

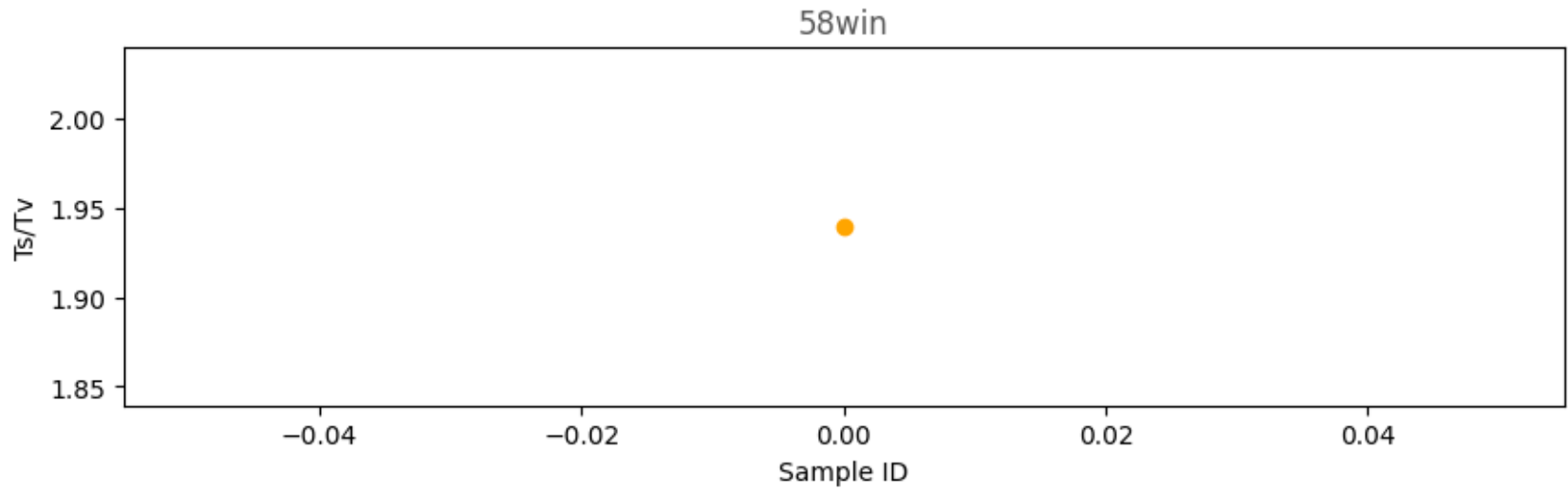
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
58win	4,035,374	1.94	1.94	934,846	–	0	0
* frameshift ratio: out/(out+in)							

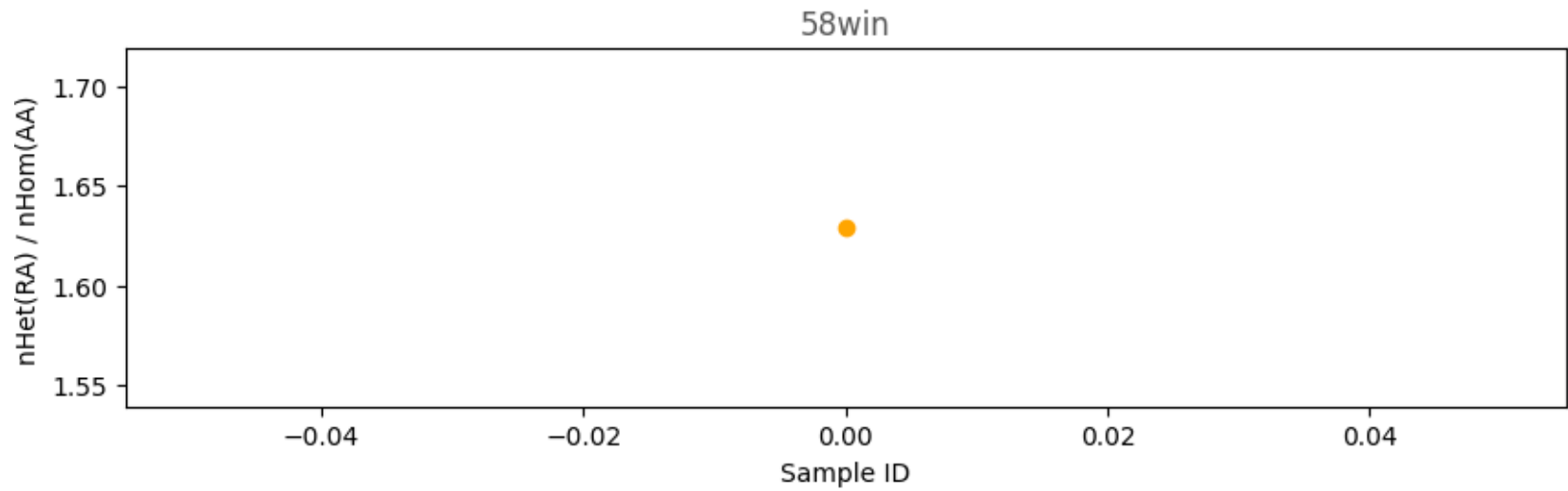
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
58win	62.0%	1.91	66.1%	91,582	1,781

- 58win .. /ngc/projects2/gm/data/archive/2022/variants/snv/58wingonm-103892826107-Normal_B
 lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01043-220727_A01176_BHNV7KDSX3-EXT_LAB
 KA_NGCWGS-NGCWGS04720_snv_germline_raw.haplotype_caller.vcf.gz

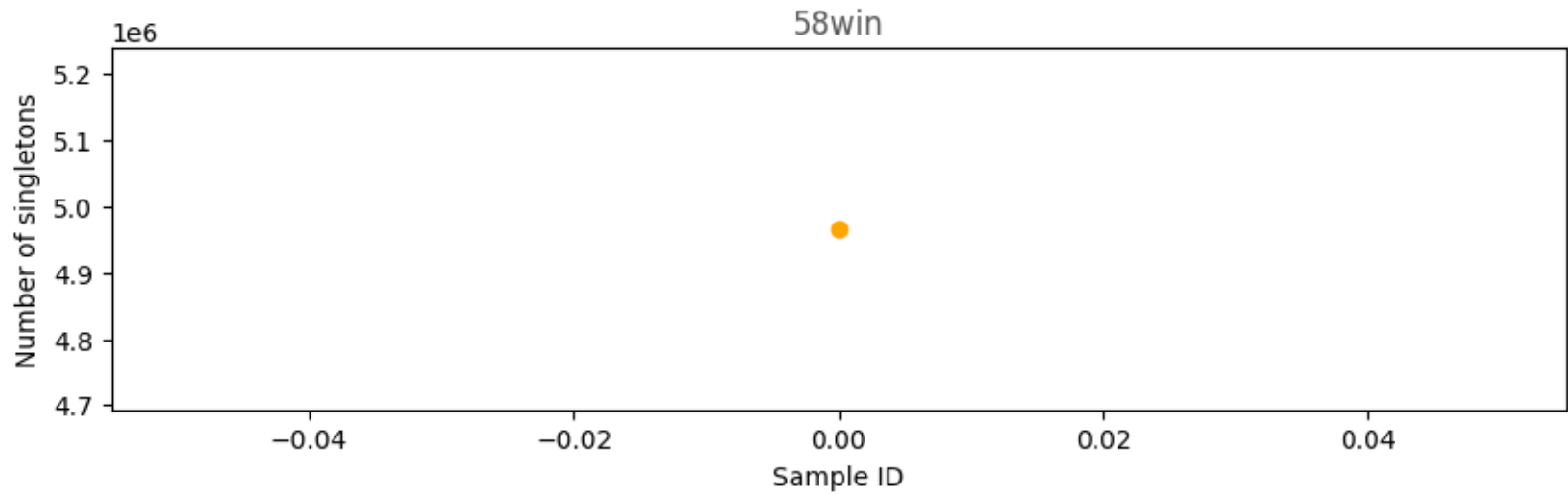
Ts/Tv by sample



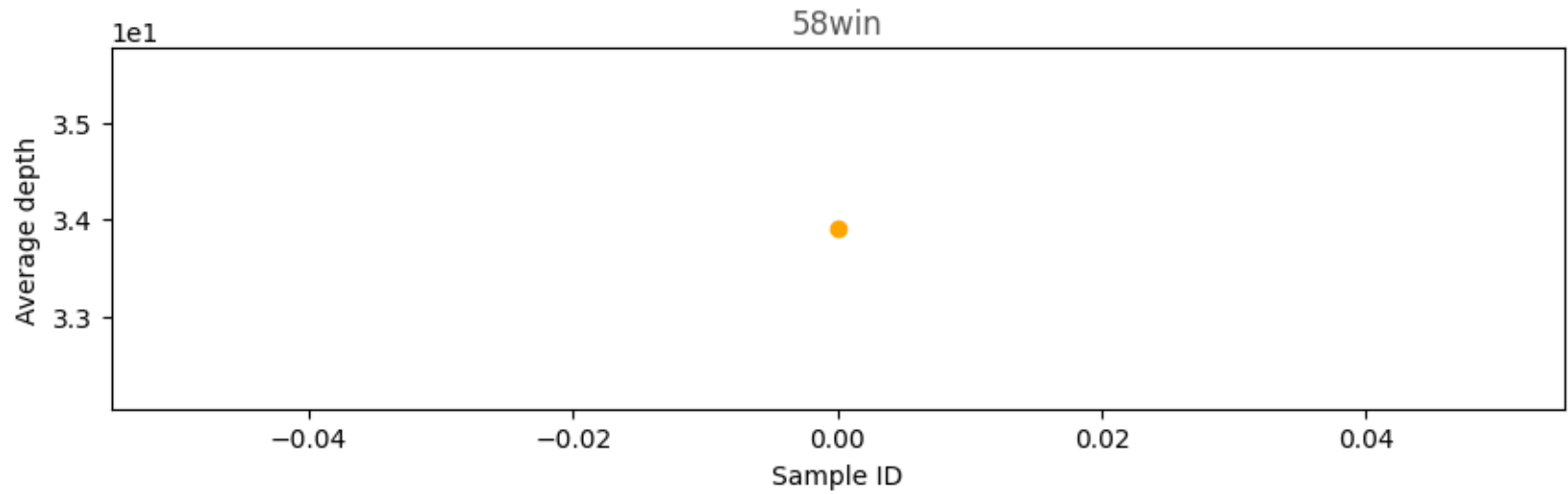
Hets vs non-ref Homs by sample



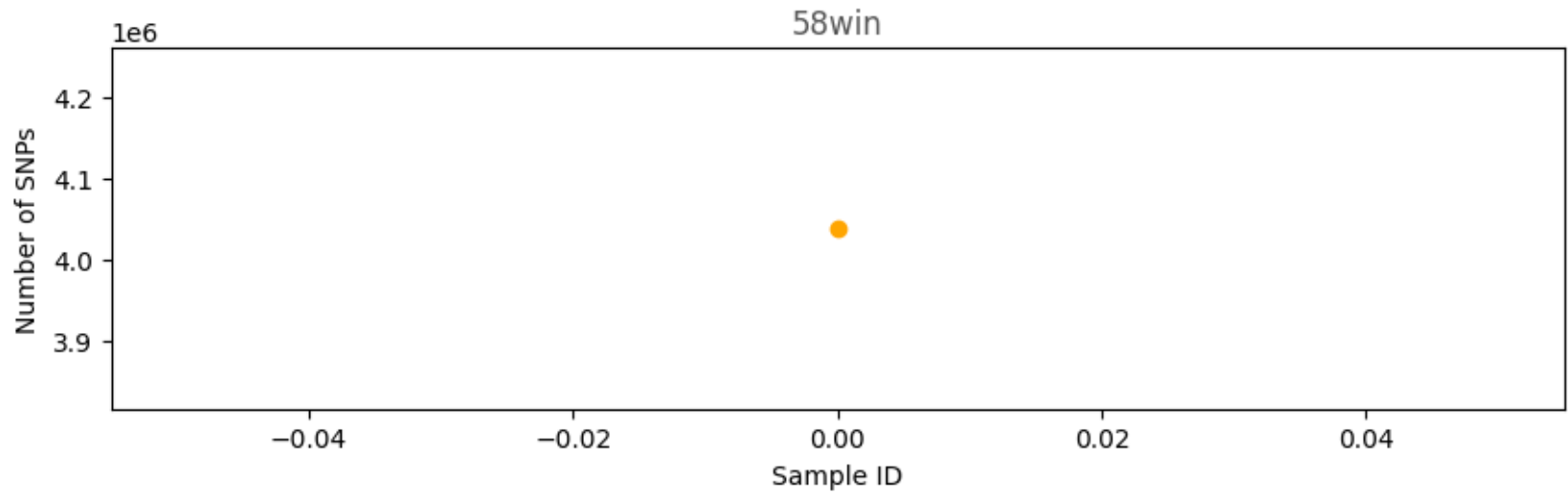
Singletons by sample (hets and homs)



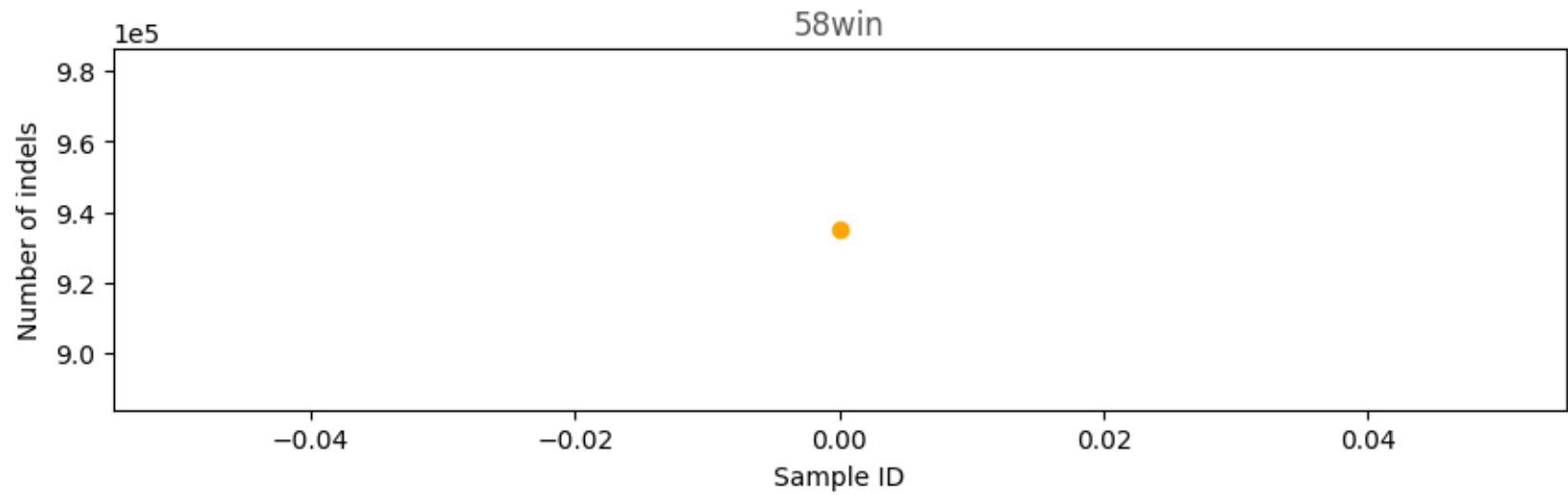
Average depth by sample



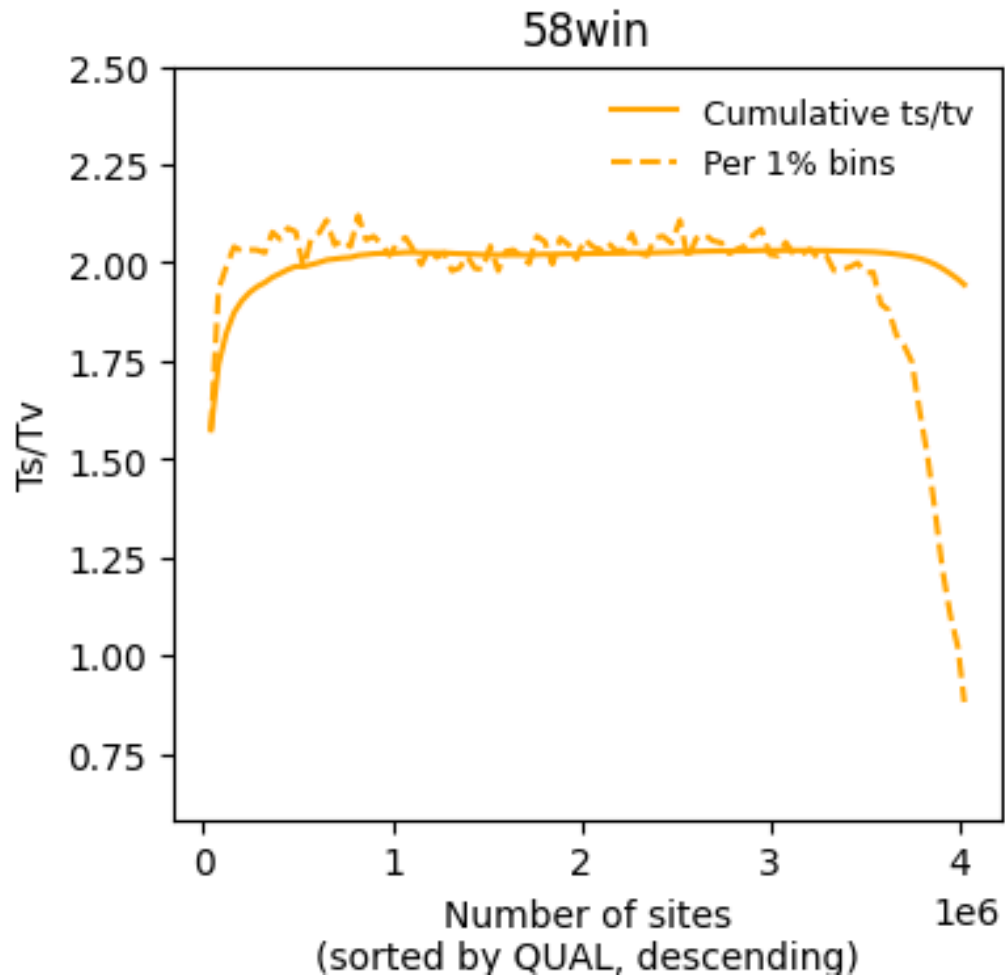
Number of SNPs by sample



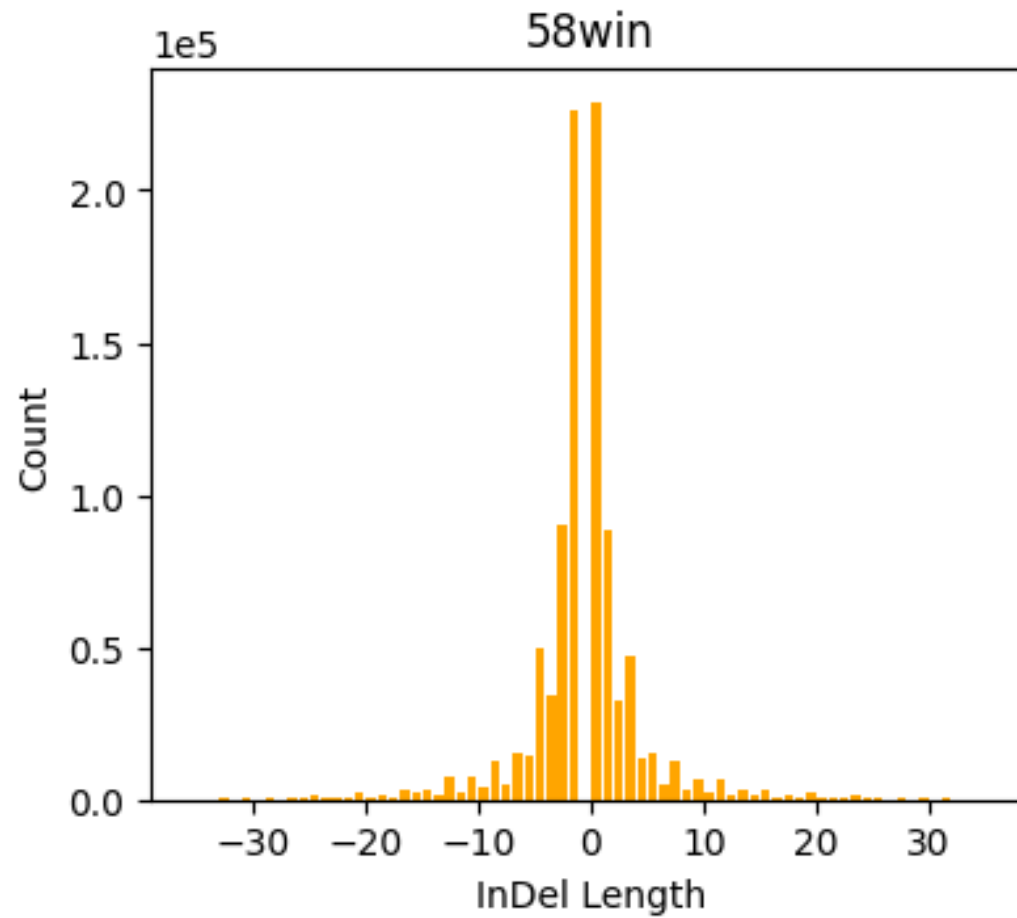
Number of indels by sample



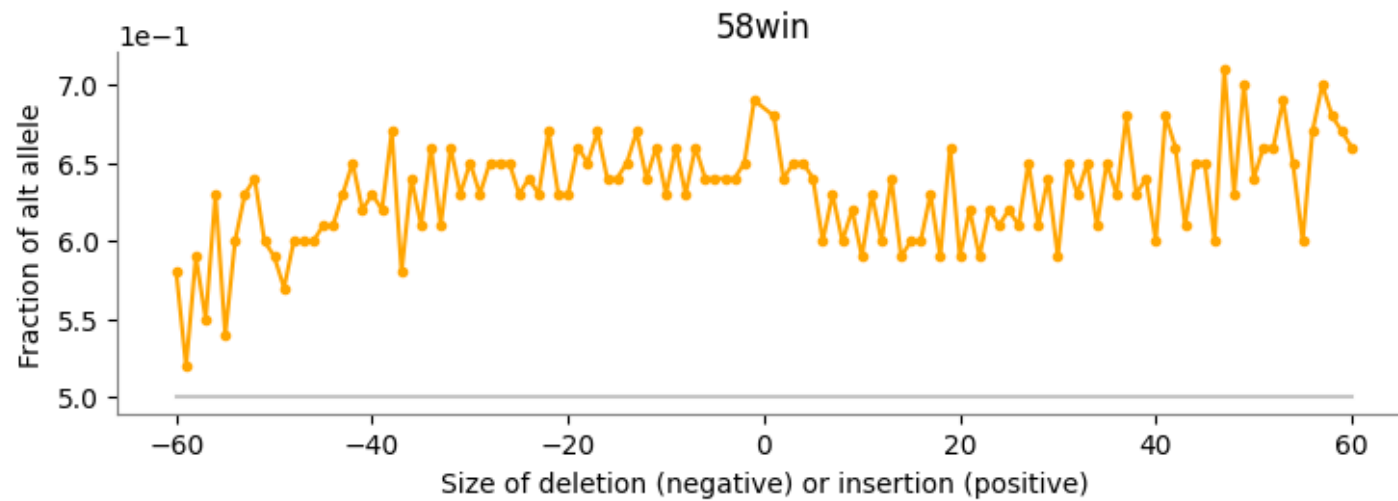
Ts/Tv stratified by QUAL



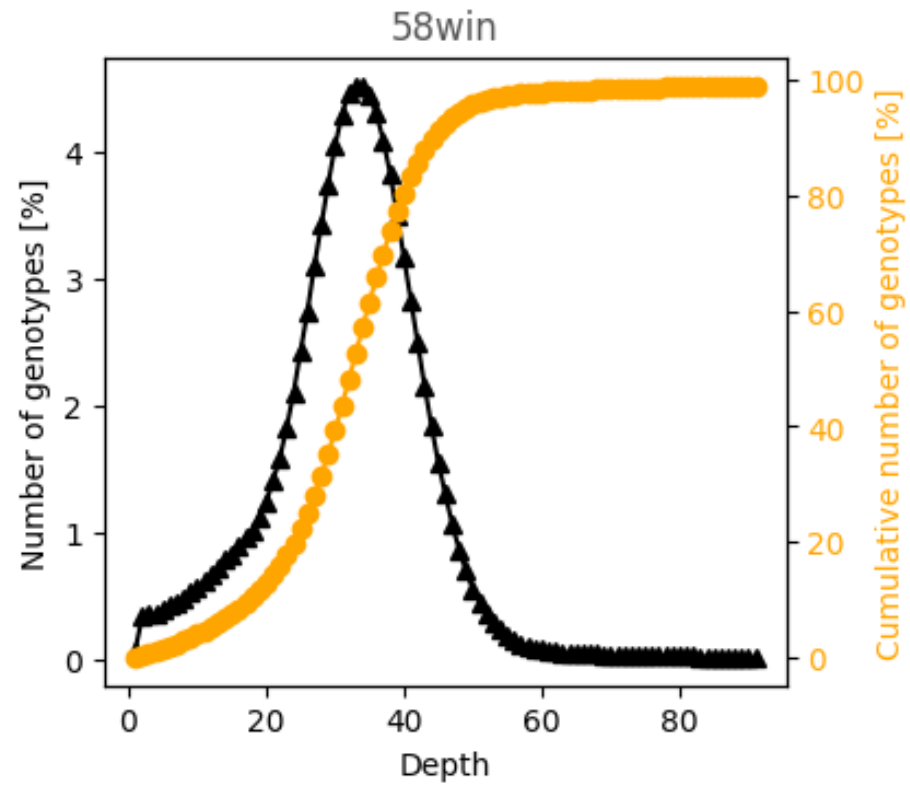
Indel distribution



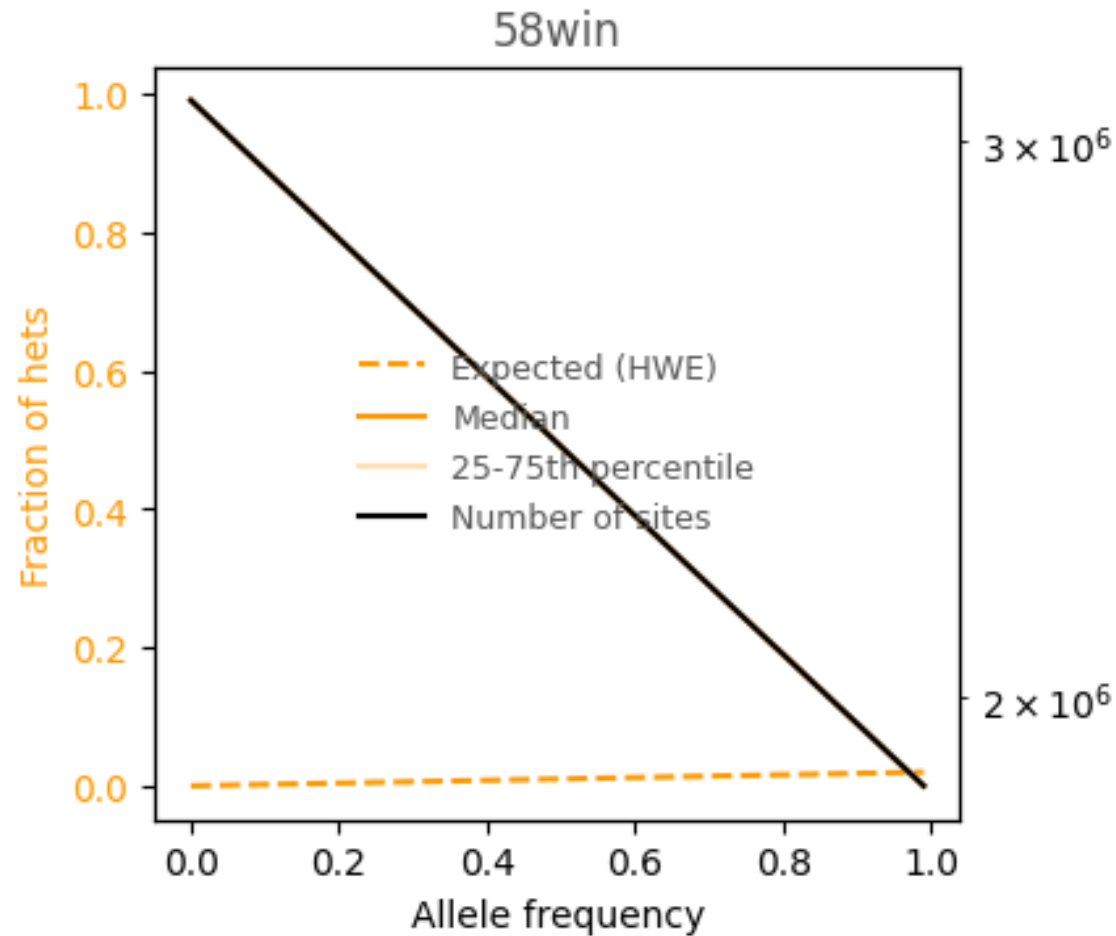
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

