Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
06qut	4,817,325	1.94	1.94	1,113,927	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
06qut	72.4%	1.92	78.3%	143,893	2,780

- 06qut .. `/ngc/projects2/gm/data/archive/2022/variants/snv/06qutsmef-110299556526-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-221026_A01961_AH3TGFDSX5-RHGM_LABKA_WGS- WGS05981_snv_germline_raw.haplotype_caller.vcf.gz`
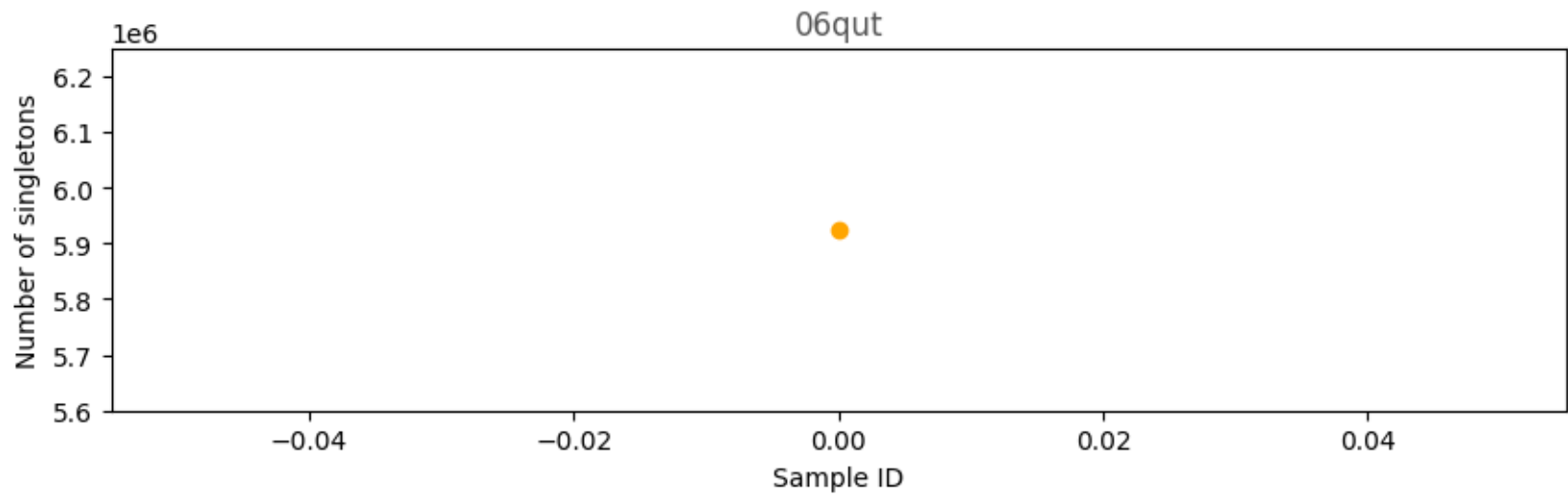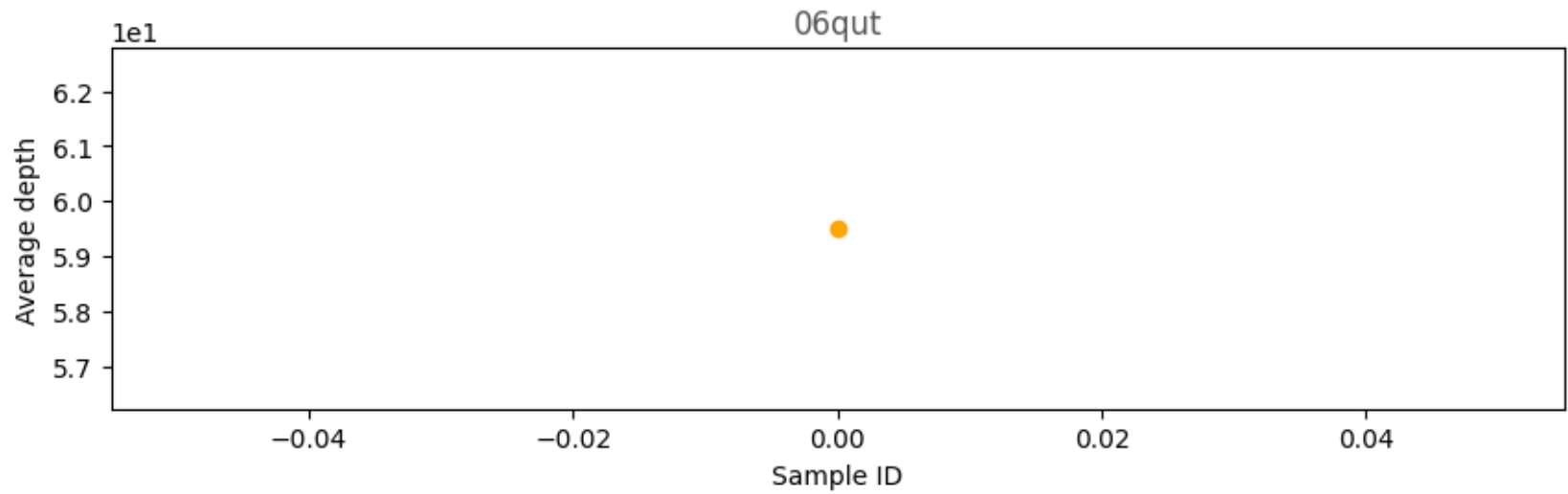
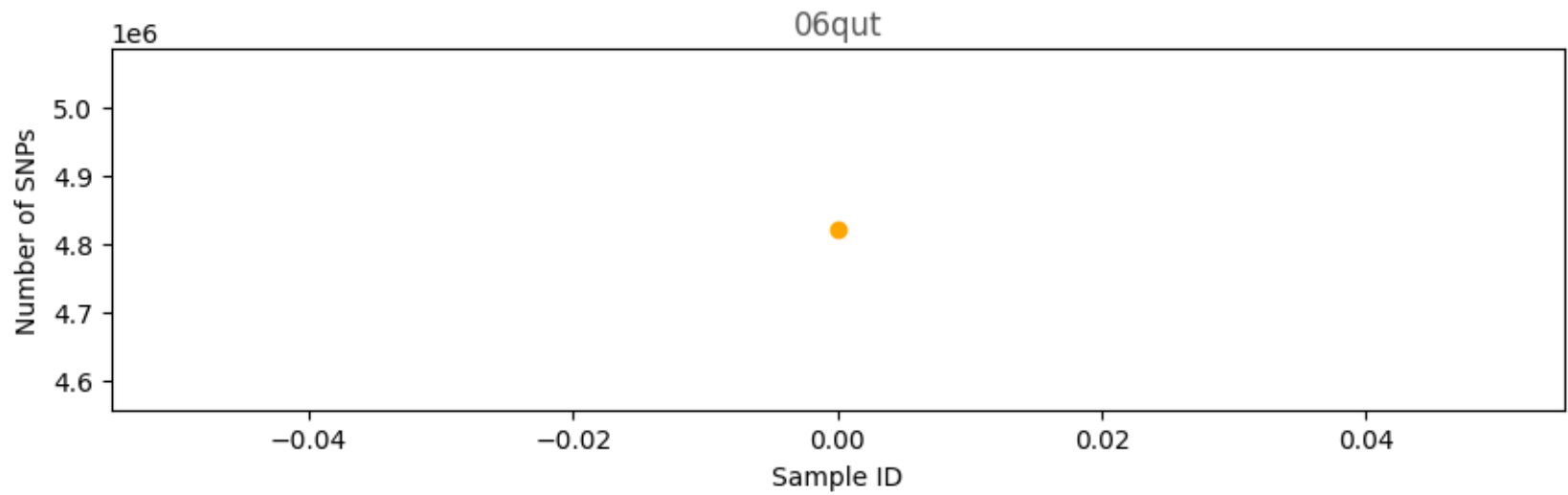# Ts/Tv by sample
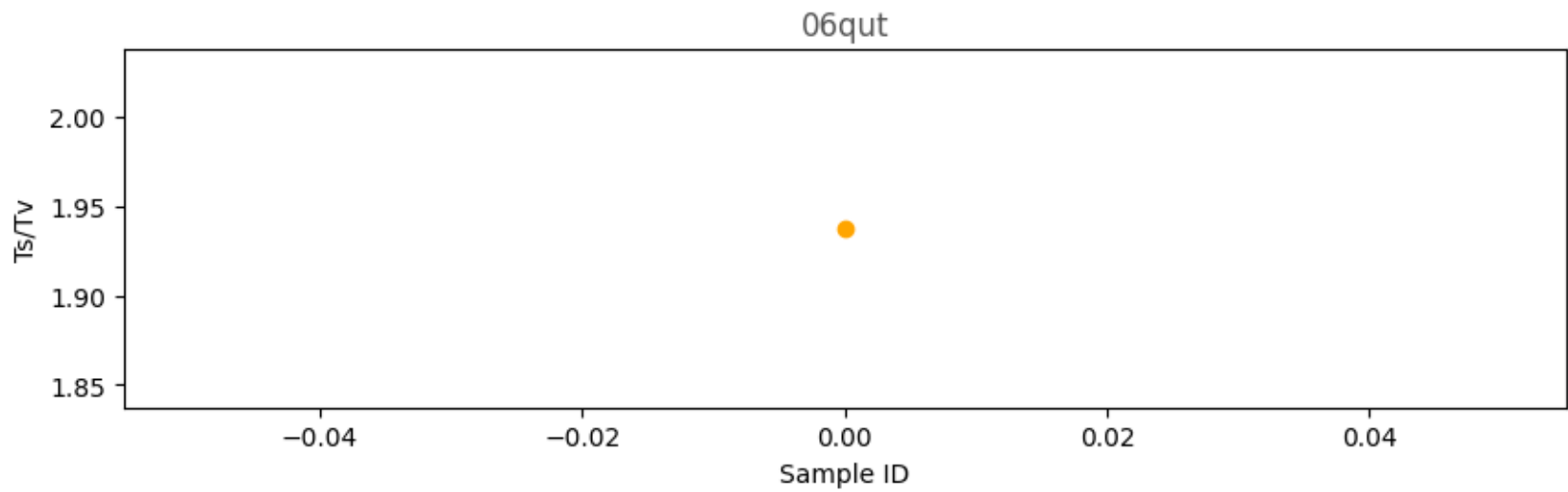
Hets vs non-ref Homs by sample
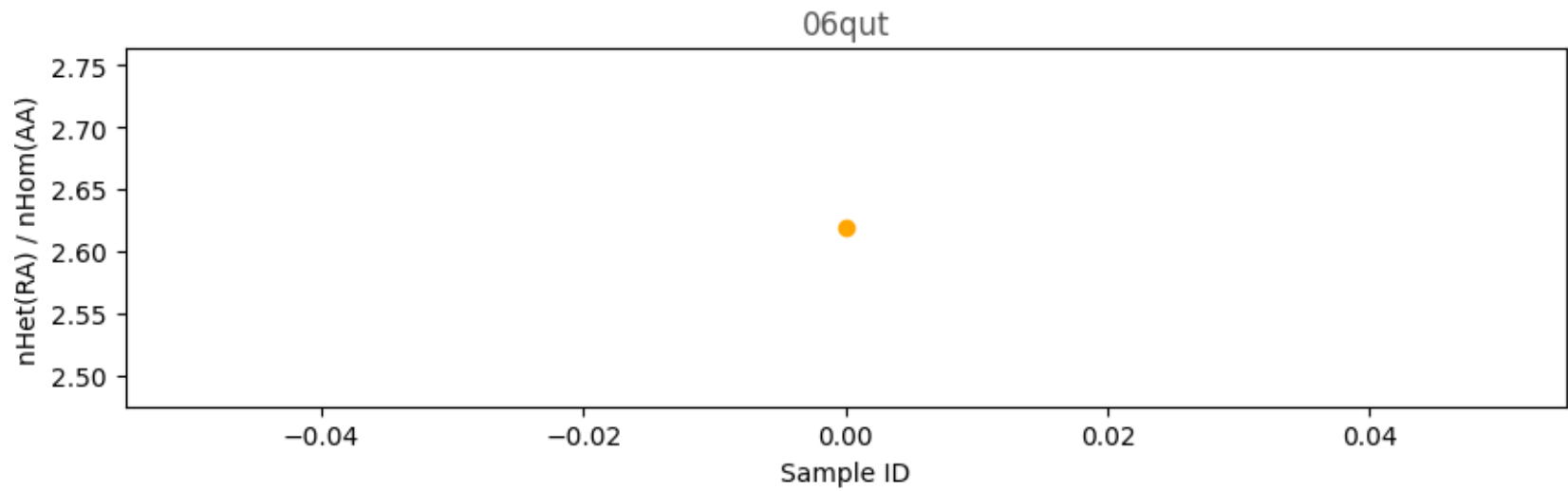
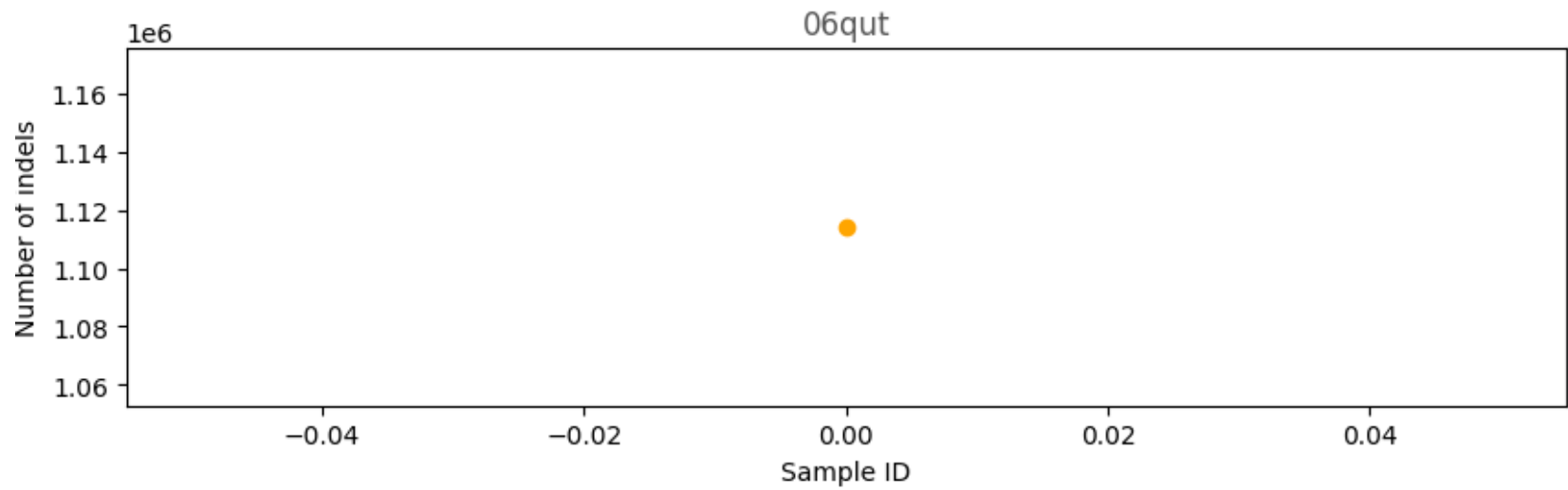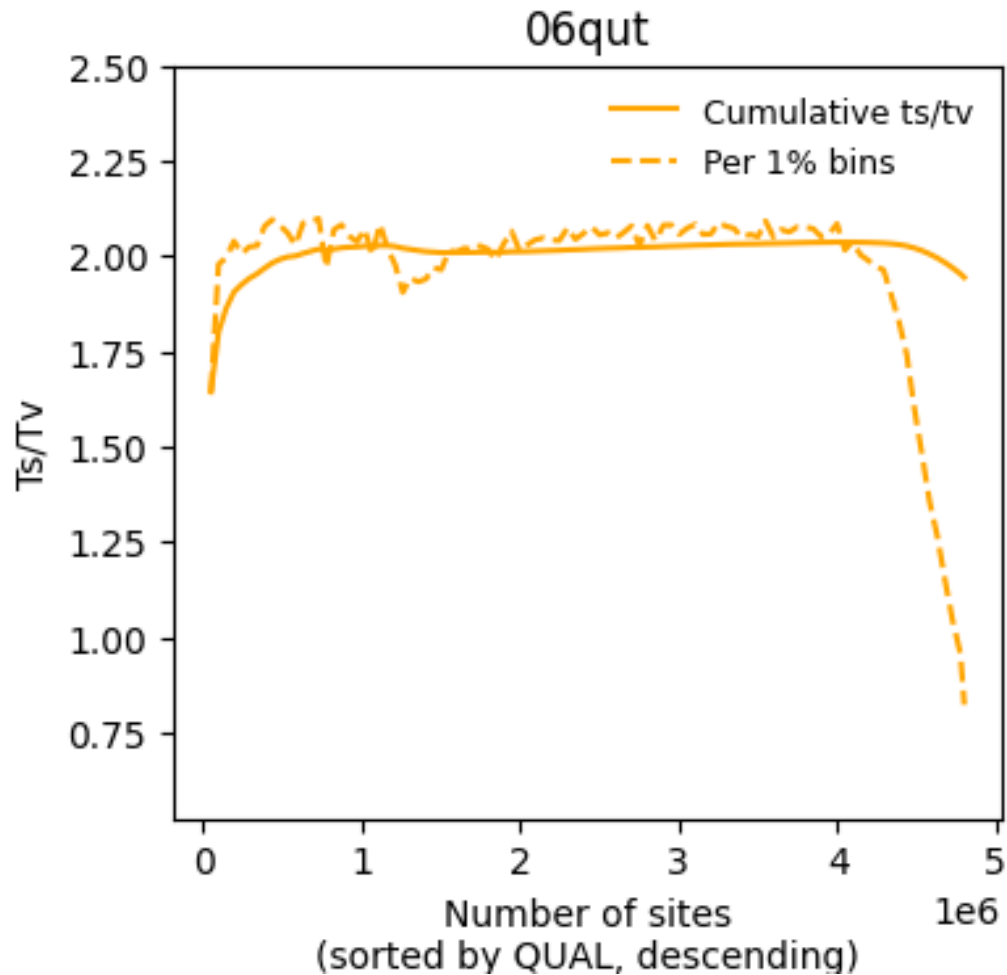# Singletons by sample (hets and homs)
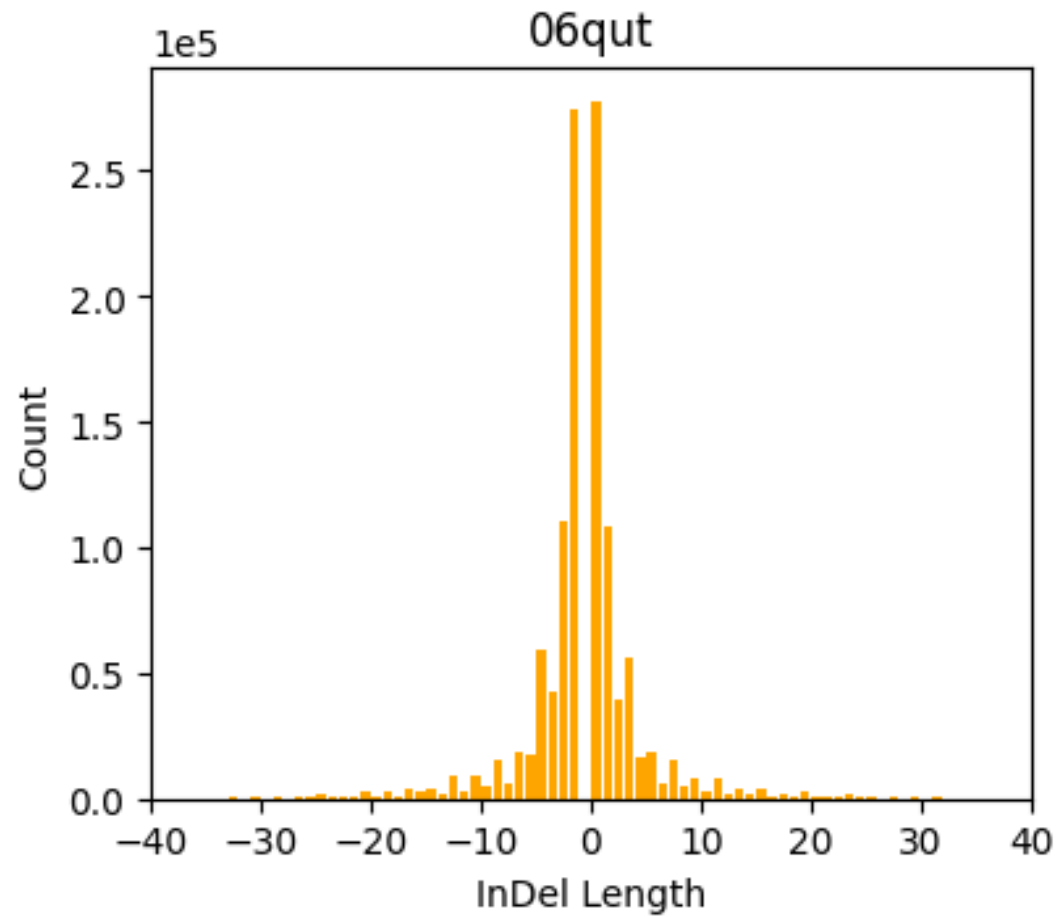
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

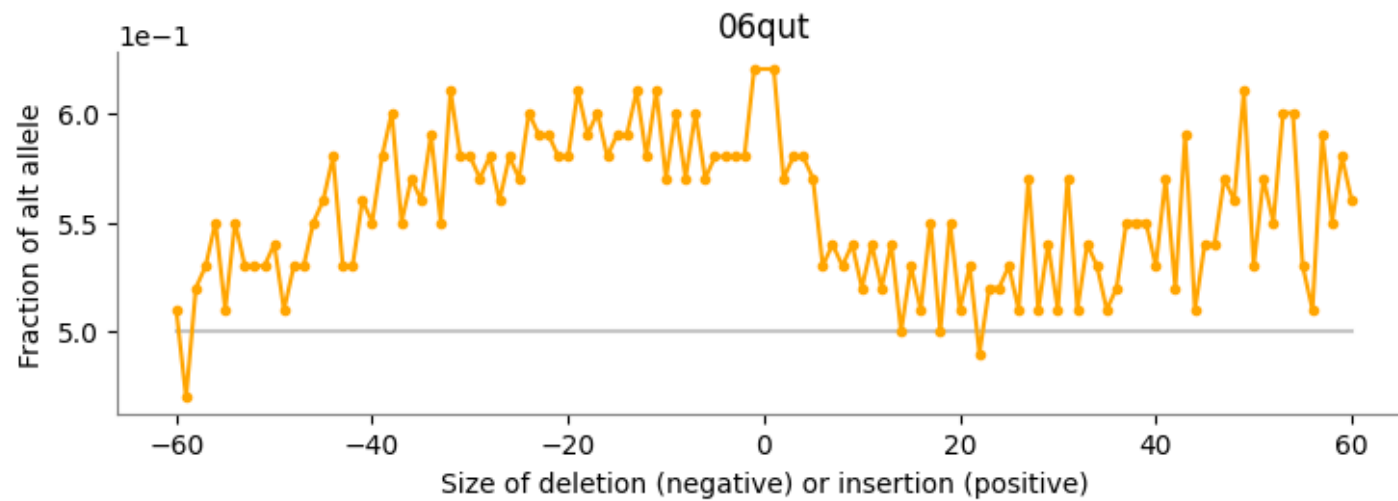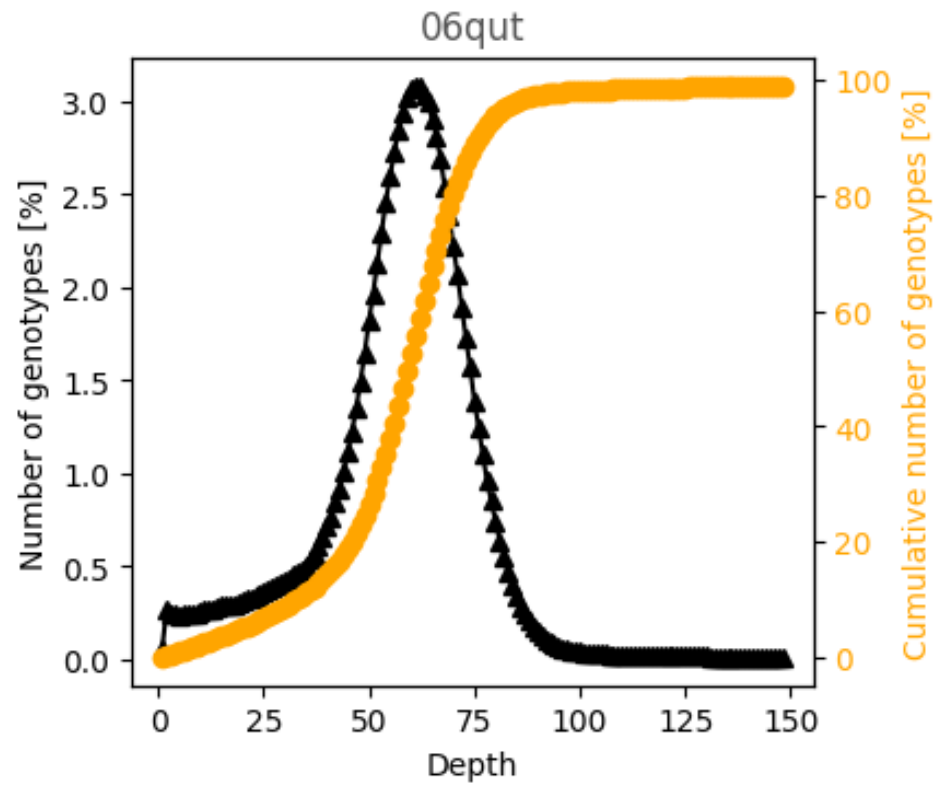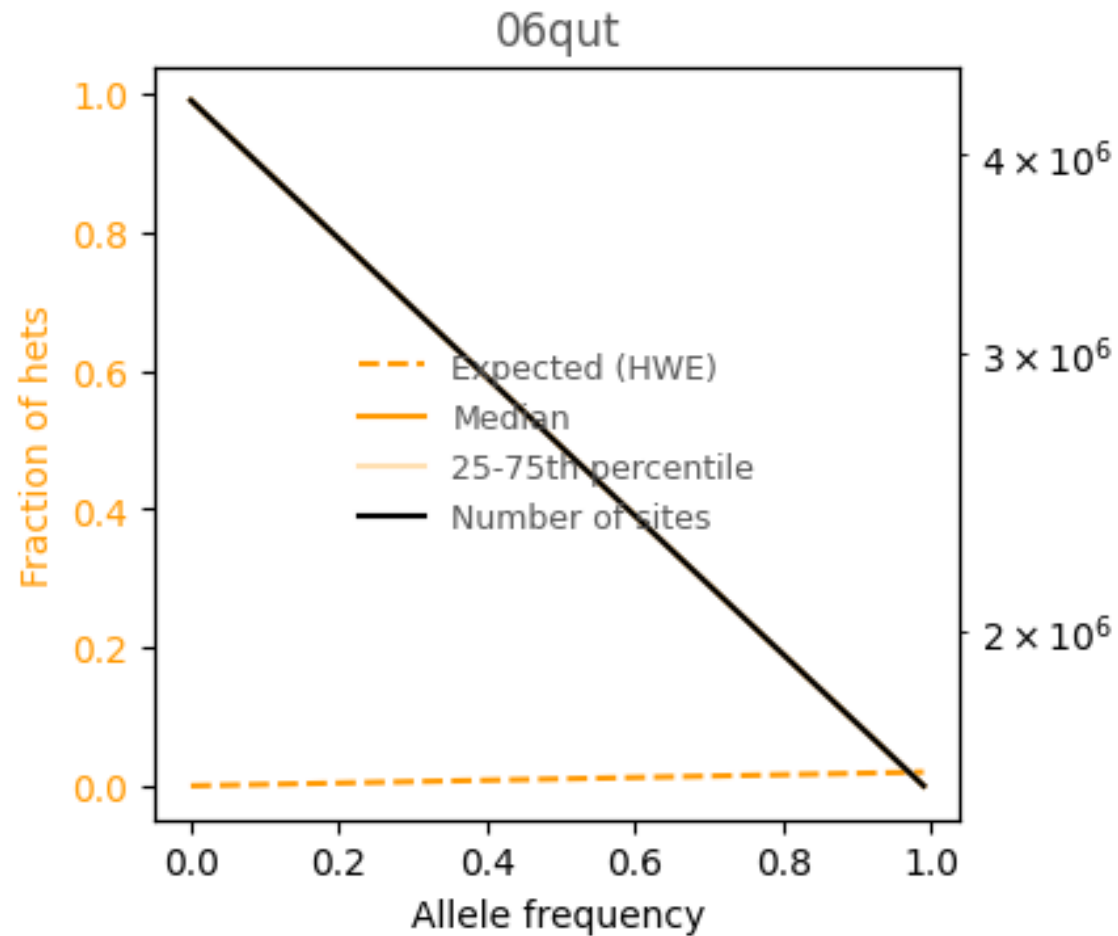# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types