

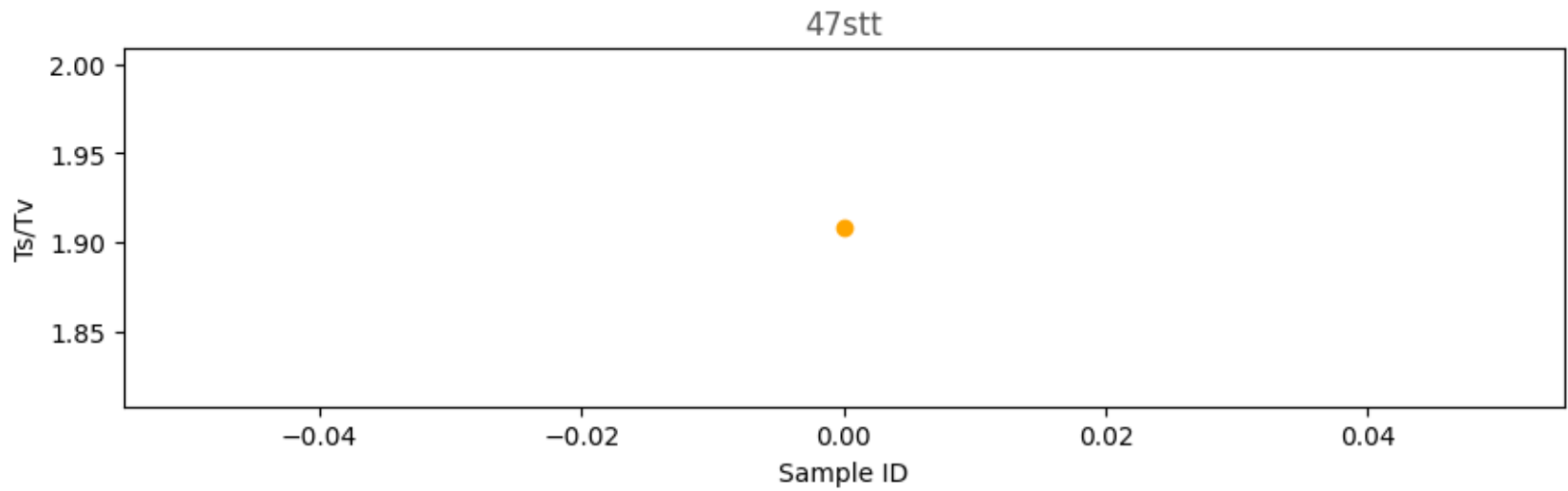
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
47stt	4,155,211	1.91	1.91	976,698	–	0	0
* frameshift ratio: out/(out+in)							

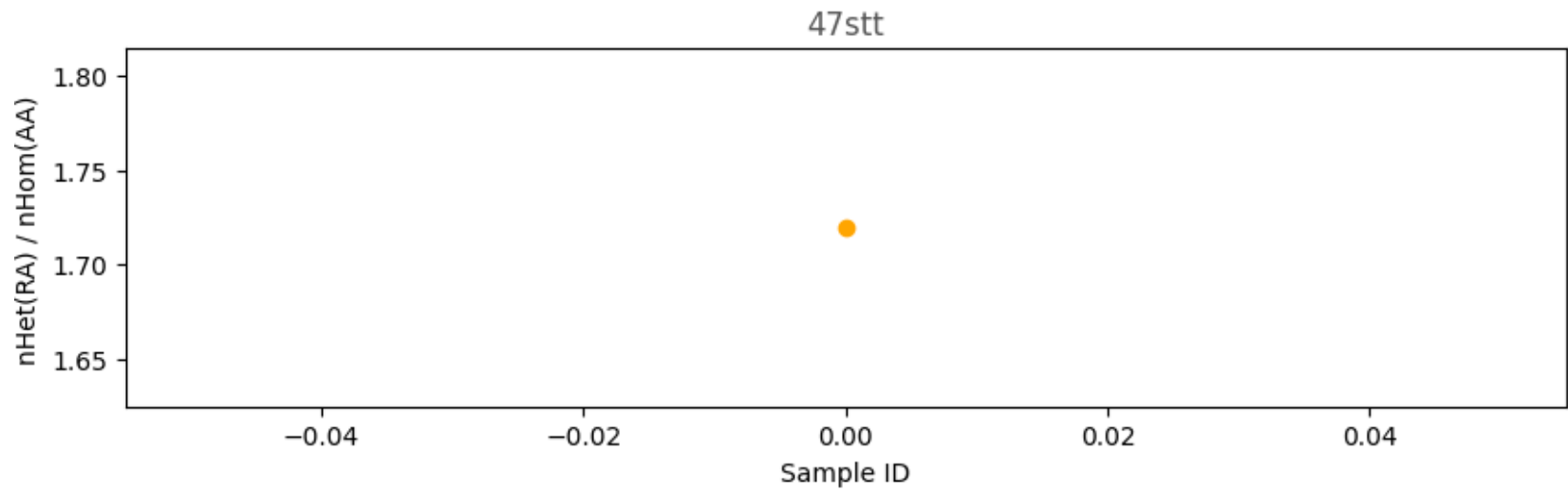
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
47stt	63.2%	1.87	69.3%	114,164	2,277

- 47stt .. /ngc/projects2/gm/data/archive/2022/variants/snv/47stttwif-103843543229-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220304_A01176_AHCFKFDSX3-RHGM_LABKA_WGS-WGS03936_snv_germline_raw.haplotype_caller.vcf.gz

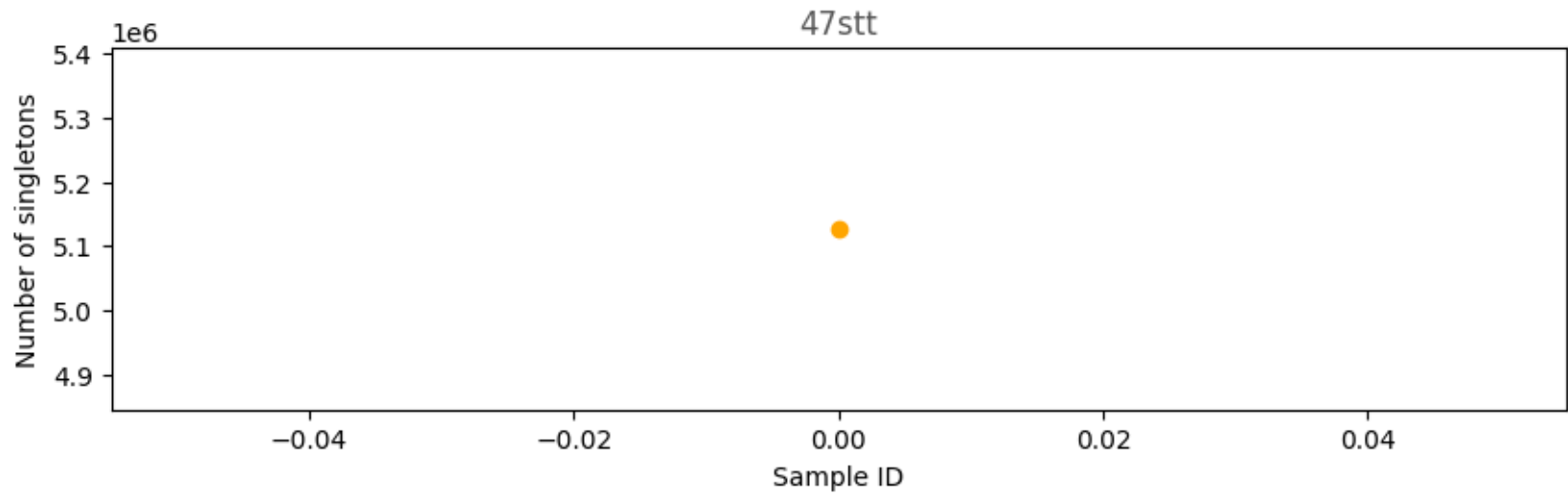
Ts/Tv by sample



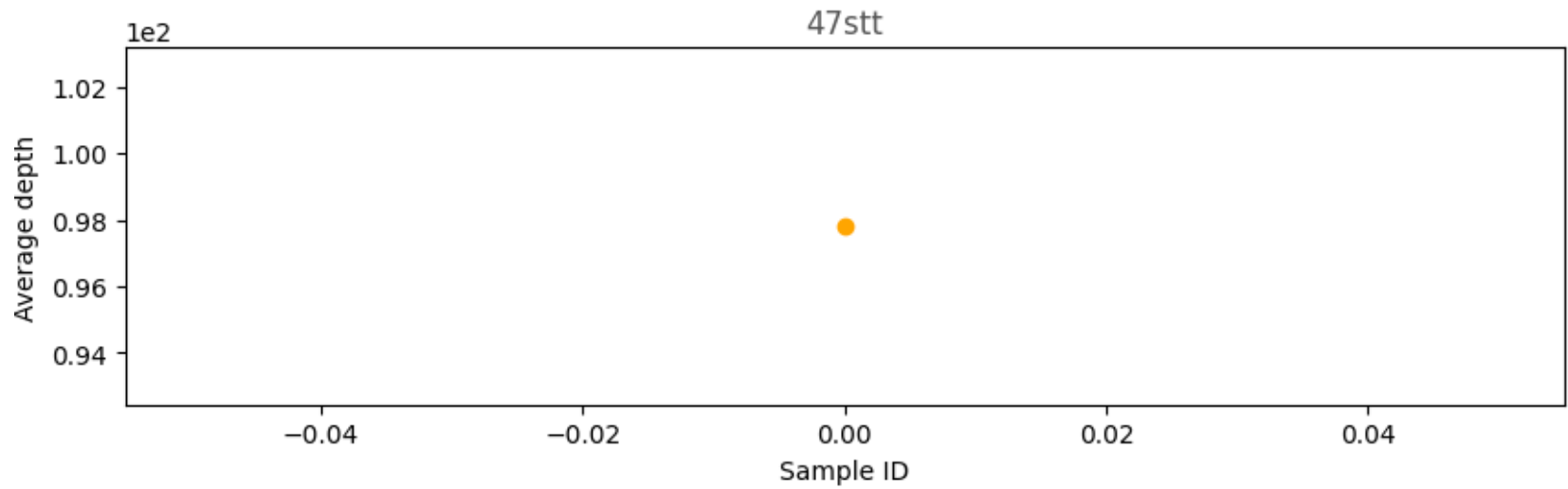
Hets vs non-ref Homs by sample



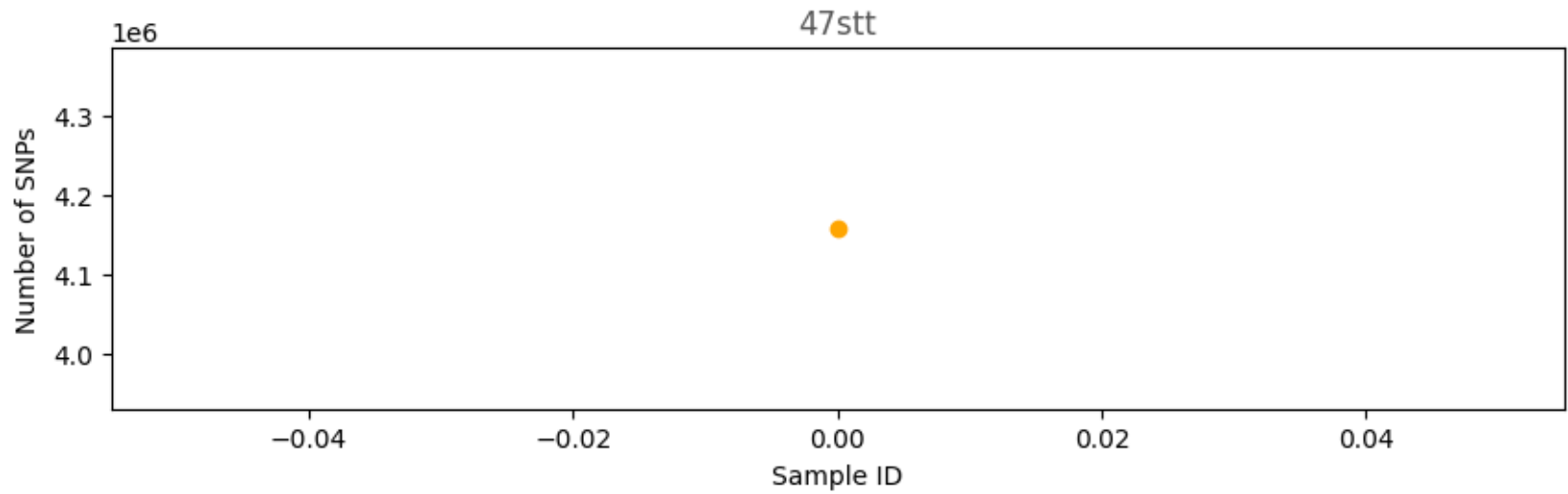
Singletons by sample (hets and homs)



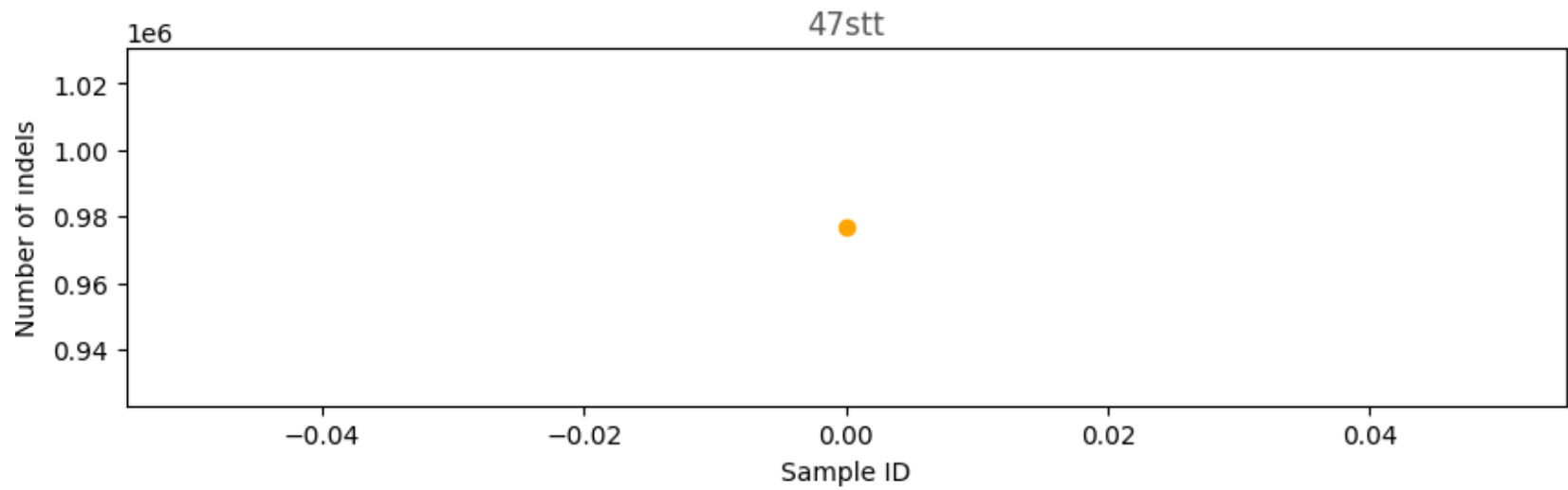
Average depth by sample



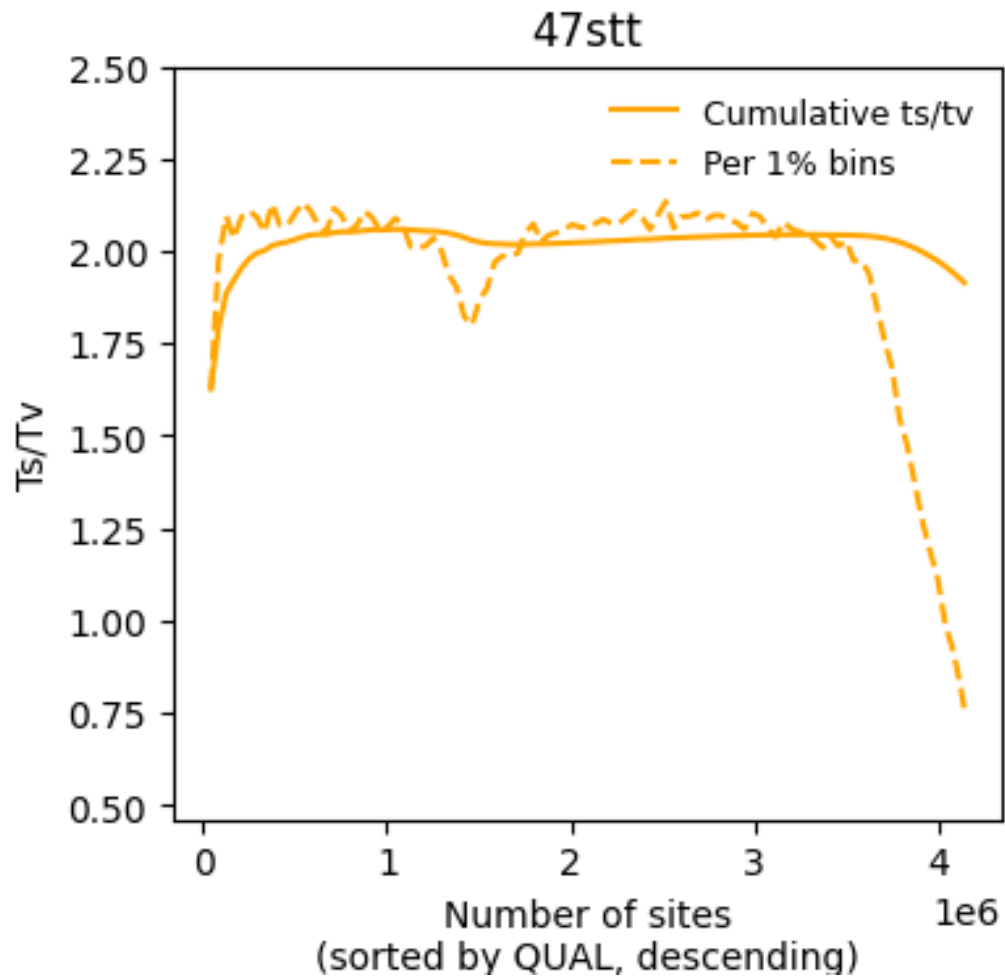
Number of SNPs by sample



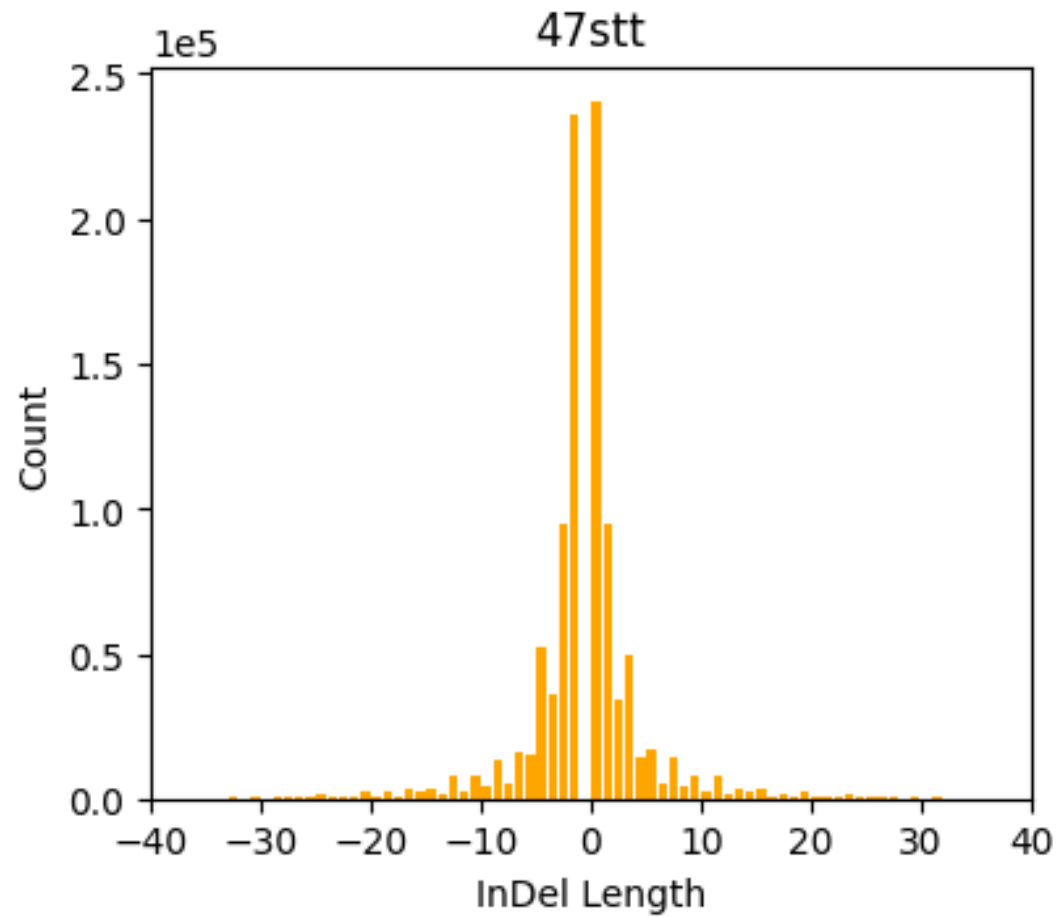
Number of indels by sample



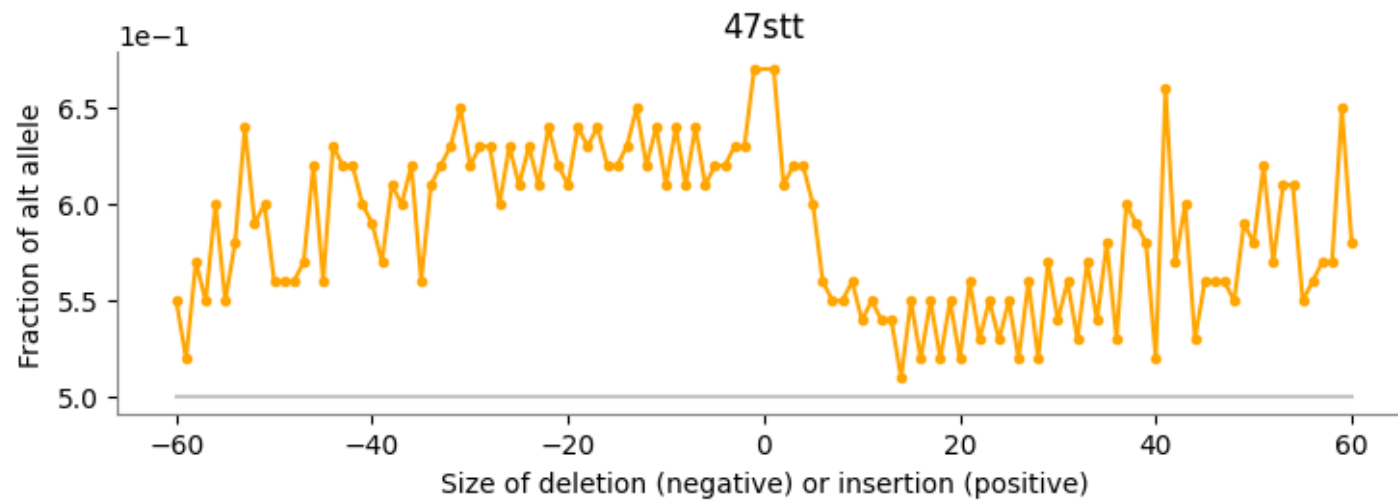
Ts/Tv stratified by QUAL



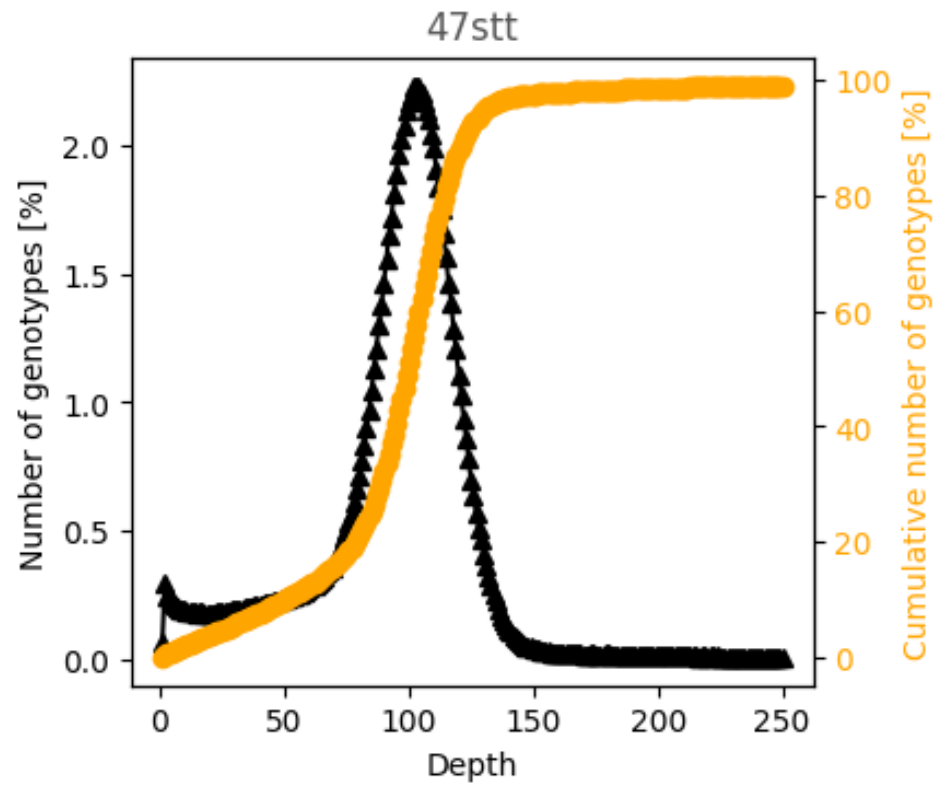
Indel distribution



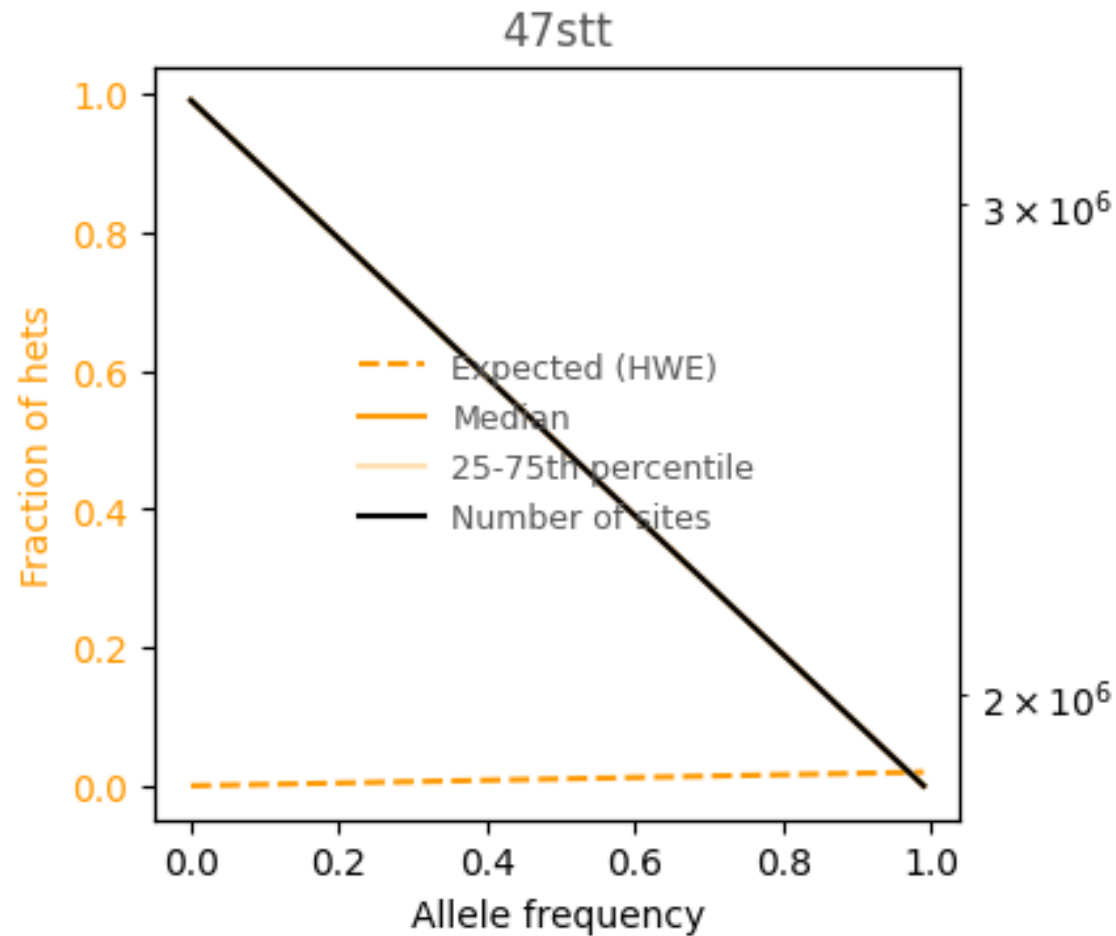
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

