

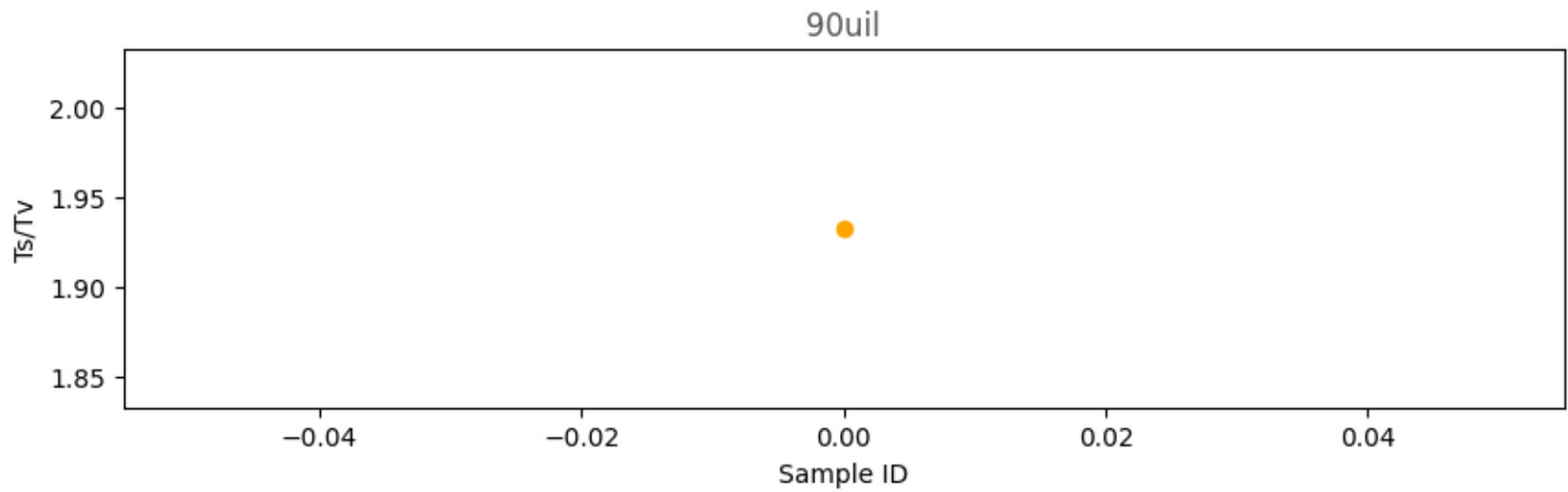
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90uil	4,055,894	1.93	1.94	938,525	–	0	0
* frameshift ratio: out/(out+in)							

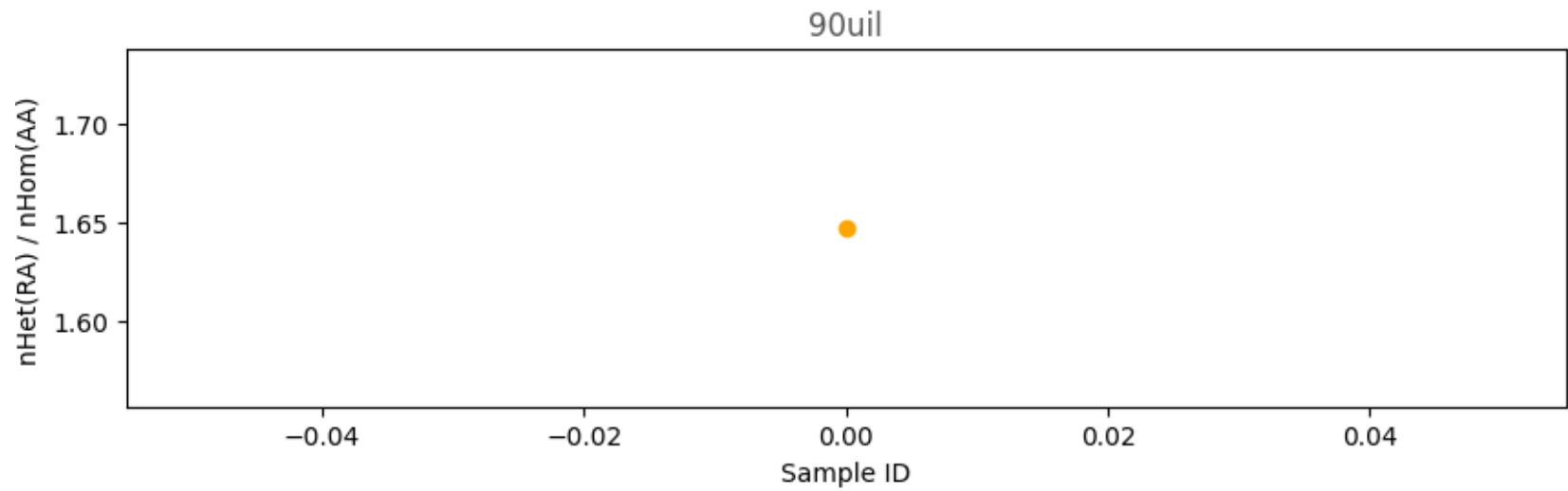
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90uil	62.2%	1.90	66.6%	95,581	1,921

- 90uil .. /ngc/projects2/gm/data/archive/2022/variants/snv/90uilerdm-103893002965-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220720_A00559_AHMHLDX3-RHGM_LABKA_WGSA_KUT-WGSAKUT04651_22RKG017238x01_snv_germline_raw.haplotype_caller.vcf.gz

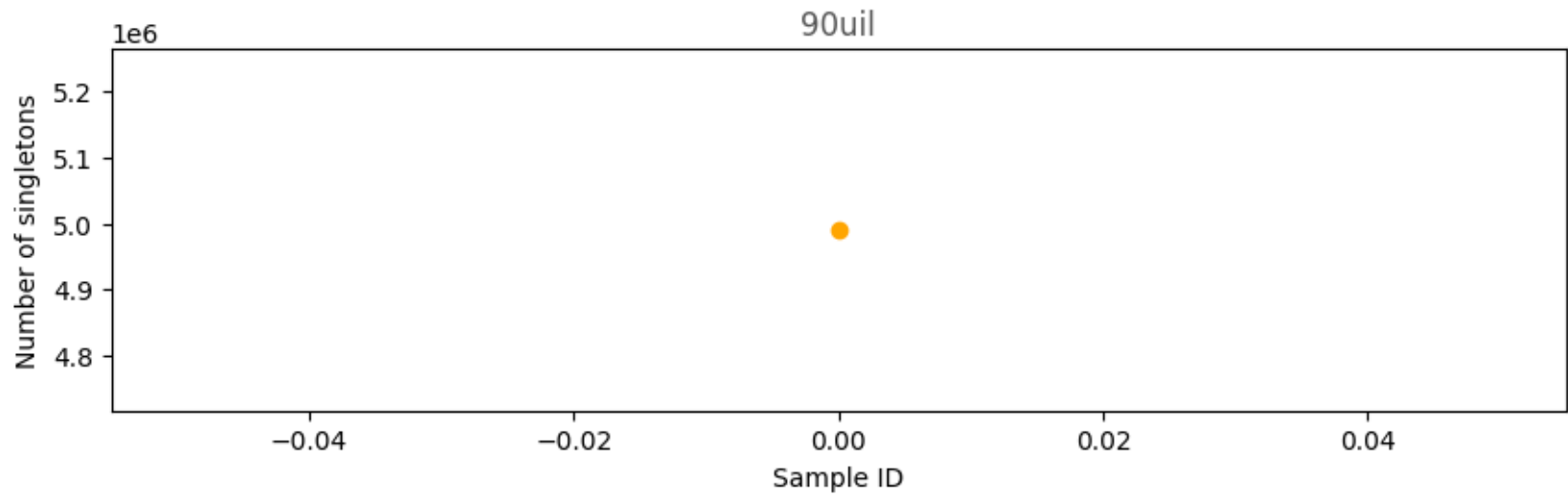
Ts/Tv by sample



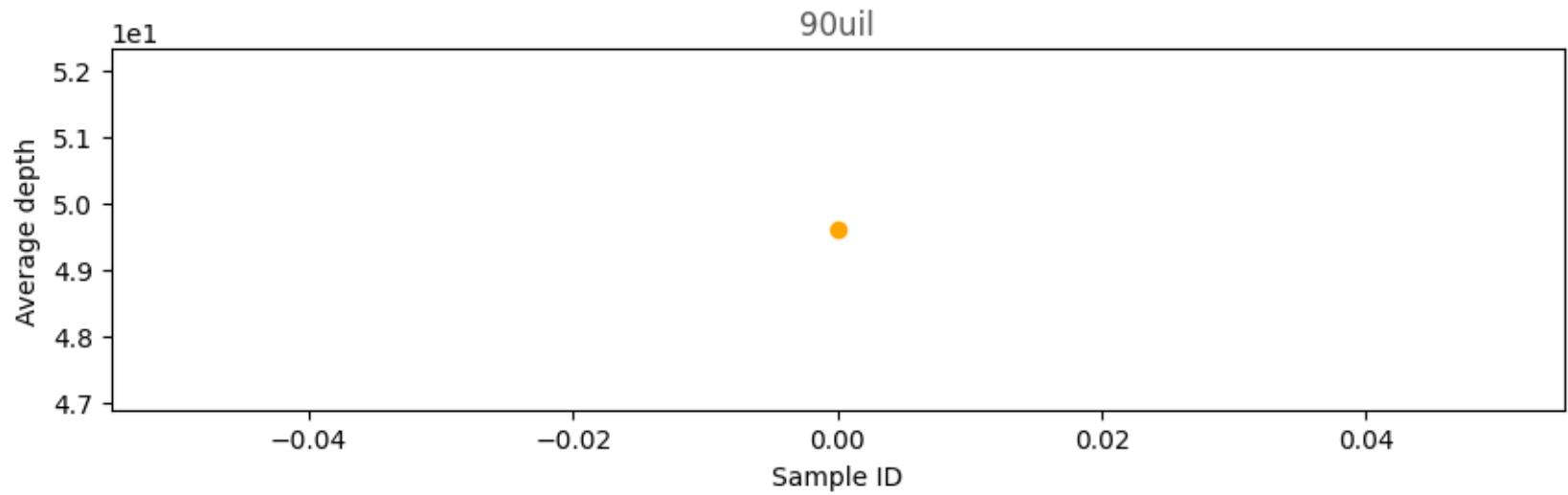
Hets vs non-ref Homs by sample



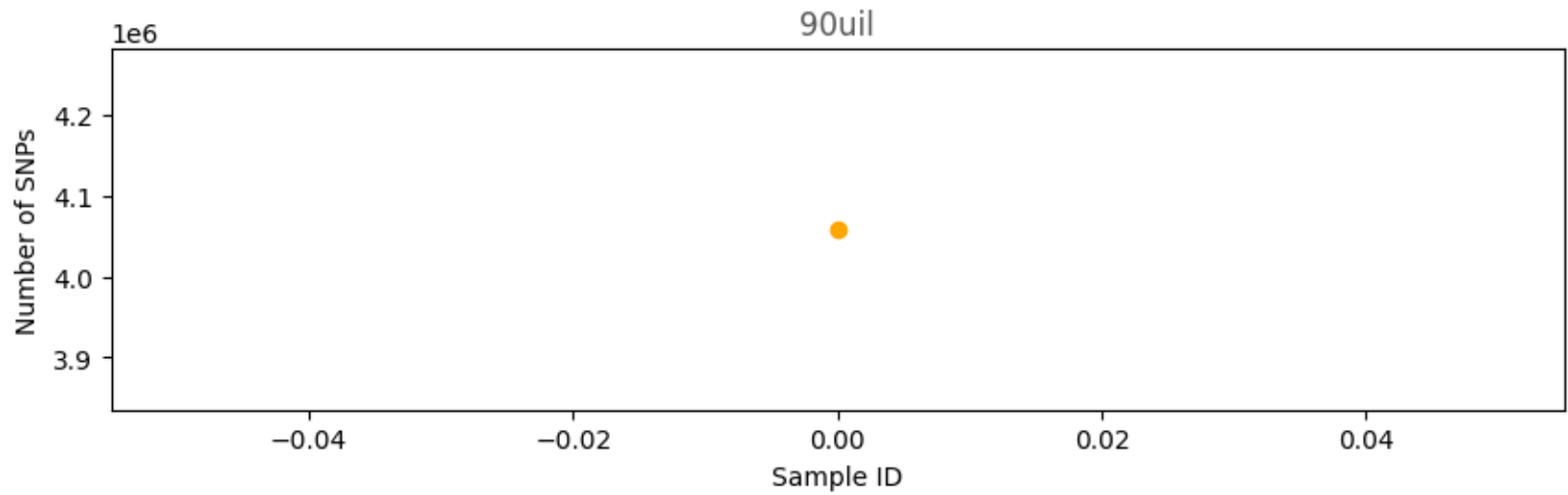
Singletons by sample (hets and homs)



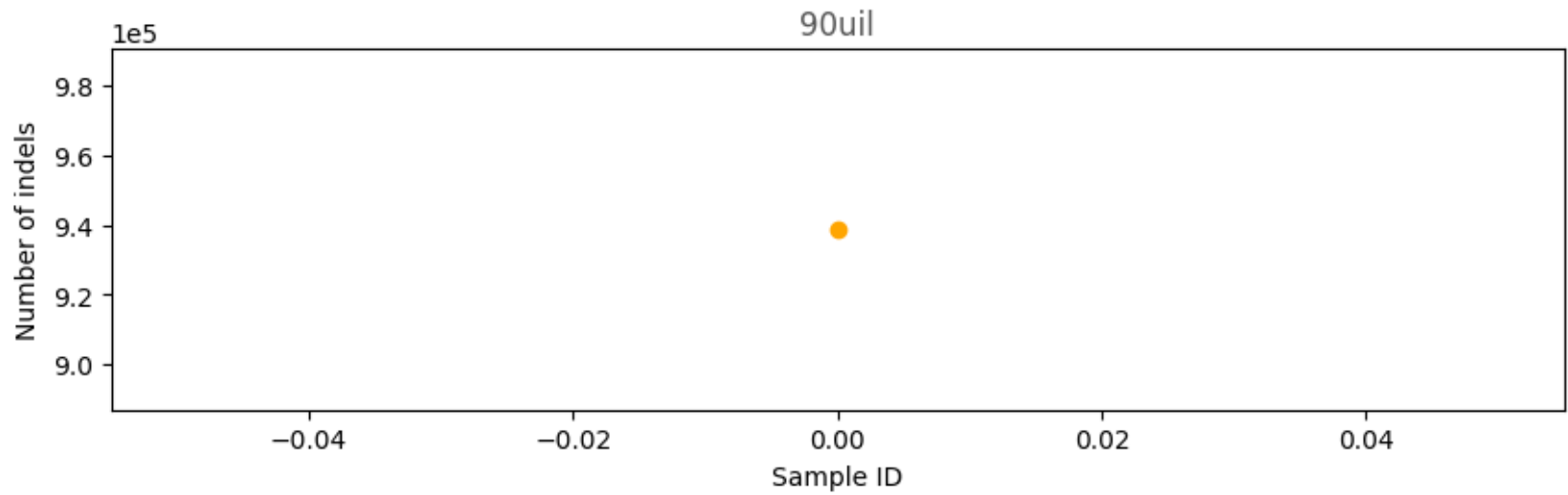
Average depth by sample



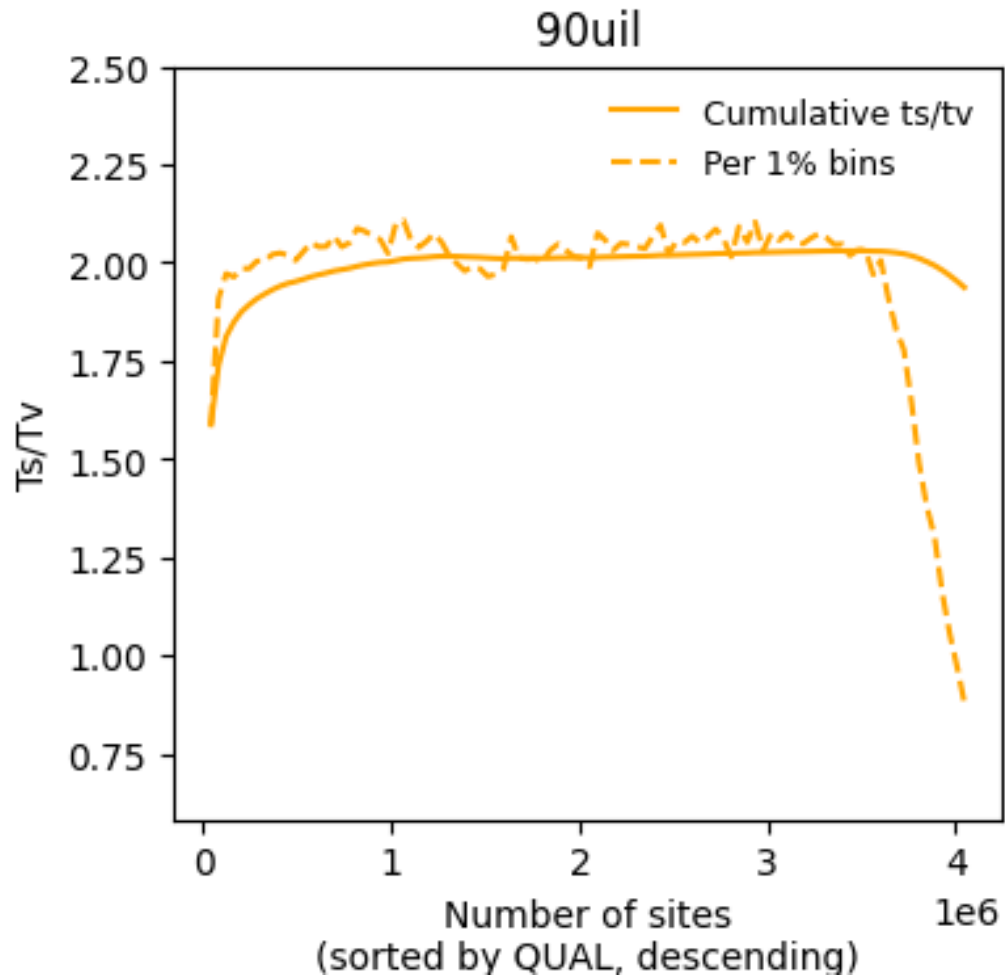
Number of SNPs by sample



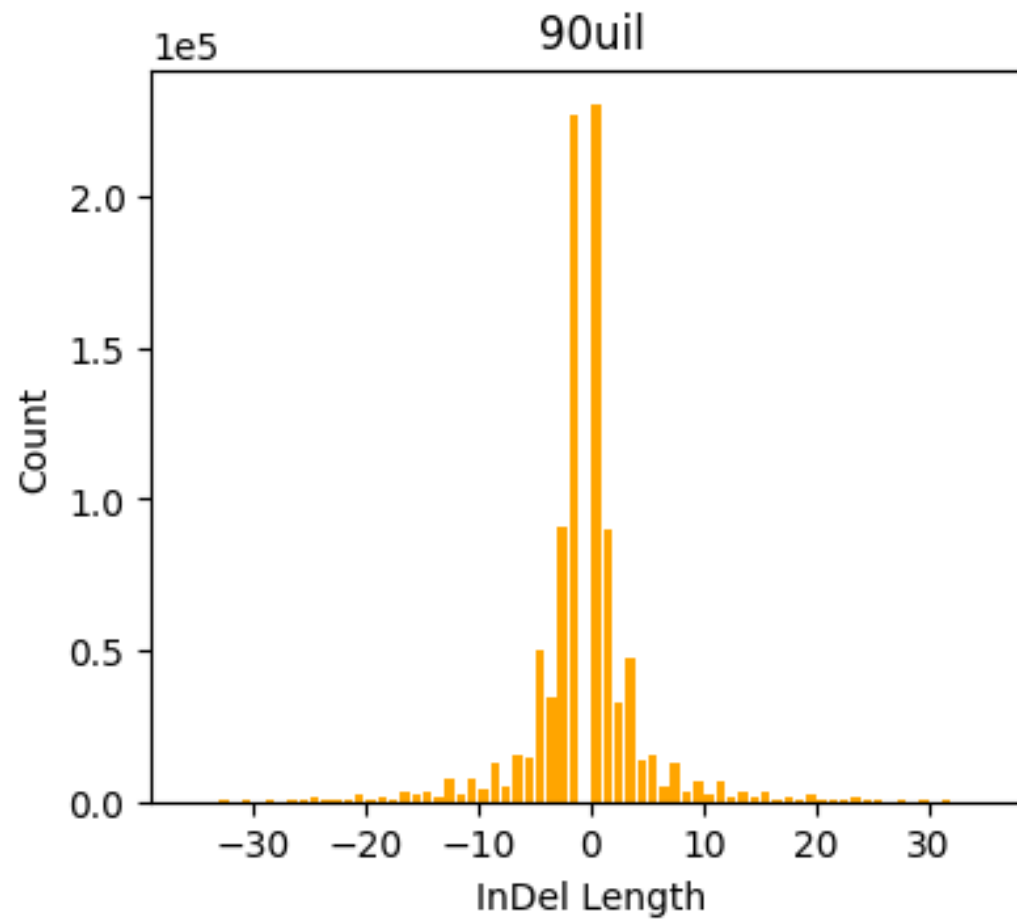
Number of indels by sample



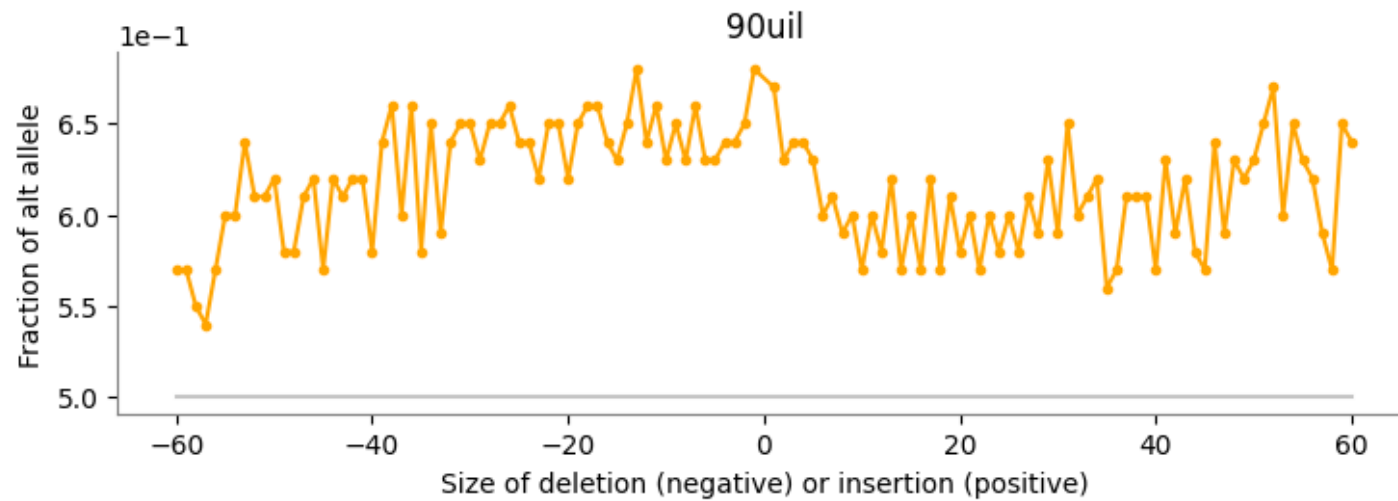
Ts/Tv stratified by QUAL



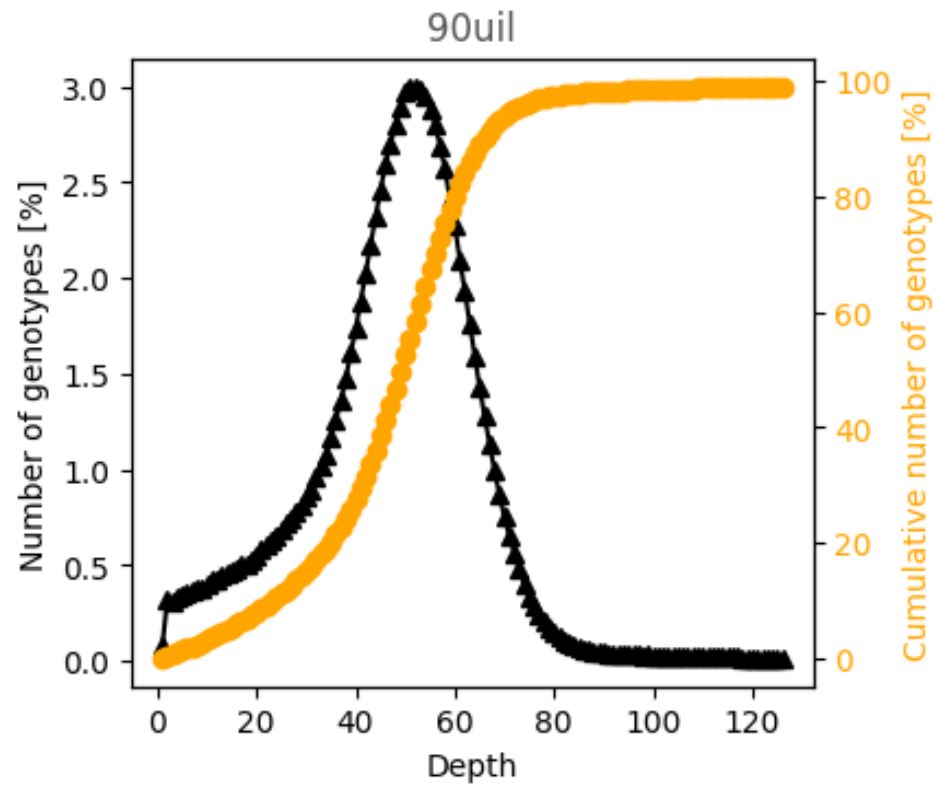
Indel distribution



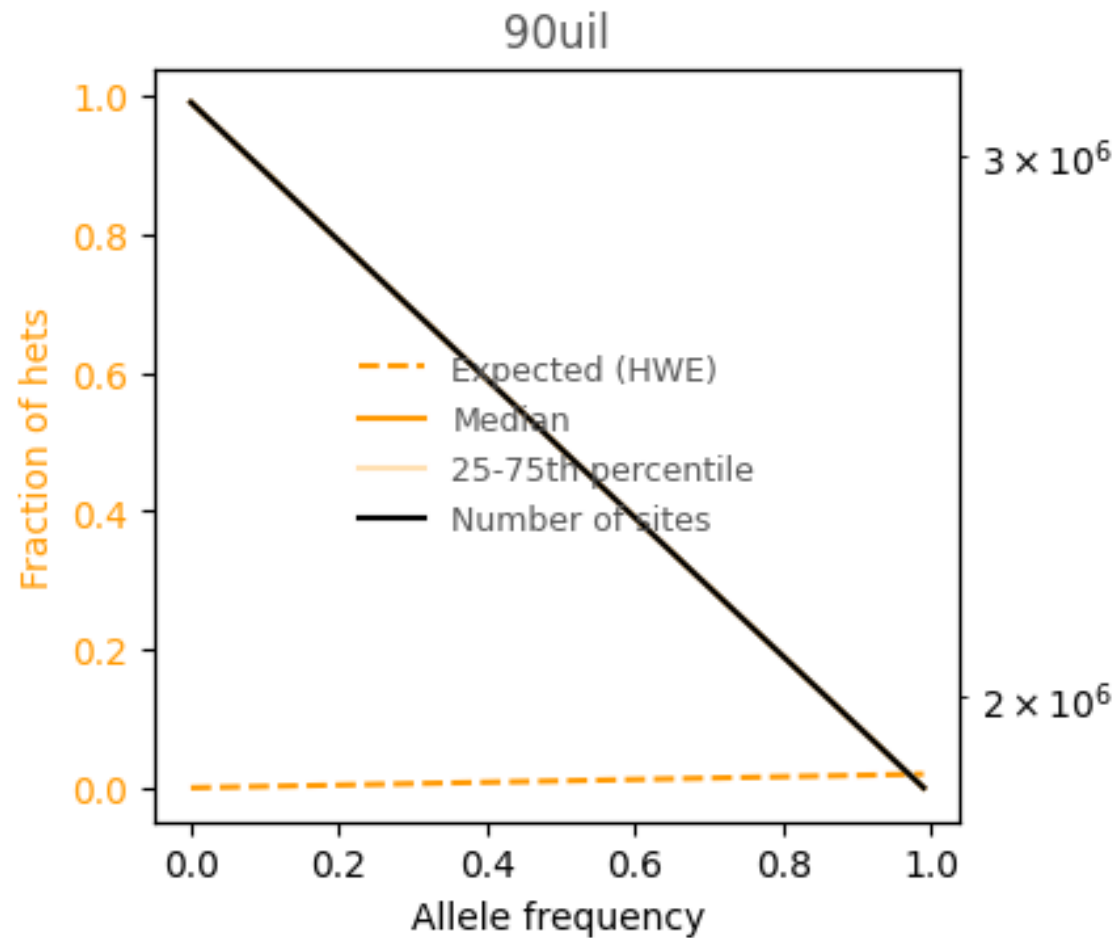
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

