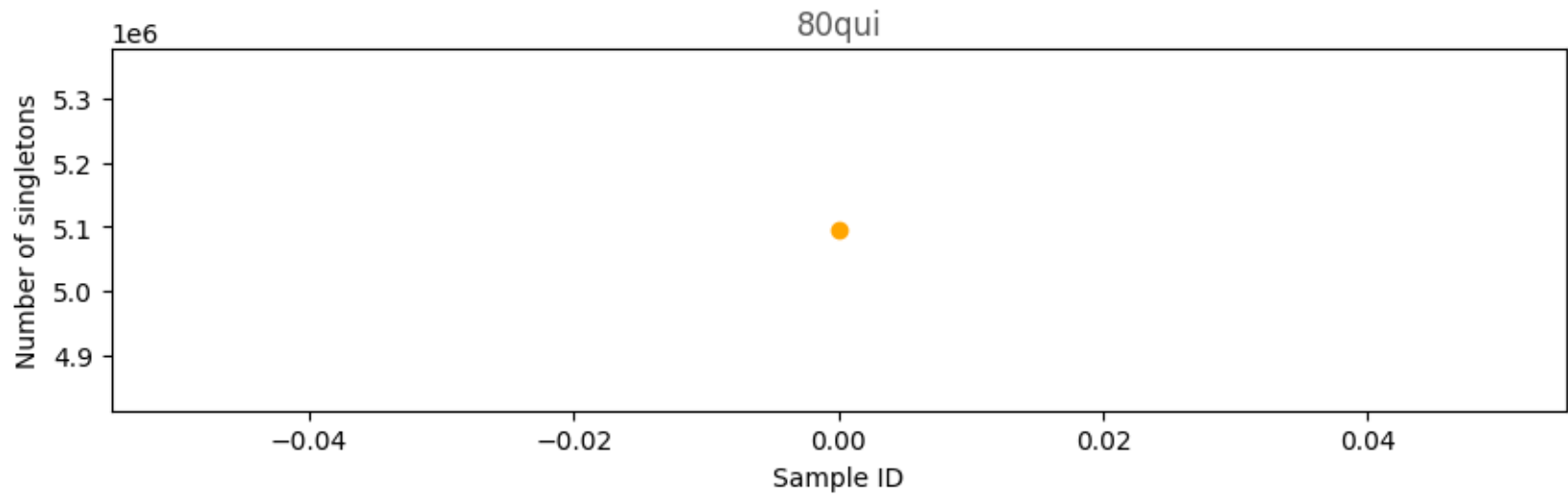
Ts/Tv by sample



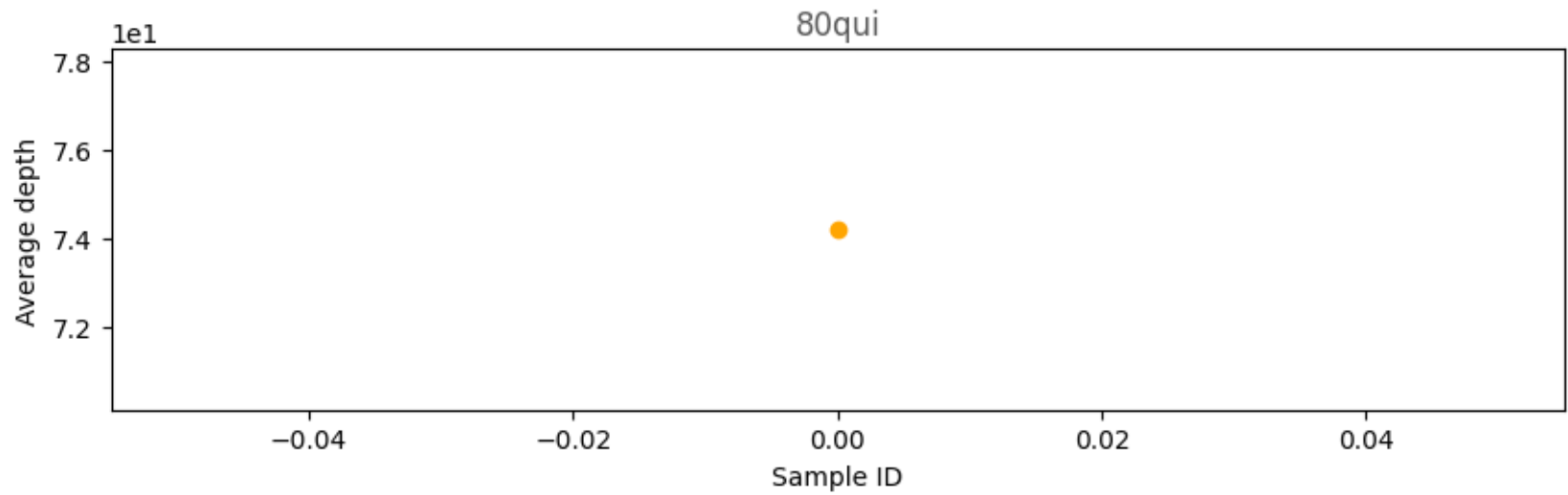
Hets vs non-ref Homs by sample



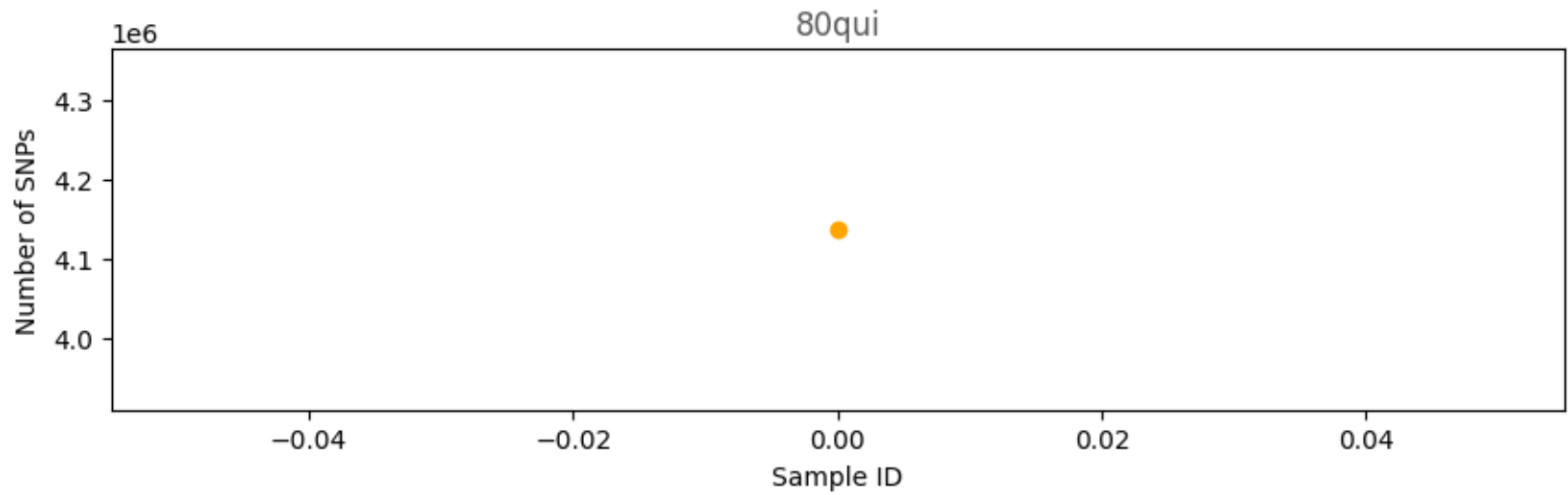
Singletons by sample (hets and homs)



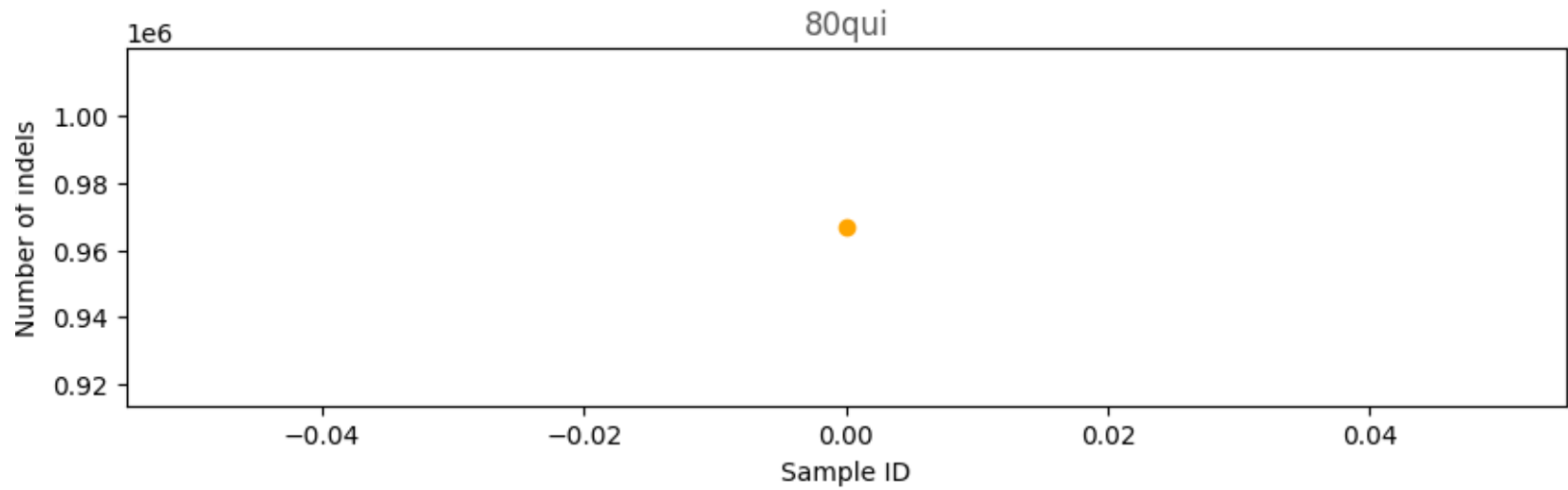
Average depth by sample



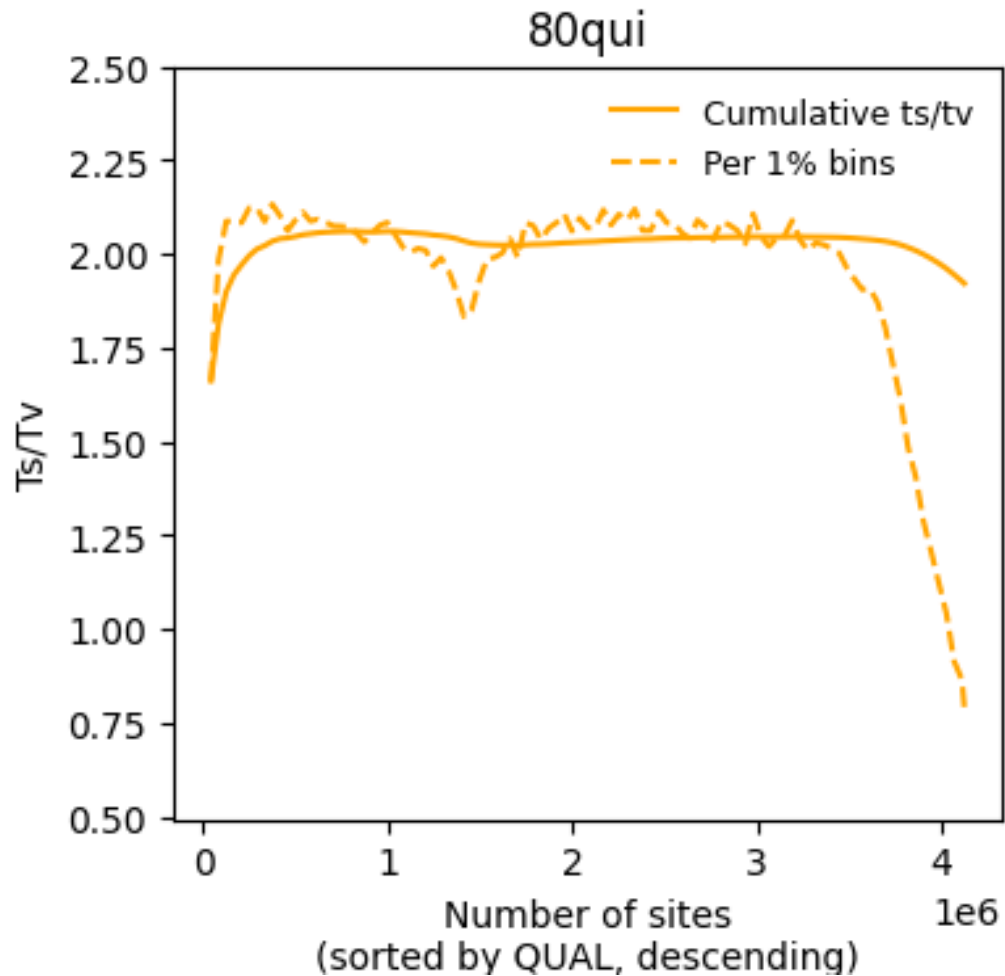
Number of SNPs by sample



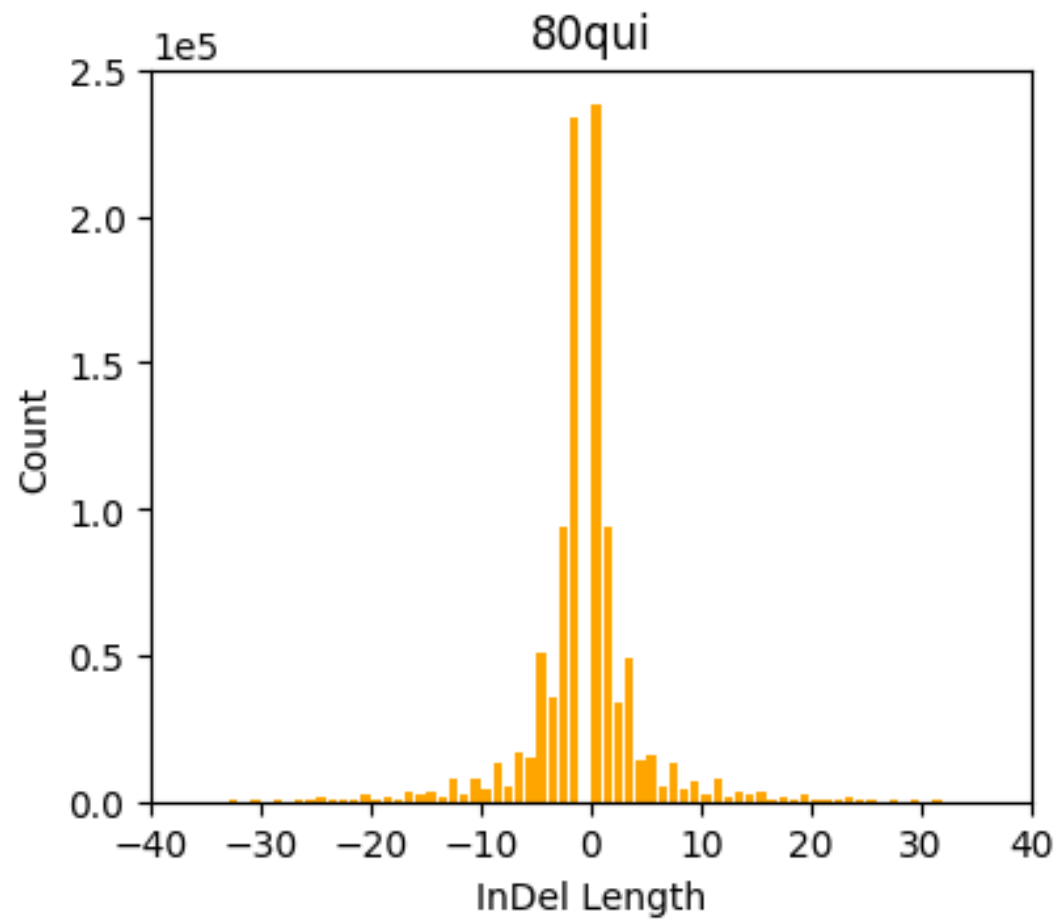
Number of indels by sample



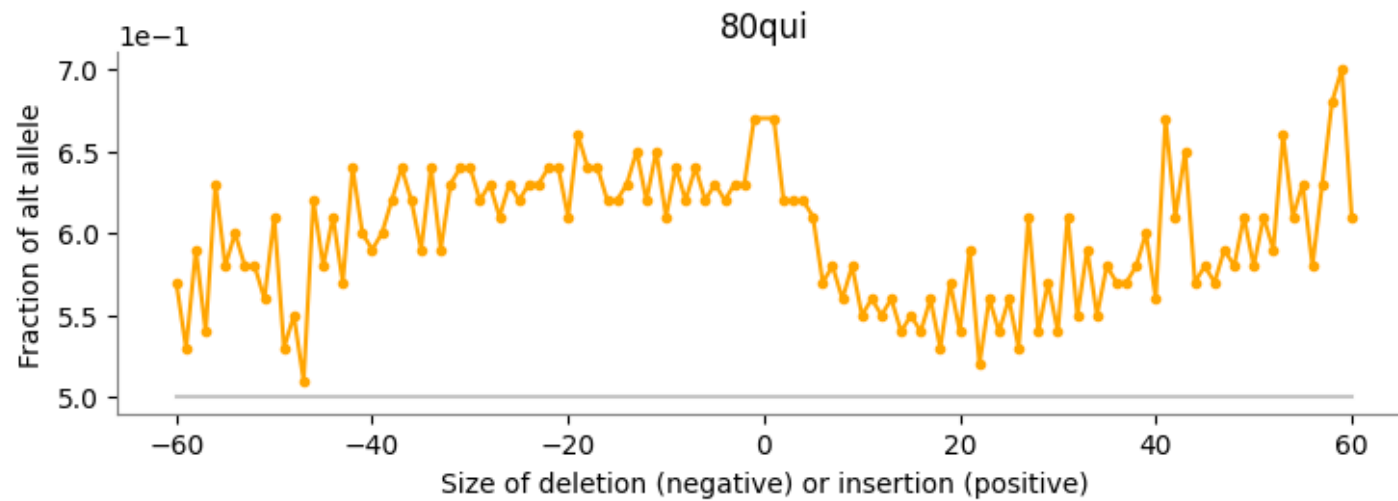
Ts/Tv stratified by QUAL



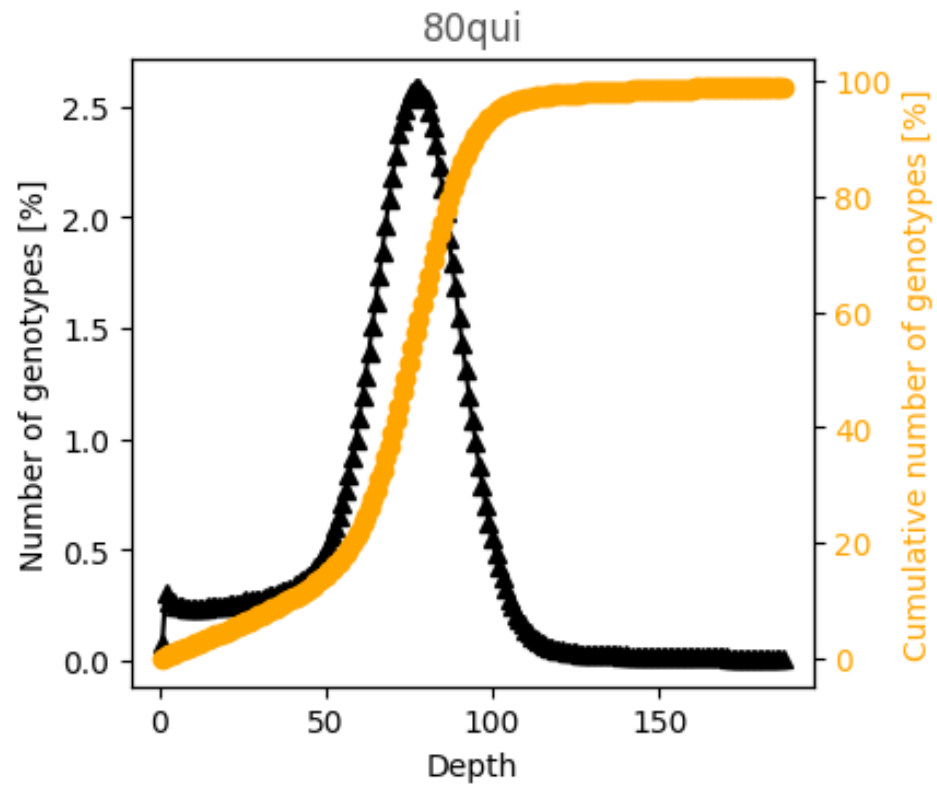
Indel distribution



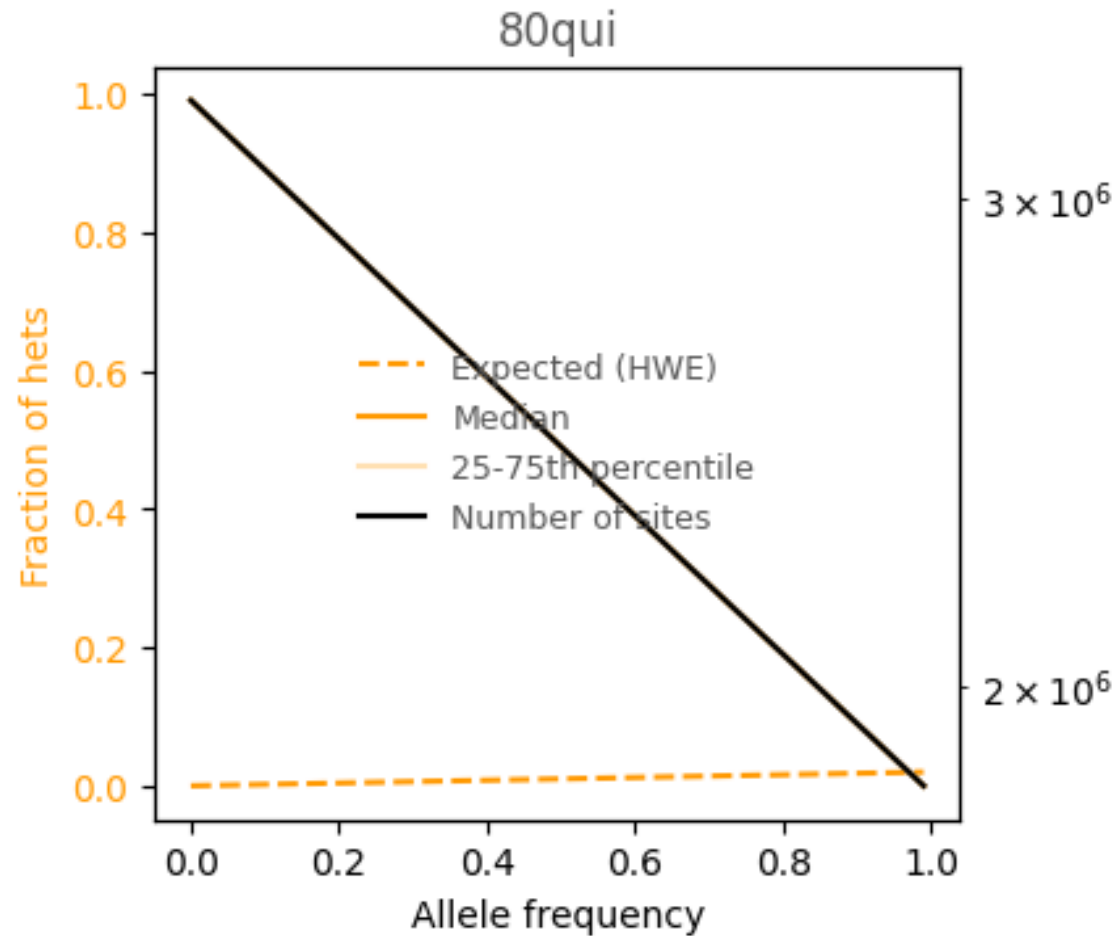
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

