Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
61cht	4,027,457	1.93	1.94	930,948	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
61cht	61.7%	1.90	65.7%	91,616	2,024

- 61cht .. `/ngc/projects2/gm/data/archive/2022/variants/snv/61chtbecm-111115085852-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220225_A01411_AH2CHTDSX3-RHGM_LABKA_WGS-` `WGS03869_snv_germline_raw.haplotype_caller.vcf.gz`
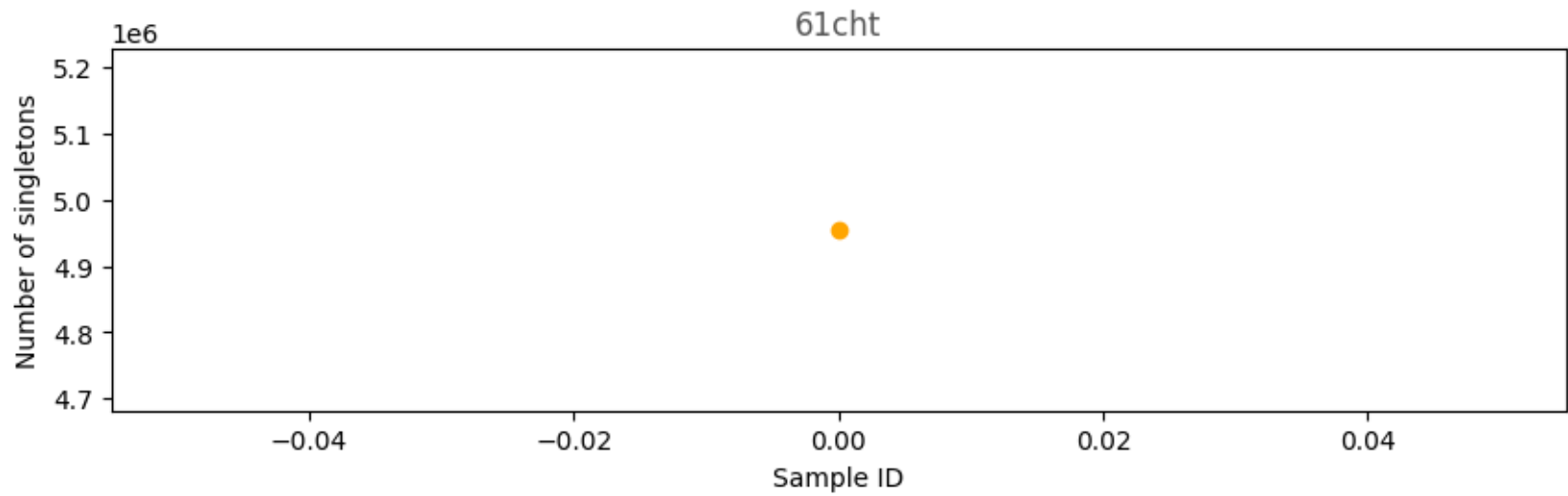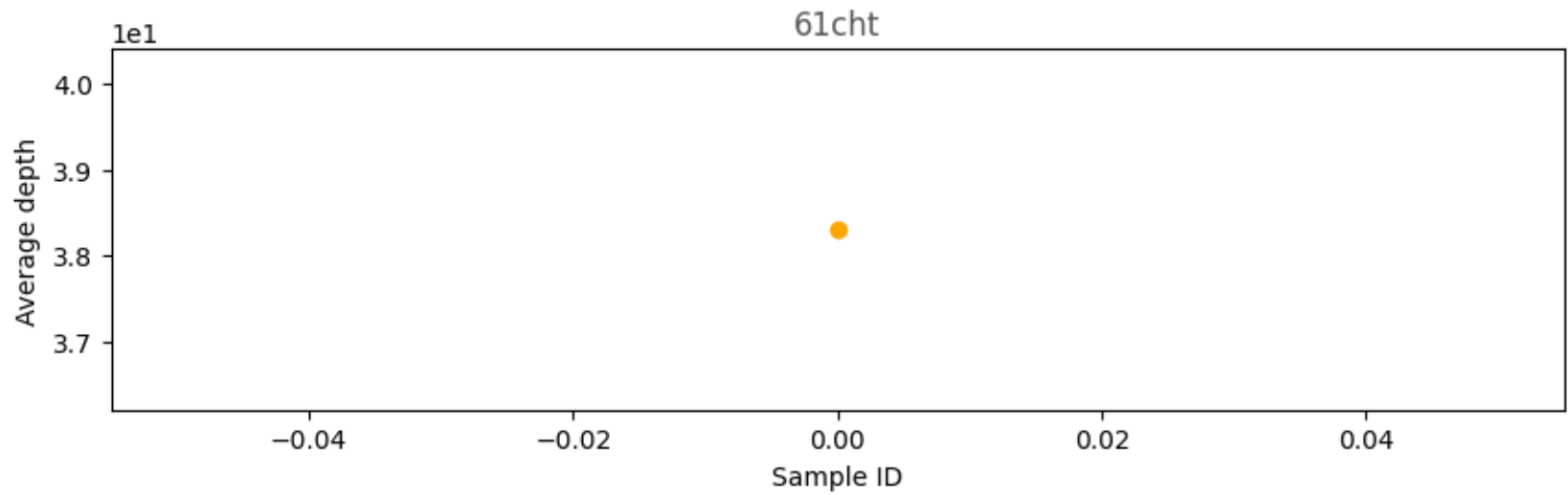
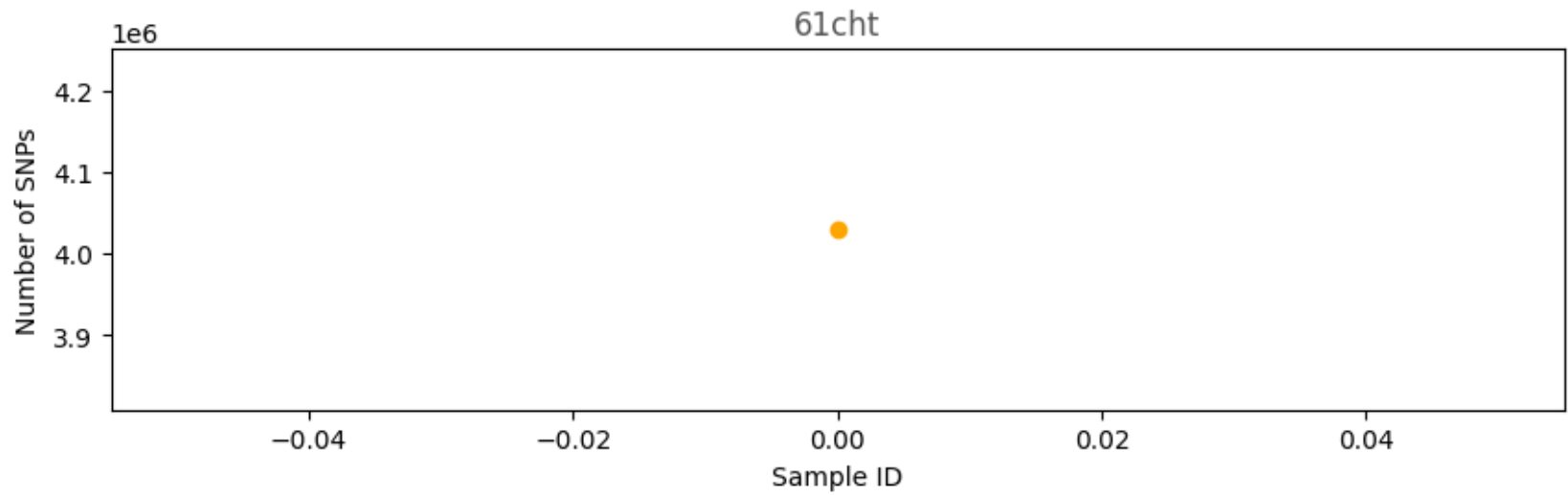# Ts/Tv by sample
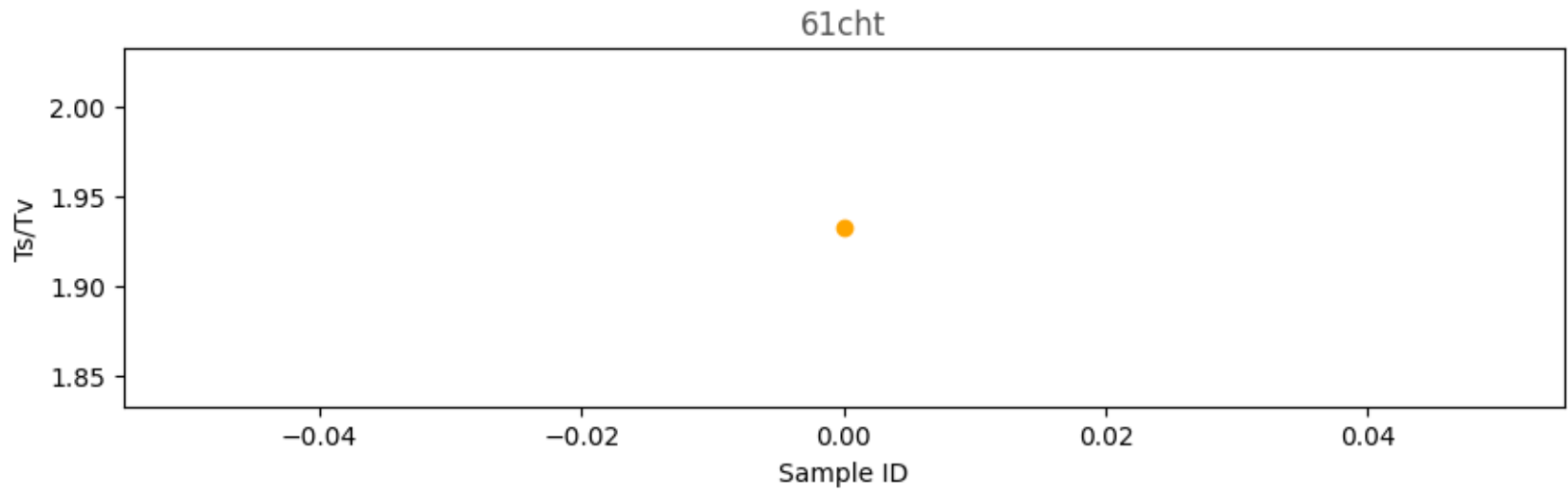
Hets vs non-ref Homs by sample
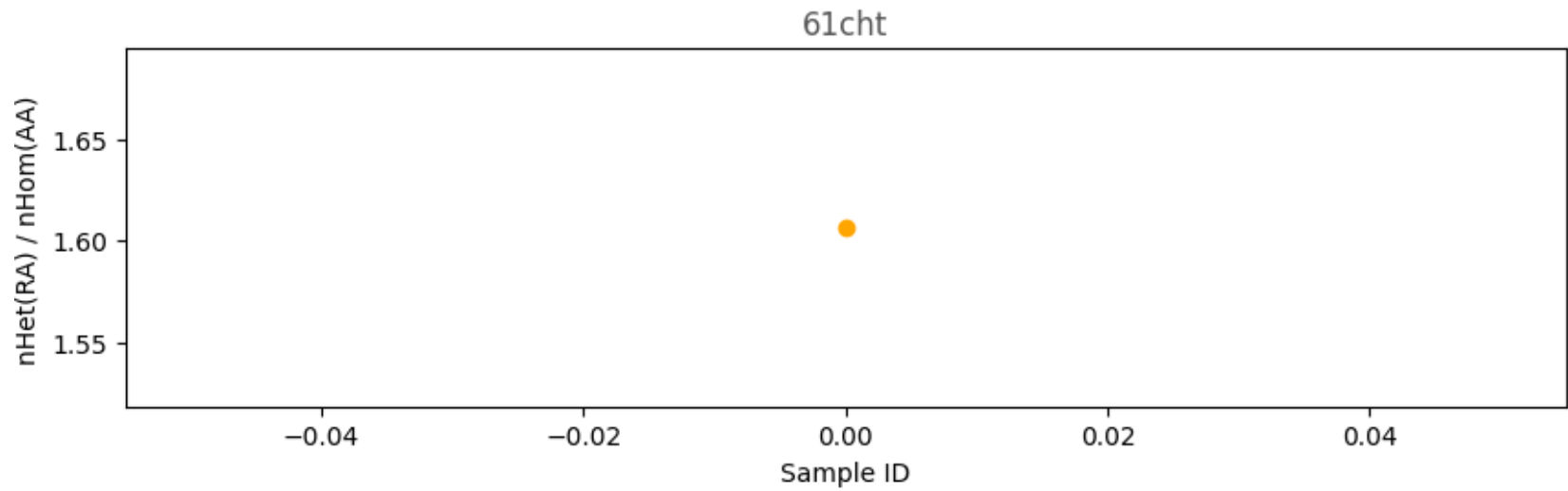
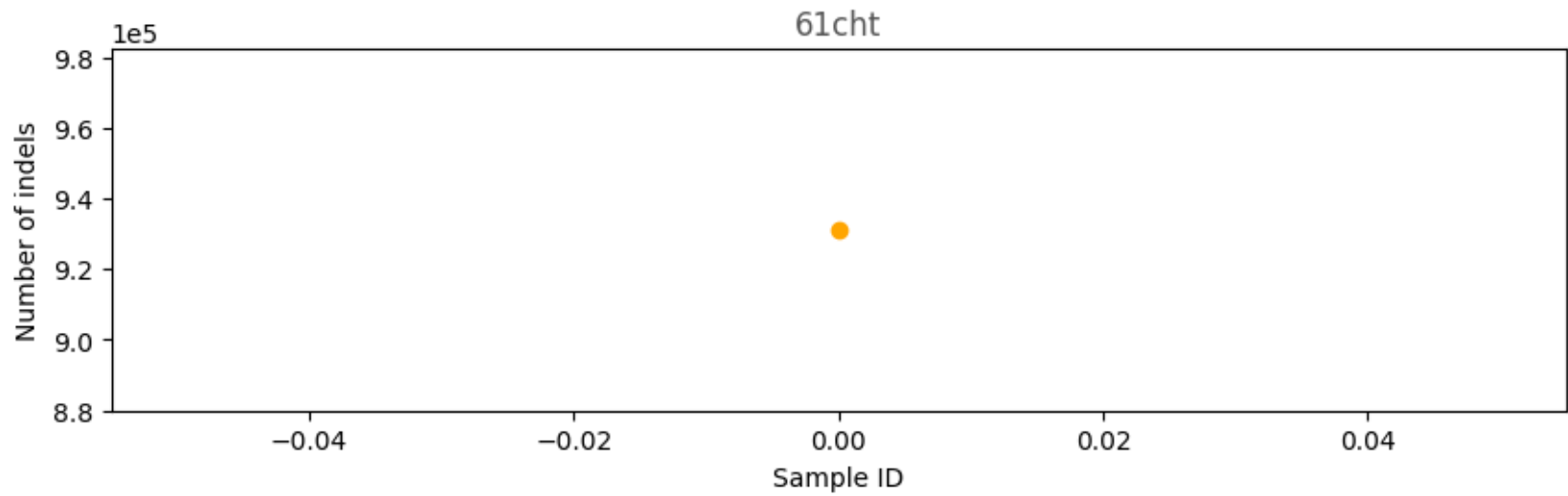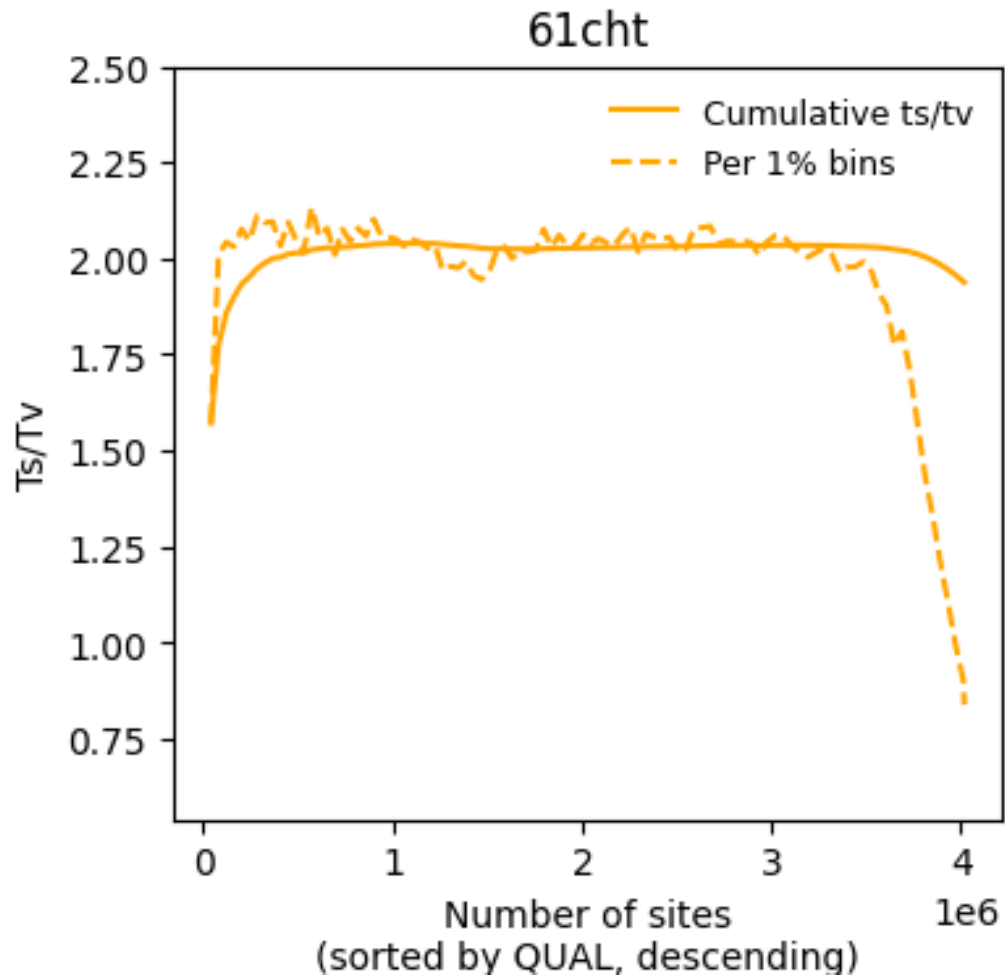# Singletons by sample (hets and homs)
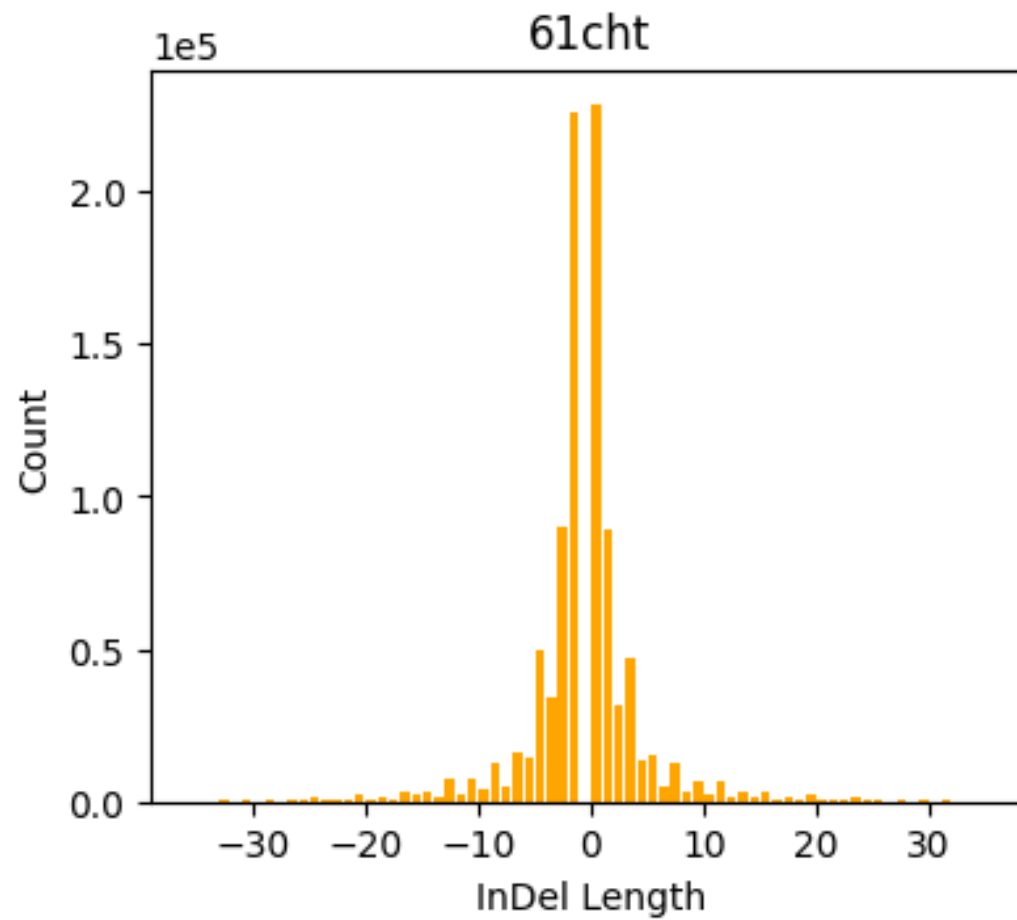
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

8

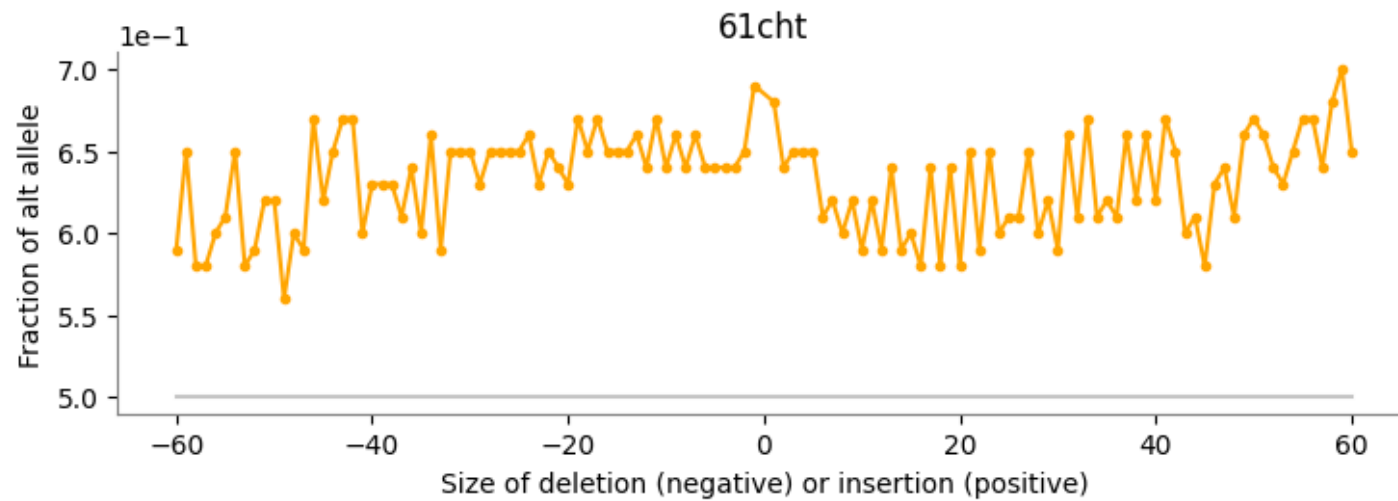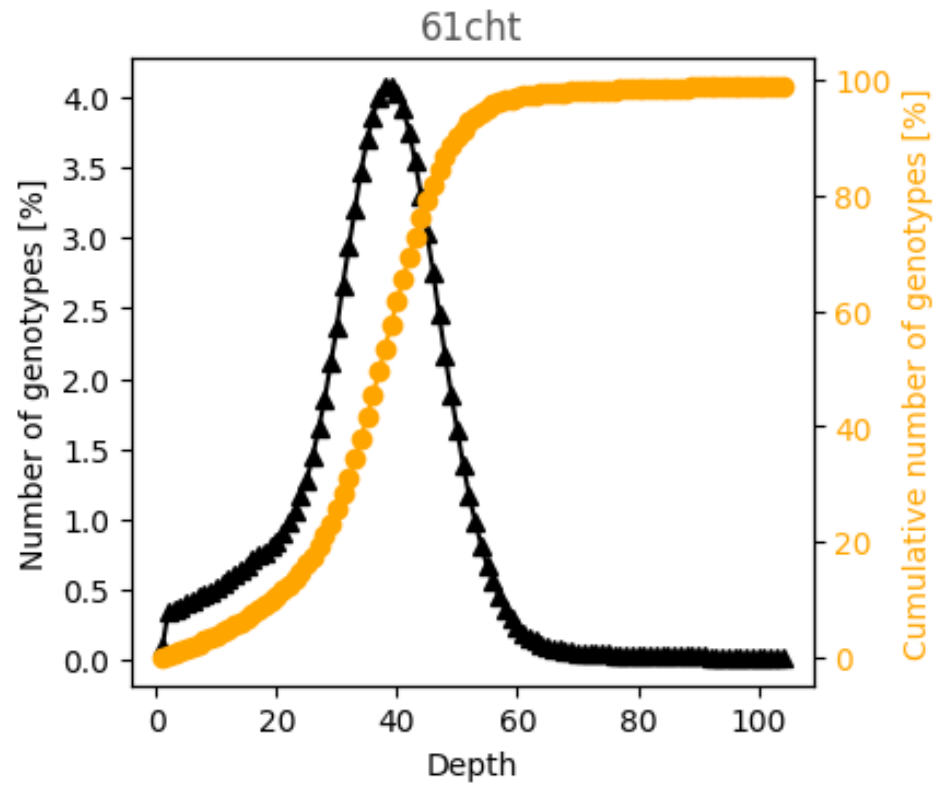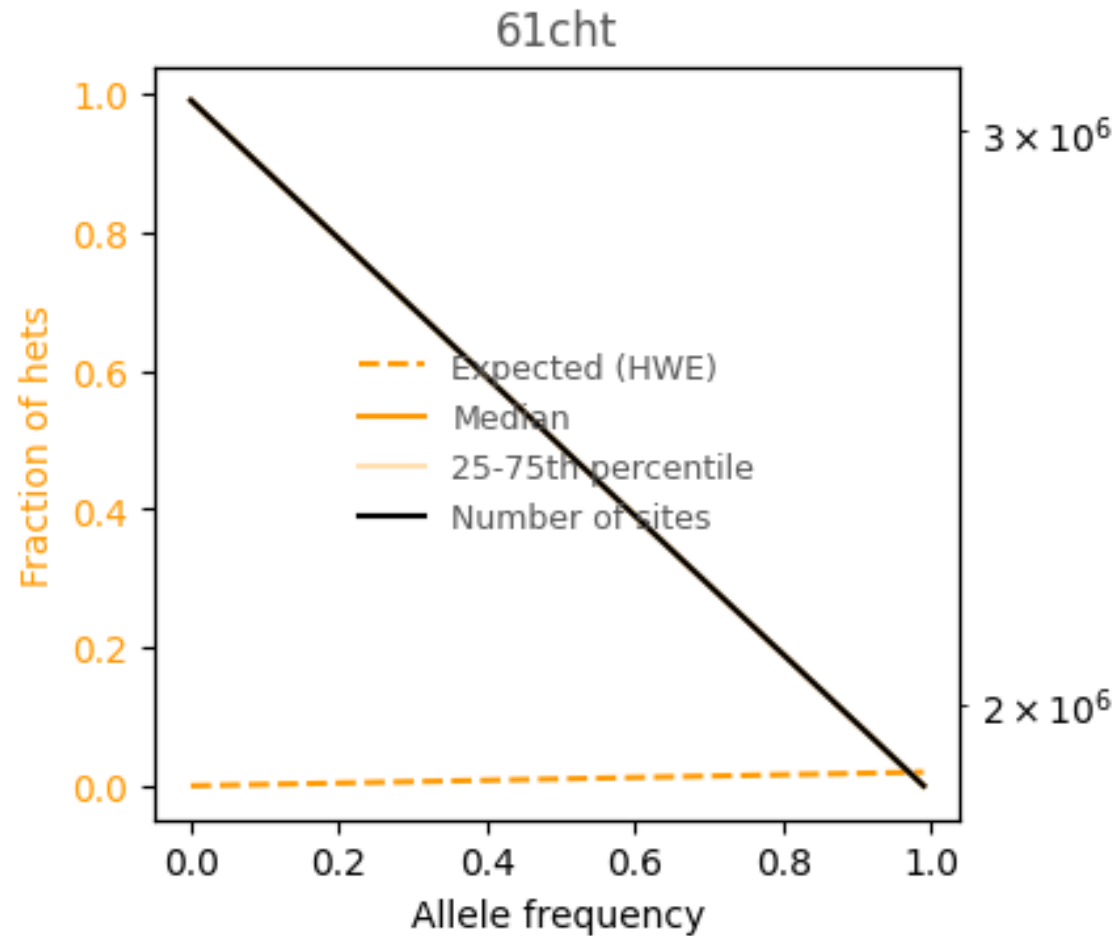# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types