

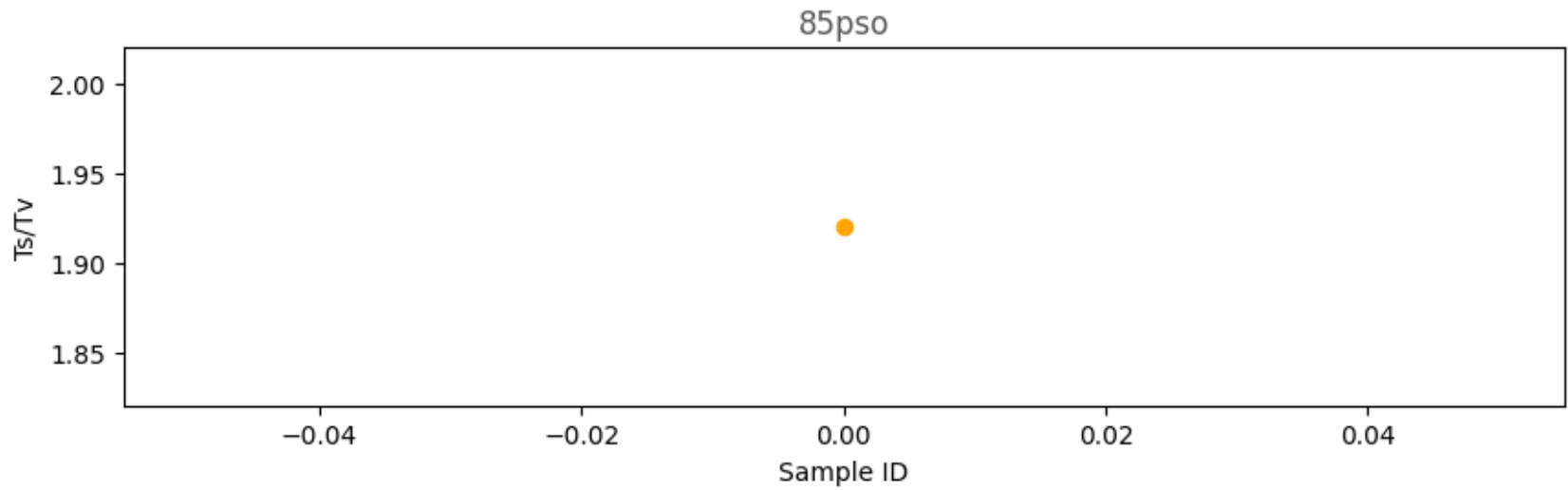
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
85pso	4,115,001	1.92	1.93	964,034	–	0	0
* frameshift ratio: out/(out+in)							

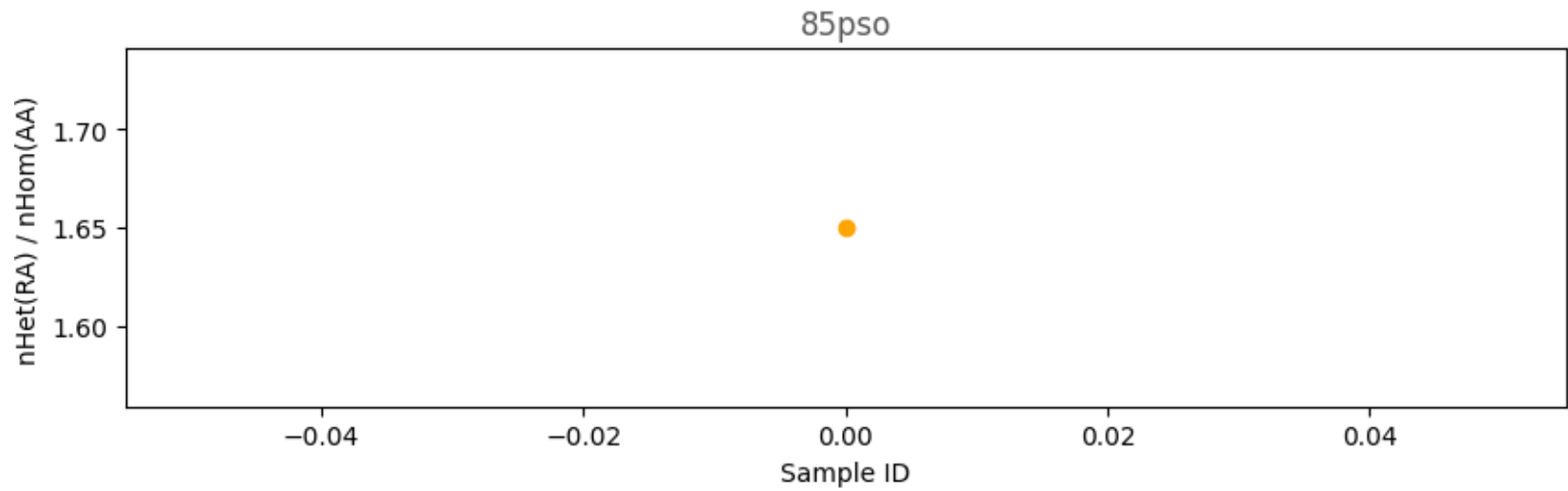
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
85pso	62.3%	1.88	67.4%	106,068	2,155

- 85pso .. /ngc/projects2/gm/data/archive/2022/variants/snv/85psofttm-103854313939-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00712-220407_A01176_BHGT17DSX3-EXT_LAB
 KA_NGCWGS-NGCWGS04114_22RKG003147x01_snv_germline_raw.haplotype_caller.vcf.gz

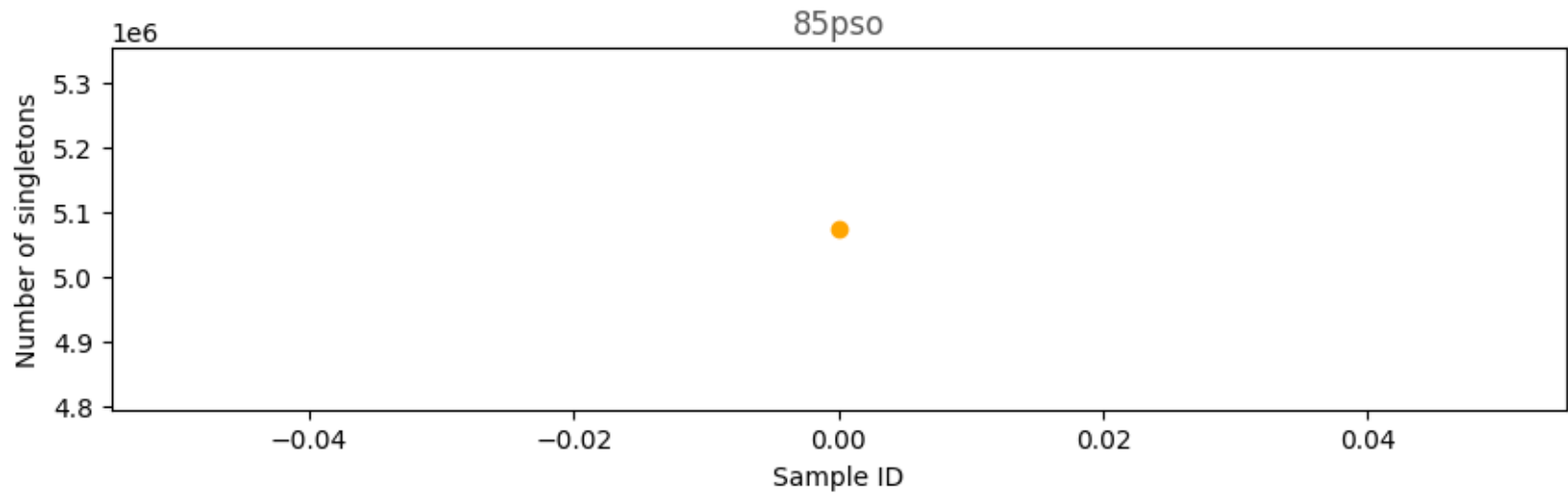
Ts/Tv by sample



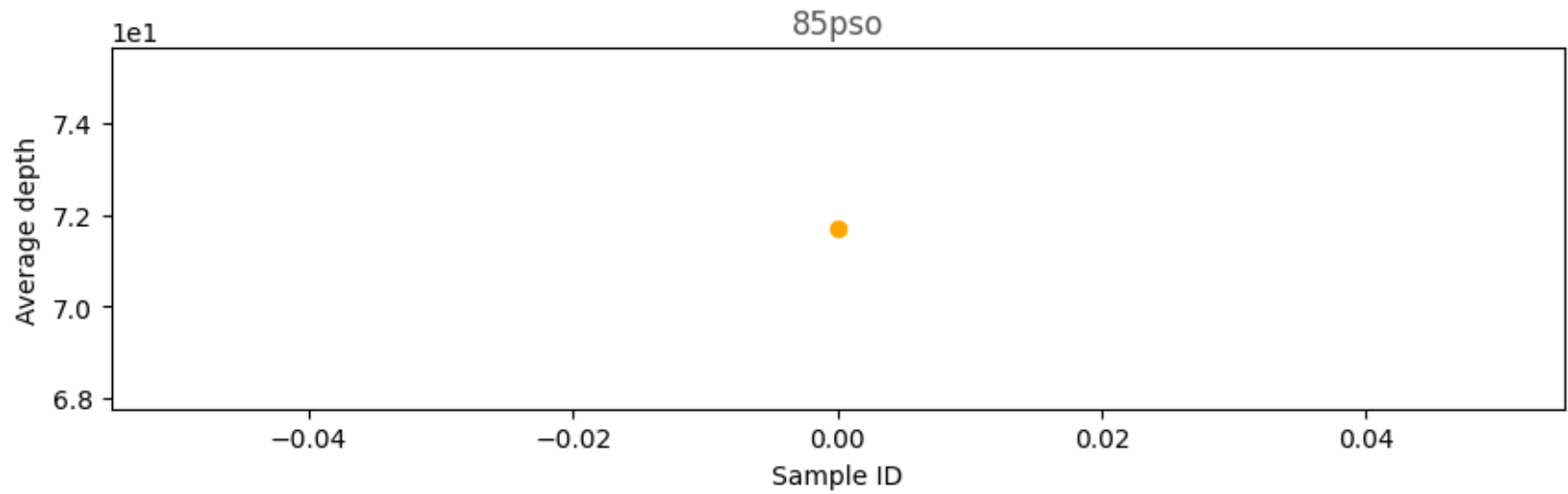
Hets vs non-ref Homs by sample



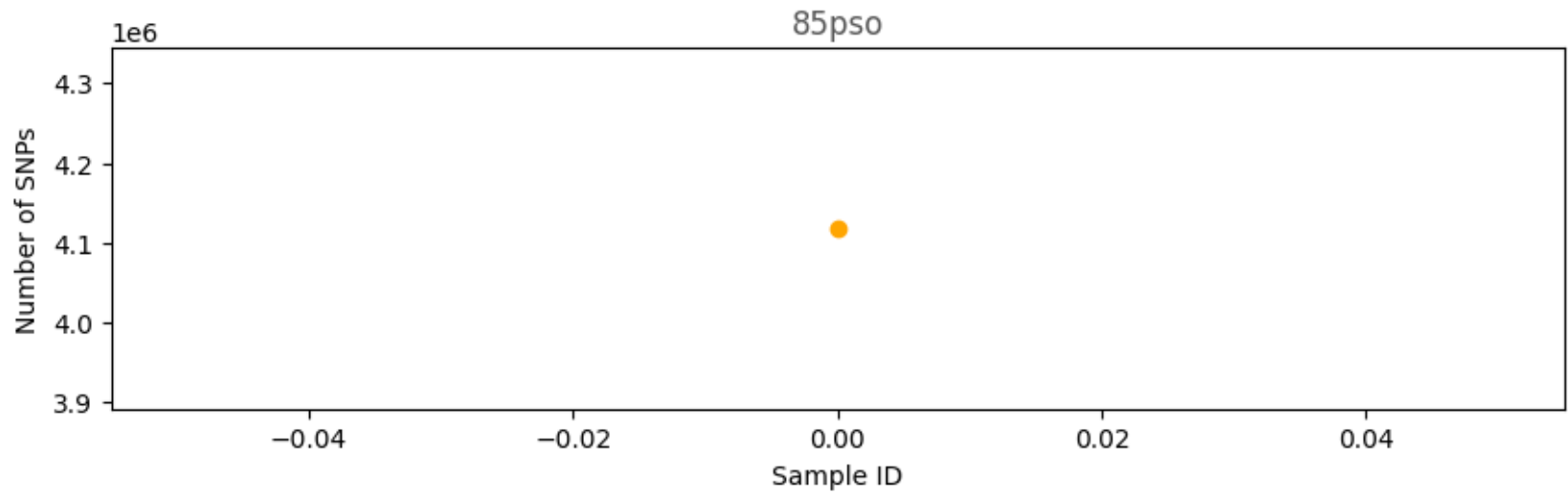
Singletons by sample (hets and homs)



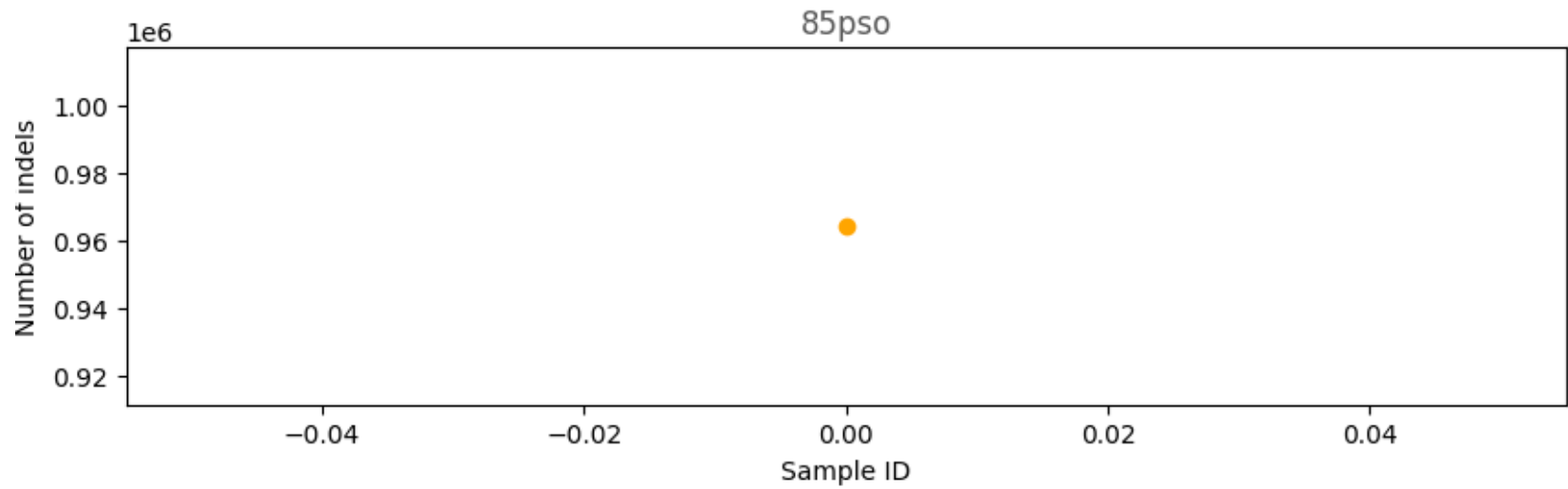
Average depth by sample



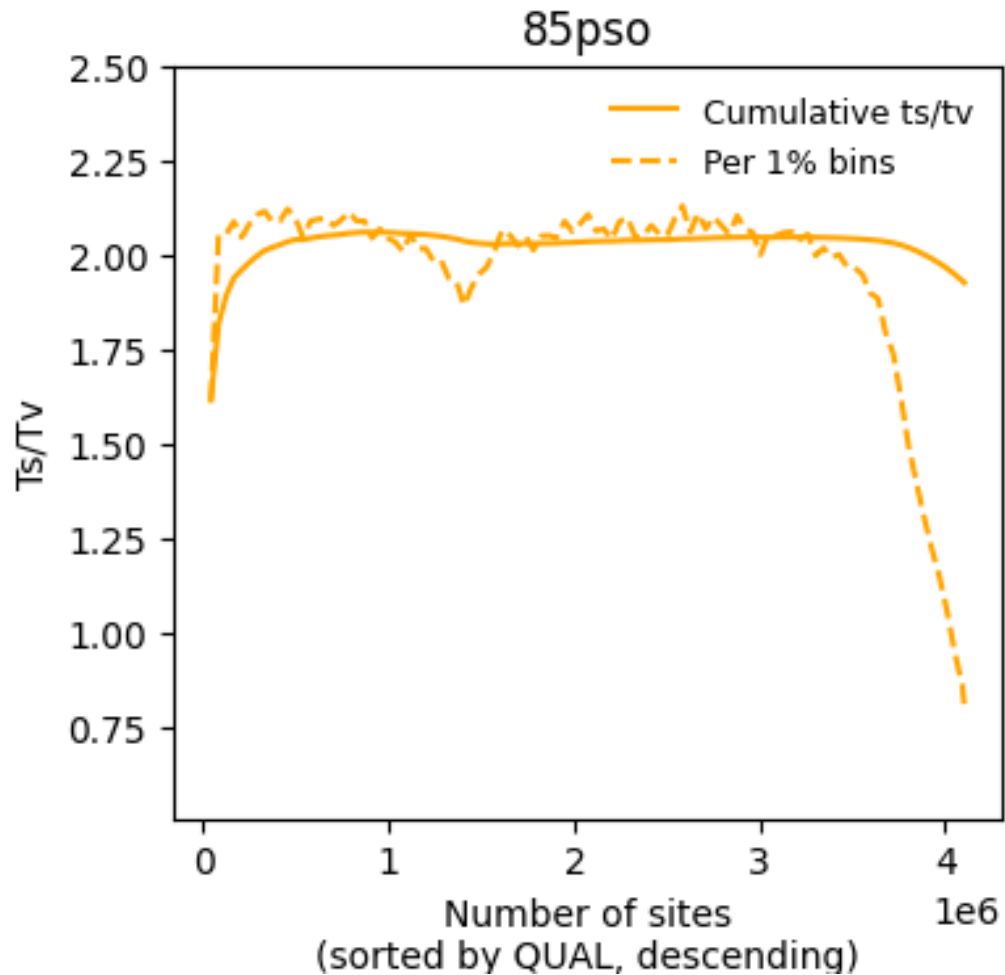
Number of SNPs by sample



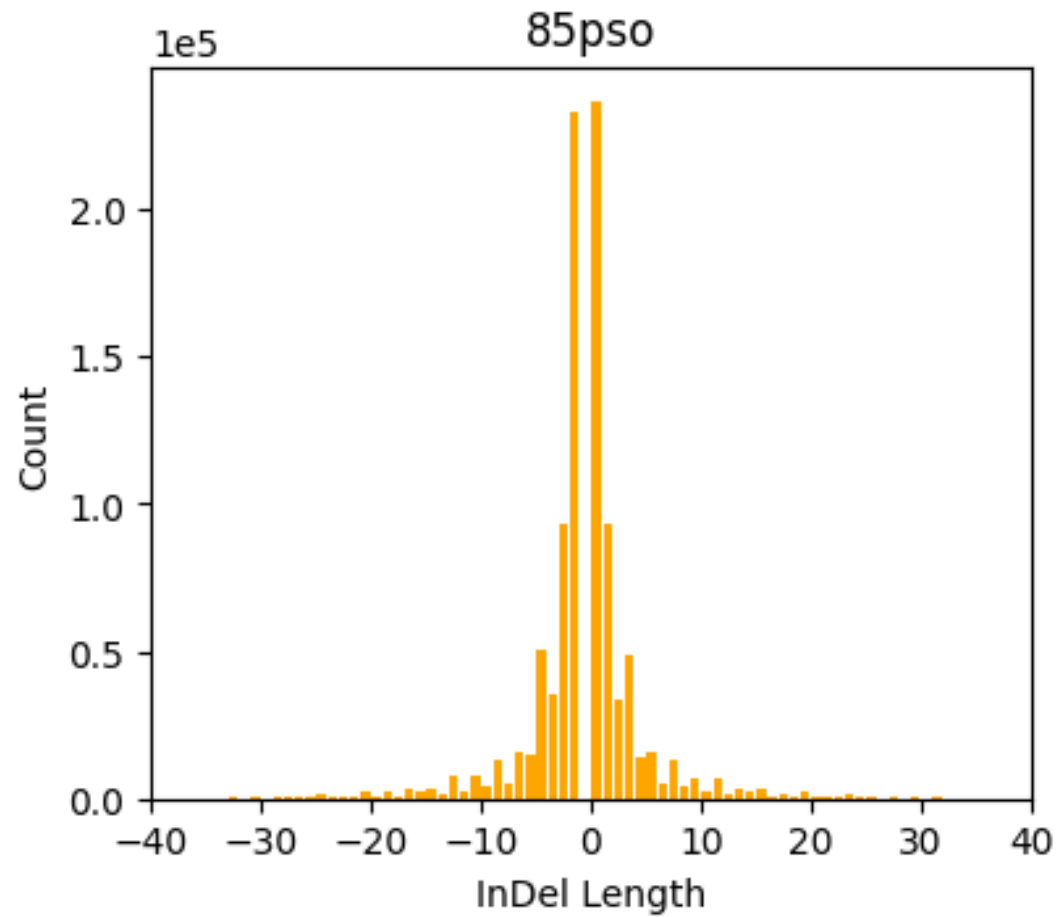
Number of indels by sample



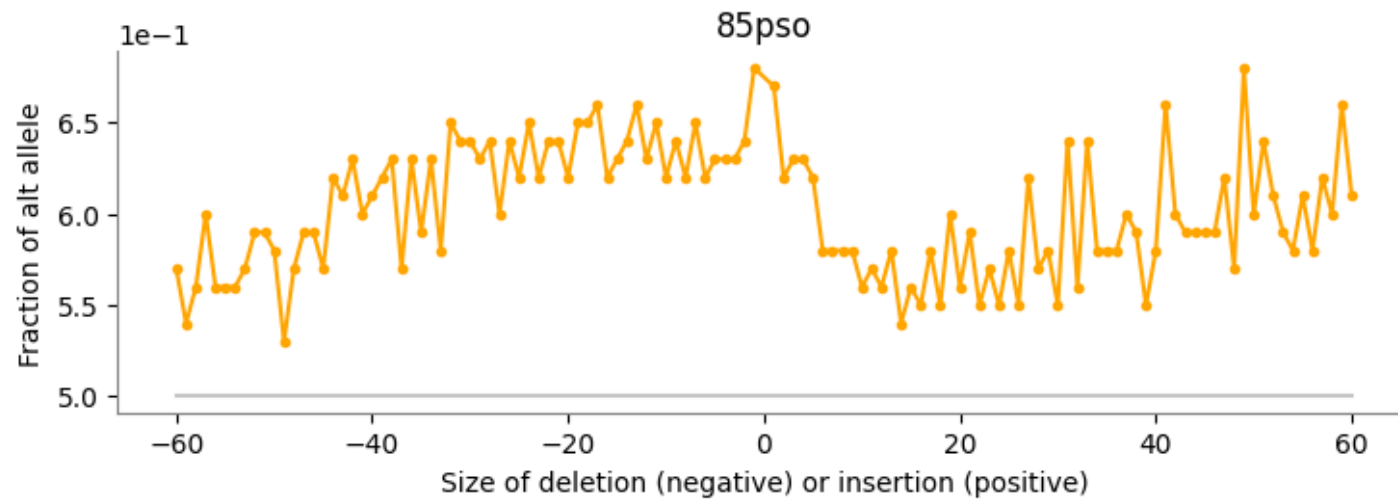
Ts/Tv stratified by QUAL



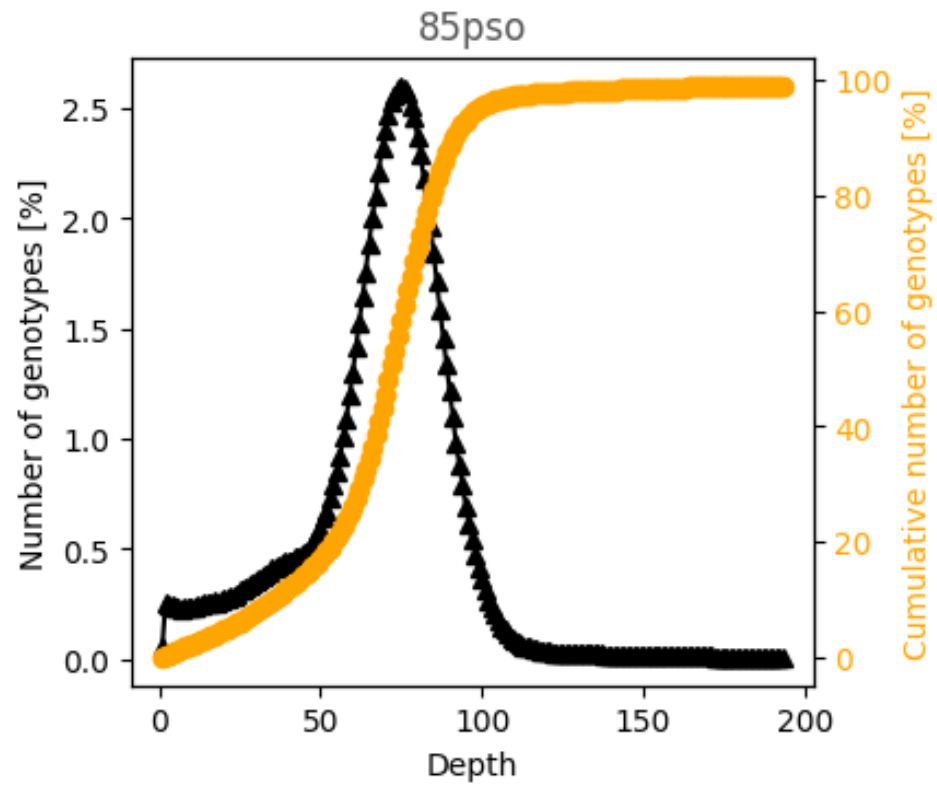
Indel distribution



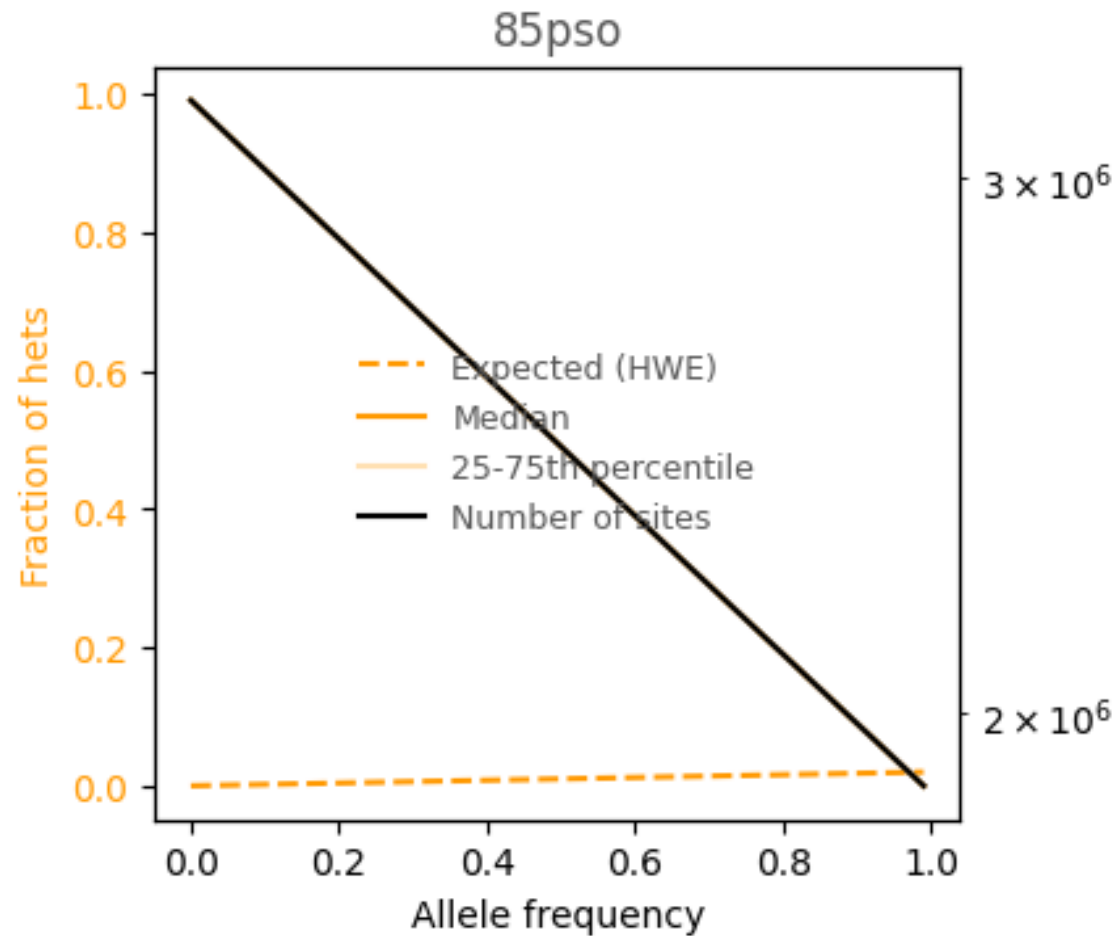
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

