

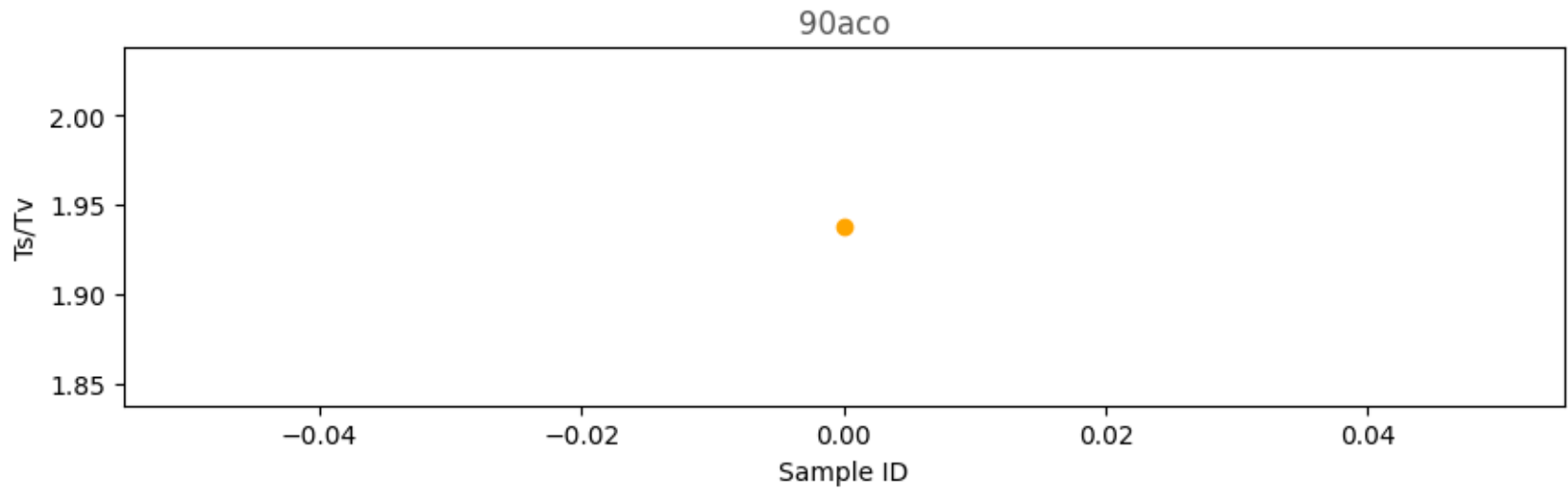
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90aco	4,208,662	1.94	1.94	976,077	–	0	0
* frameshift ratio: out/(out+in)							

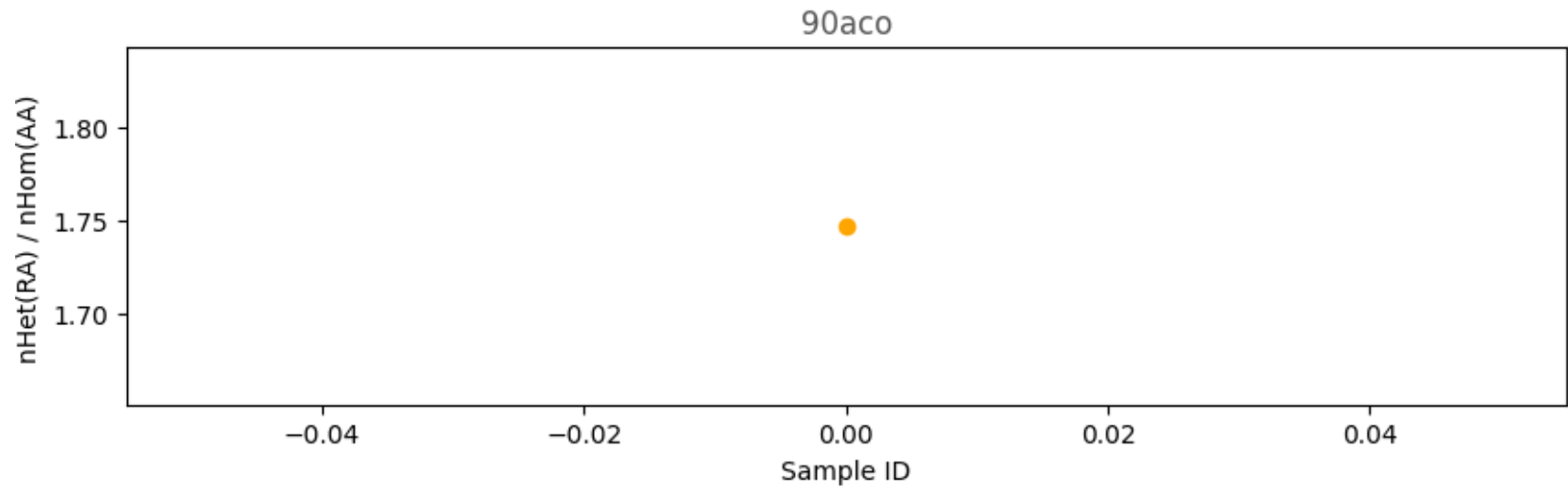
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90aco	63.6%	1.91	68.2%	104,659	1,931

- 90aco .. /ngc/projects2/gm/data/archive/2022/variants/snv/90acouthm-103876135804-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00858-220609_A00559_AHJ5LMDSX3-EXT_LAB_KA_NGCWGS-NGCWGS04410_22RKG012054x01_snv_germline_raw.haplotype_caller.vcf.gz

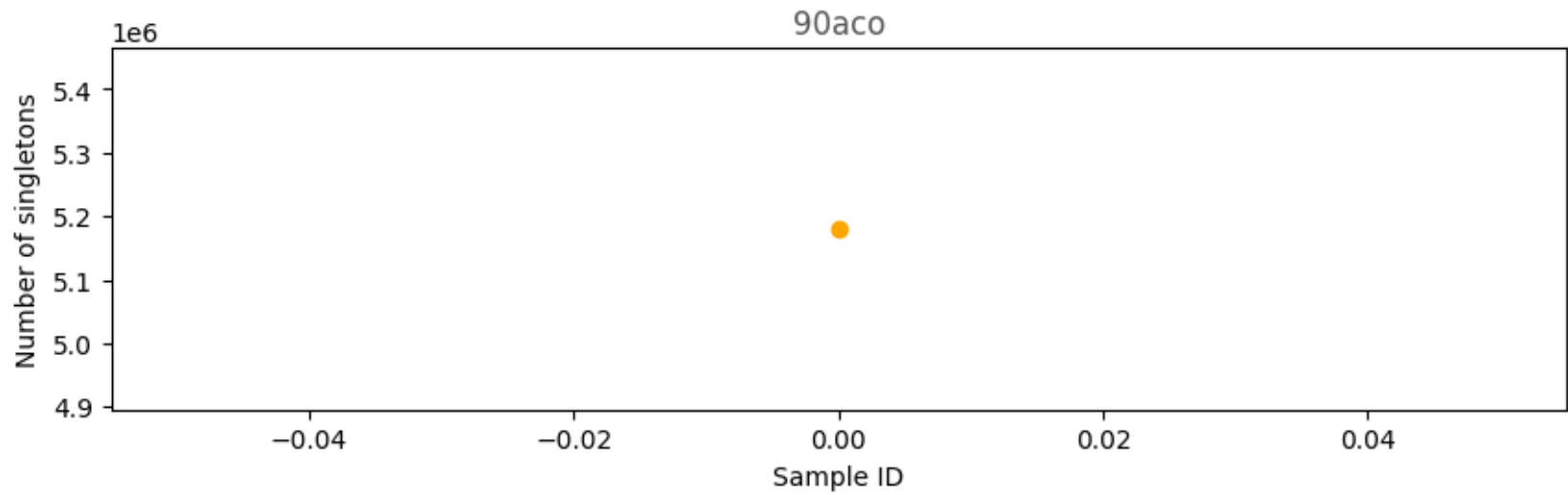
Ts/Tv by sample



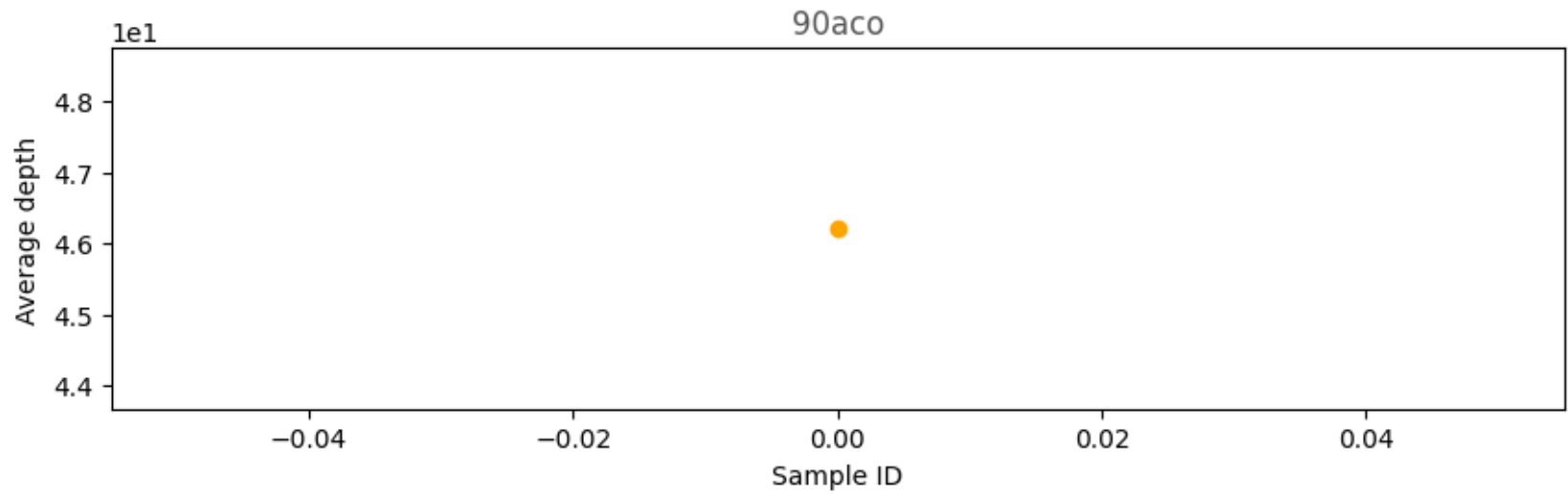
Hets vs non-ref Homs by sample



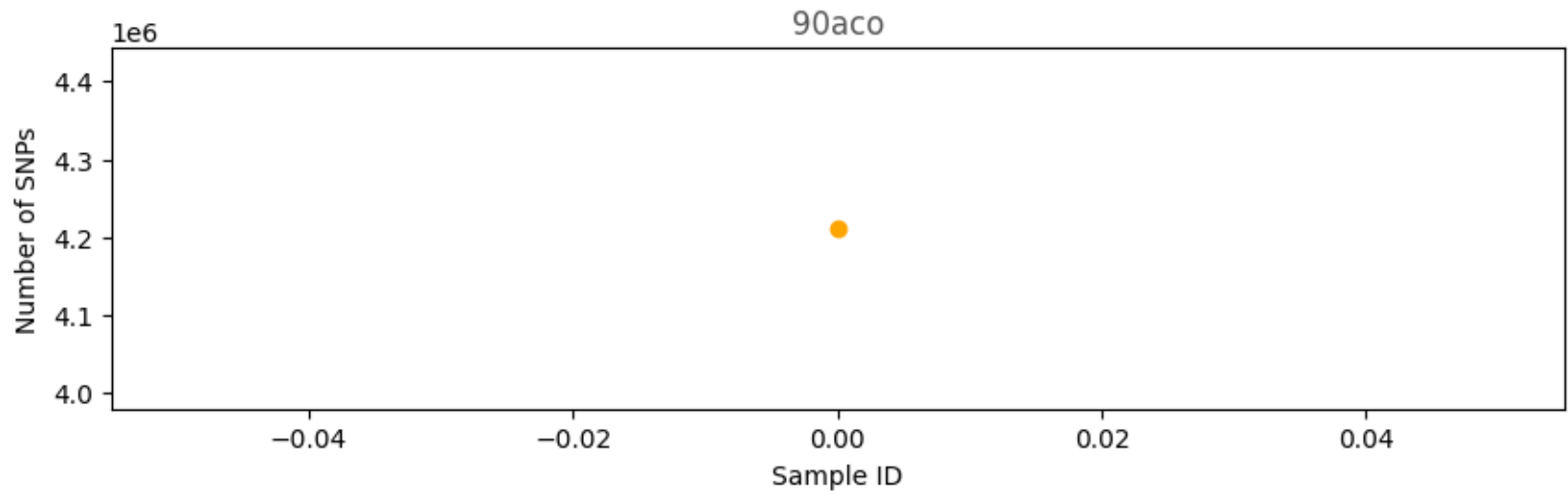
Singletons by sample (hets and homs)



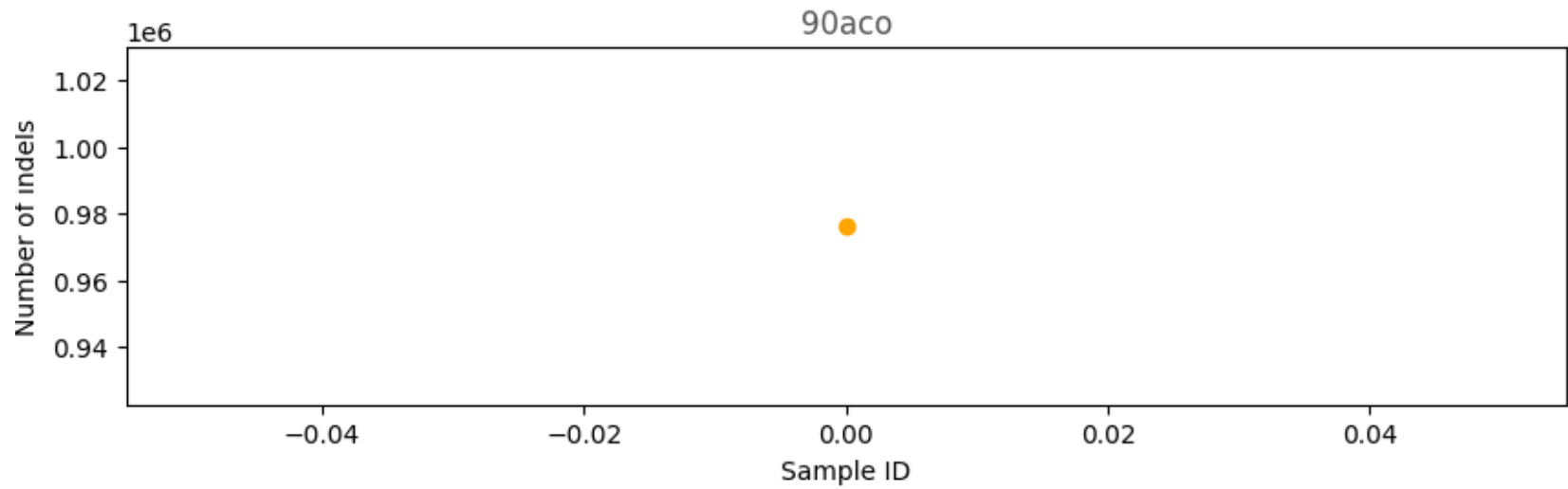
Average depth by sample



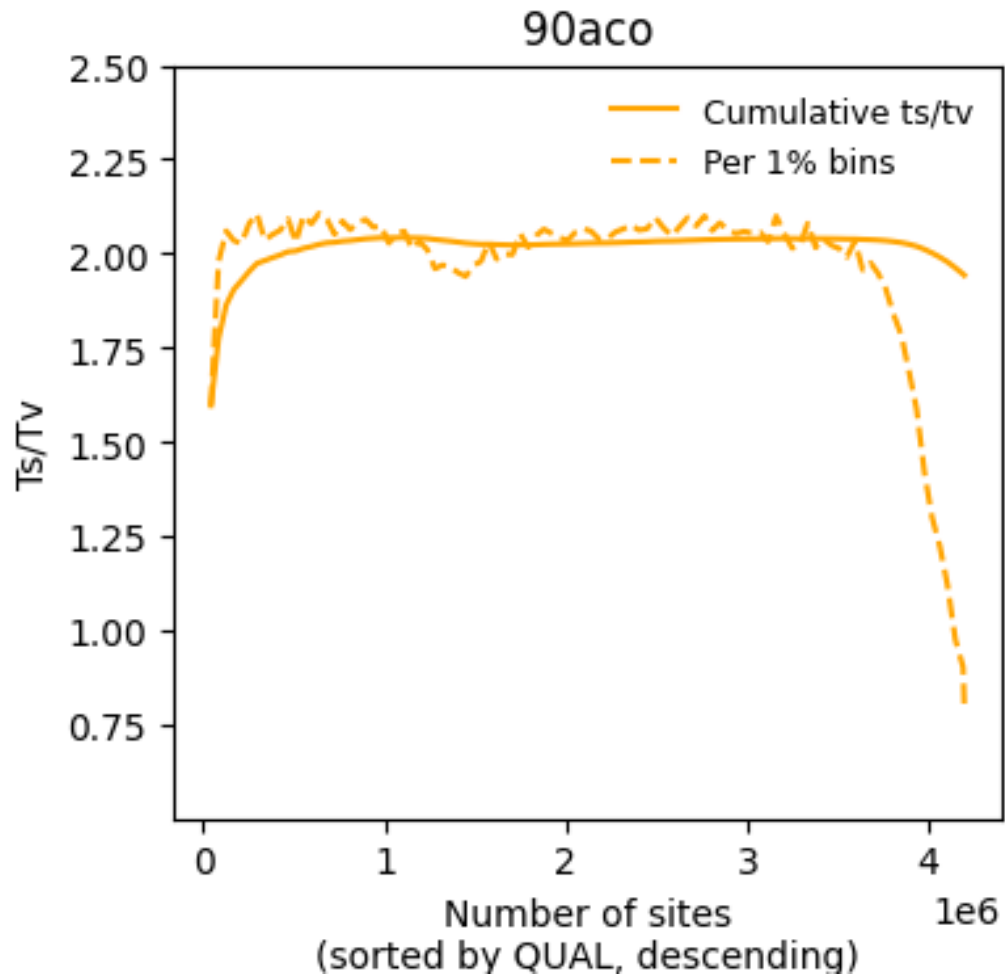
Number of SNPs by sample



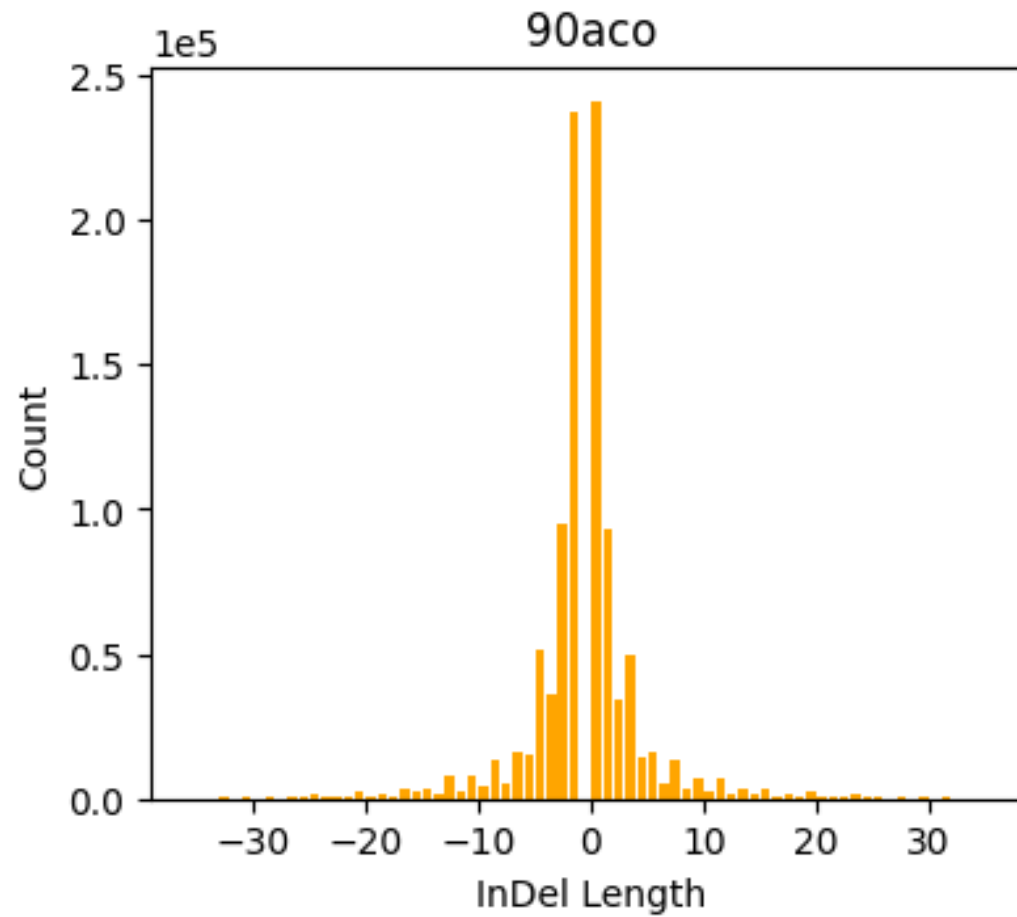
Number of indels by sample



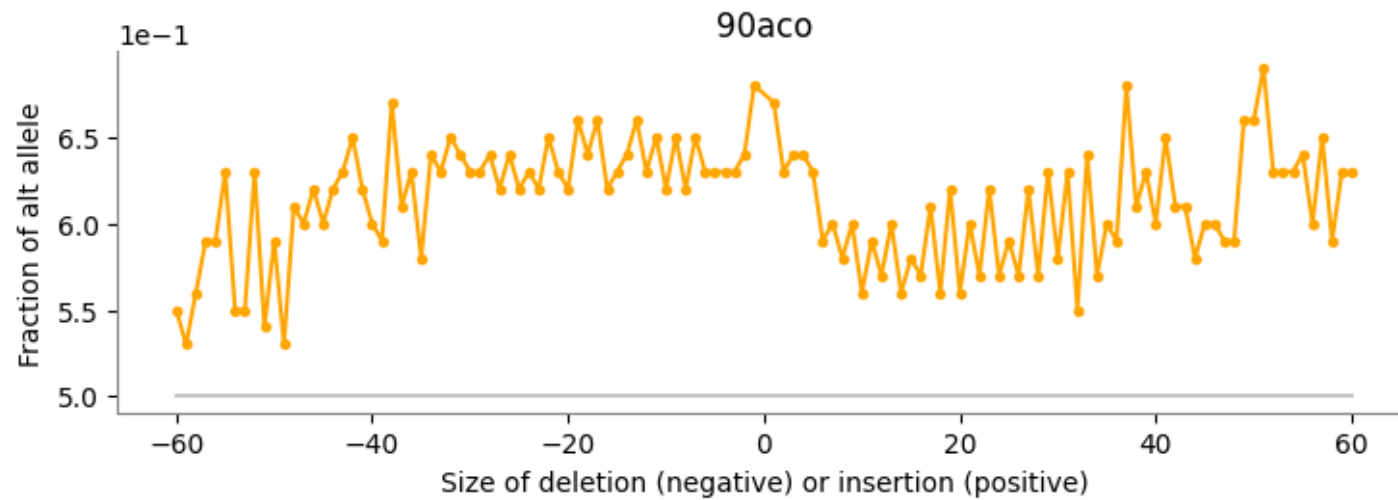
Ts/Tv stratified by QUAL



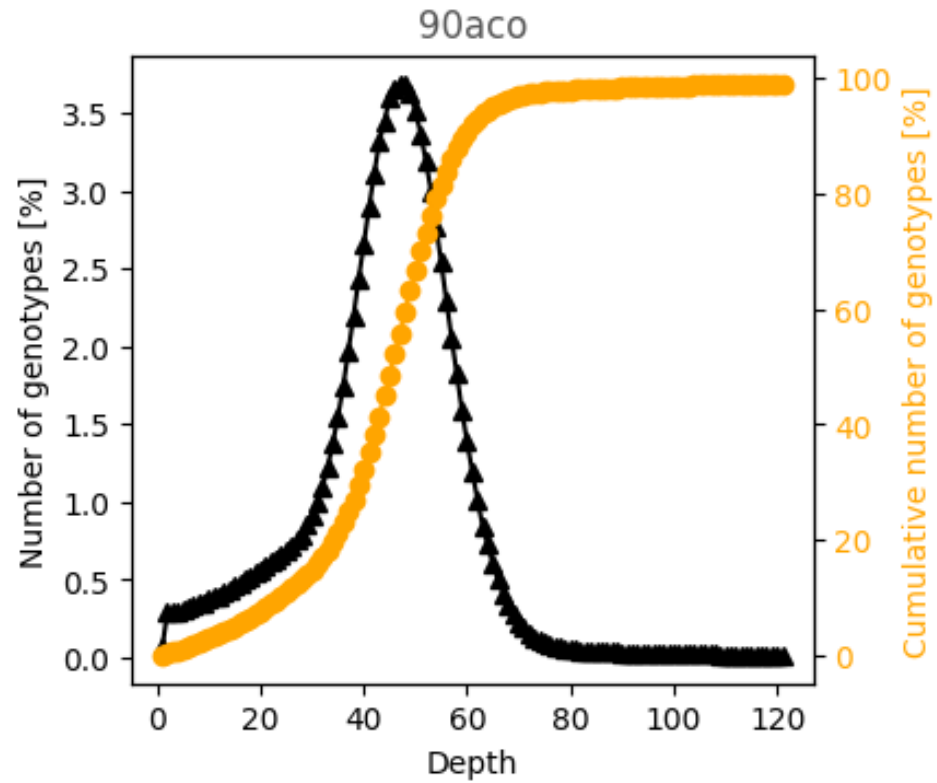
Indel distribution



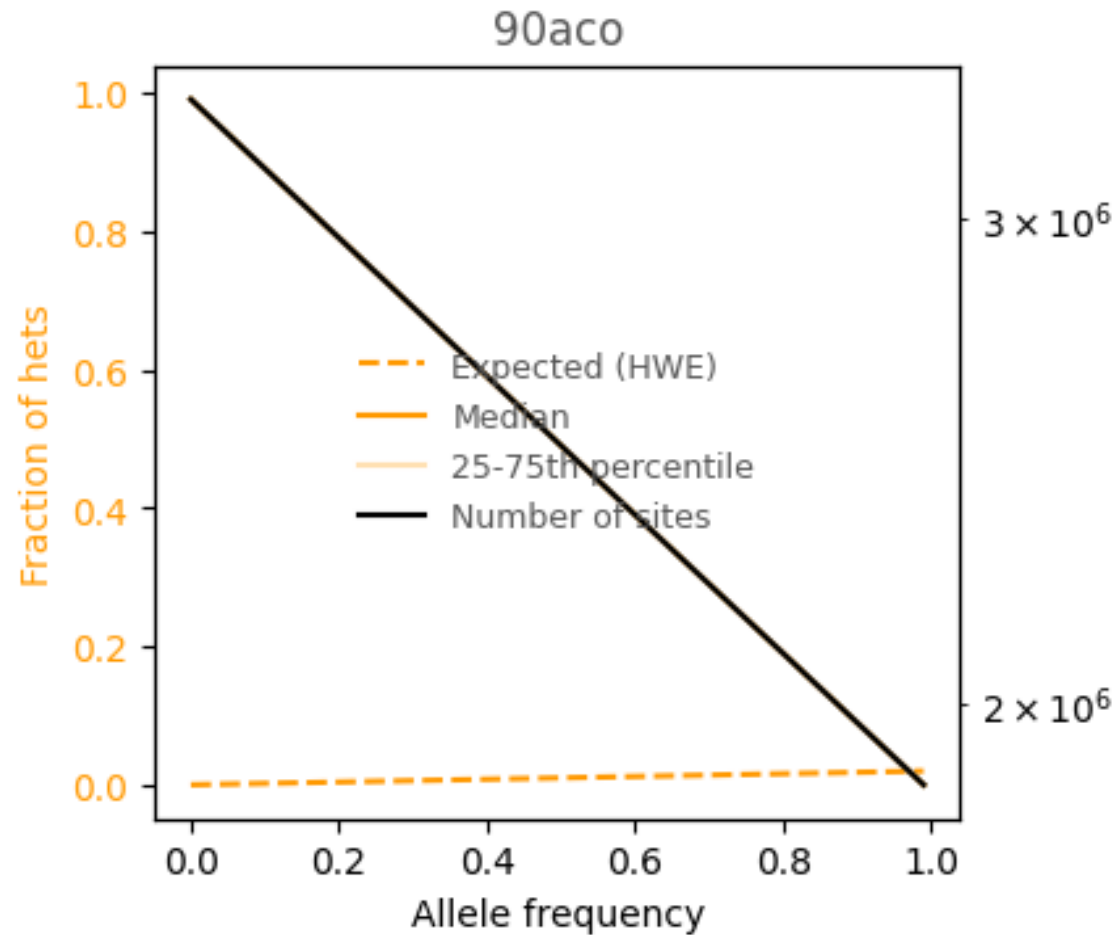
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

