Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
80err	4,152,438	1.93	1.93	968,740	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
80err	63.2%	1.89	68.3%	105,838	2,238

- 80err .. `/ngc/projects2/gm/data/archive/2022/variants/snv/80errefuf-103776936493-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00312-210921_A01411_BHJTJJDSX2-EXT_LABKA_NGCWGS-NGCWGS00205_21RKG021208x01_snv_germline_raw.haplotype_caller.vcf.gz`
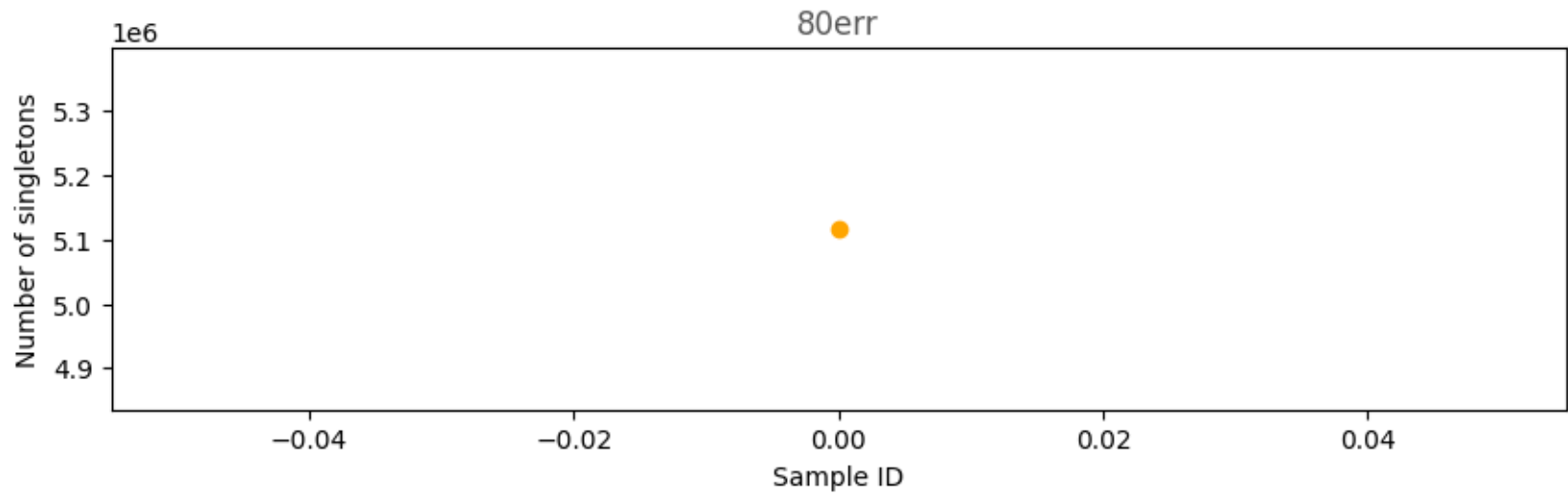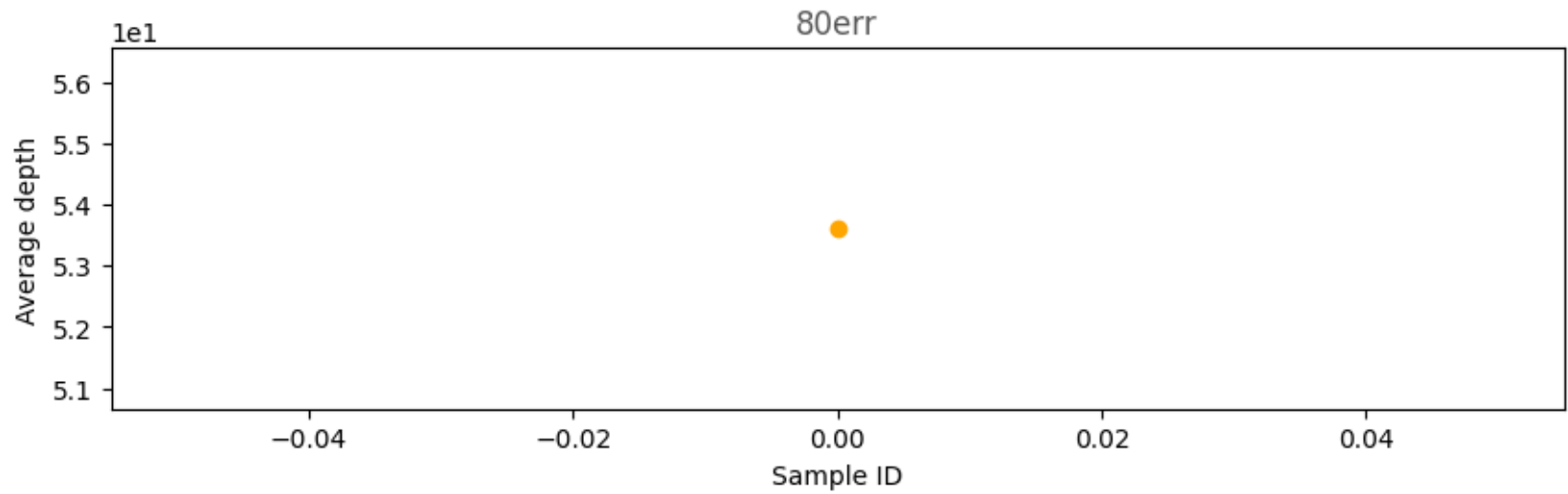
Ts/Tv by sample
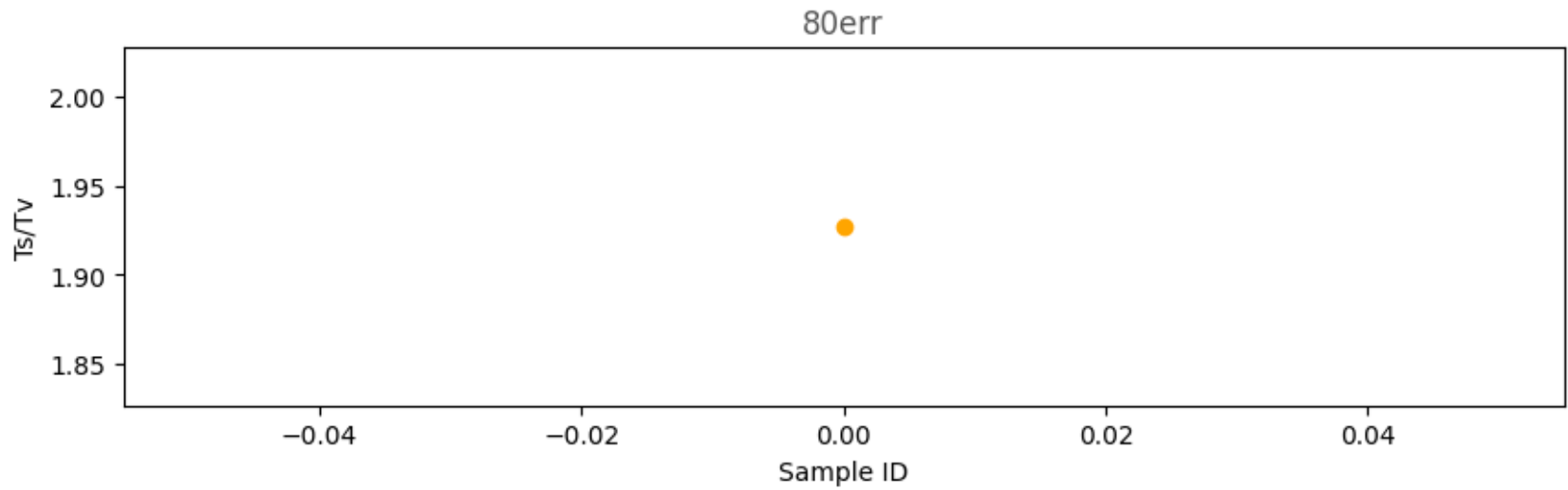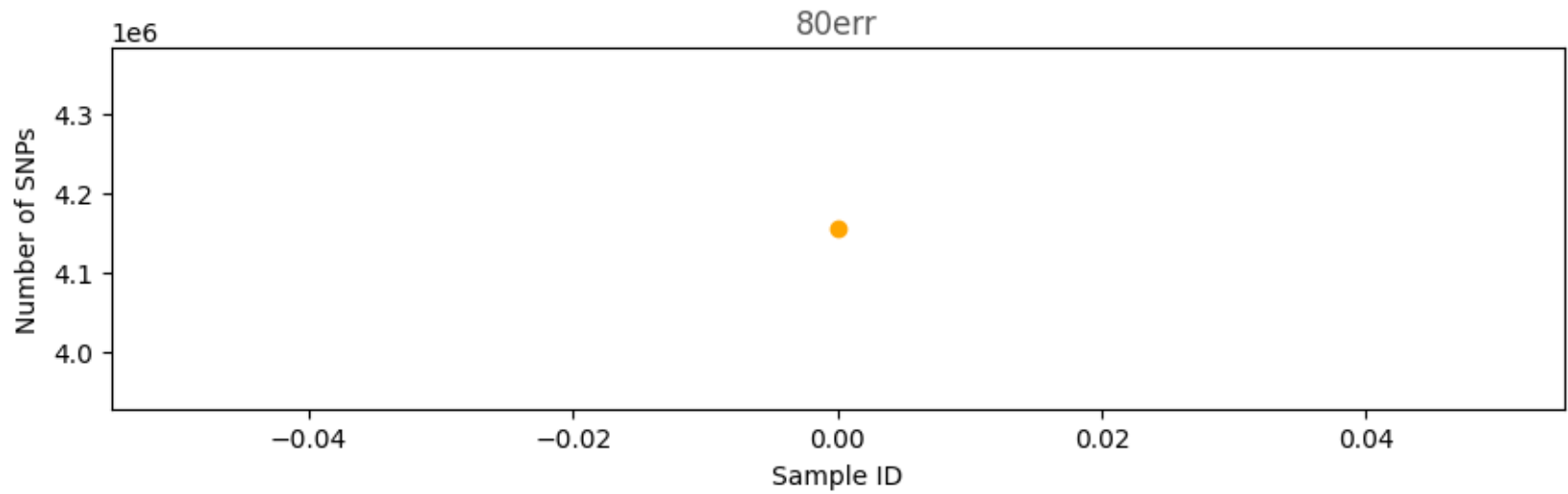
Hets vs non-ref Homs by sample
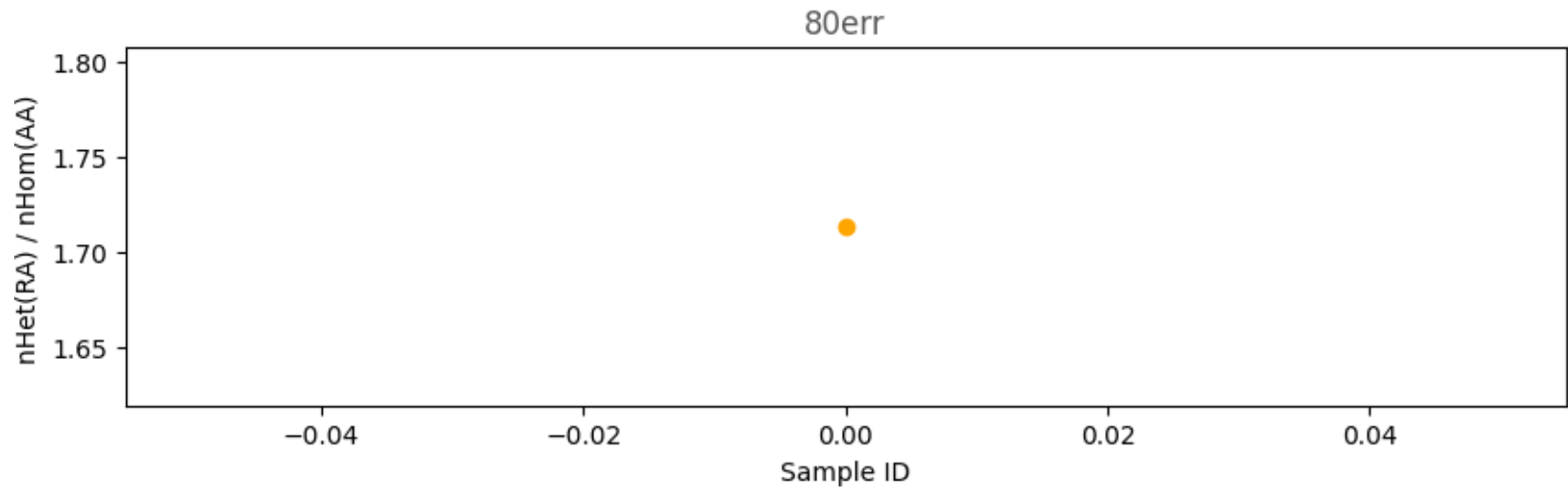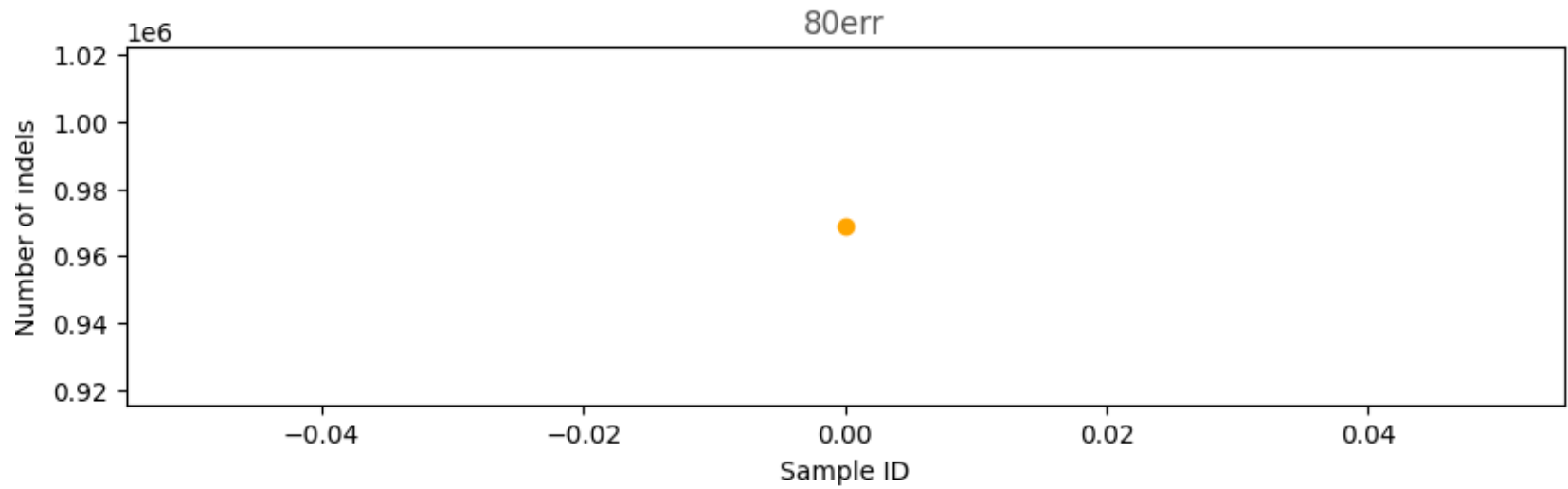
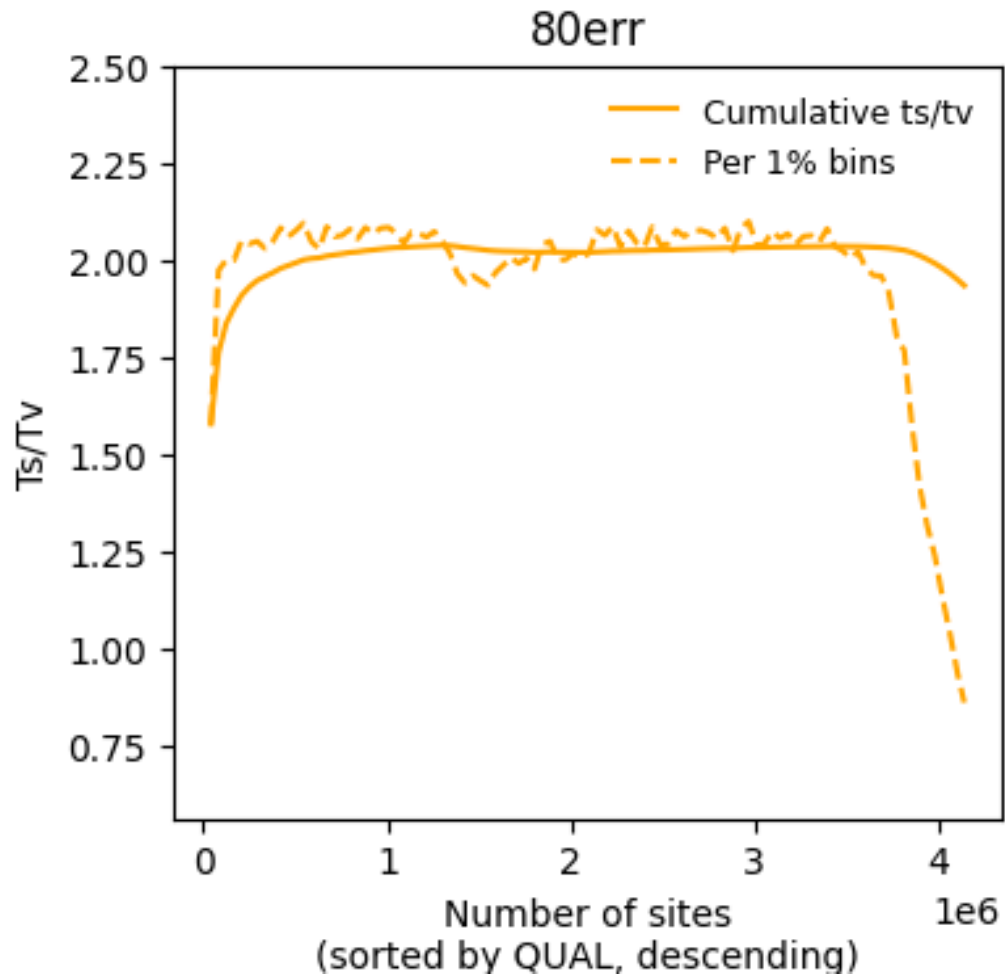# Singletons by sample (hets and homs)

Average depth by sample

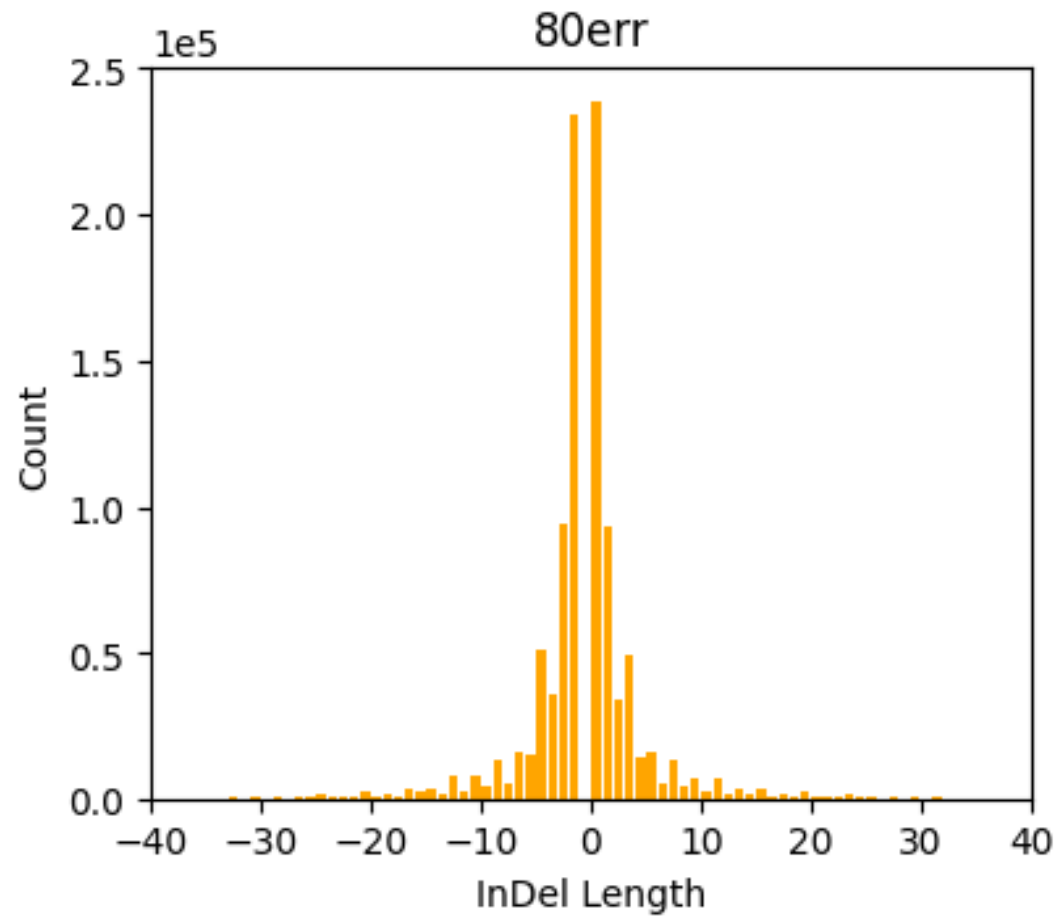# Number of SNPs by sample

Number of indels by sample
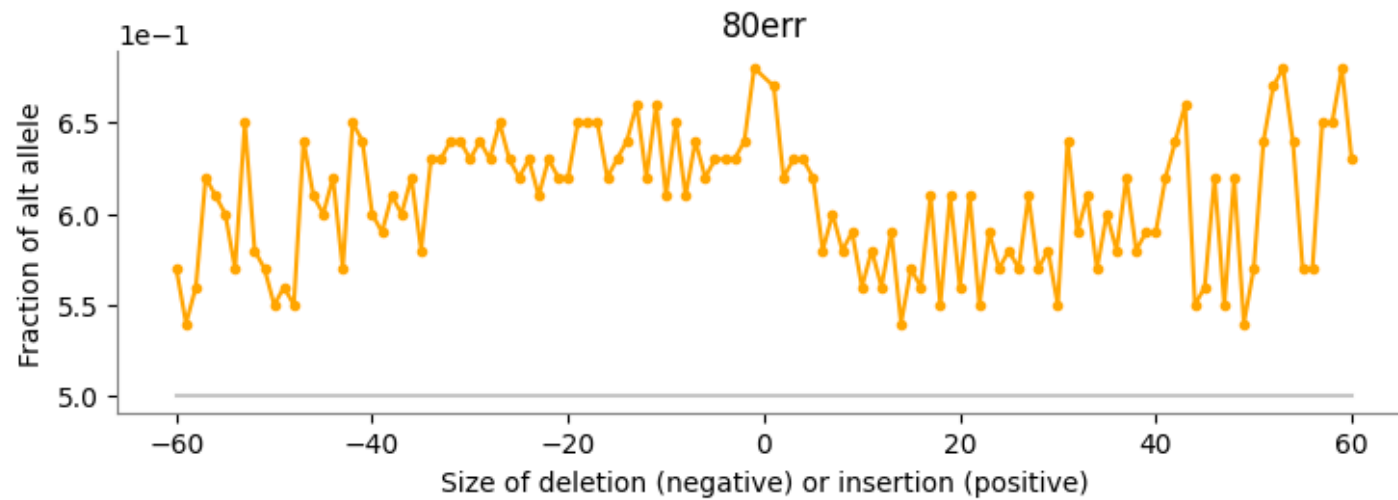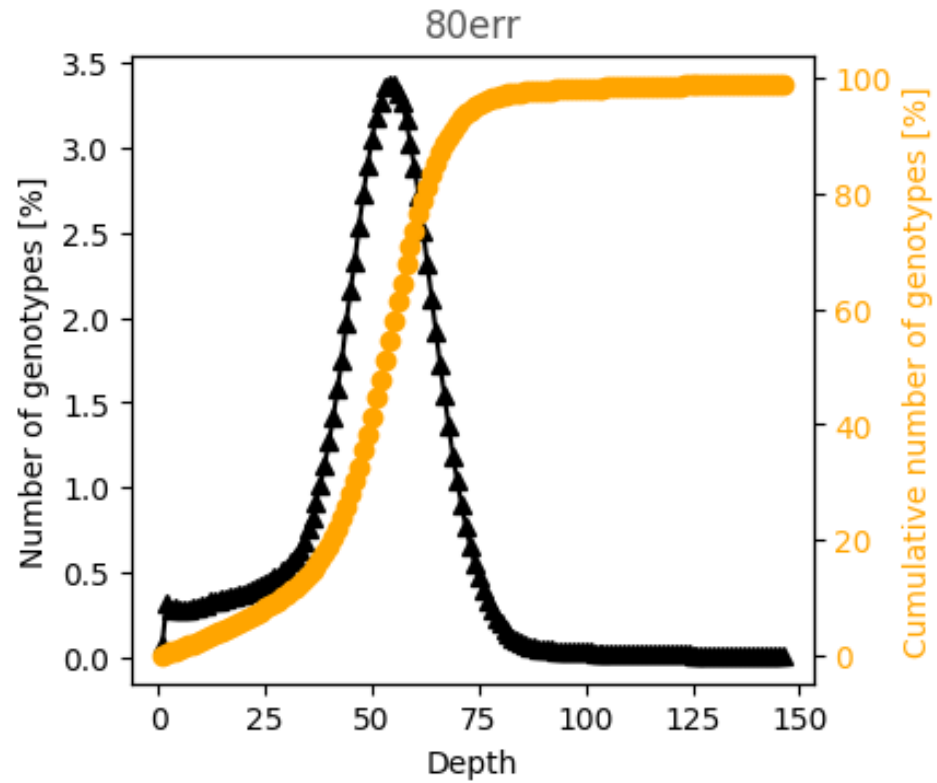
Ts/Tv stratified by QUAL
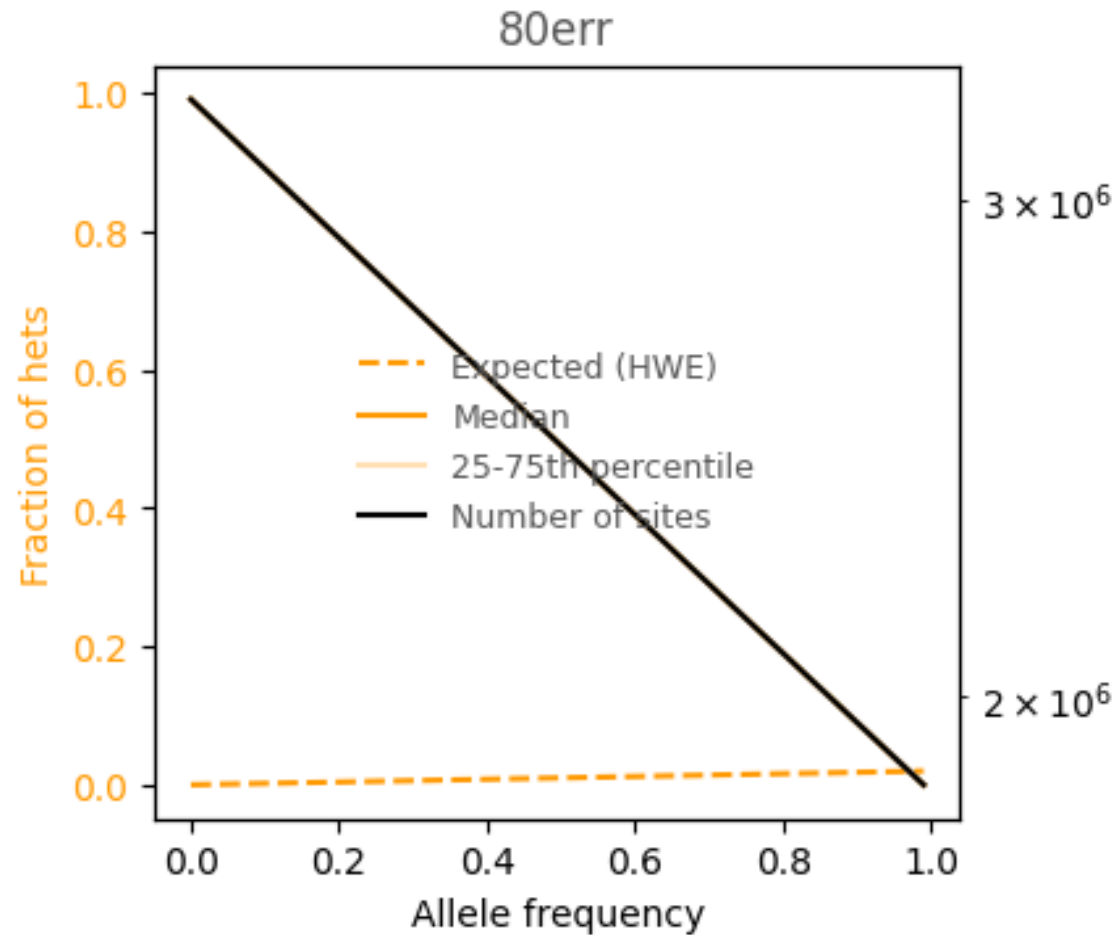

8

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types