

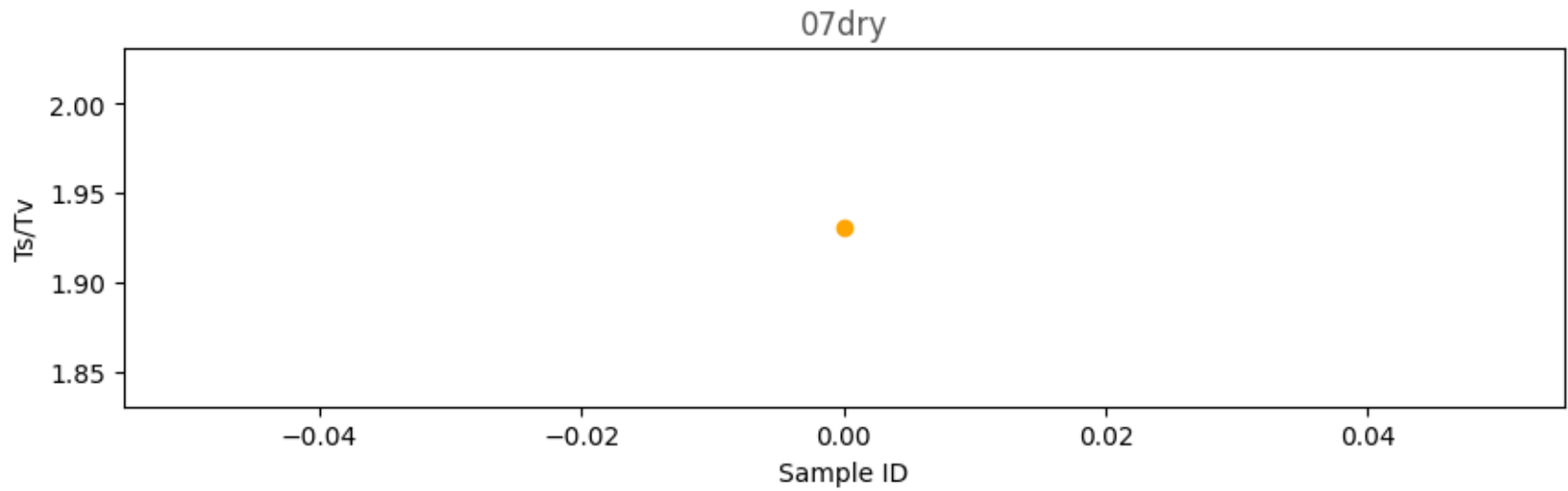
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
07dry	4,080,747	1.93	1.94	949,923	–	0	0
* frameshift ratio: out/(out+in)							

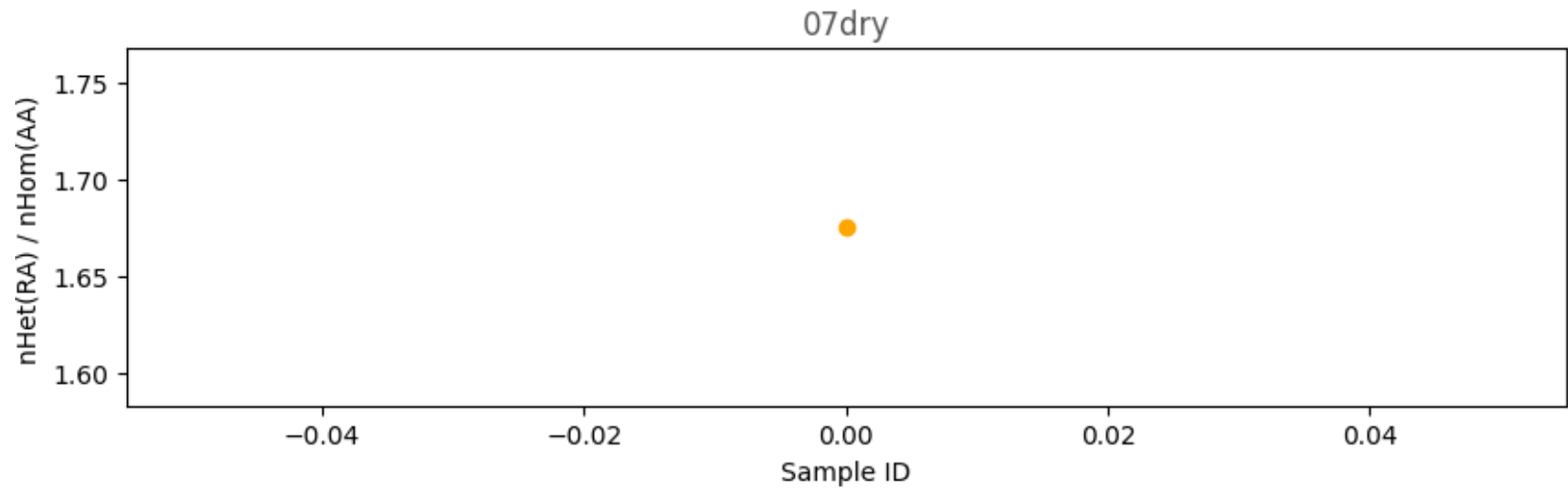
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
07dry	62.6%	1.90	67.3%	99,614	2,124

- 07dry .. /ngc/projects2/gm/data/archive/2022/variants/snv/07drysesm-103880904892-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01555-220916_A01411_BHYNLVDSX3-EXT_LAB
KA_NGCWGS-NGCWGS05356_22RKG014509_snv_germline_raw.haplotype_caller.vcf.gz

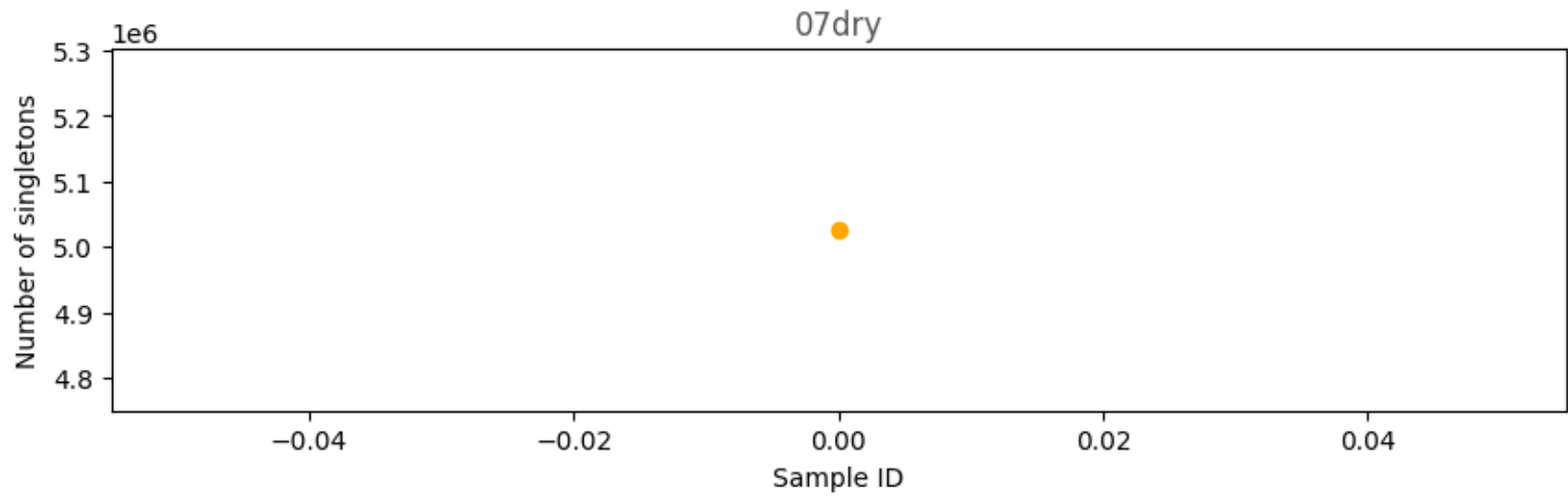
Ts/Tv by sample



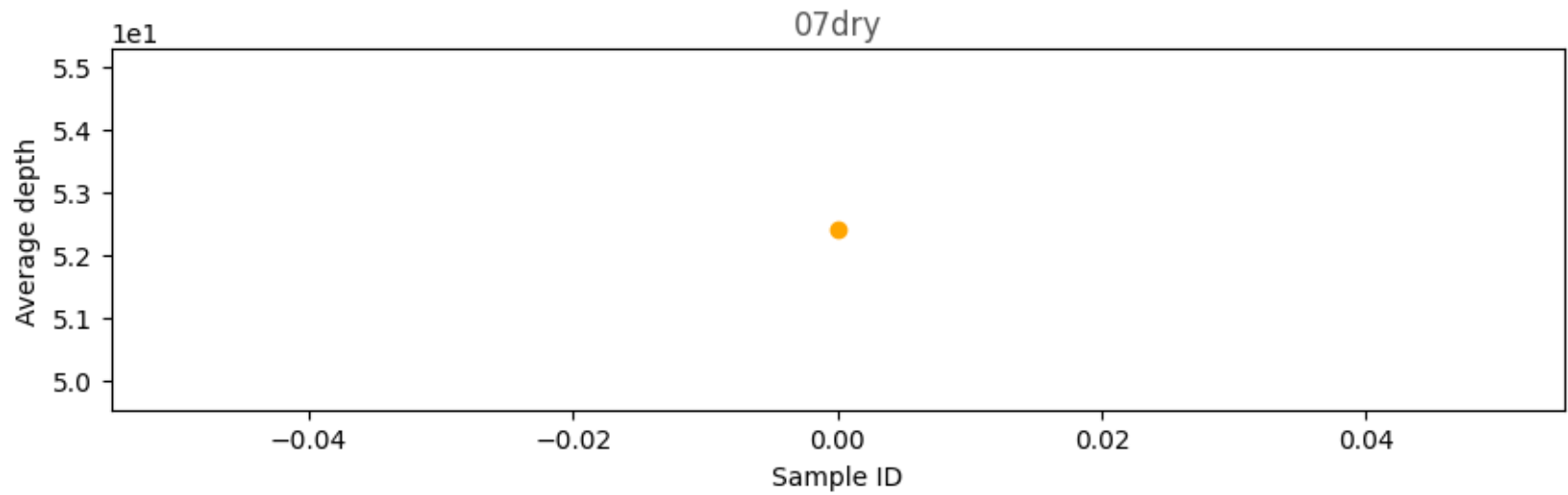
Hets vs non-ref Homs by sample



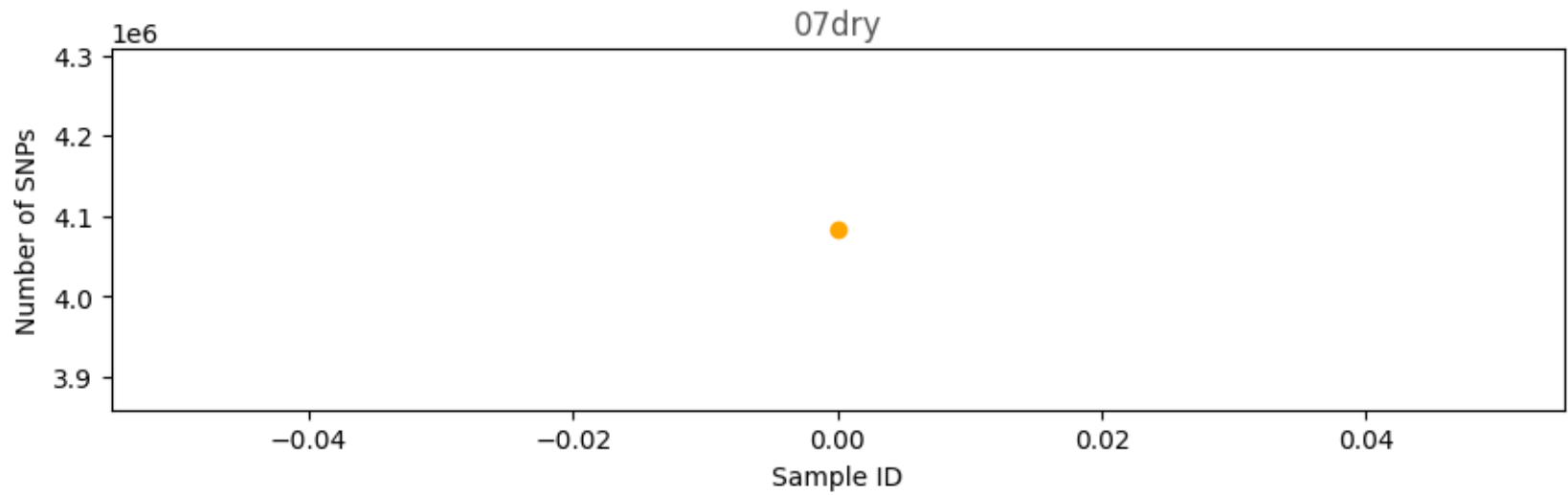
Singletons by sample (hets and homs)



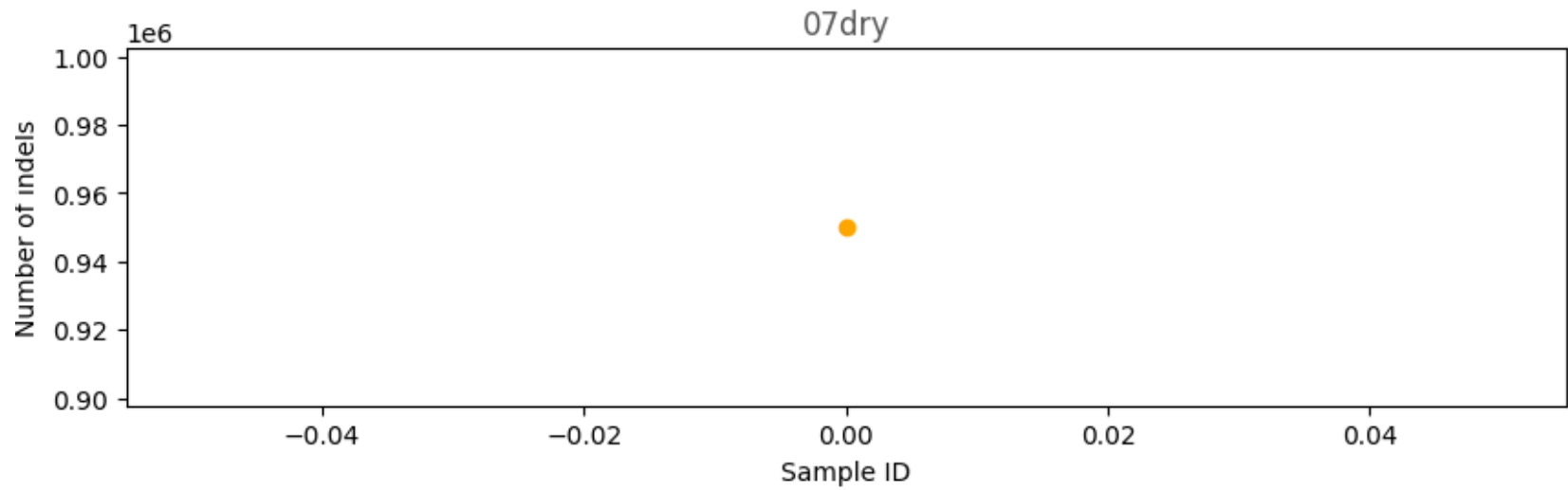
Average depth by sample



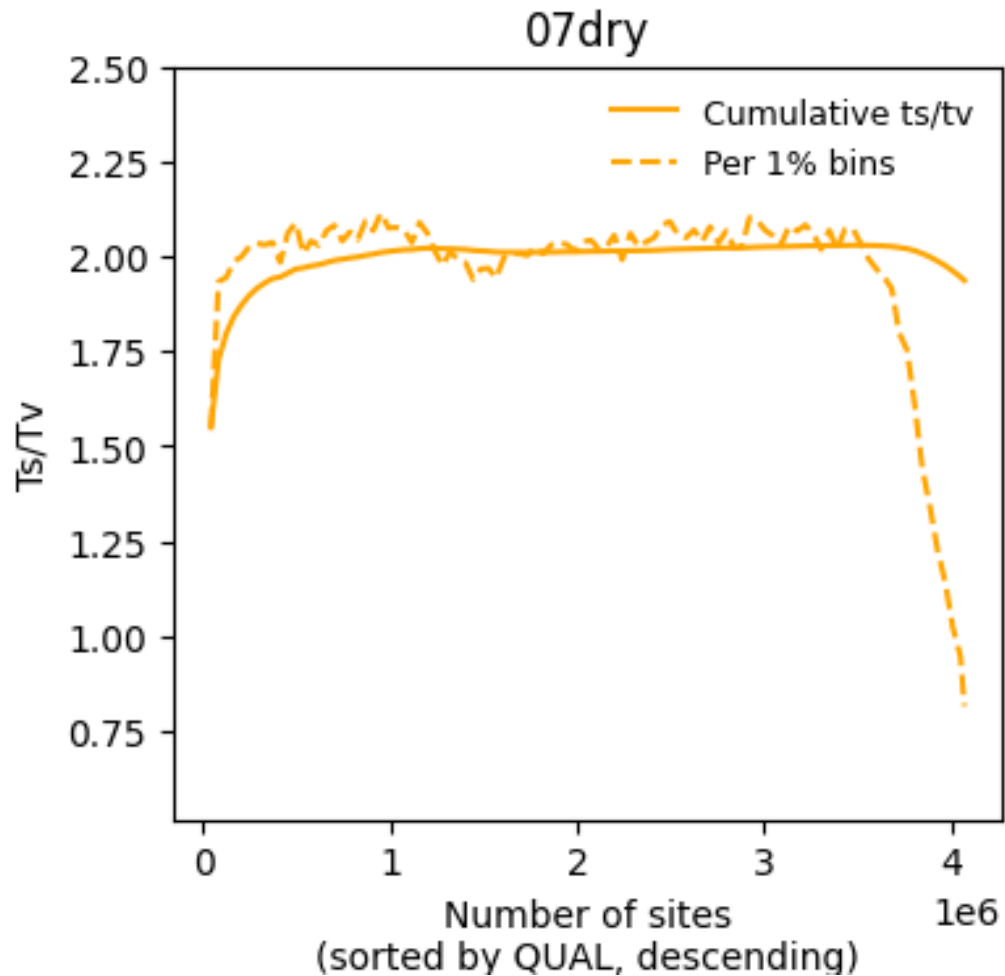
Number of SNPs by sample



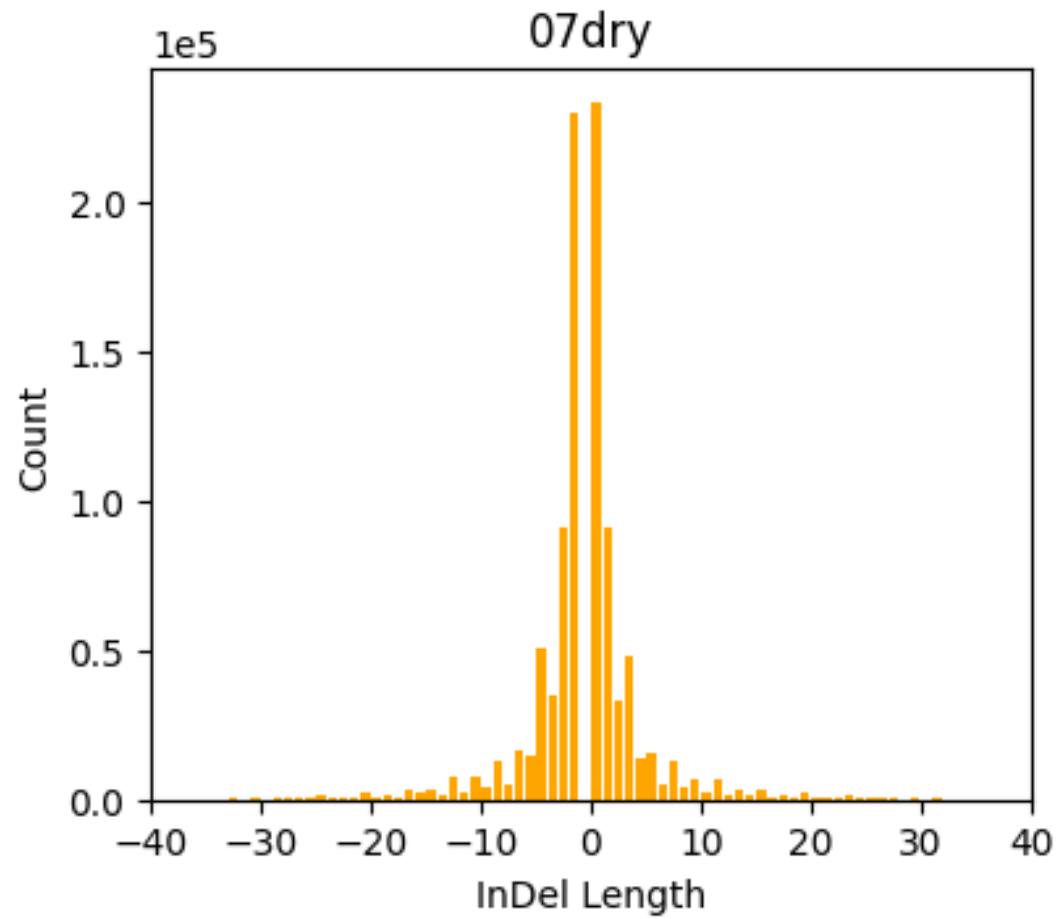
Number of indels by sample



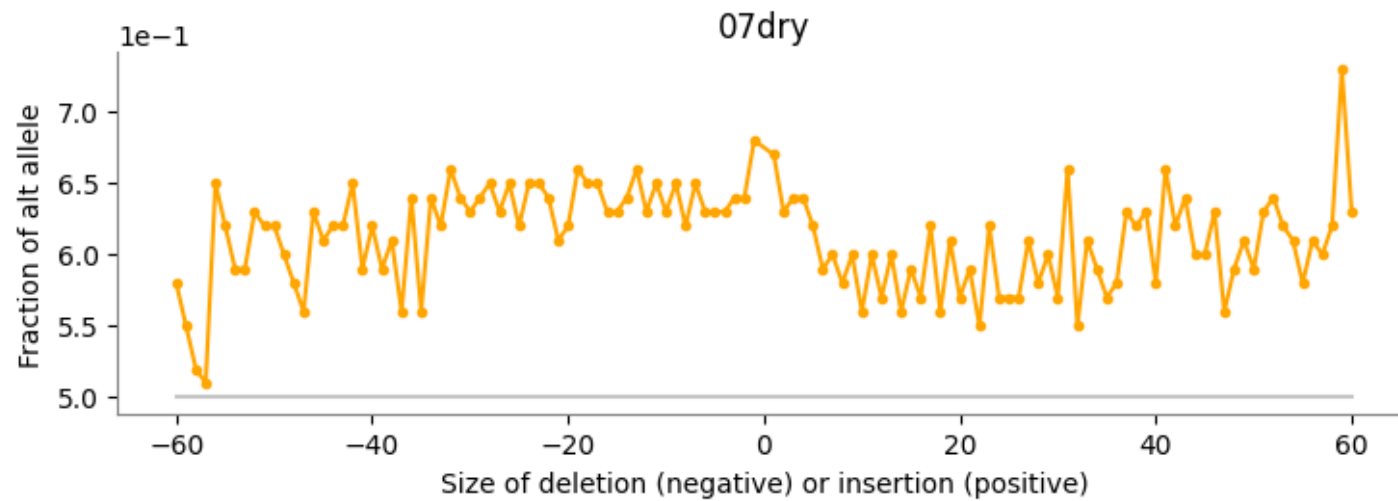
Ts/Tv stratified by QUAL



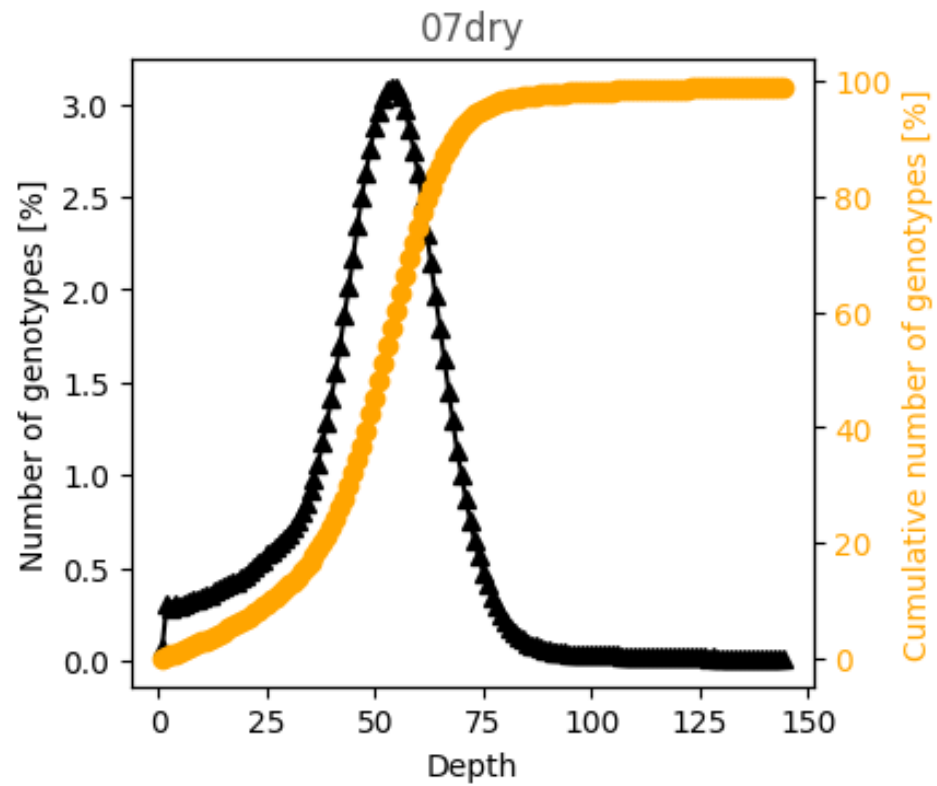
Indel distribution



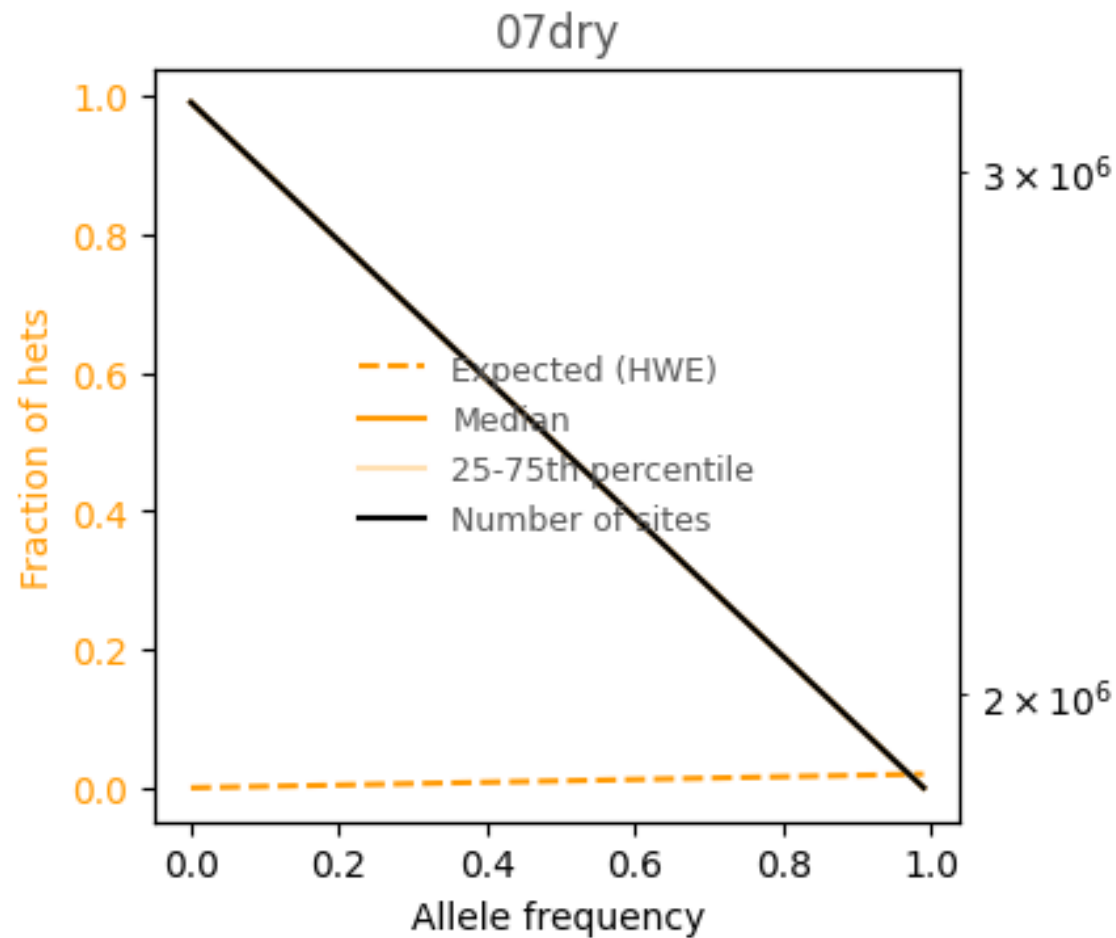
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

