

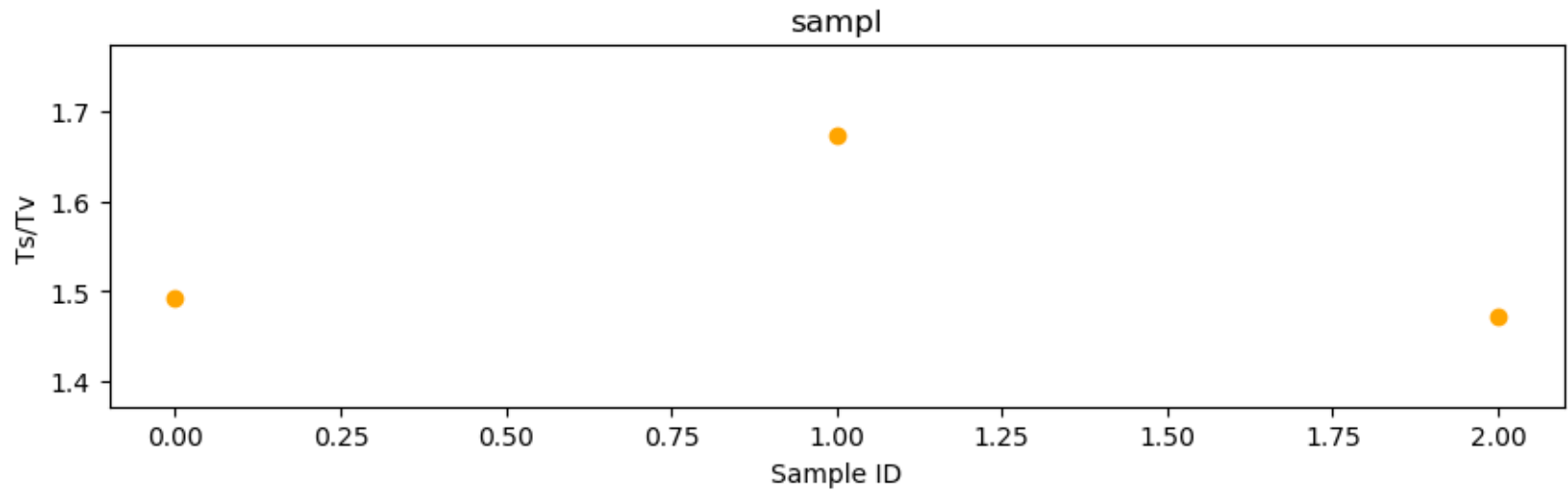
Summary Numbers

	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
sampl	471	1.53	1.53	36	–	0	0
* frameshift ratio: out/(out+in)							

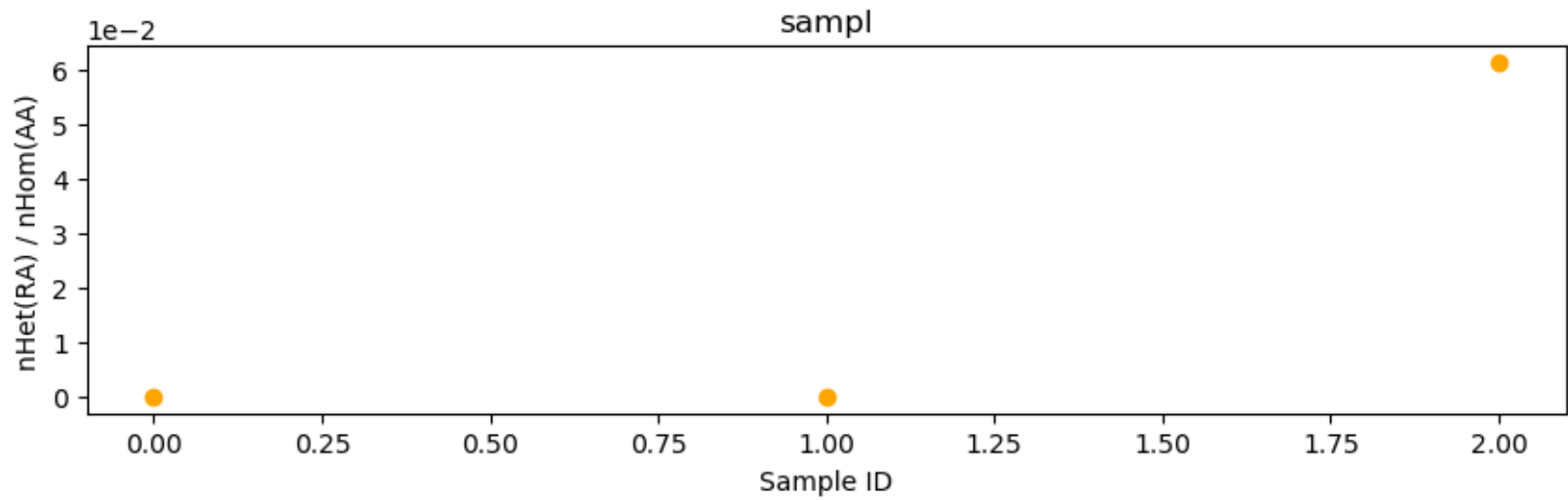
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
sampl	2.1%	2.33	0.0%	0	0

- `sampl .. samples.vcf.gz`

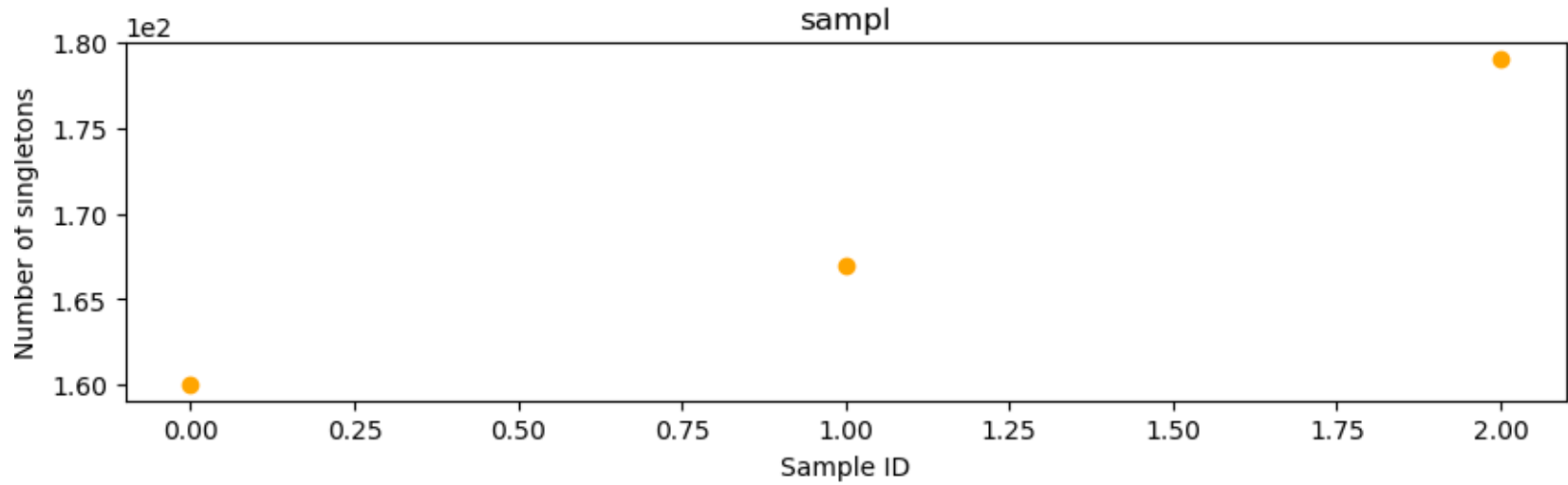
Ts/Tv by sample



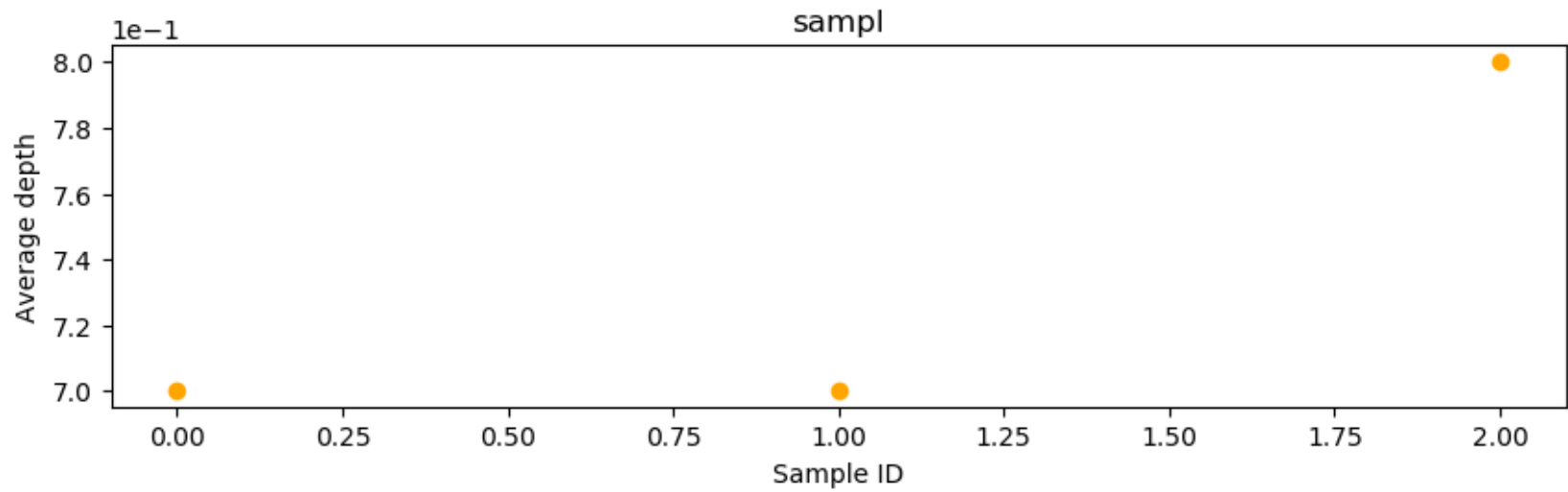
Hets vs non-ref Homs by sample



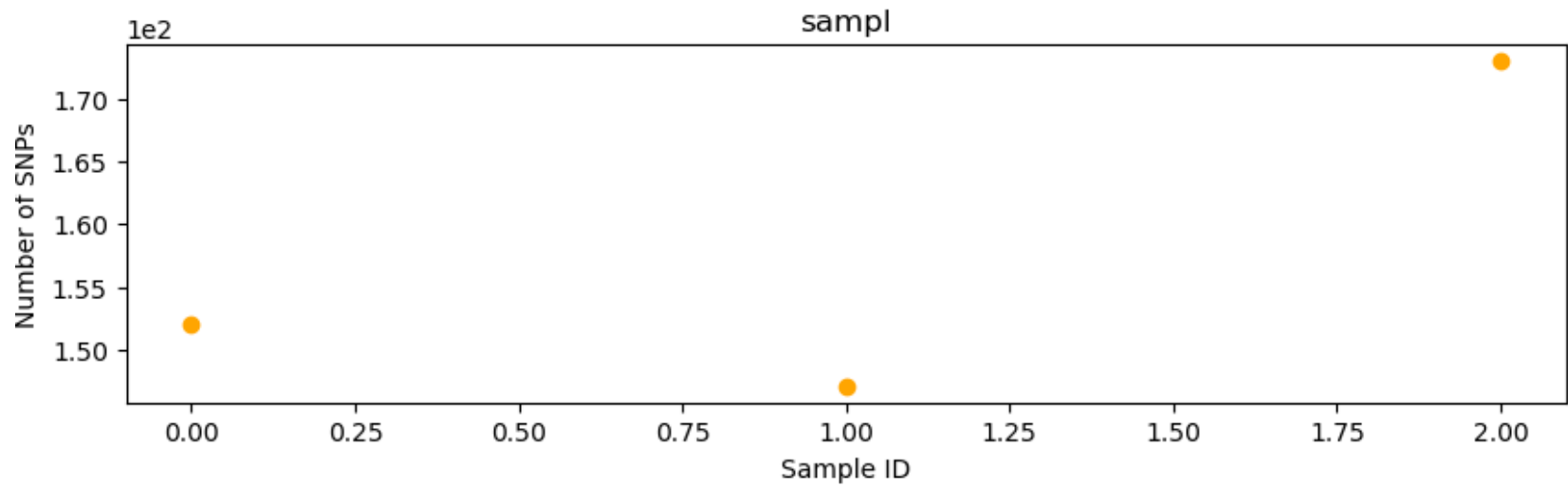
Singletons by sample (hets and homs)



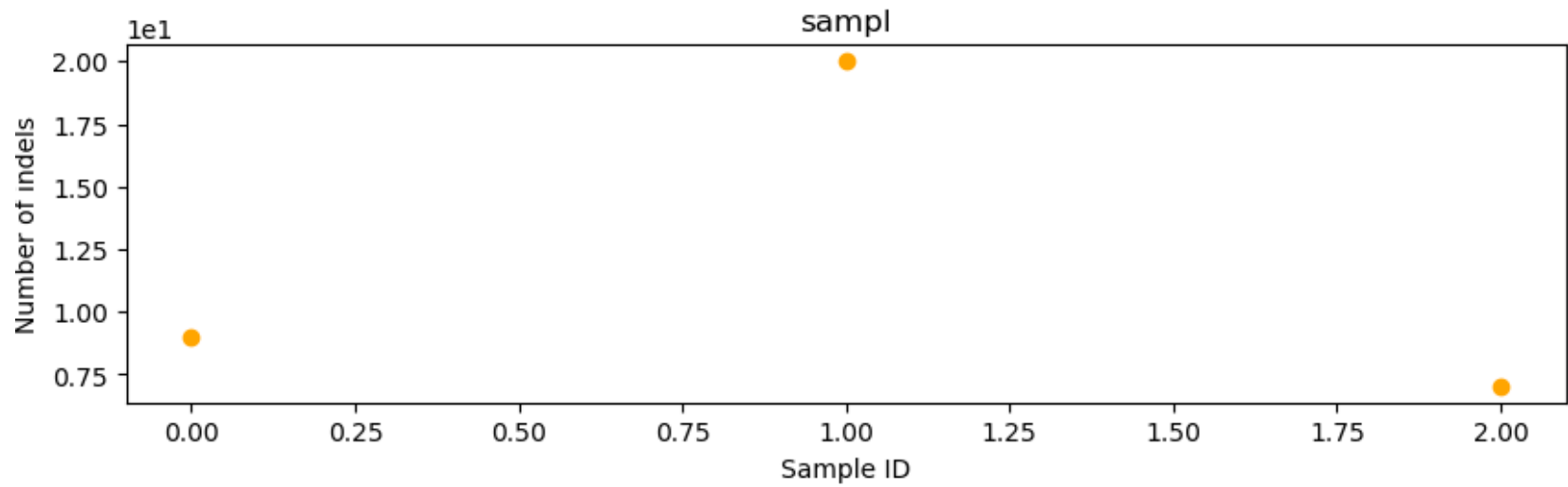
Average depth by sample



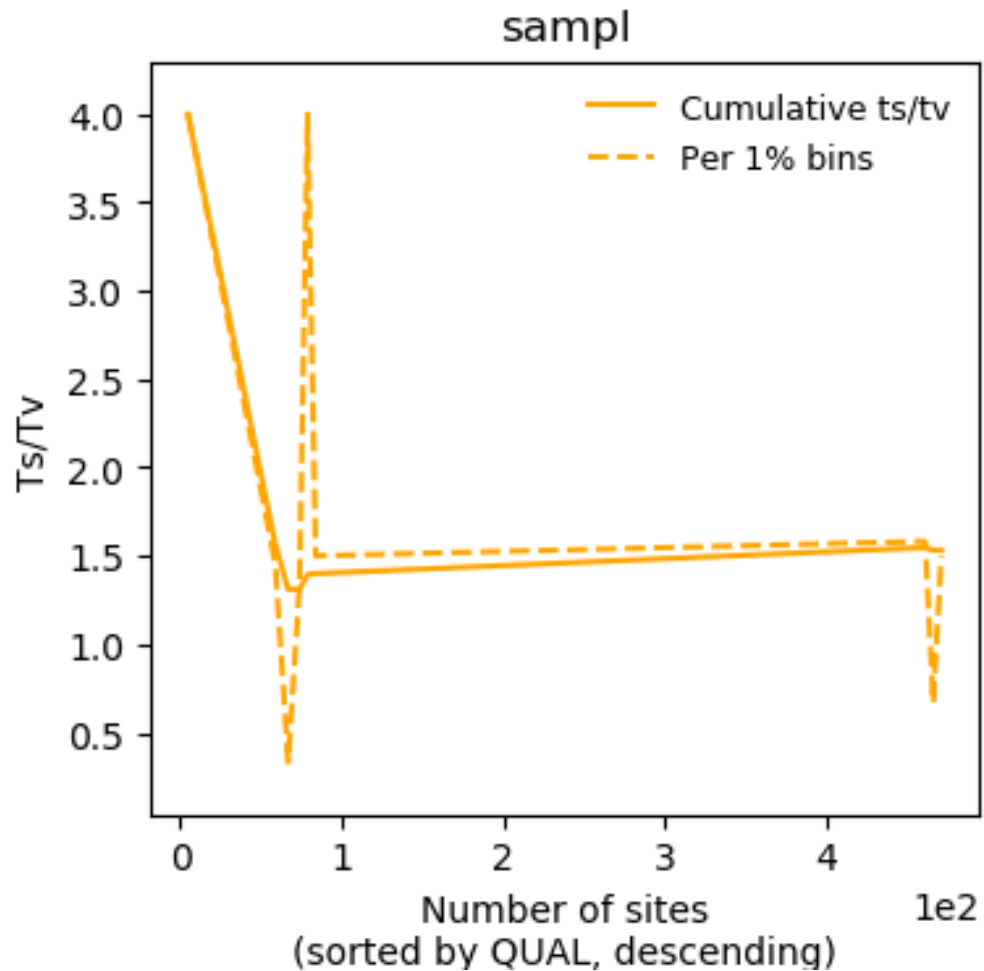
Number of SNPs by sample



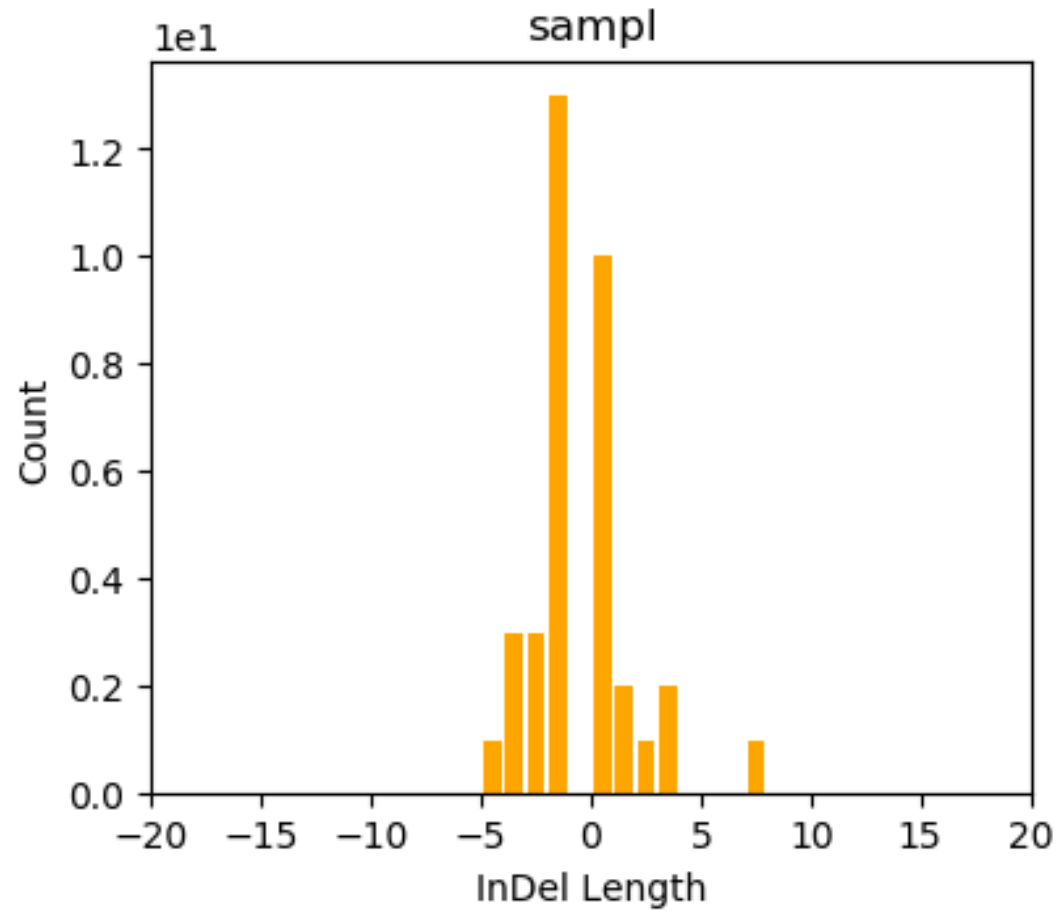
Number of indels by sample



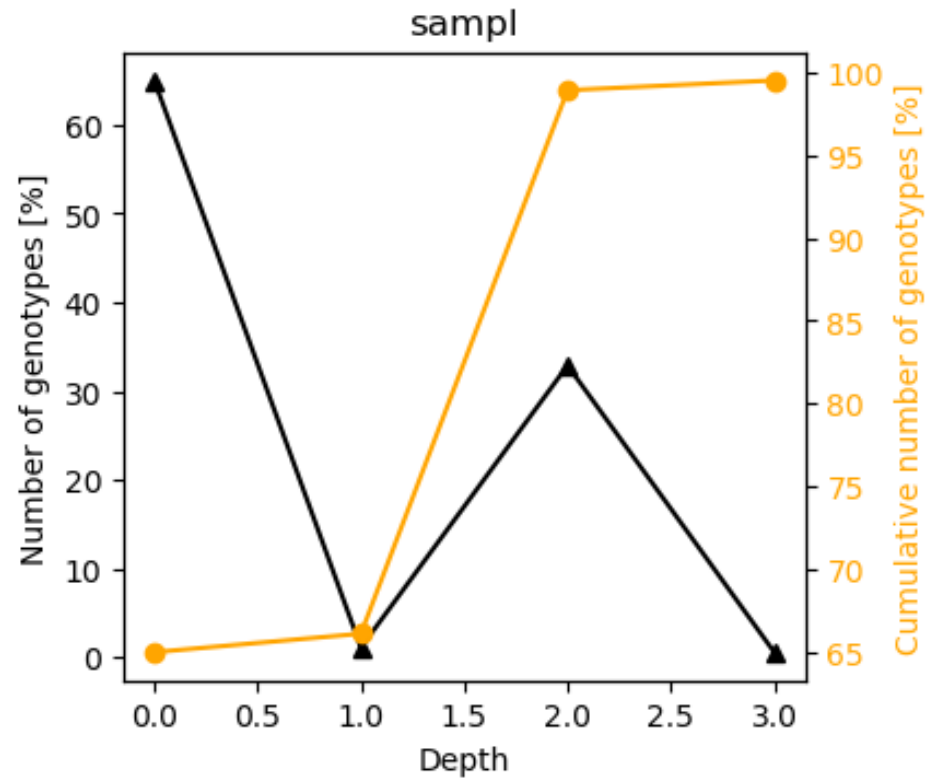
Ts/Tv stratified by QUAL



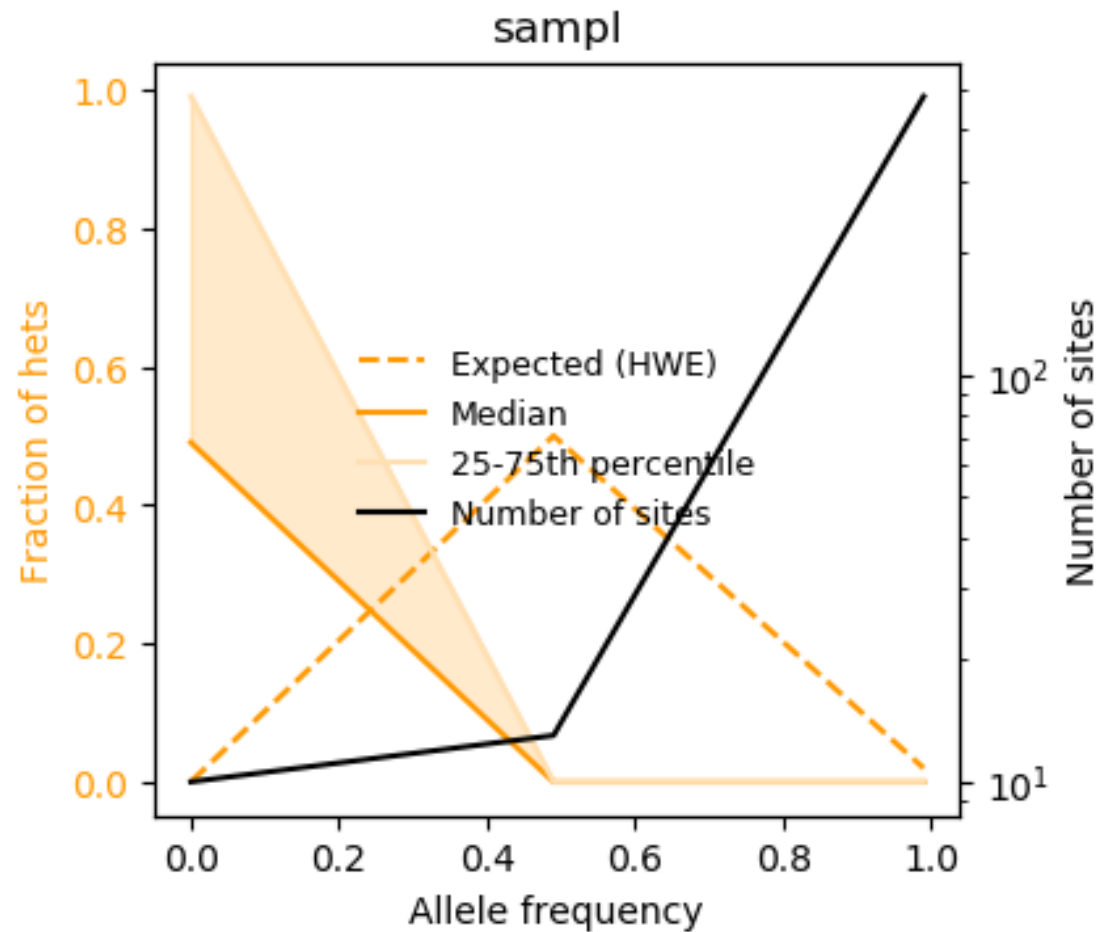
Indel distribution



Depth distribution



Number of HETs by AF



Substitution types

