Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
62con	4,061,609	1.93	1.94	942,031	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
62con	61.9%	1.91	66.2%	95,186	1,918

- 62con .. `/ngc/projects2/gm/data/archive/2022/variants/snv/62contopm-103914601028-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01779-220928_A01176_BH3FHLDSX5-EXT_LAB` `KA_NGCWGS-NGCWGS05613_22RKG009983_snv_germline_raw.haplotype_caller.vcf.gz`
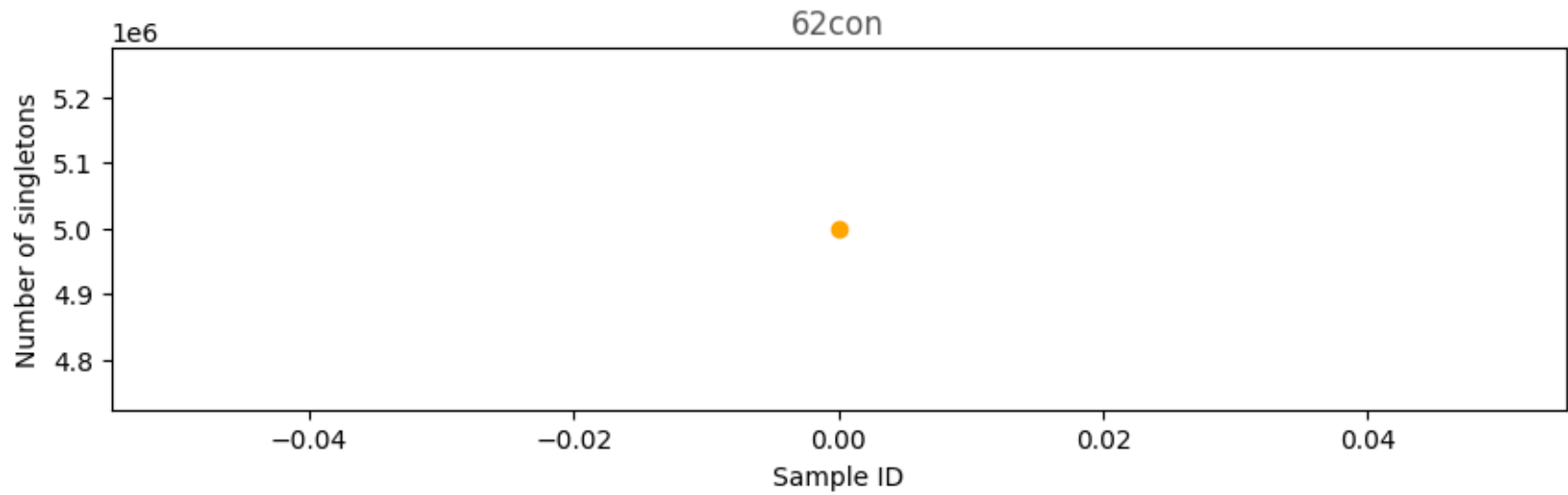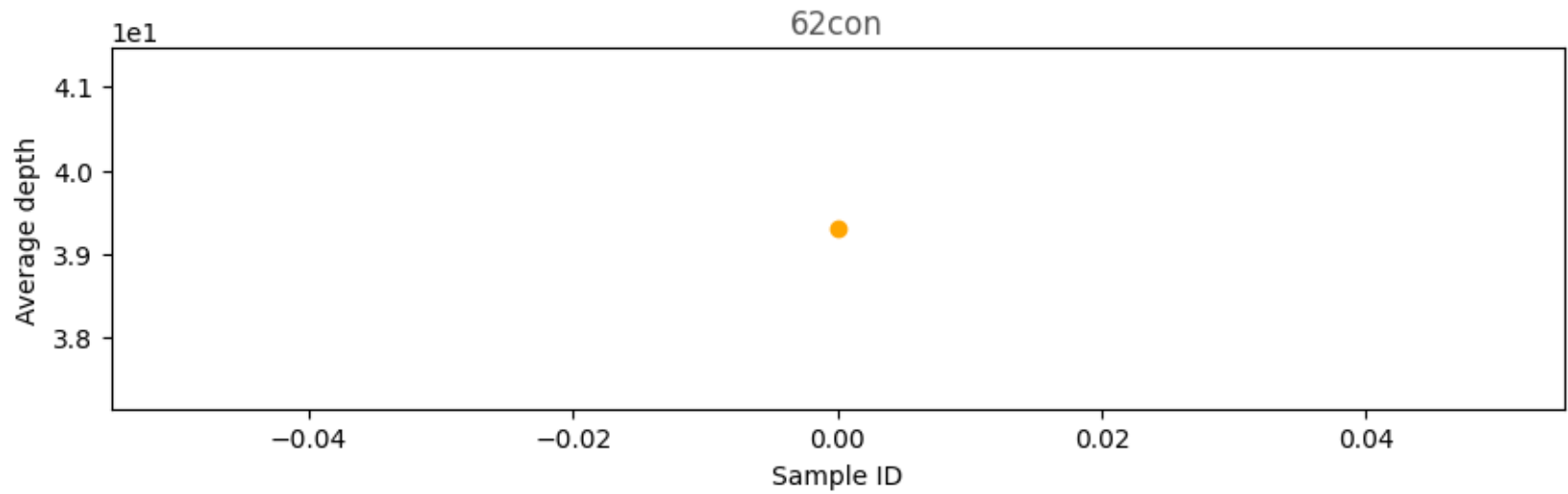
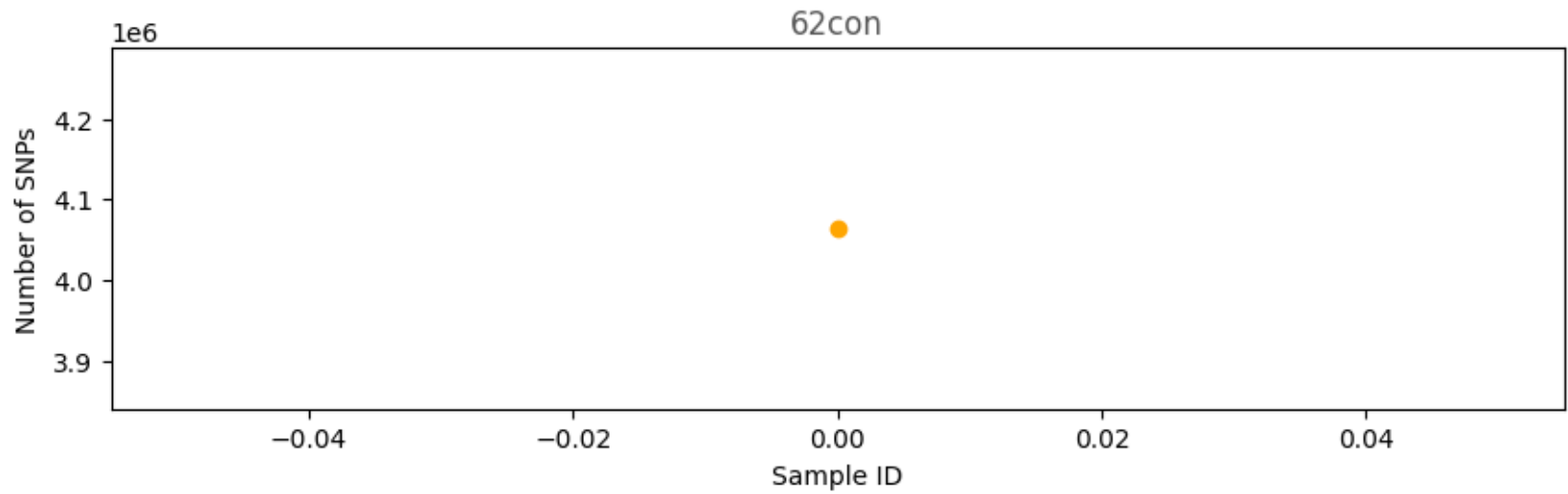# Ts/Tv by sample
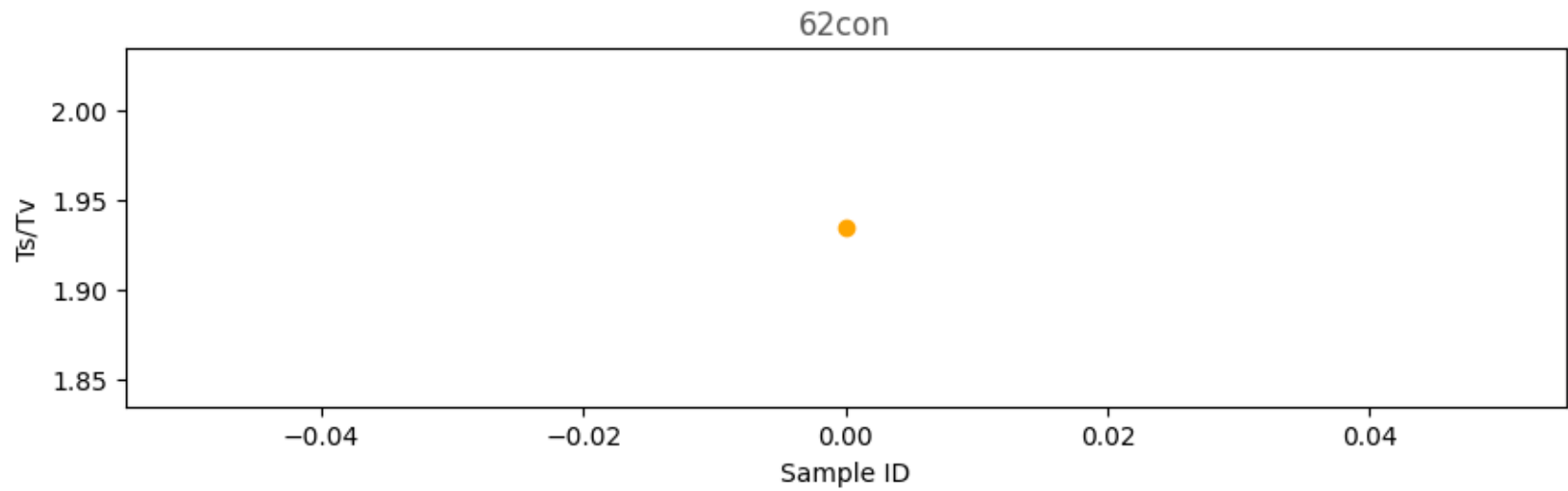
Hets vs non-ref Homs by sample
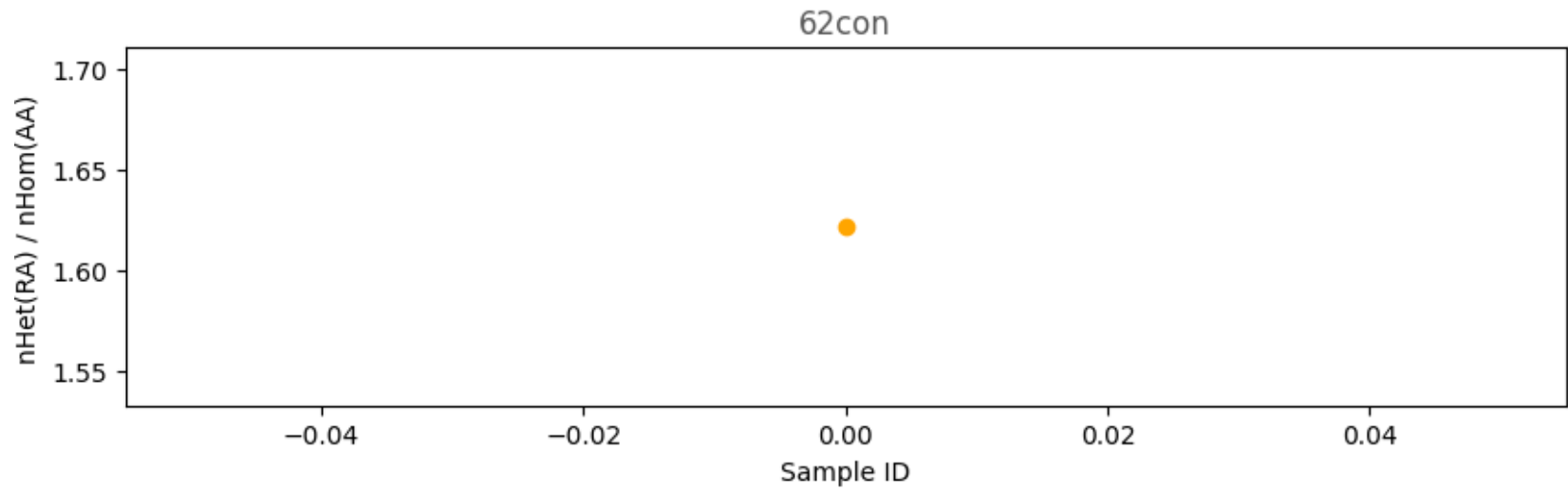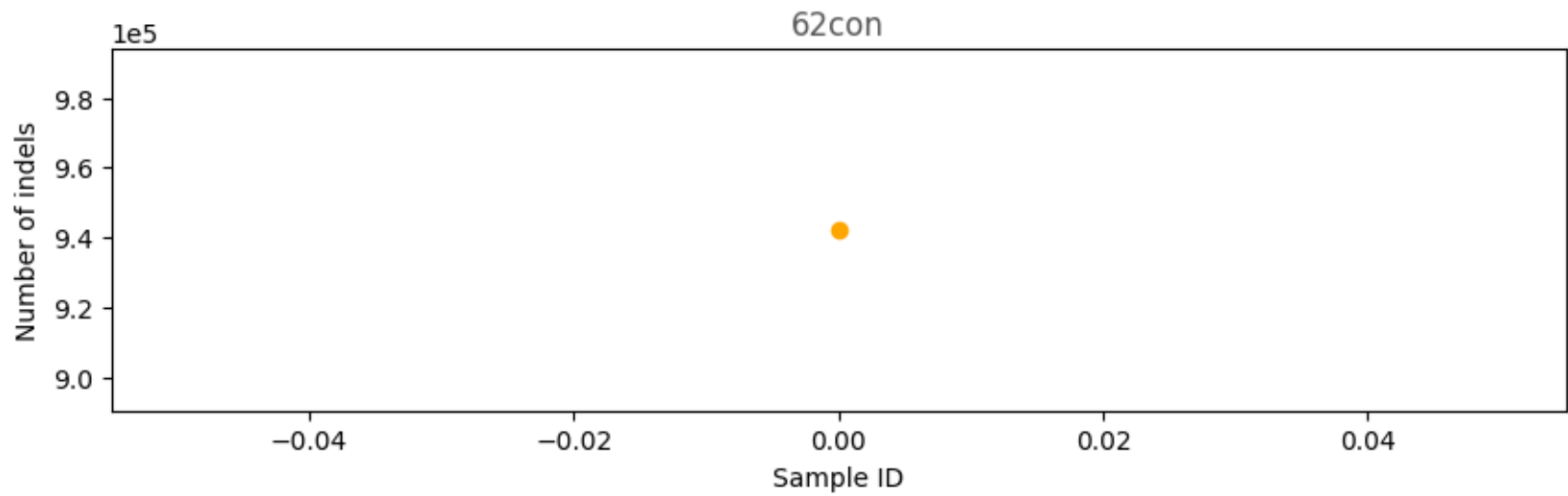
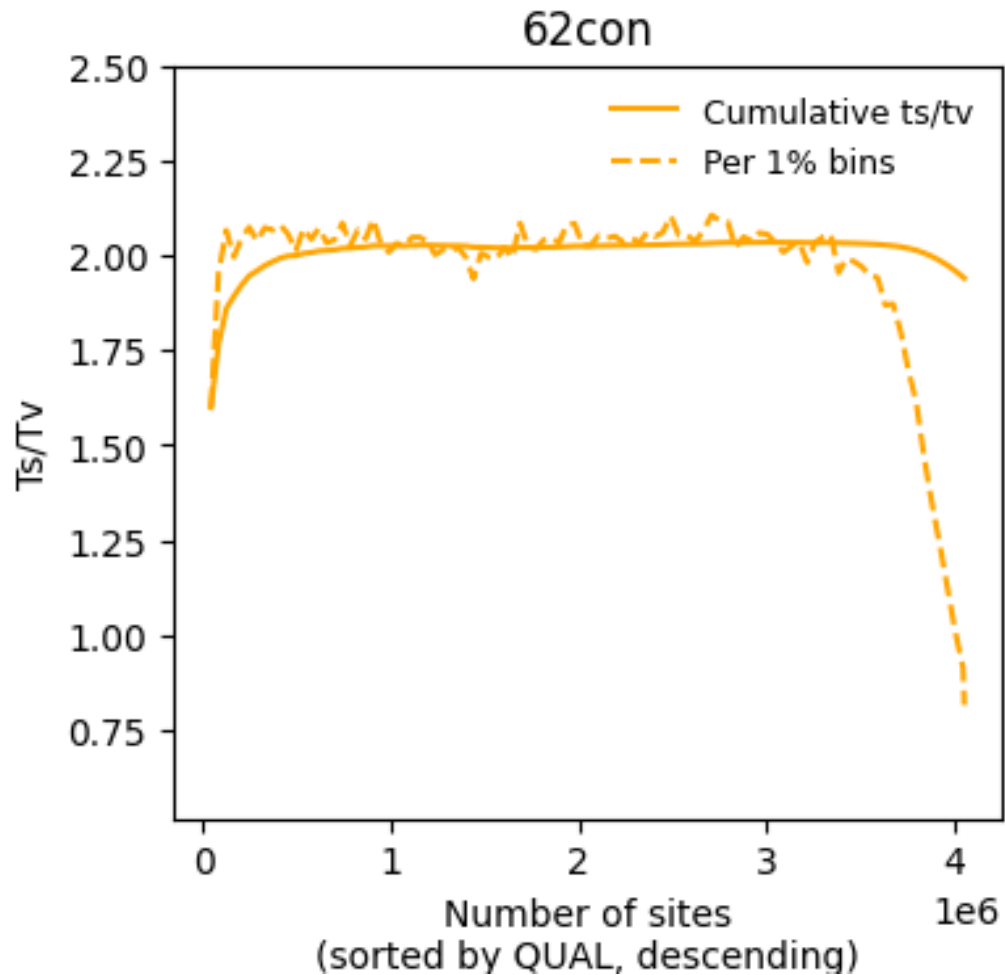# Singletons by sample (hets and homs)
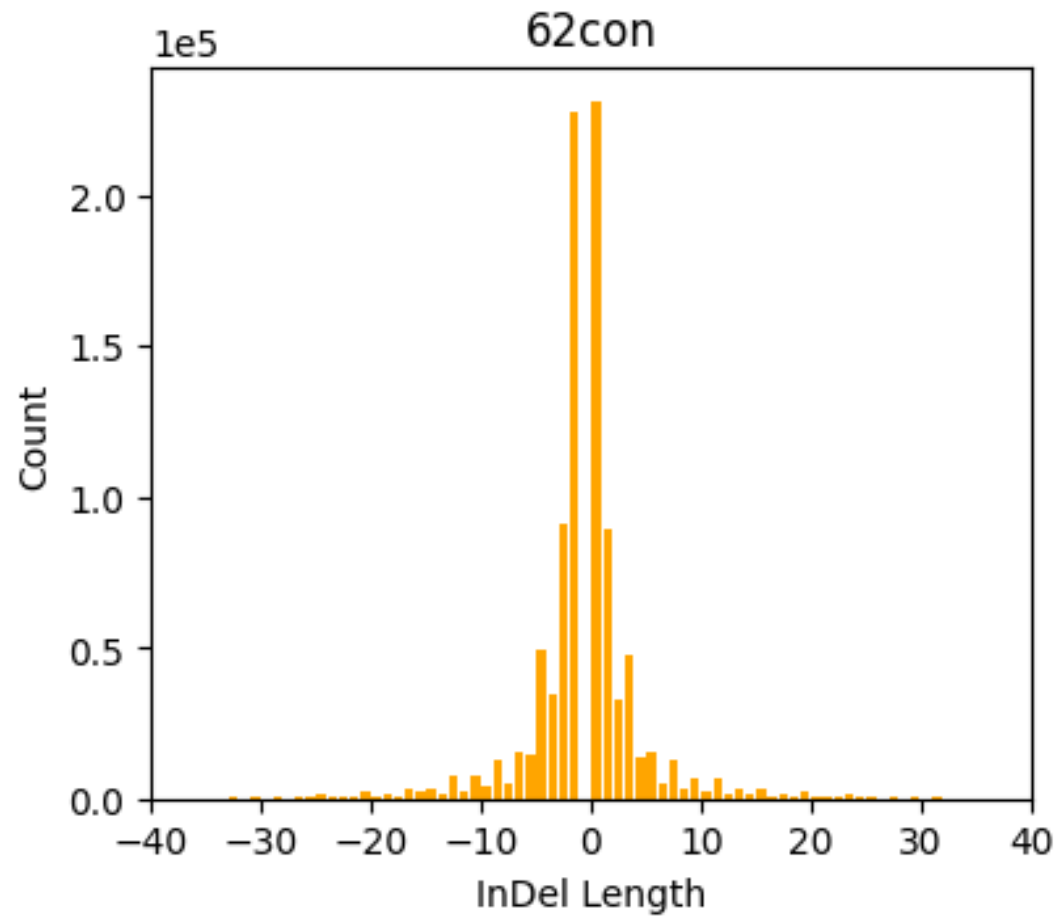
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

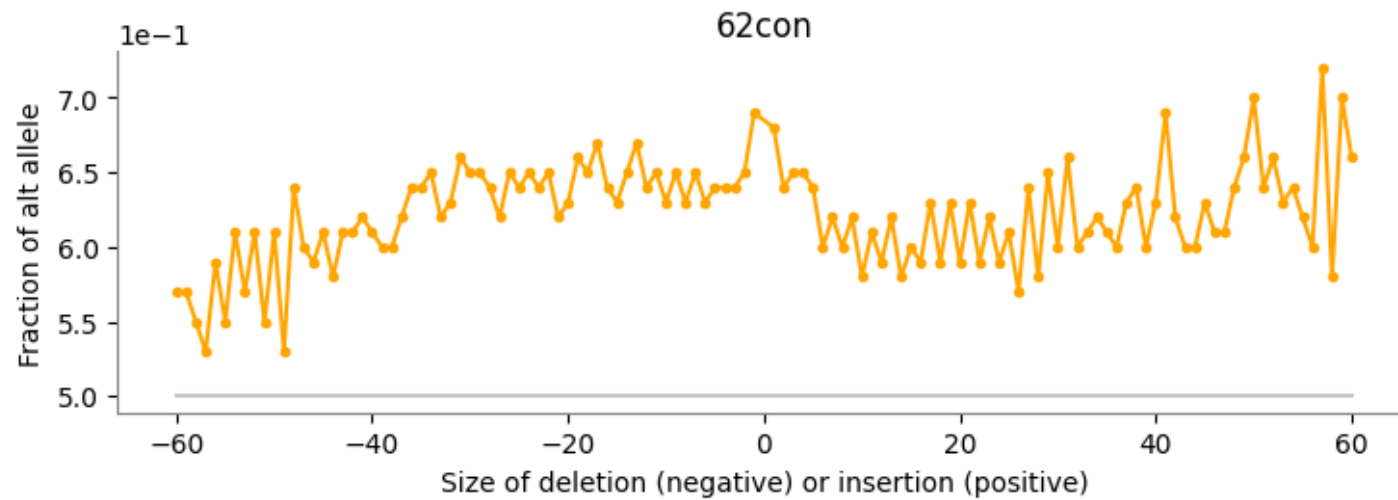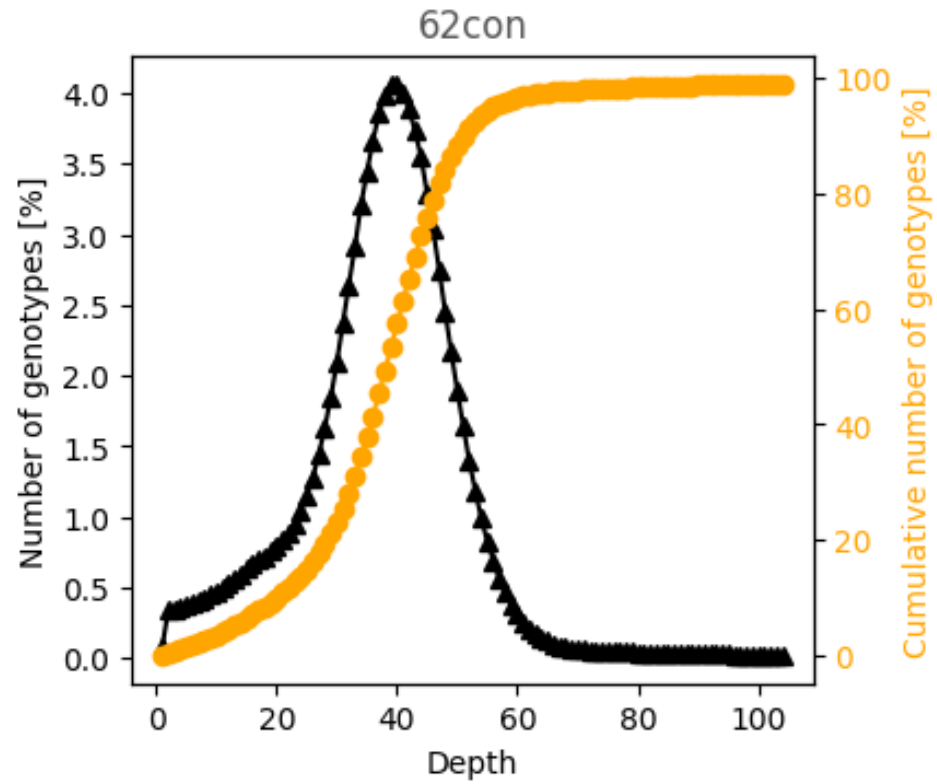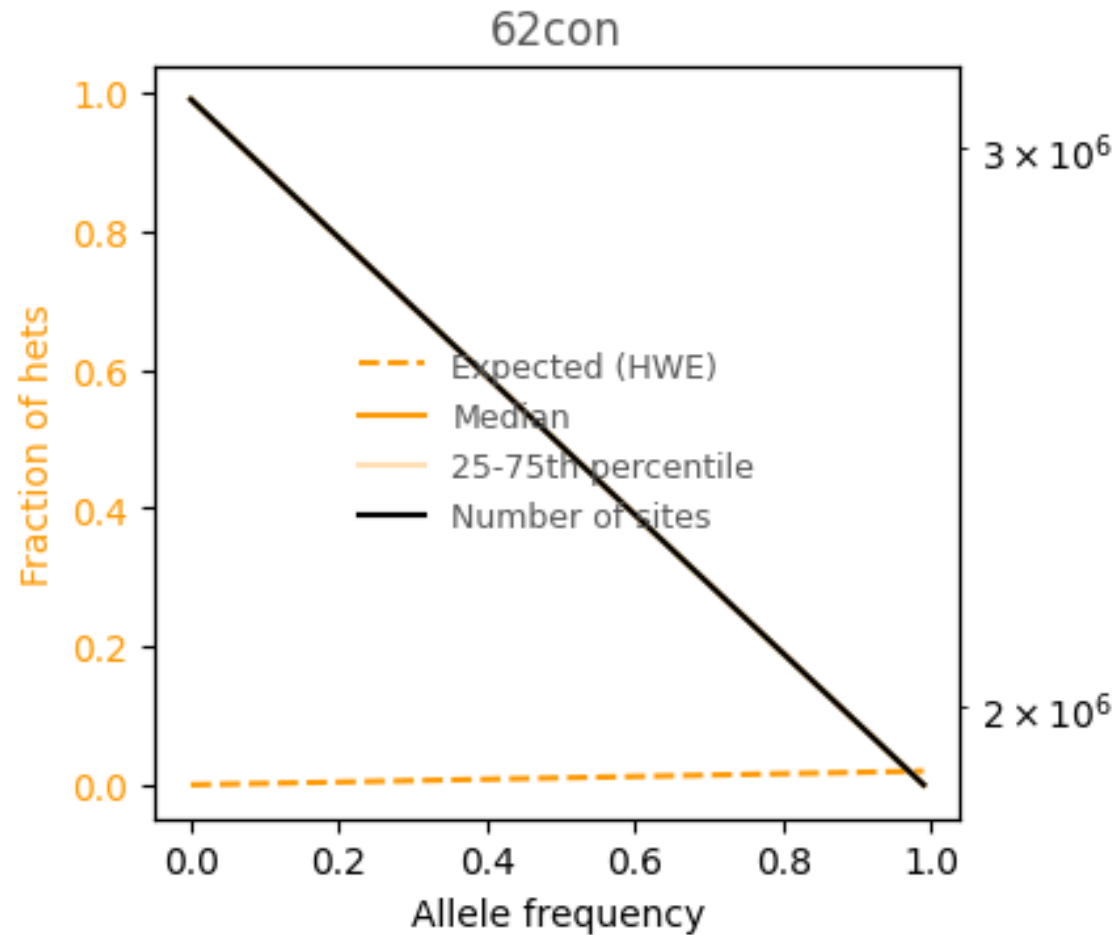# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types