Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
75ree	4,069,560	1.93	1.94	938,711	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
75ree	62.1%	1.90	66.3%	92,973	1,909

- 75ree .. `/ngc/projects2/gm/data/archive/2022/variants/snv/75reeresm-103817172170-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211215_A01466_BH2GHYDSX3-RHGM_LABKA_WGSA KUT-WGSAKUT03523_21RKG030508x01_snv_germline_raw.haplotype_caller.vcf.gz`
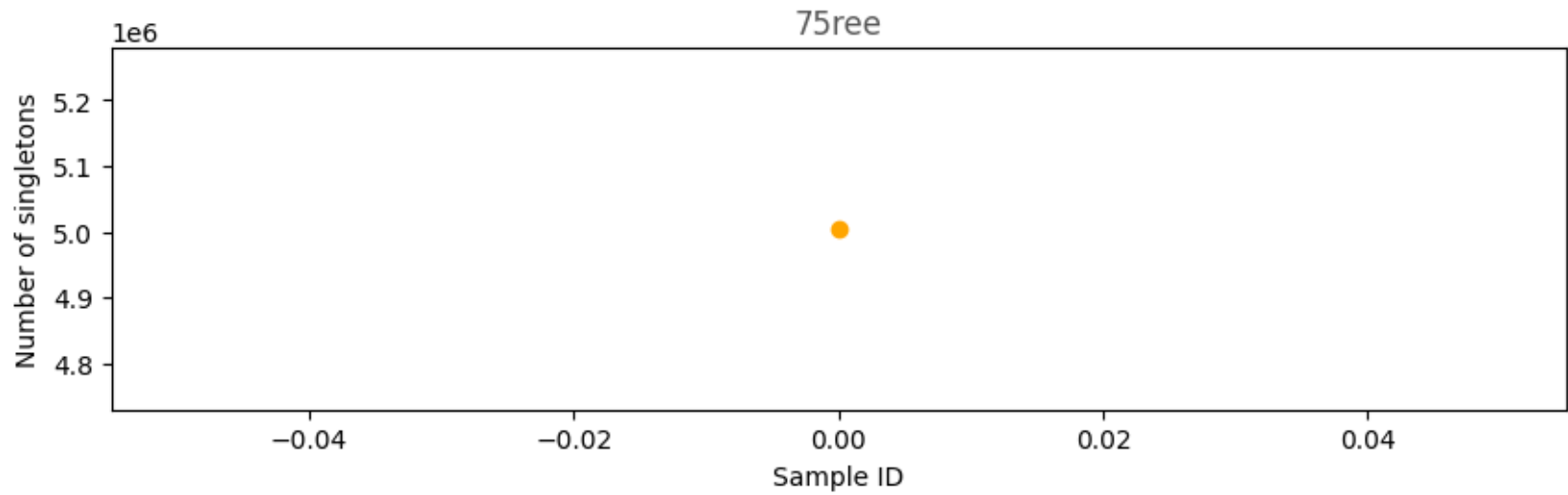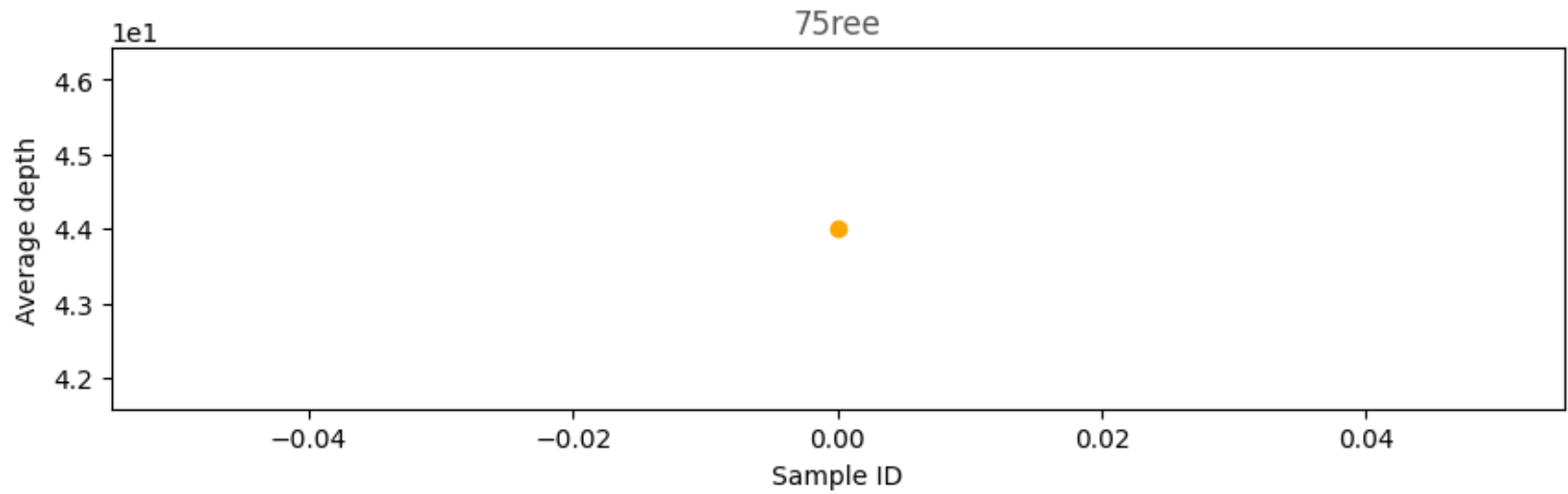
Ts/Tv by sample
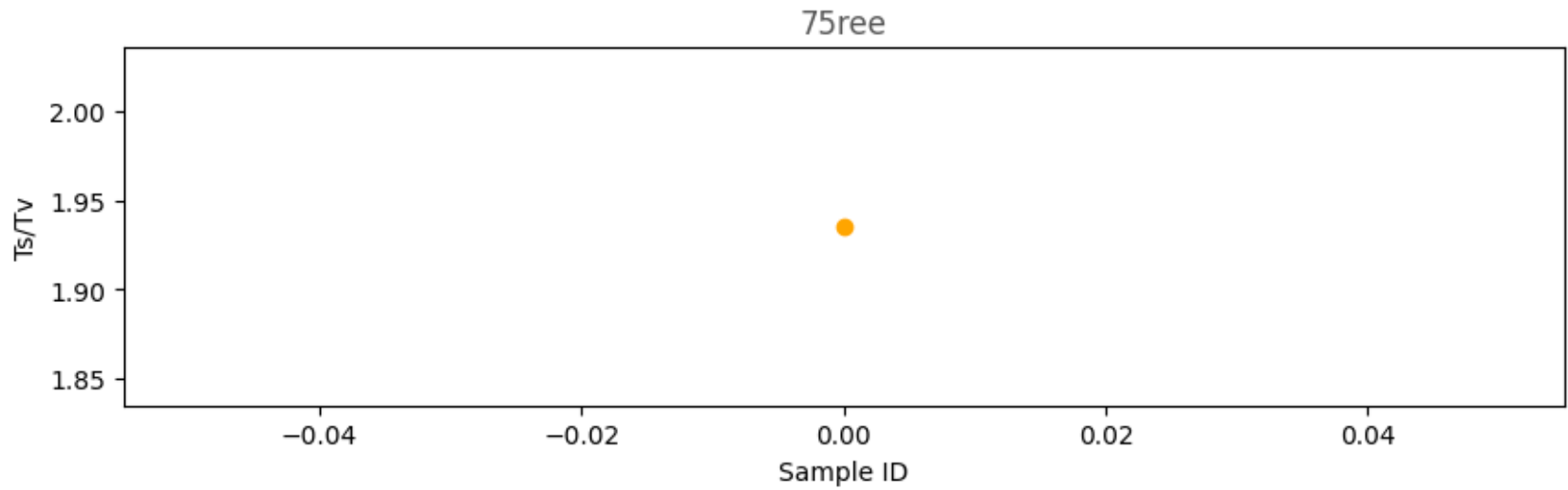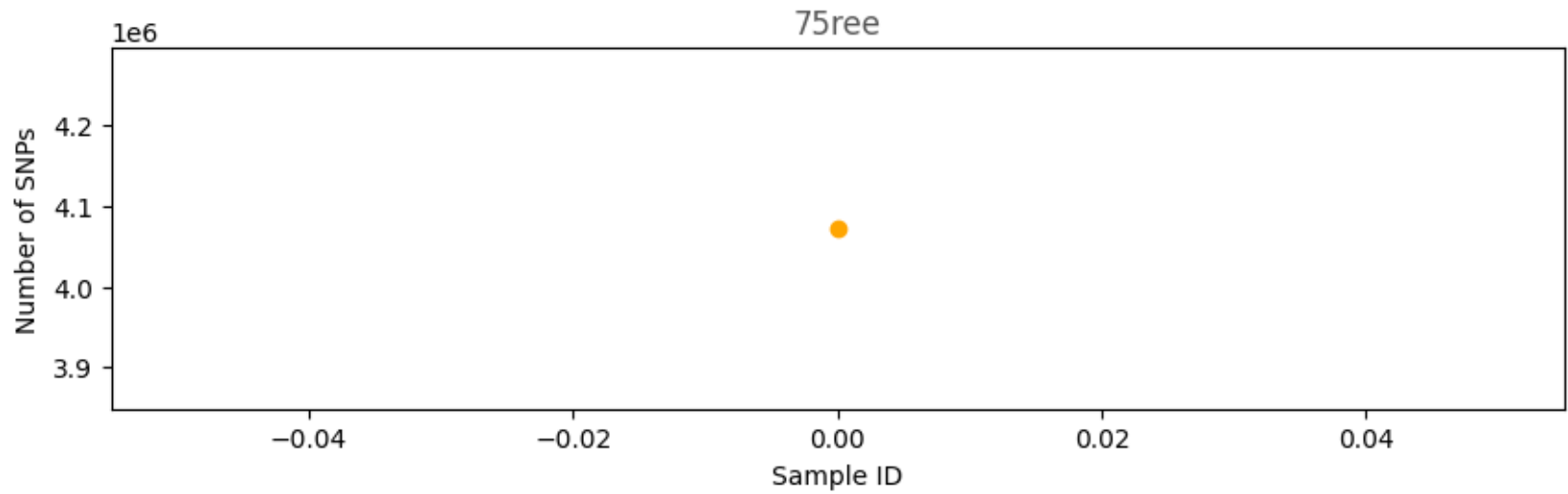
Hets vs non-ref Homs by sample
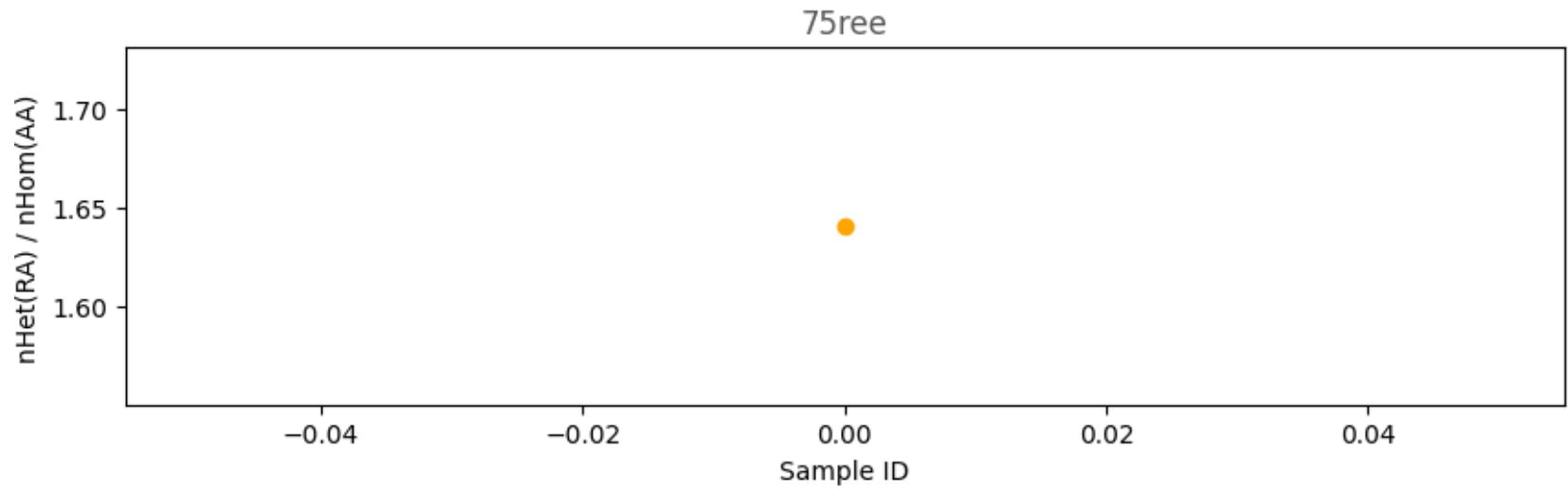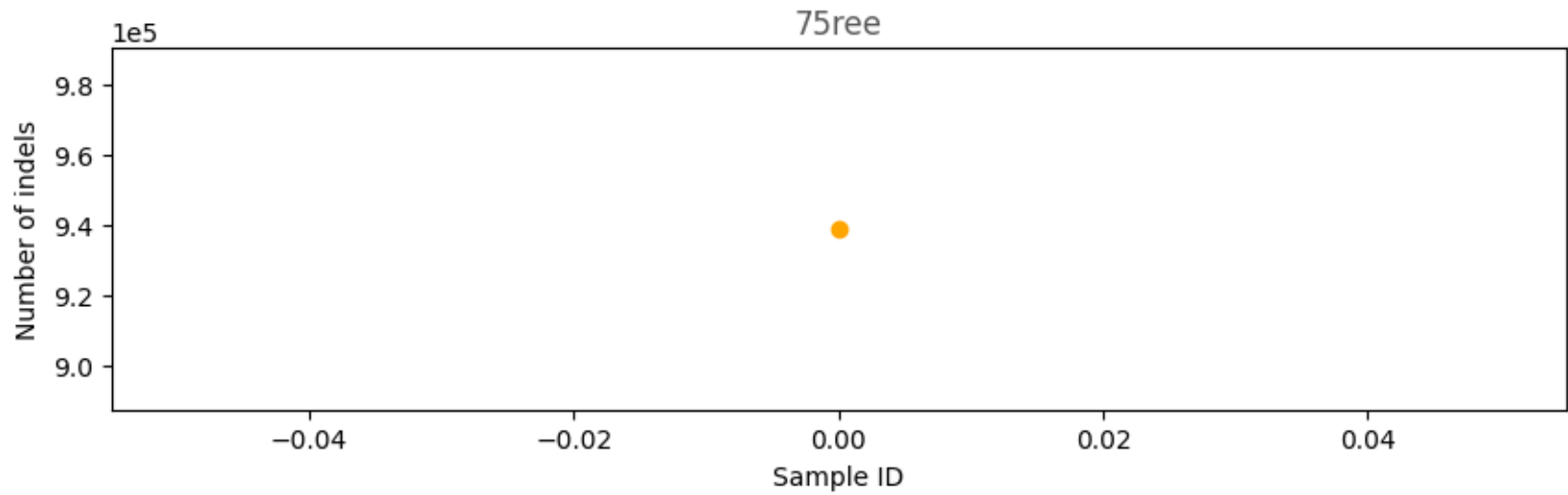
Singletons by sample (hets and homs)
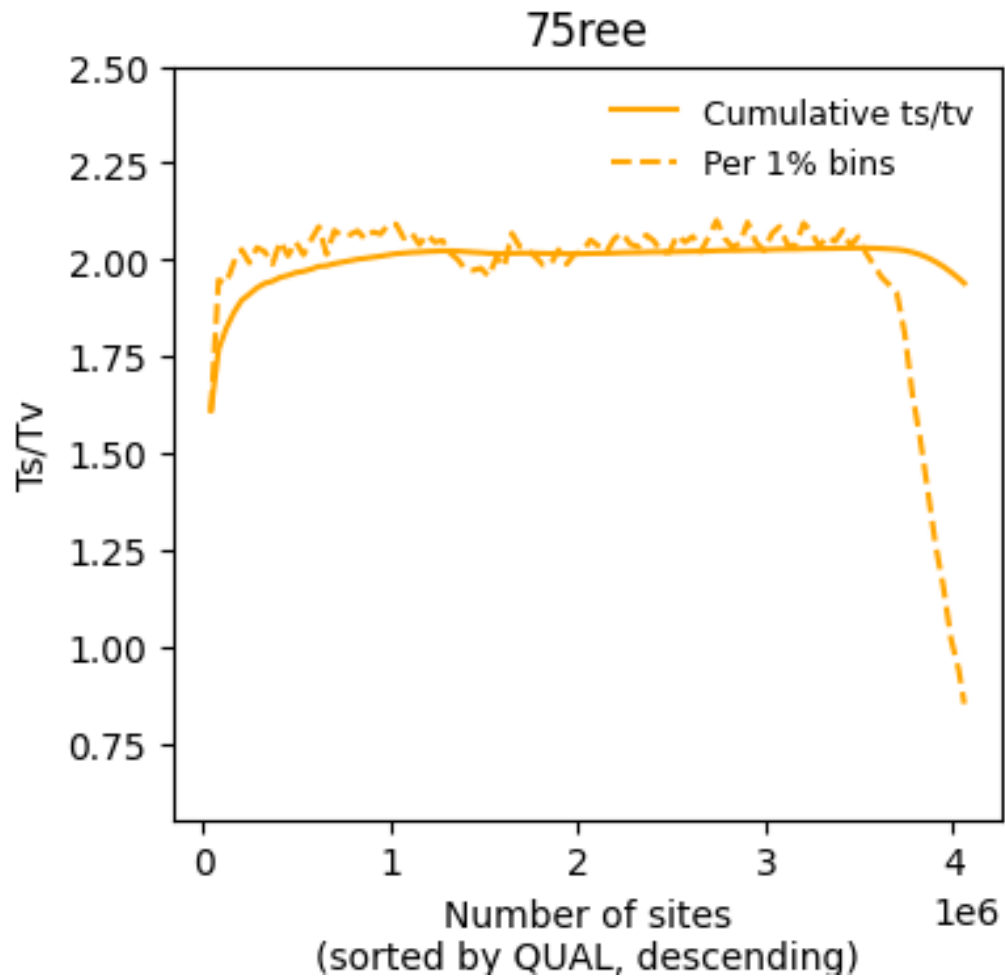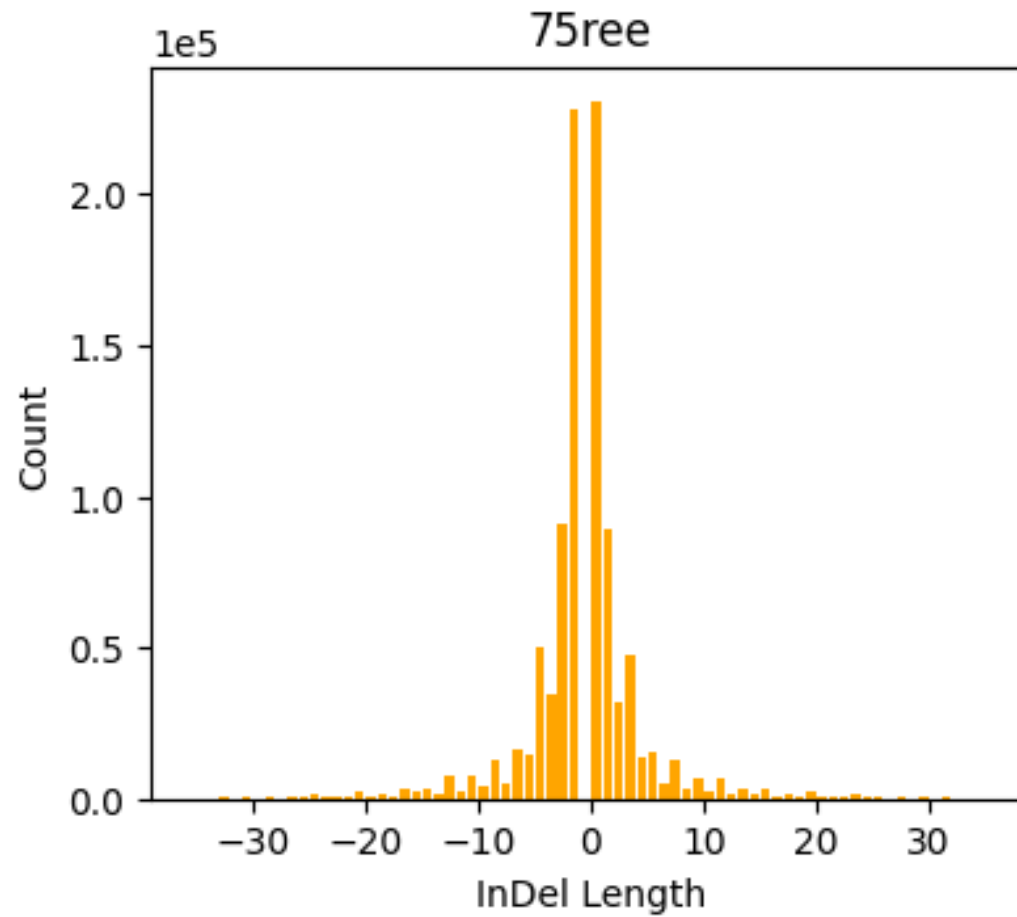
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

8

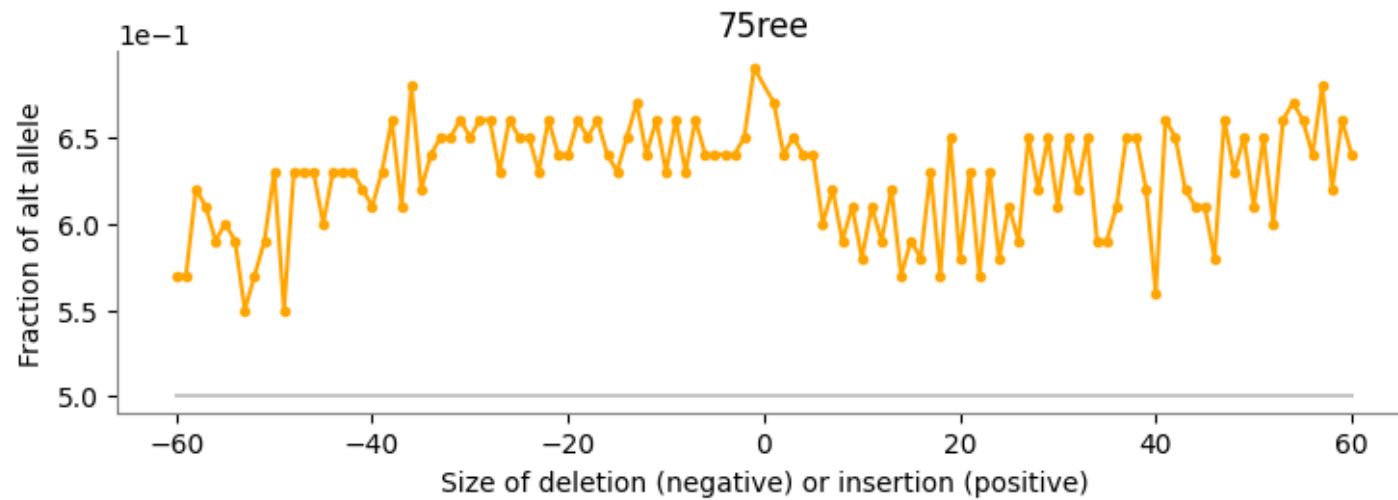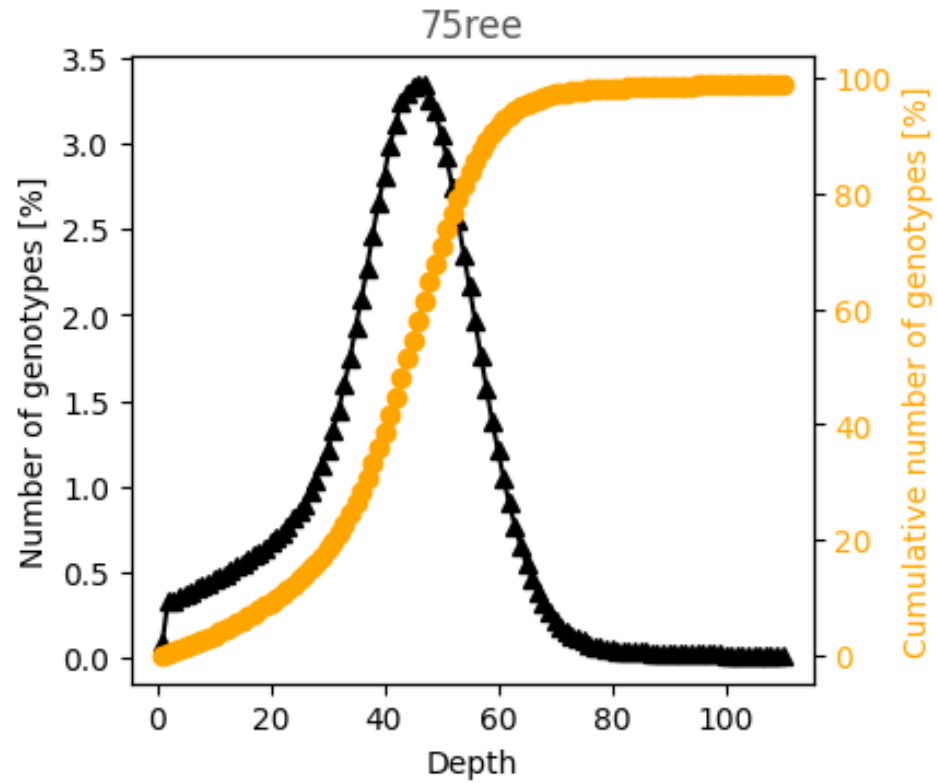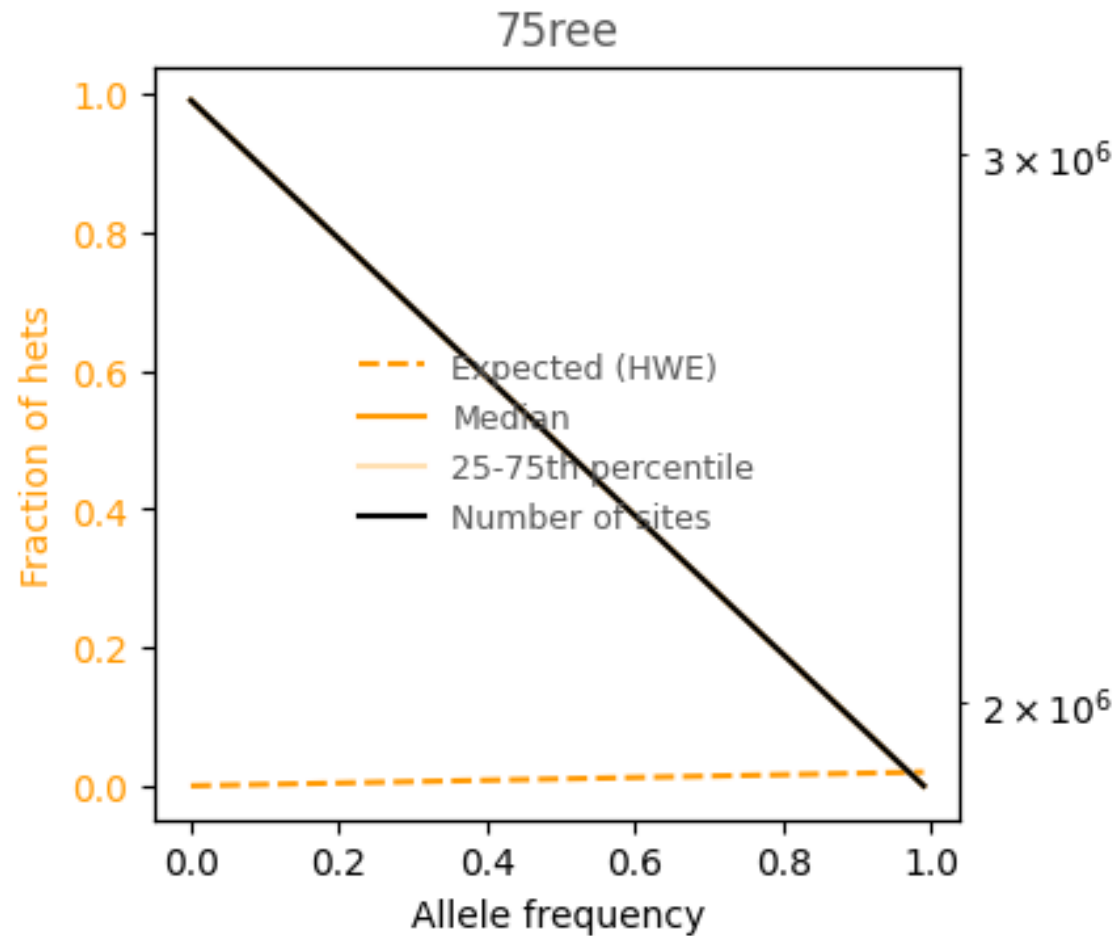# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types