

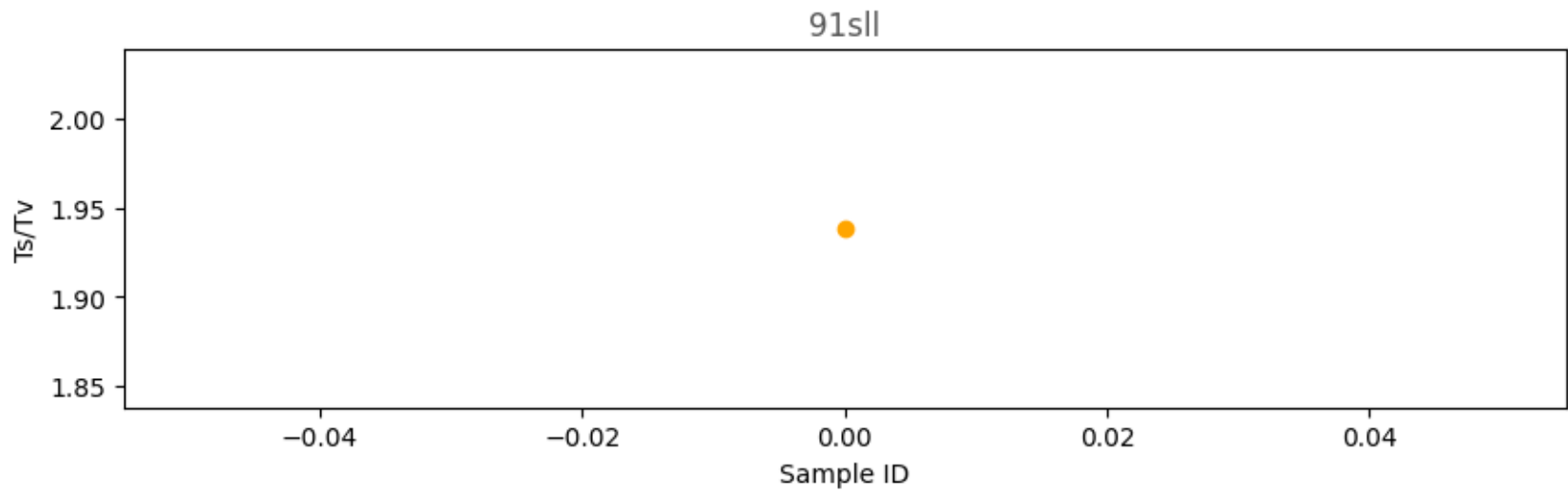
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
91sll	4,079,905	1.94	1.94	956,292	–	0	0
* frameshift ratio: out/(out+in)							

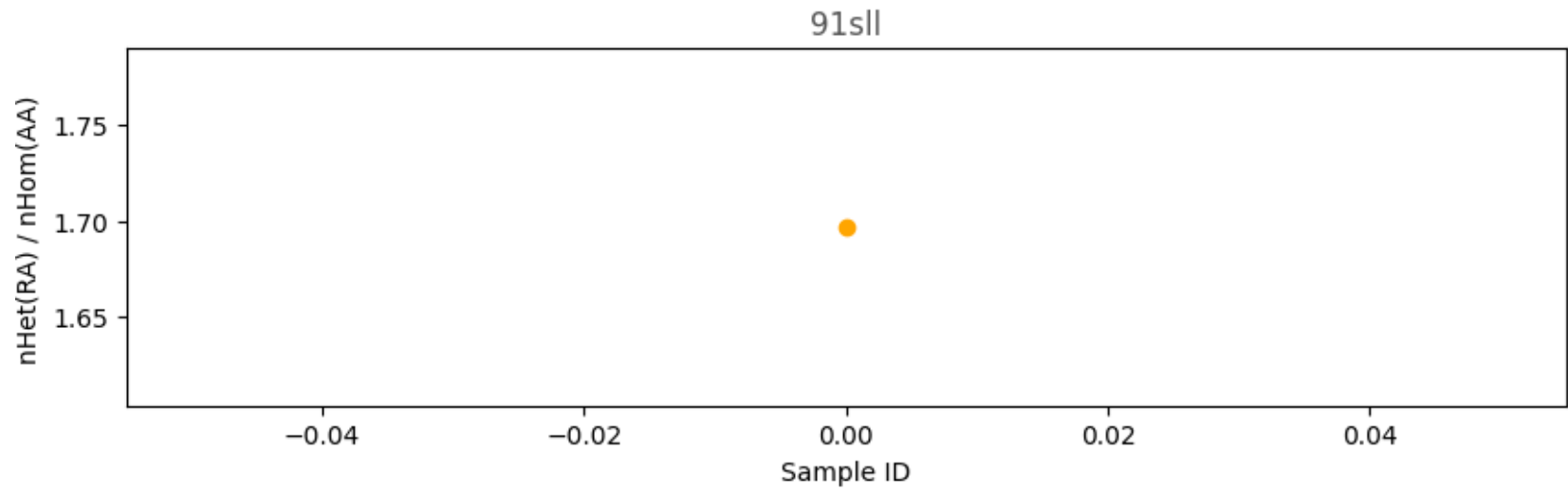
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
91sll	62.9%	1.91	68.2%	103,433	2,196

- 91sll .. /ngc/projects2/gm/data/archive/2022/variants/snv/91slllif-103794076972-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00364-211019_A01411_BHHHY3DSX2-EXT_LAB
KA_NGCWGS-NGCWGS00263_21RKG014232x01_snv_germline_raw.haplotype_caller.vcf.gz

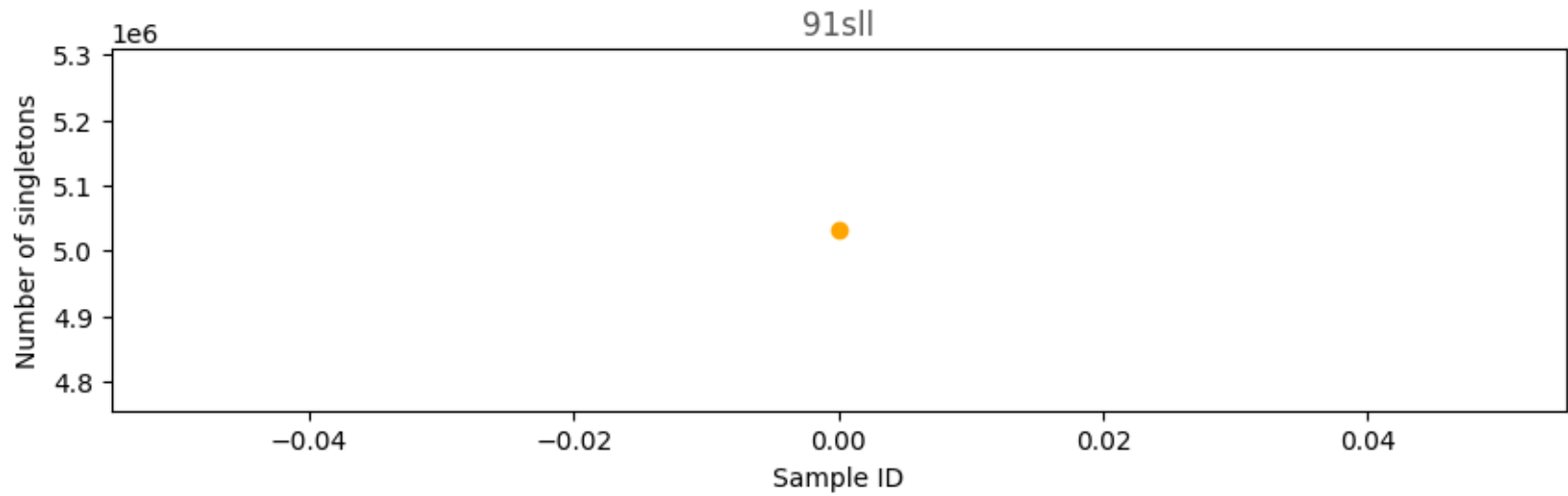
Ts/Tv by sample



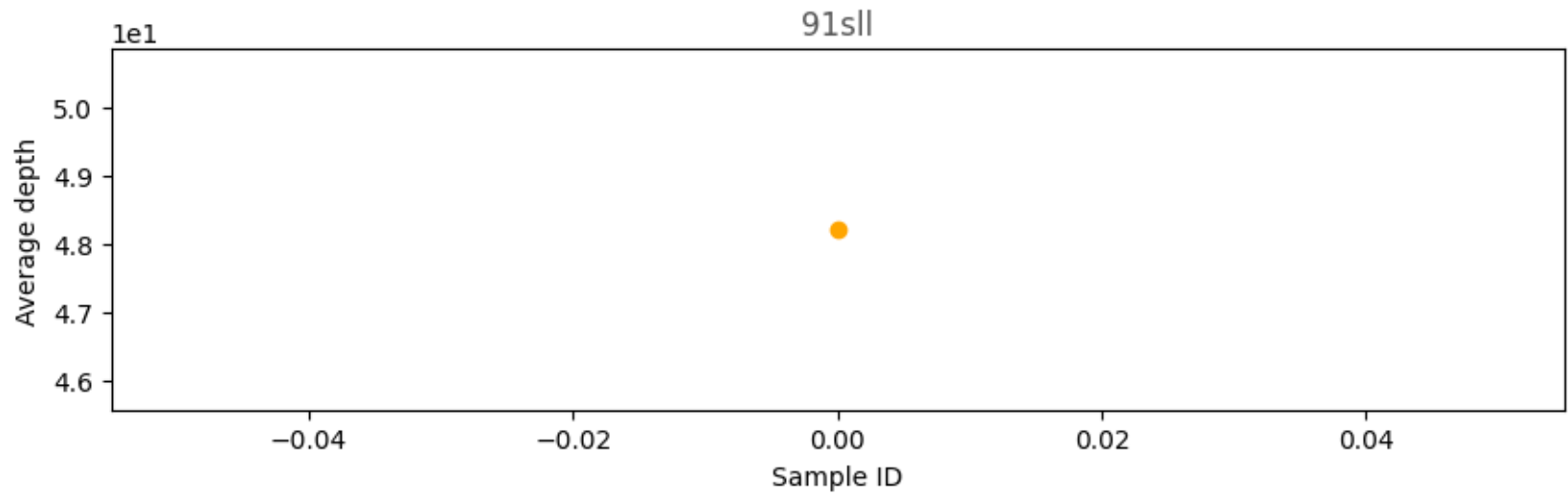
Hets vs non-ref Homs by sample



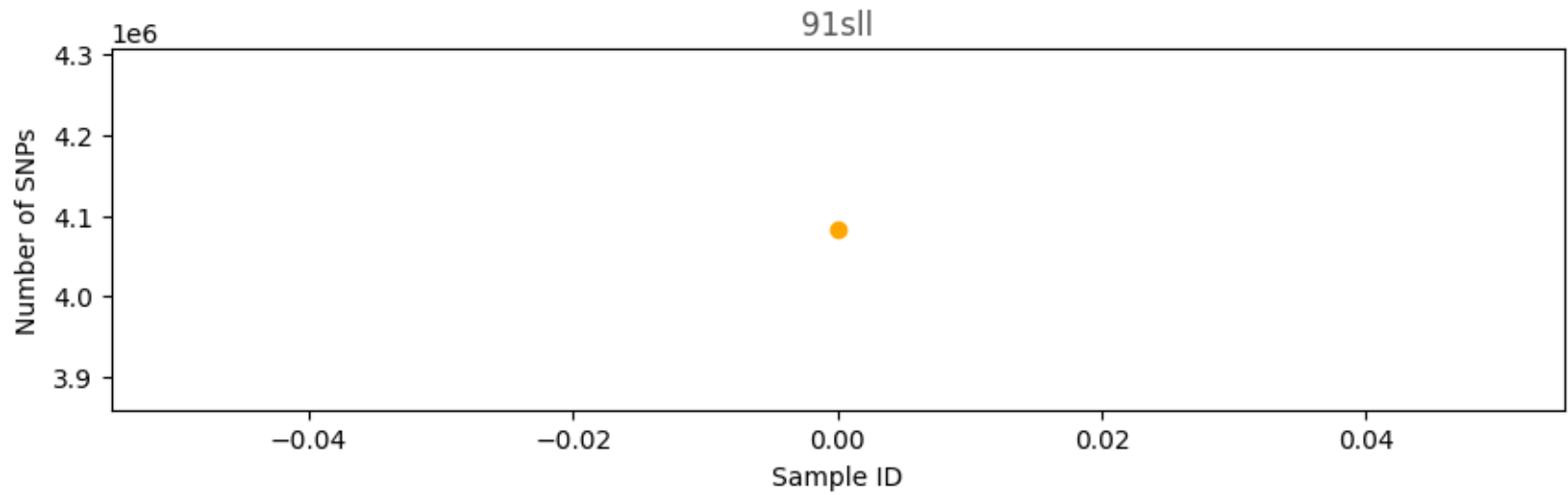
Singletons by sample (hets and homs)



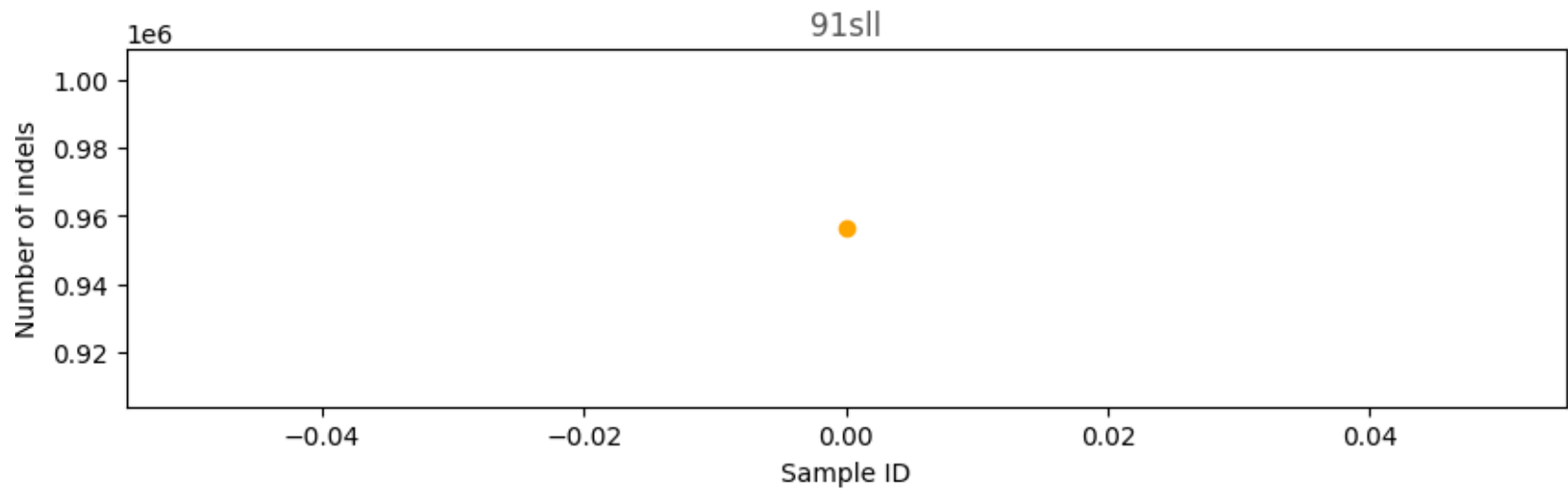
Average depth by sample



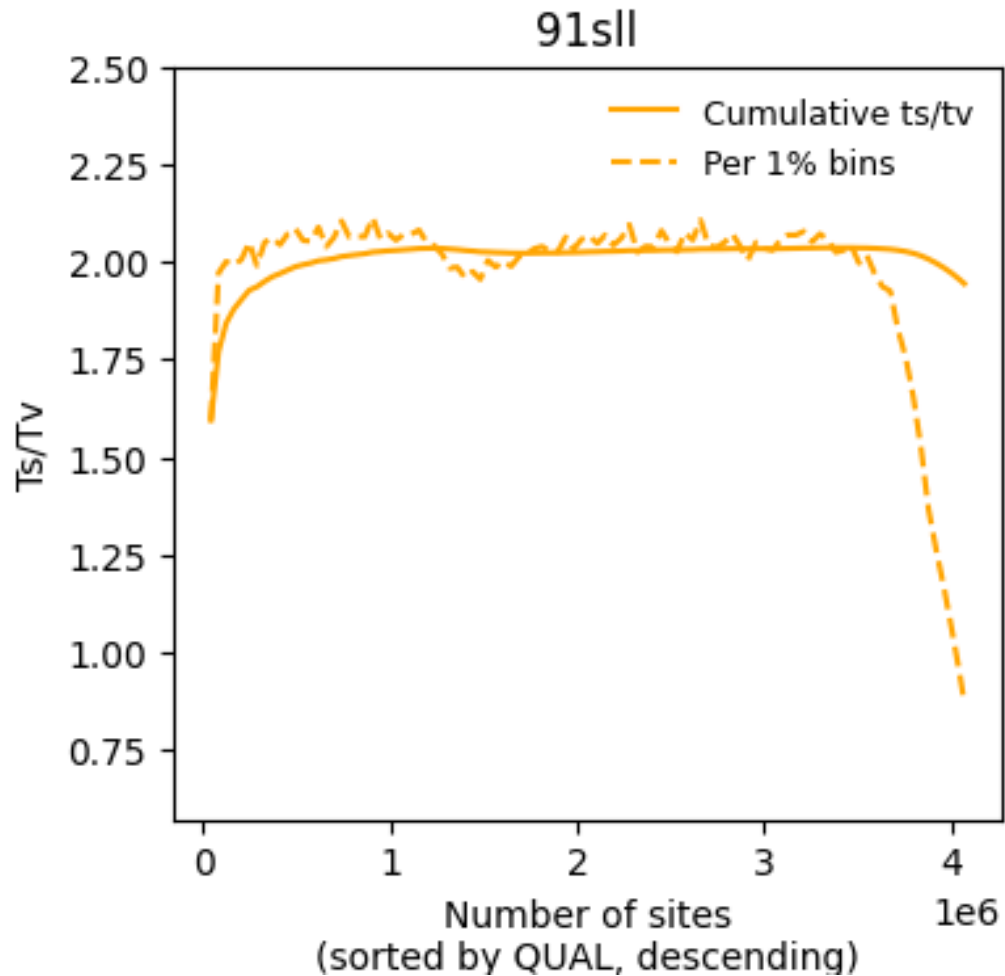
Number of SNPs by sample



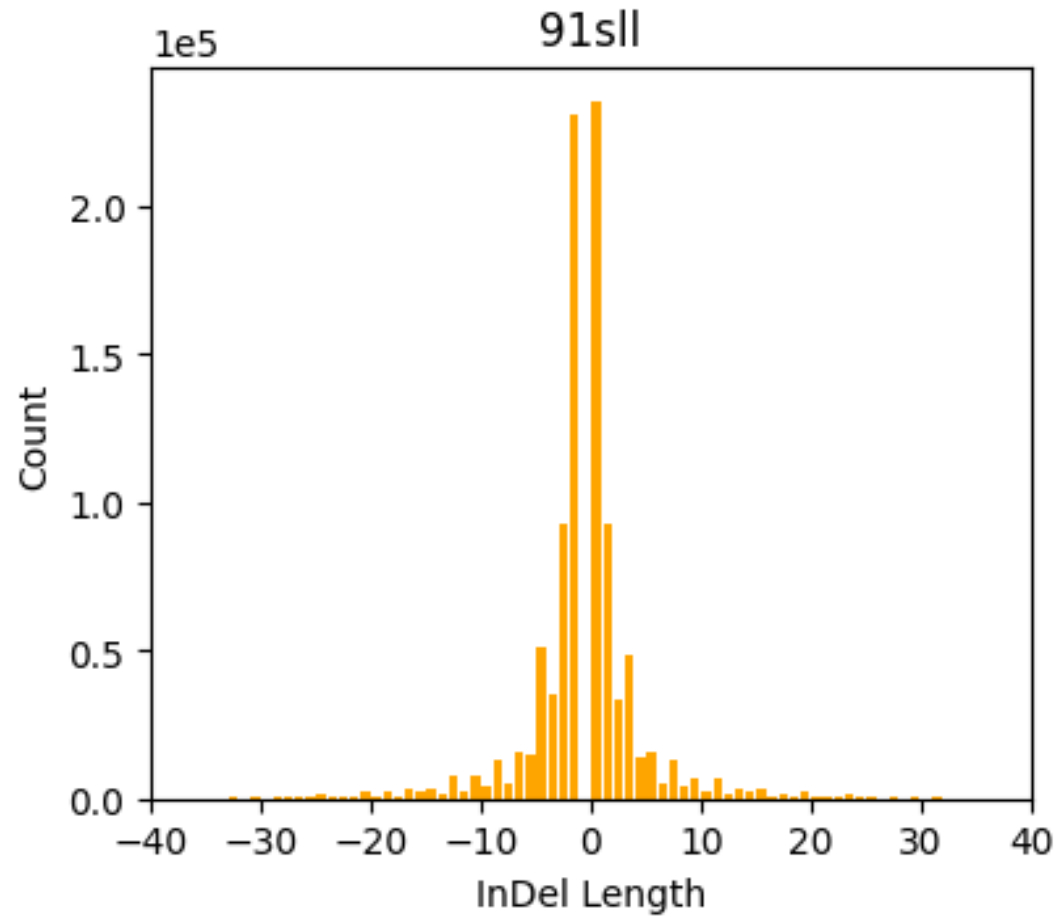
Number of indels by sample



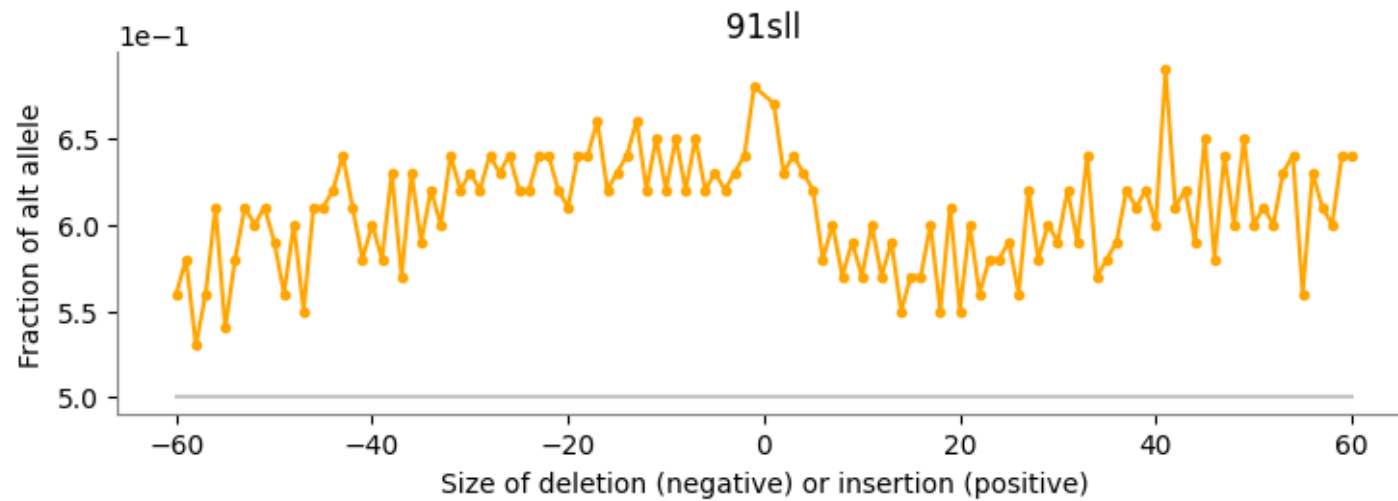
Ts/Tv stratified by QUAL



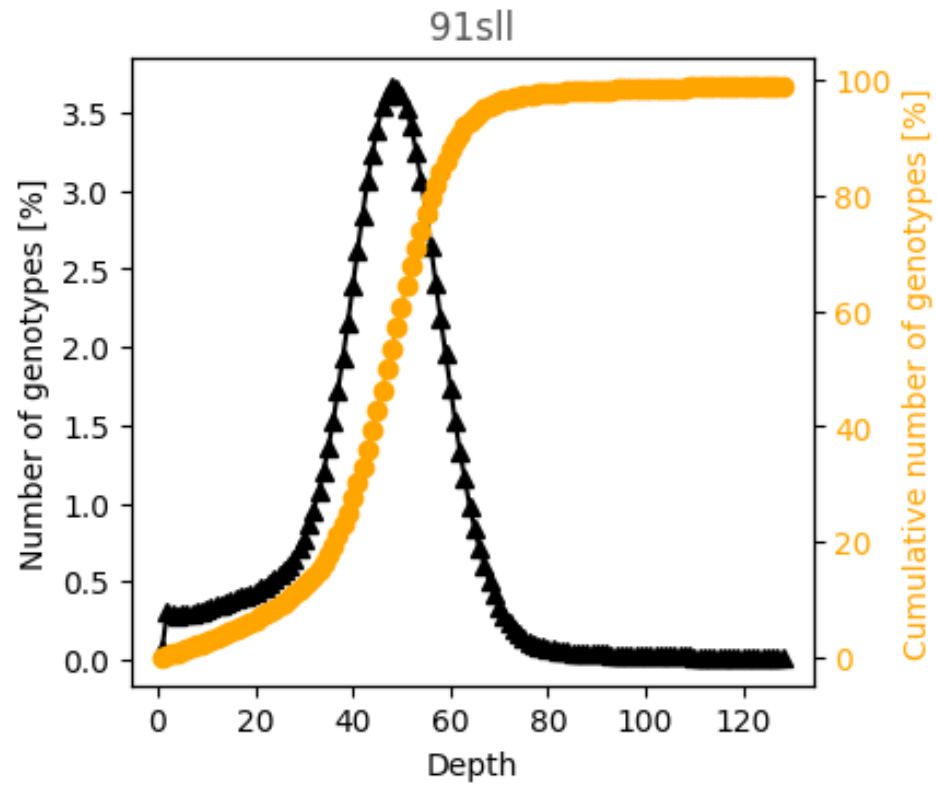
Indel distribution



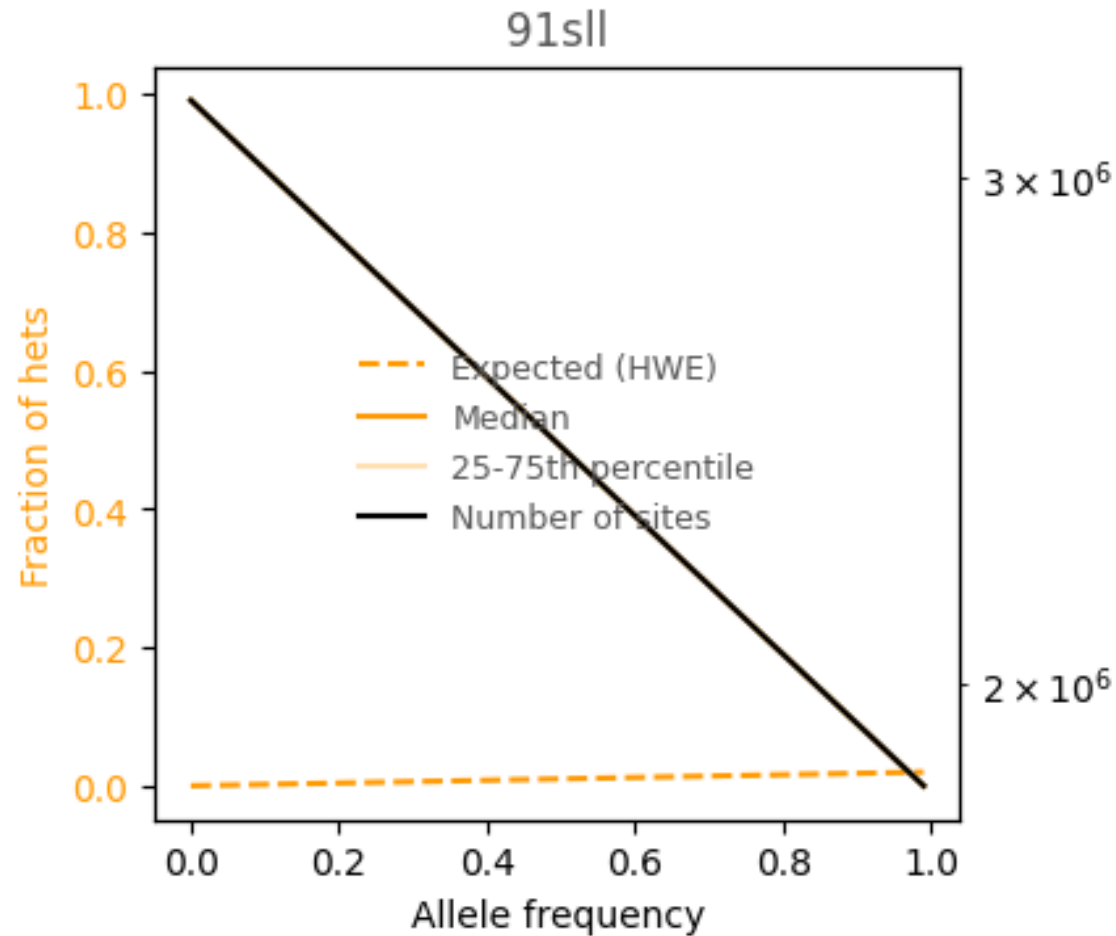
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

