

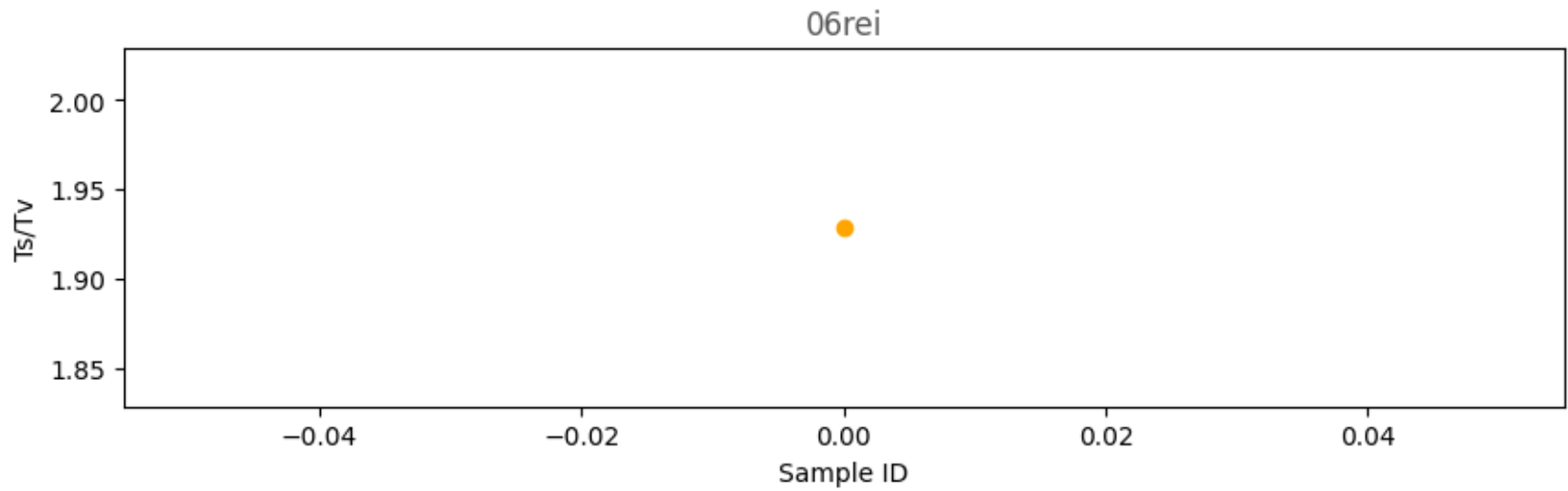
Summary Numbers

	SNPs			indels	MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	
06rei	4,061,608	1.93	1.93	942,269	–	0
* frameshift ratio: out/(out+in)						

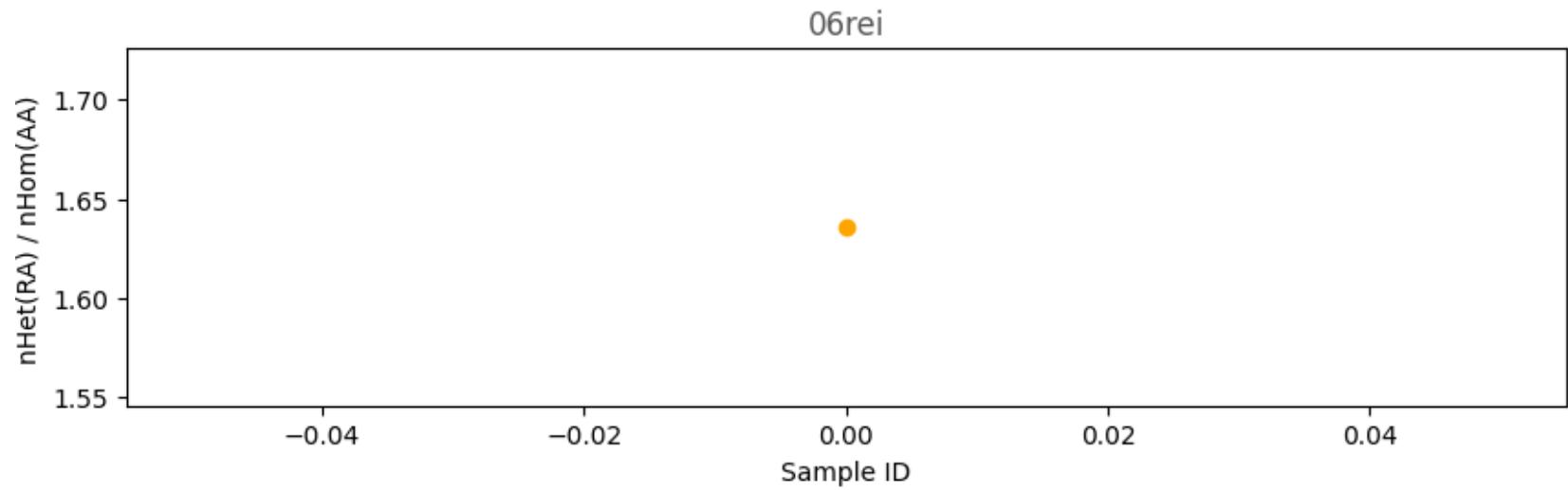
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
06rei	62.1%	1.90	66.6%	96,760	2,064

- 06rei .. /ngc/projects2/gm/data/archive/2022/variants/snv/06reilusm-103825451962-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00512-220119_A01176_BH2CT5DSX3-EXT_LAB_KA_NGCWGS-NGCWGS03685_21RKG023149x01_snv_germline_raw.haplotype_caller.vcf.gz

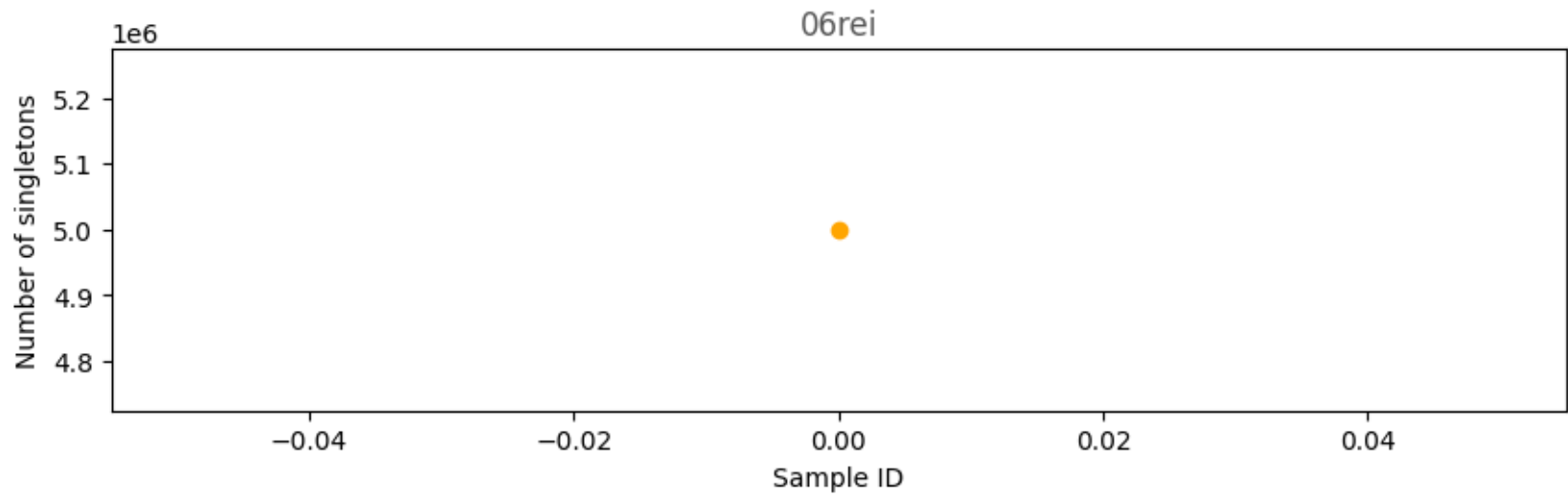
Ts/Tv by sample



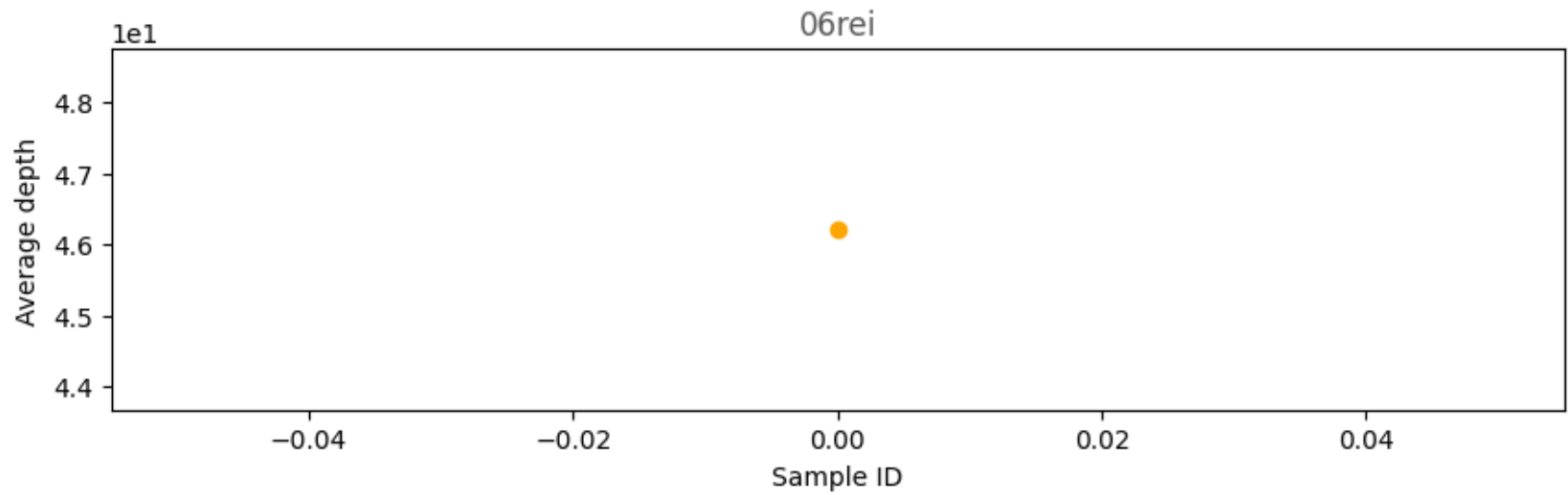
Hets vs non-ref Homs by sample



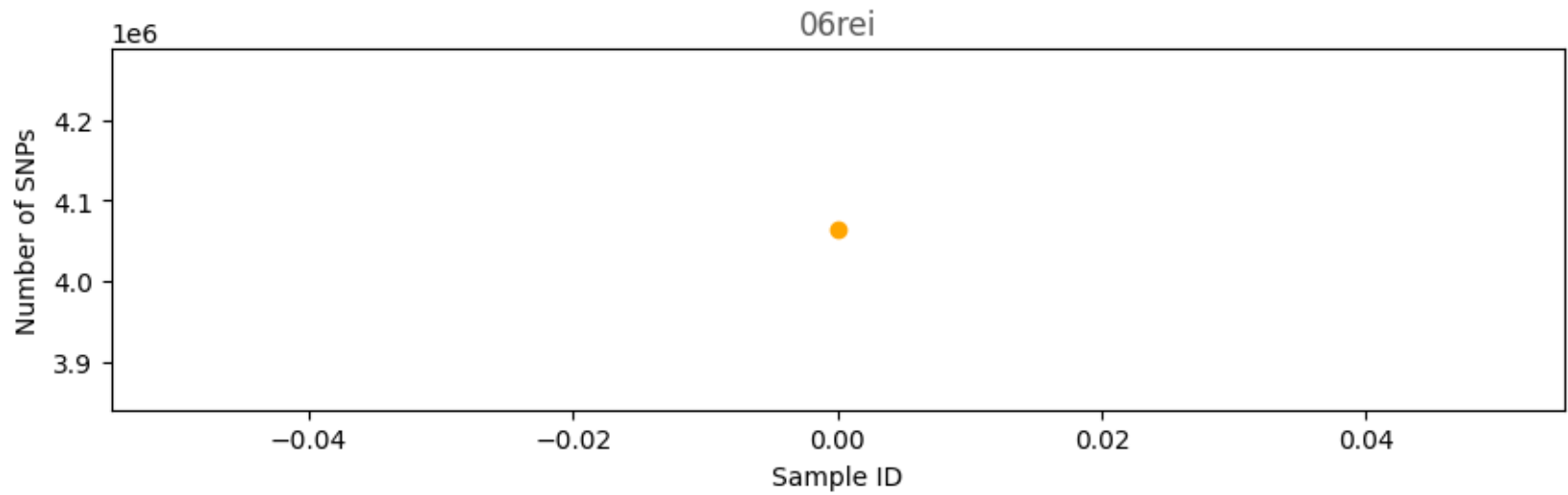
Singletons by sample (hets and homs)



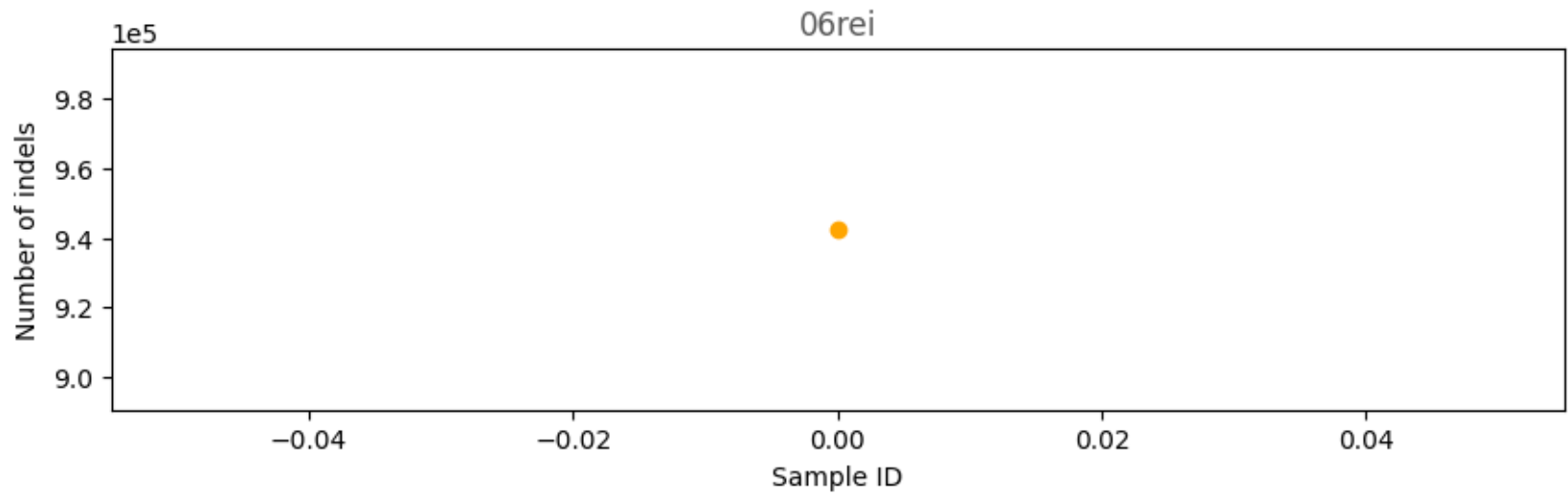
Average depth by sample



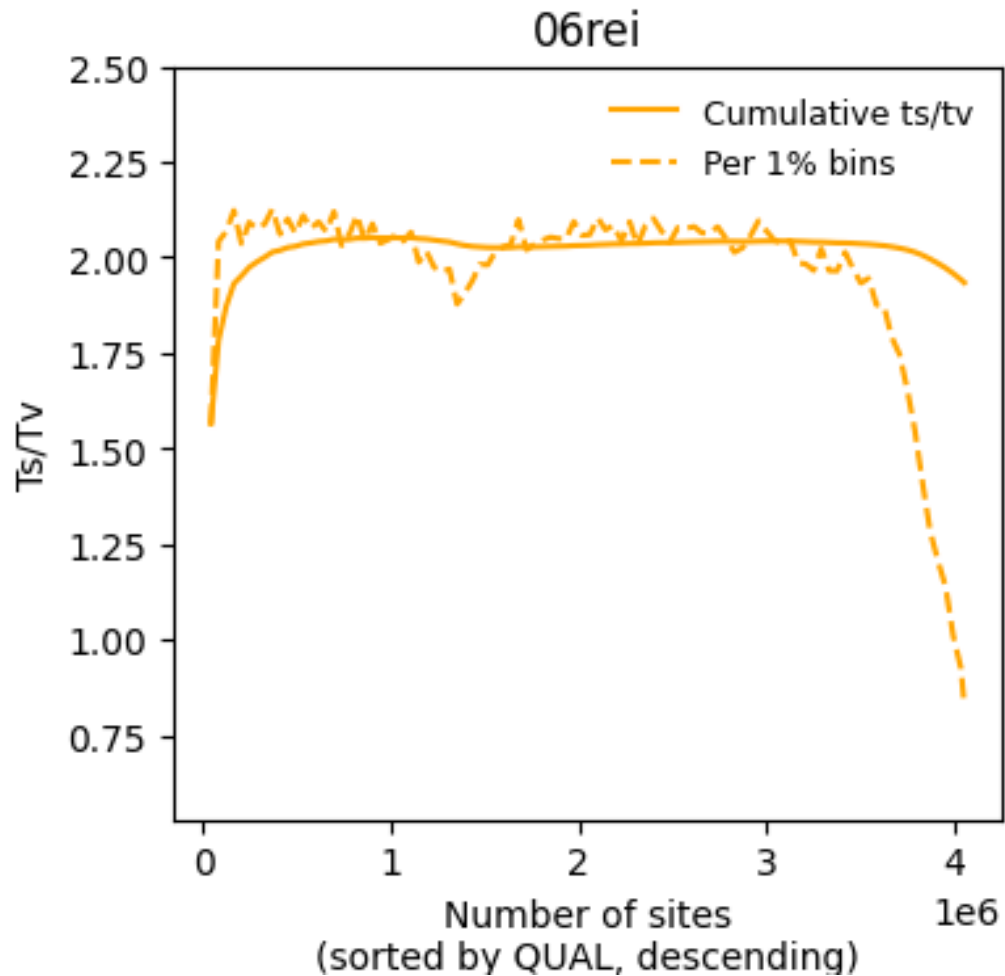
Number of SNPs by sample



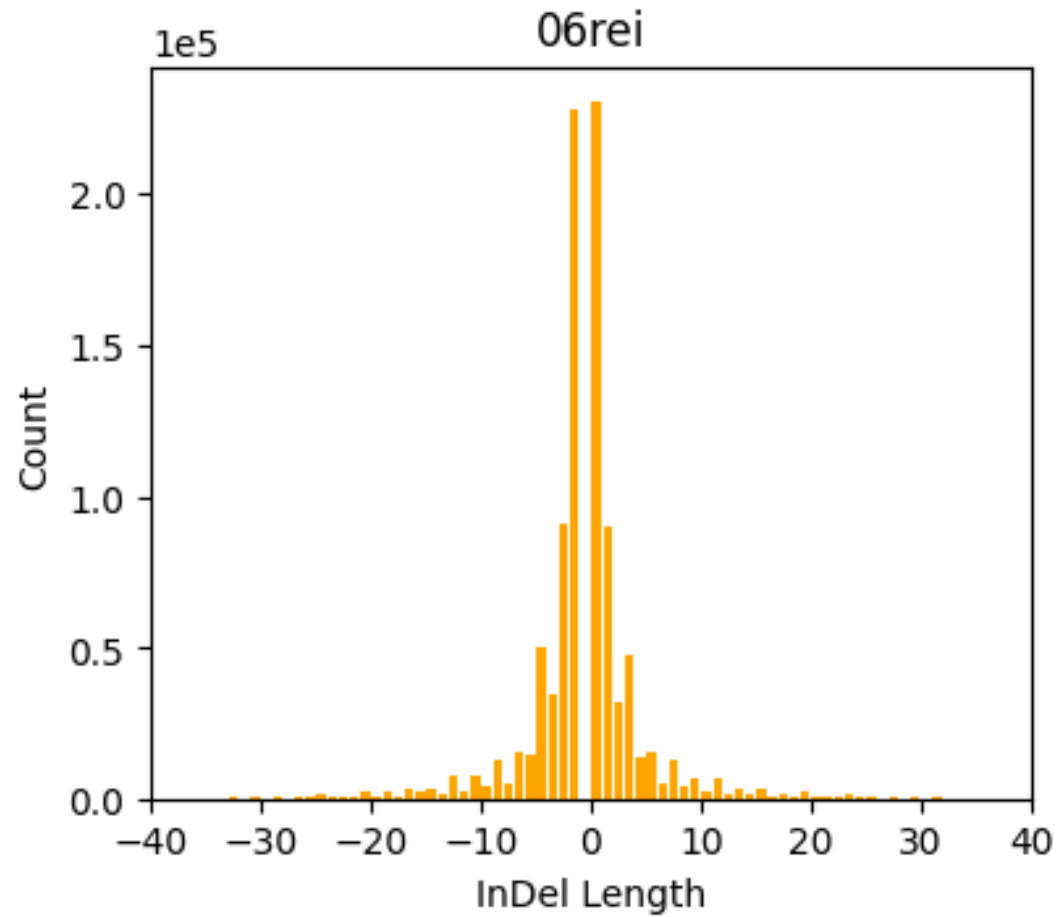
Number of indels by sample



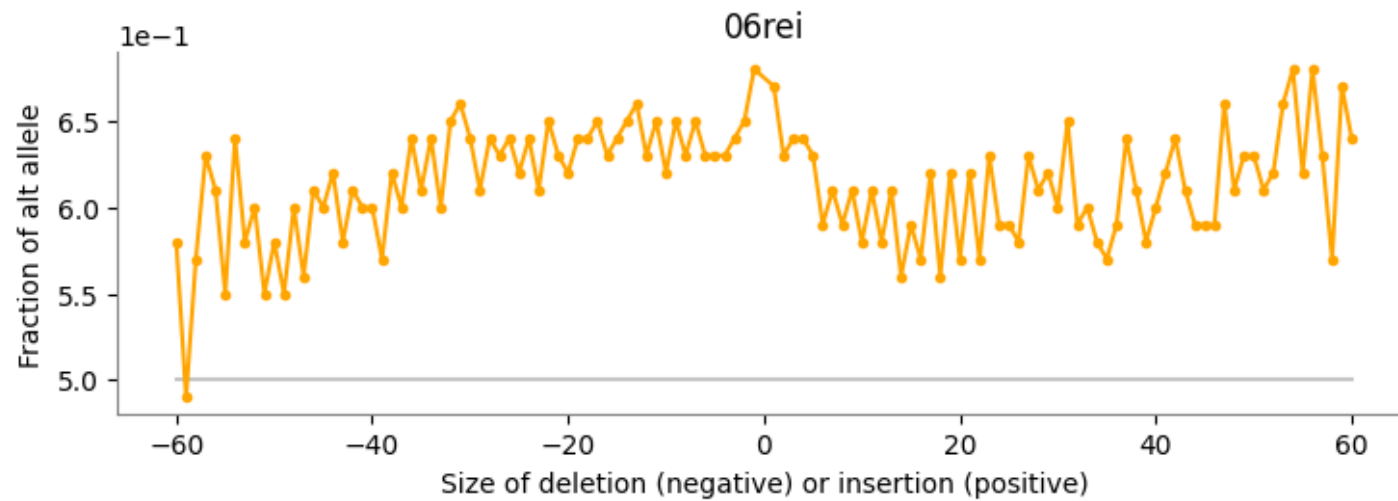
Ts/Tv stratified by QUAL



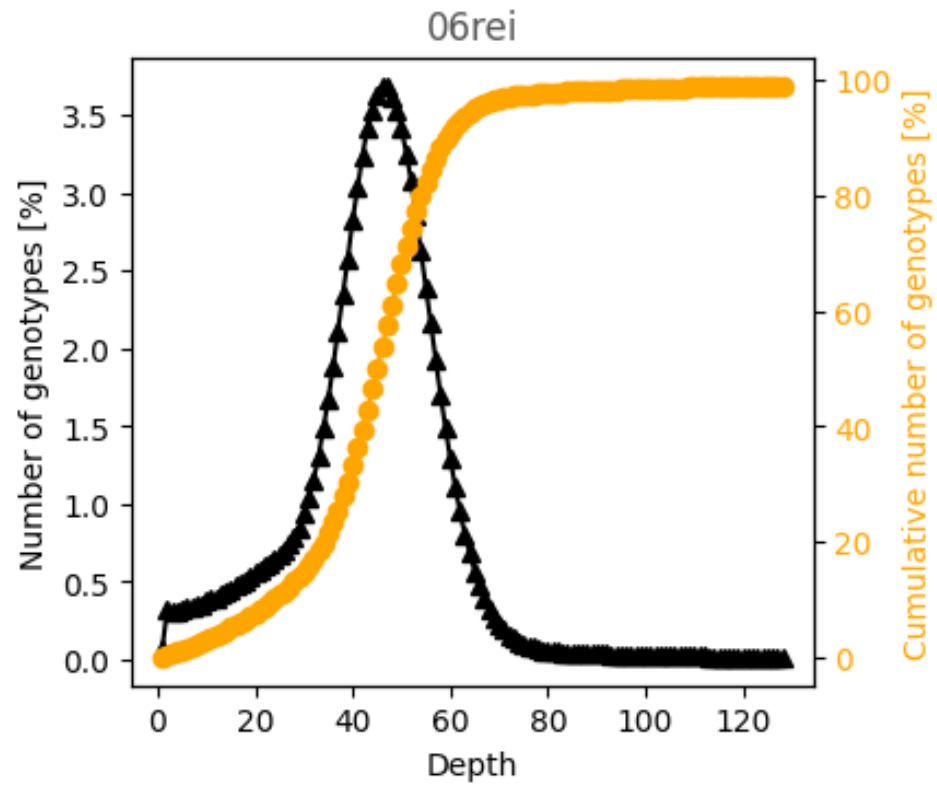
Indel distribution



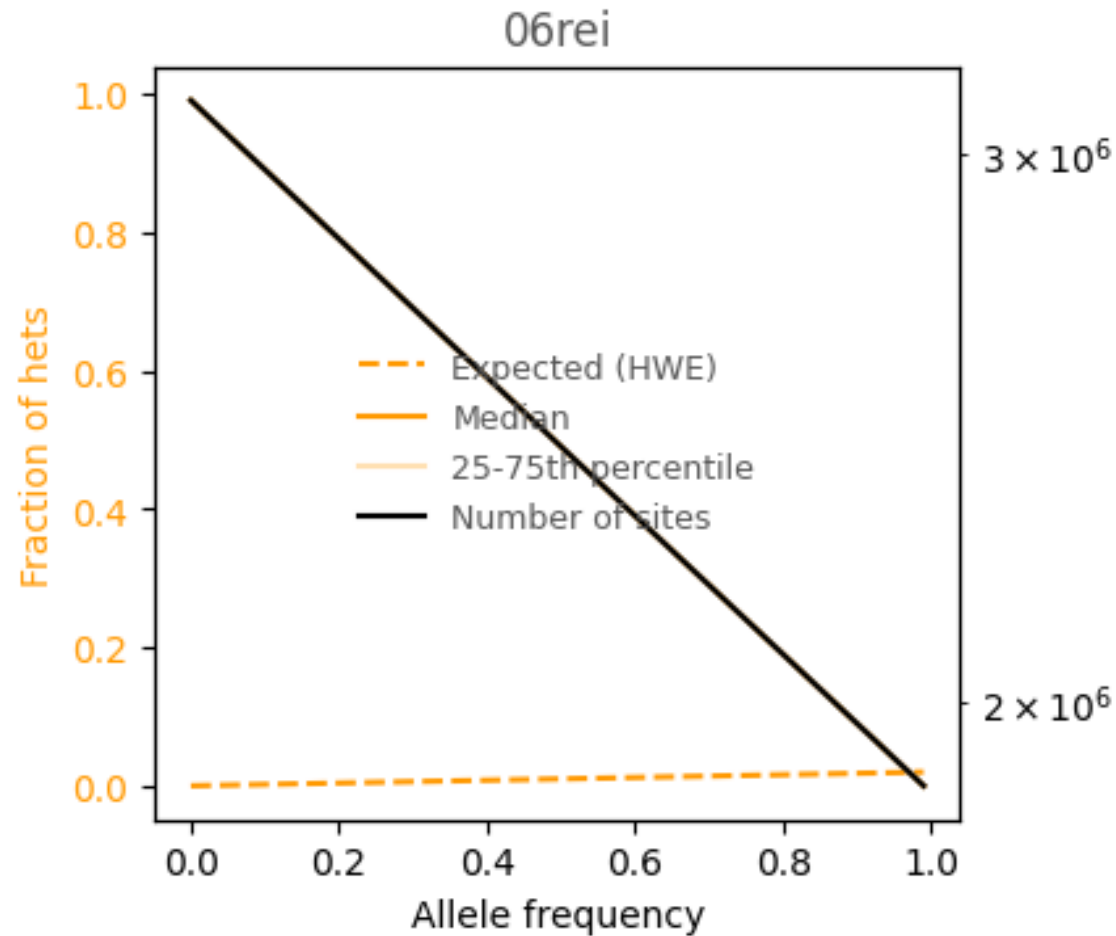
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

