

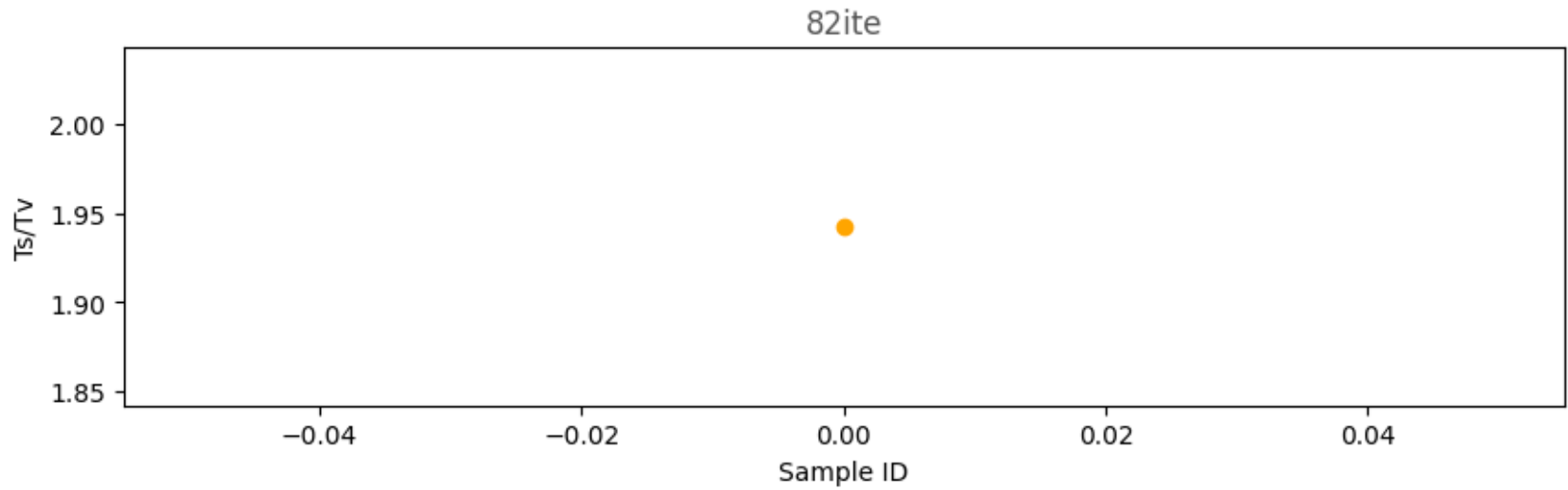
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
82ite	4,097,782	1.94	1.95	949,890	–	0	0
* frameshift ratio: out/(out+in)							

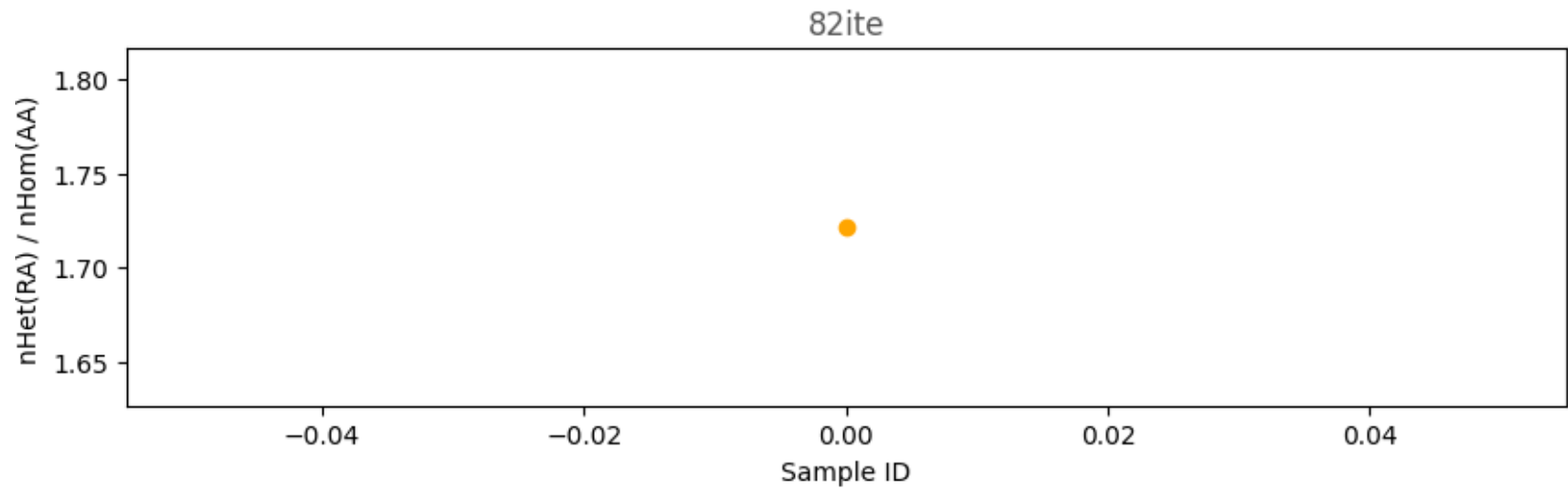
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
82ite	63.3%	1.92	67.9%	98,490	2,231

- 82ite .. /ngc/projects2/gm/data/archive/2022/variants/snv/82itemonf-103913448650-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01754-220923_A01961_AH33KNDSX5-EXT_LAB
KA_NGCWGS-NGCWGS05516_22RKG015729_snv_germline_raw.haplotype_caller.vcf.gz

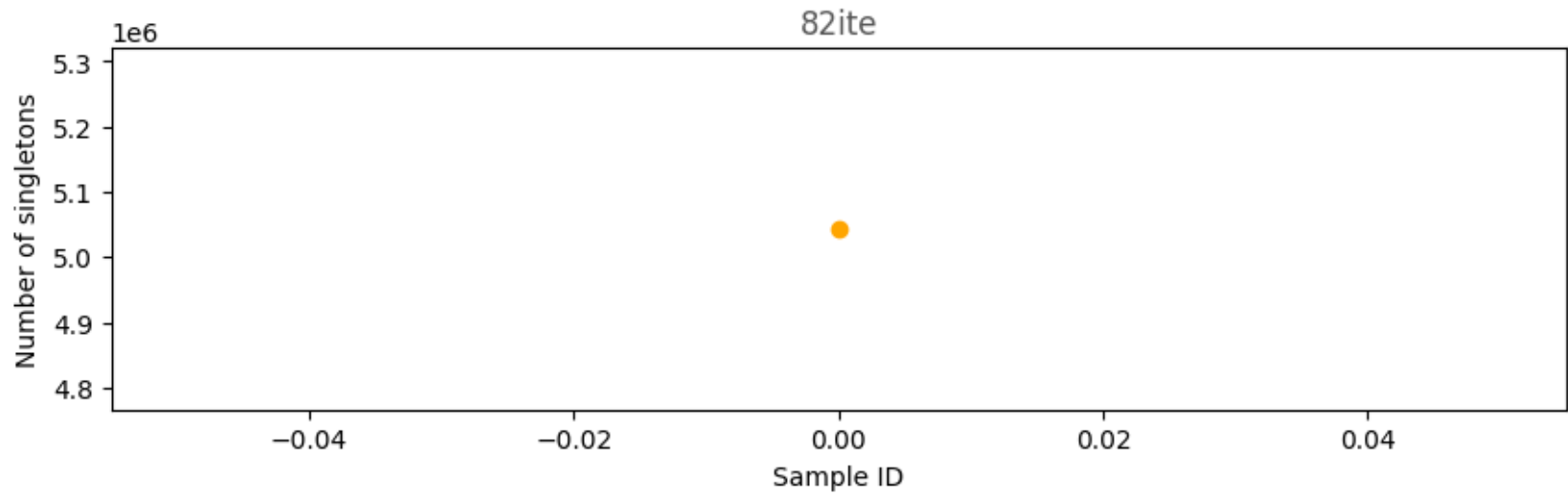
Ts/Tv by sample



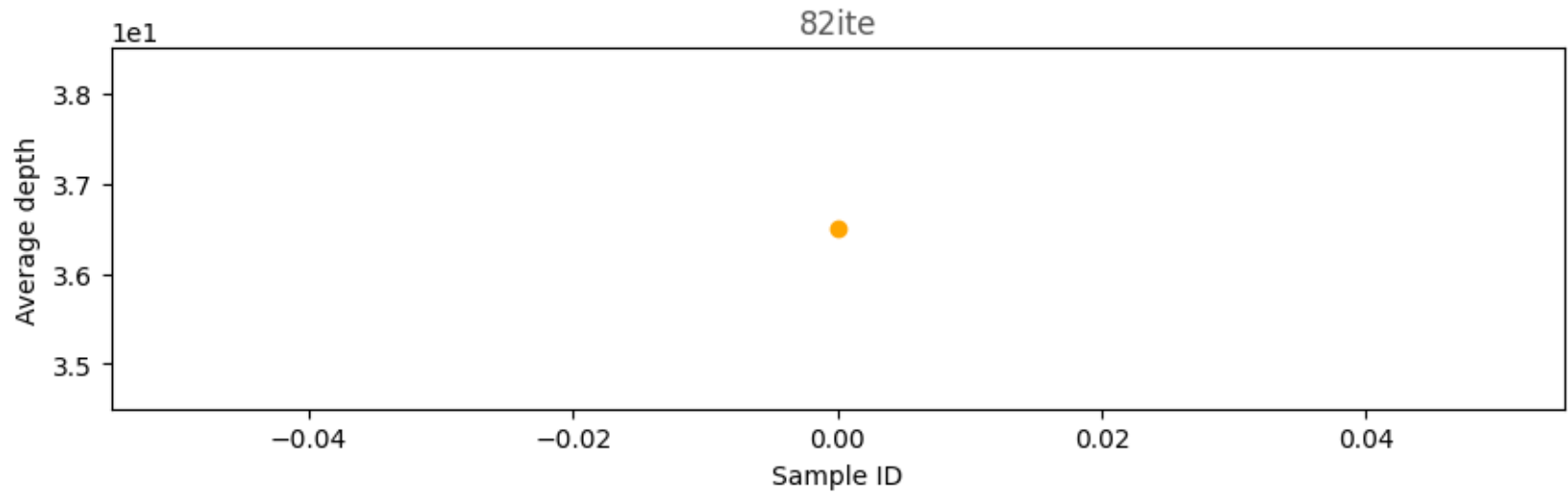
Hets vs non-ref Homs by sample



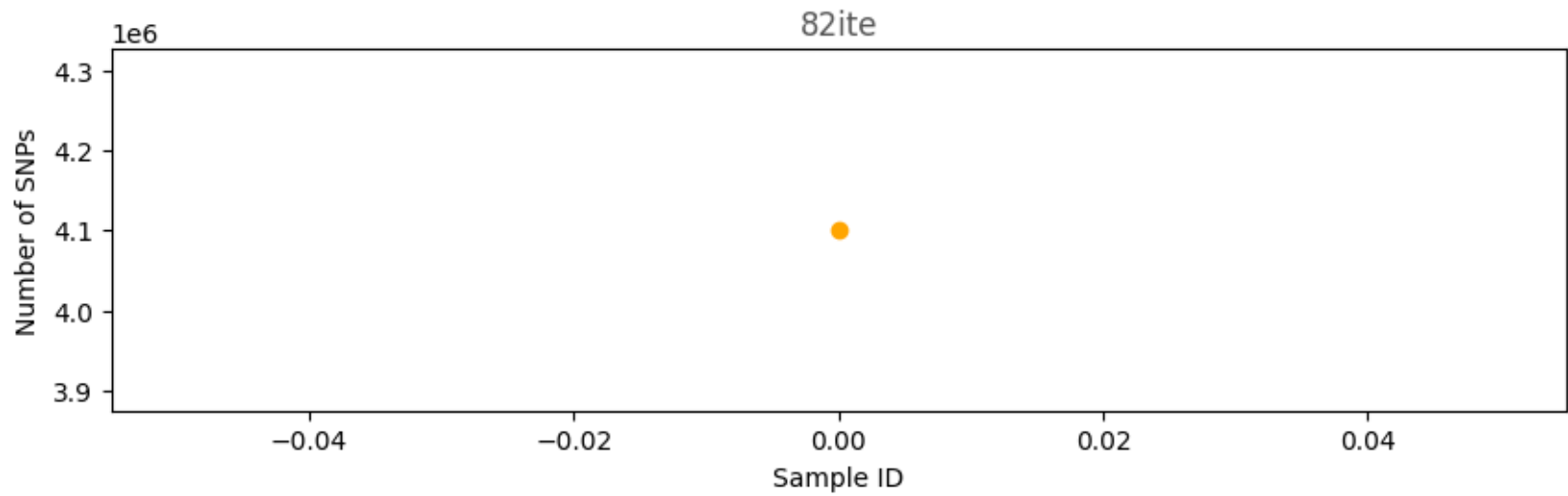
Singletons by sample (hets and homs)



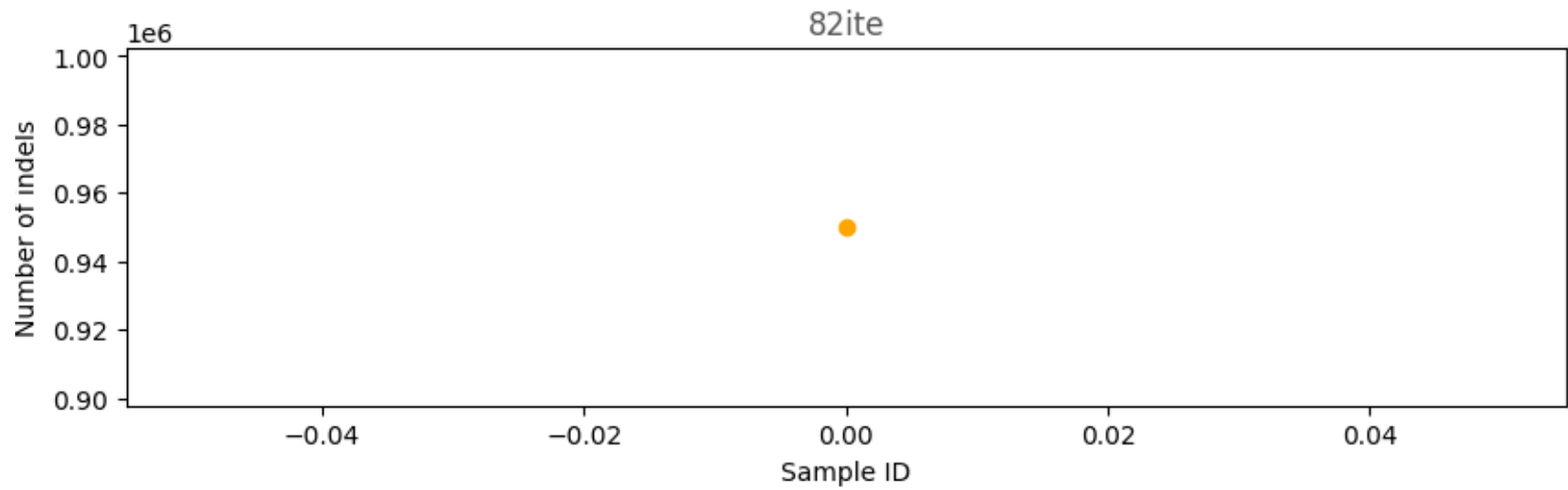
Average depth by sample



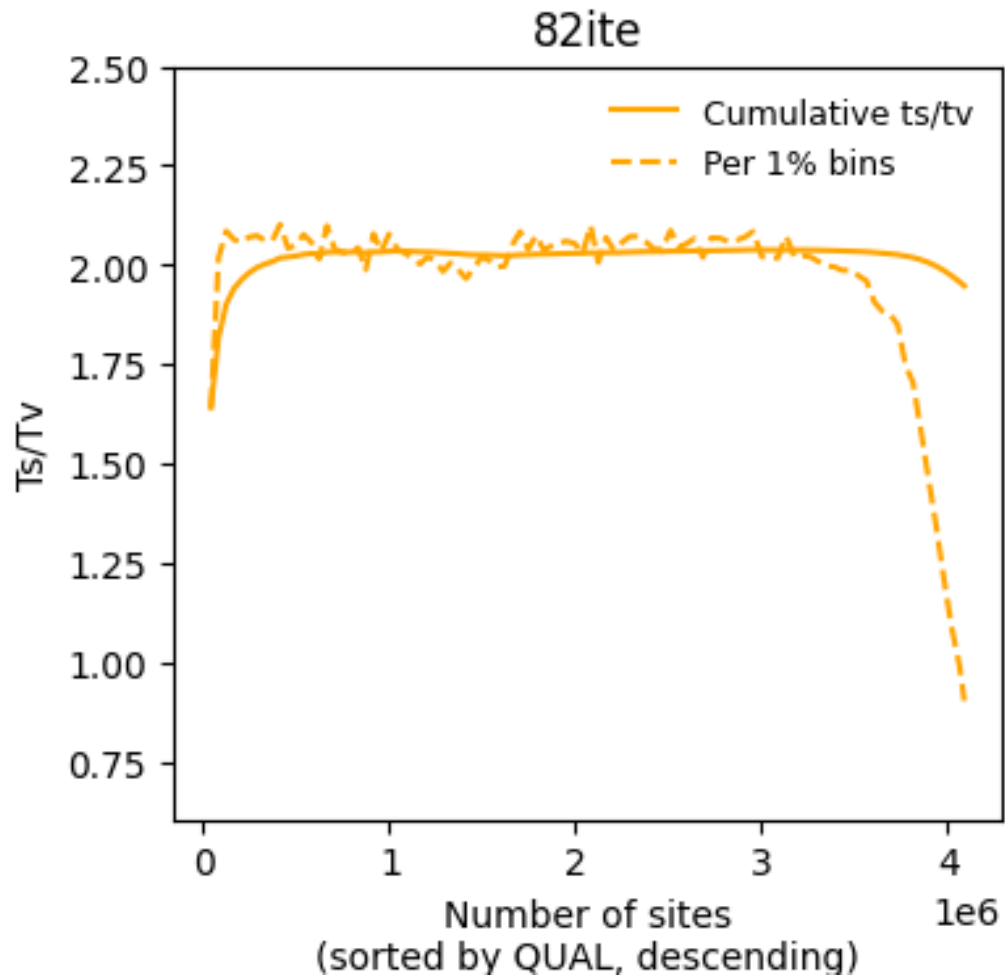
Number of SNPs by sample



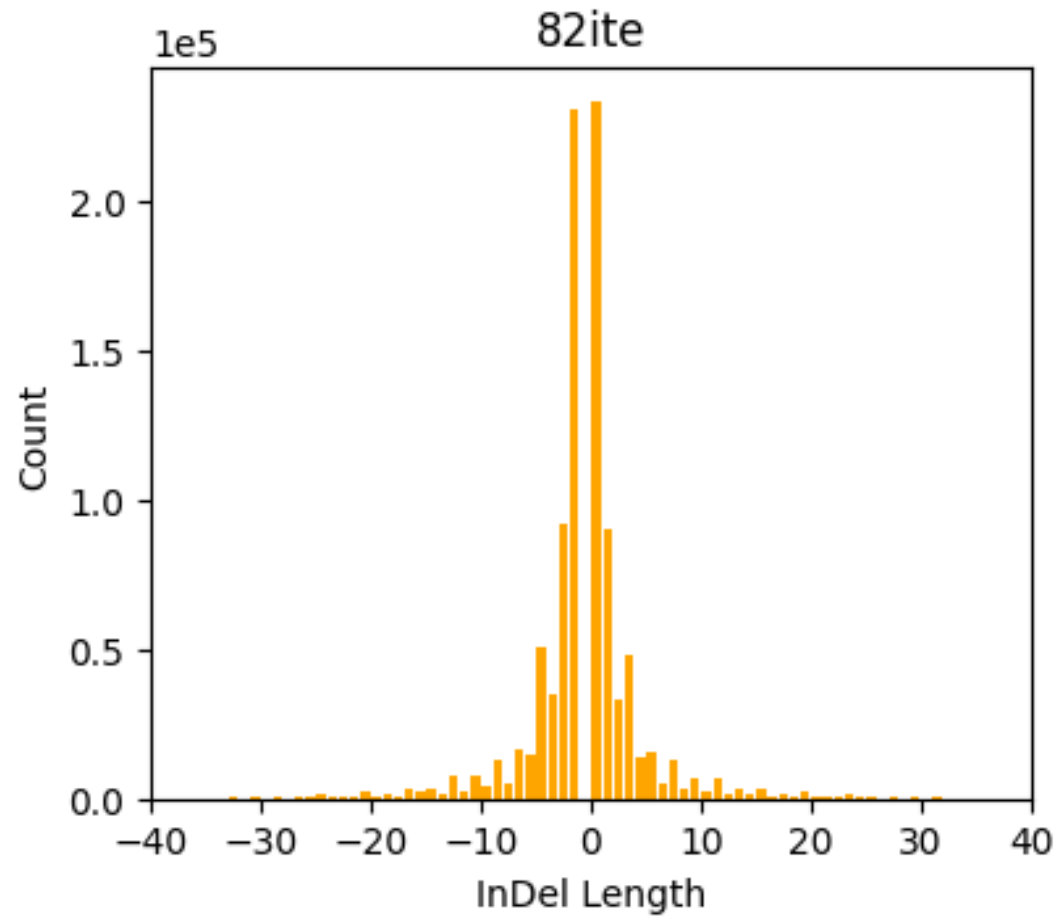
Number of indels by sample



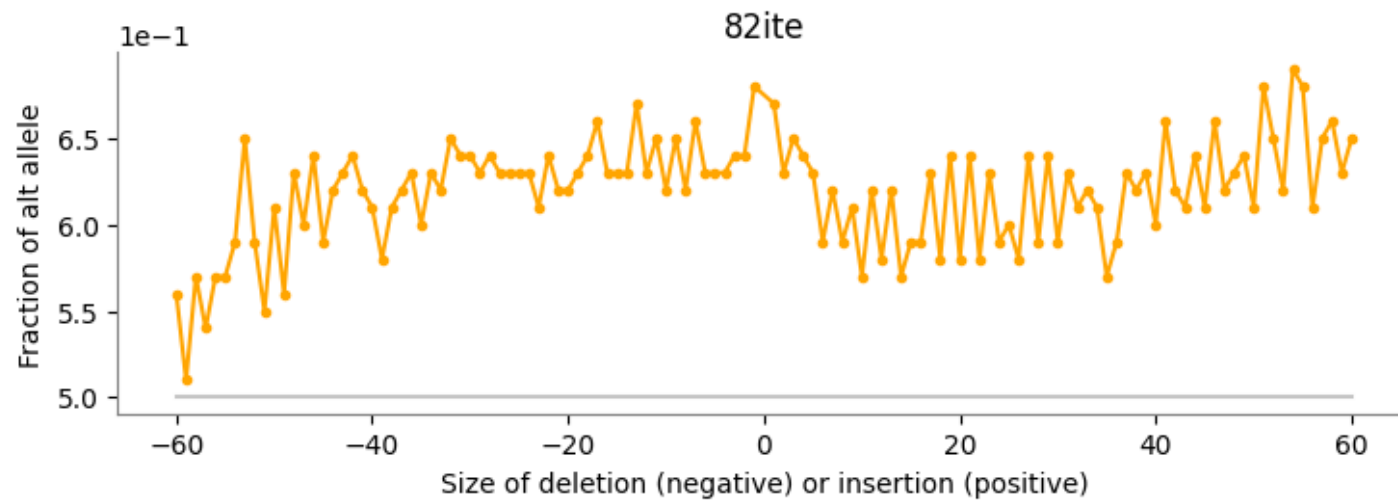
Ts/Tv stratified by QUAL



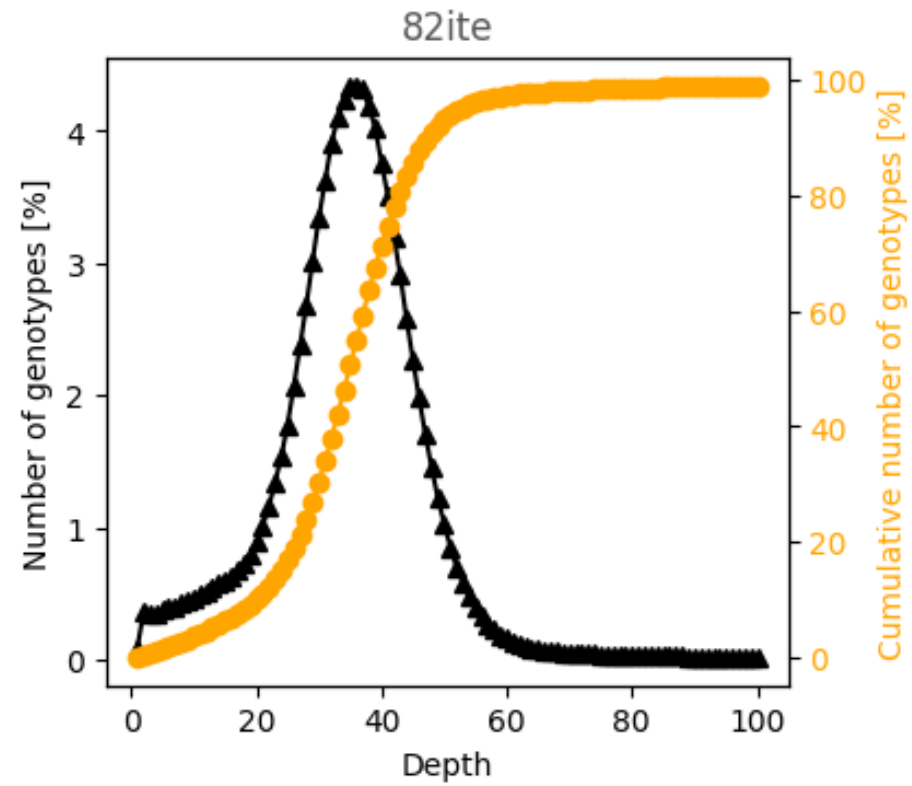
Indel distribution



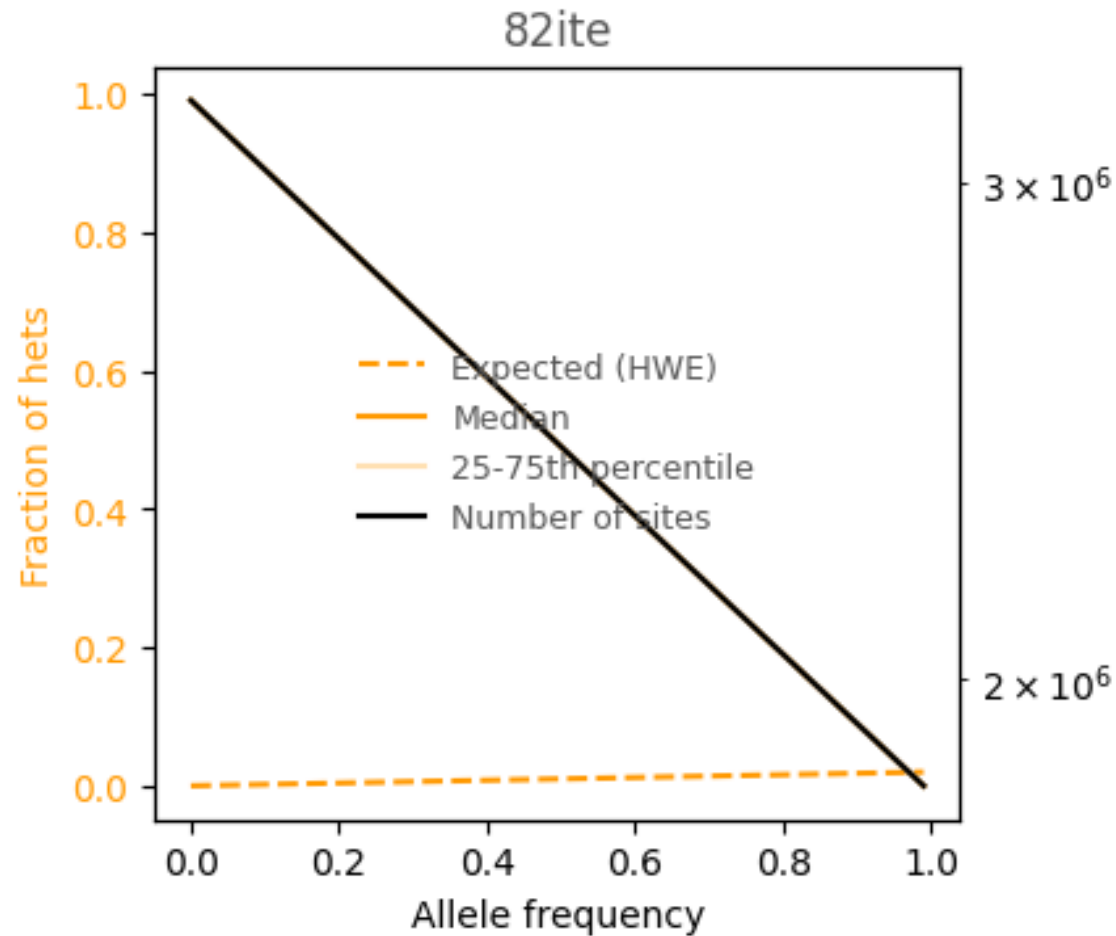
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

