Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
71cob	4,099,076	1.93	1.94	948,111	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
71cob	63.2%	1.90	68.0%	98,821	2,069

- 71cob .. `/ngc/projects2/gm/data/archive/2022/variants/snv/71cobescf-103901079620-Normal_B`
   `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01106-220817_A01411_AHNVWWDSX3-EXT_LAB`
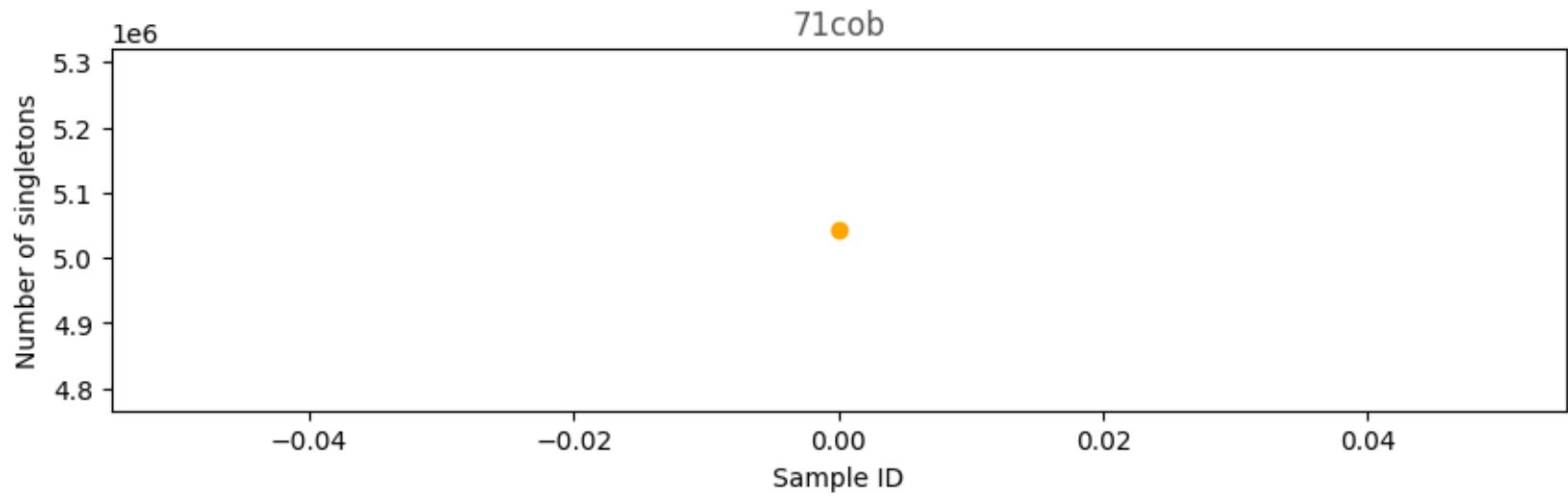   `KA_NGCWGS-NGCWGS04841_snv_germline_raw.haplotype_caller.vcf.gz`
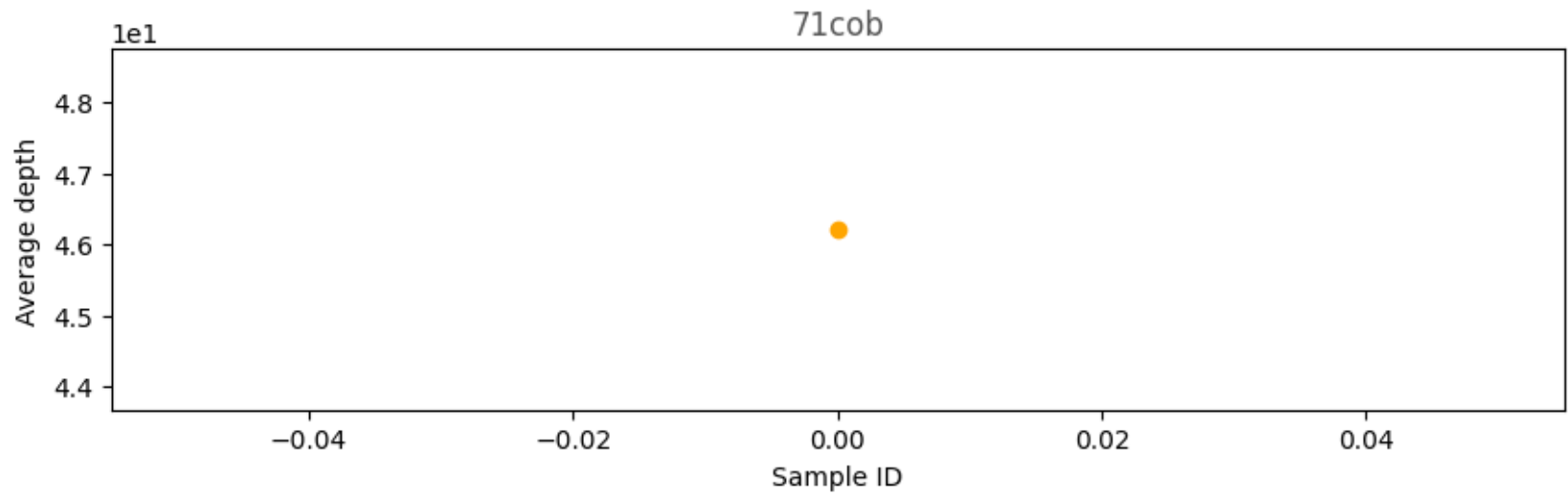
1
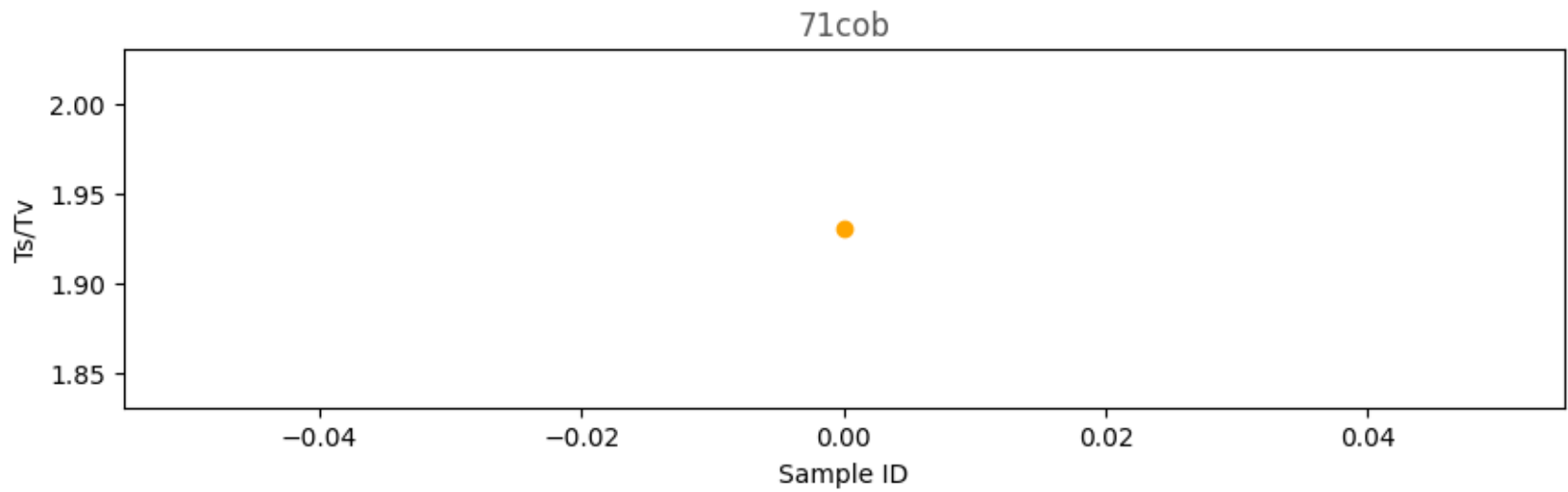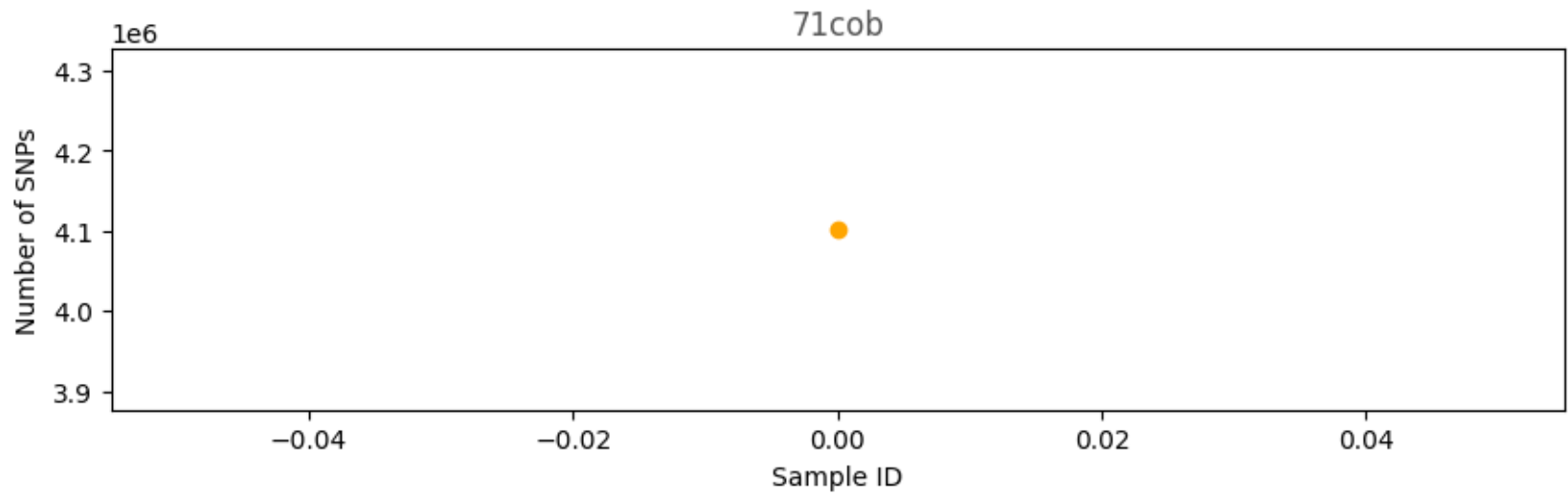
Ts/Tv by sample

Hets vs non-ref Homs by sample
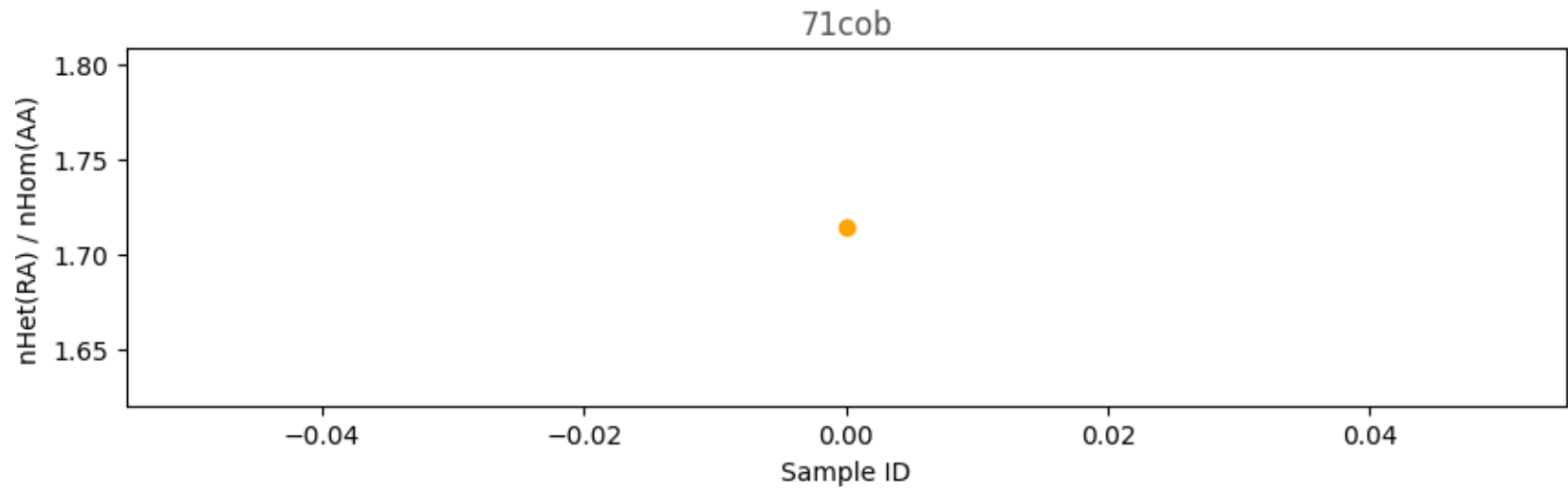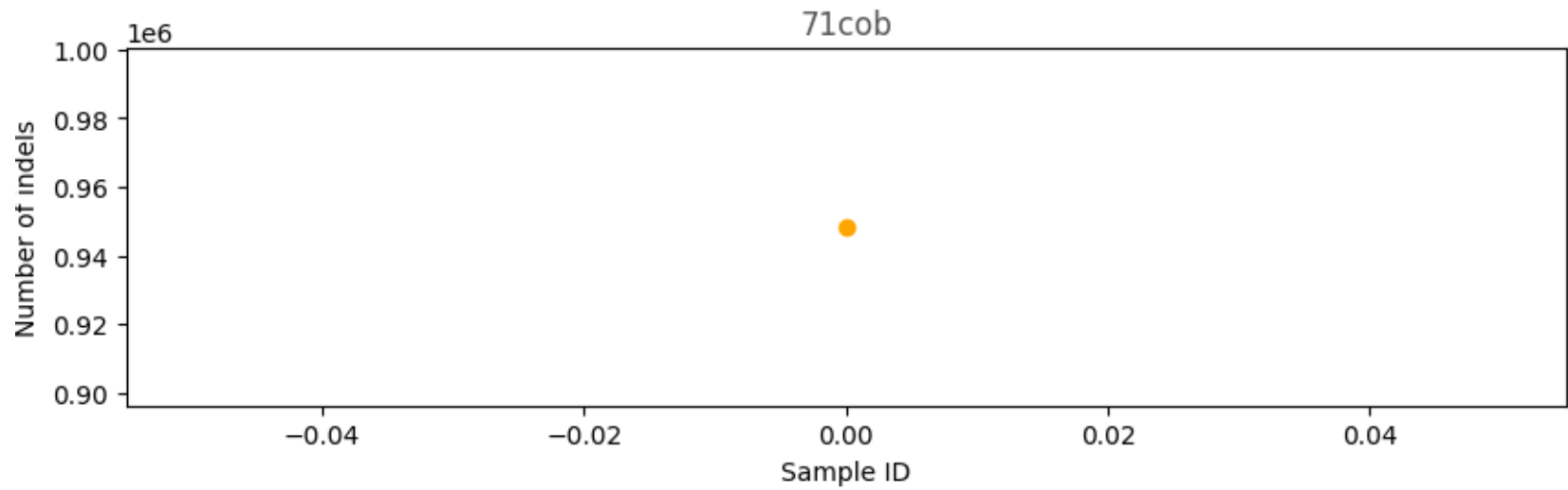
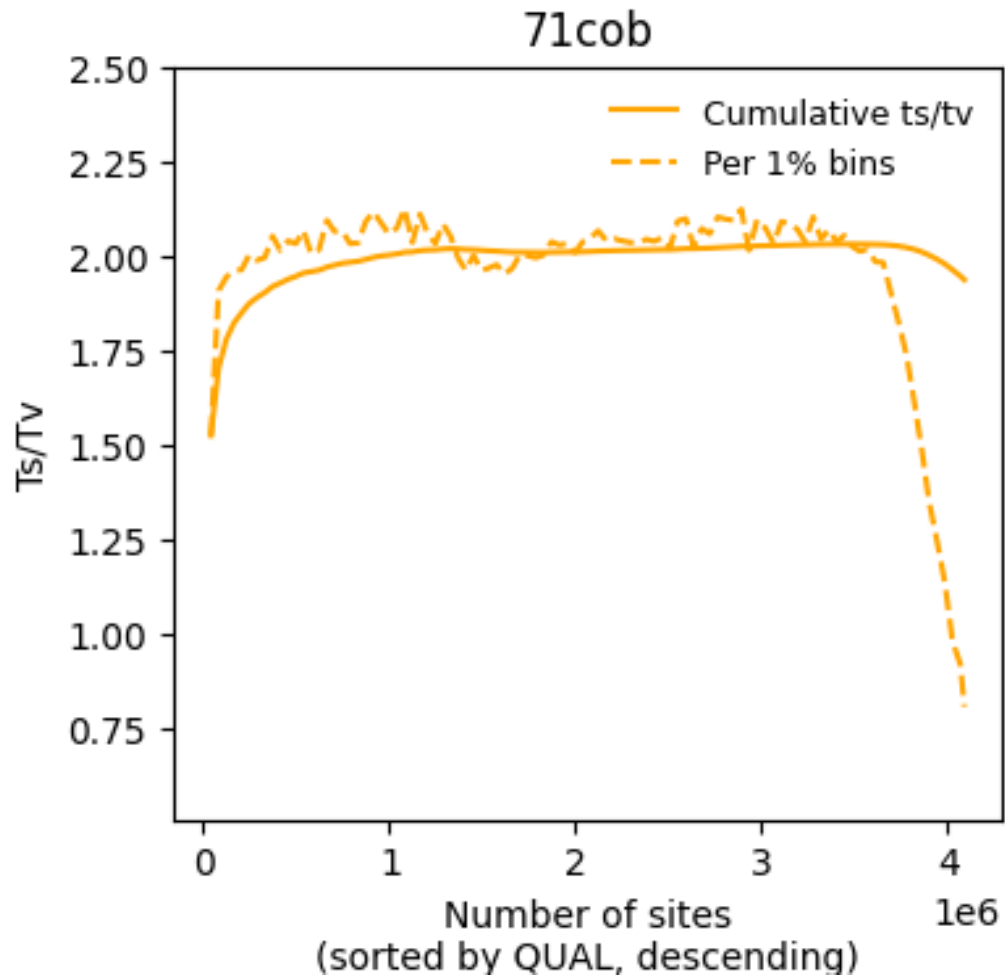# Singletons by sample (hets and homs)
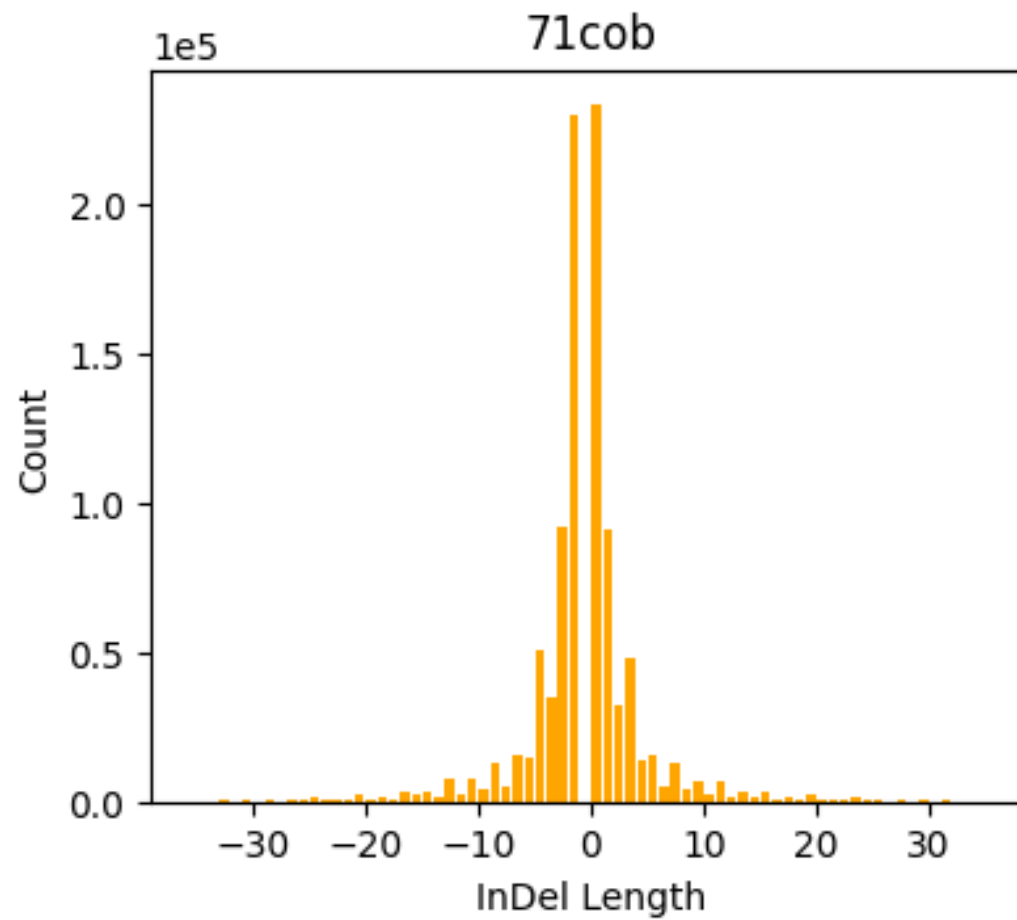
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

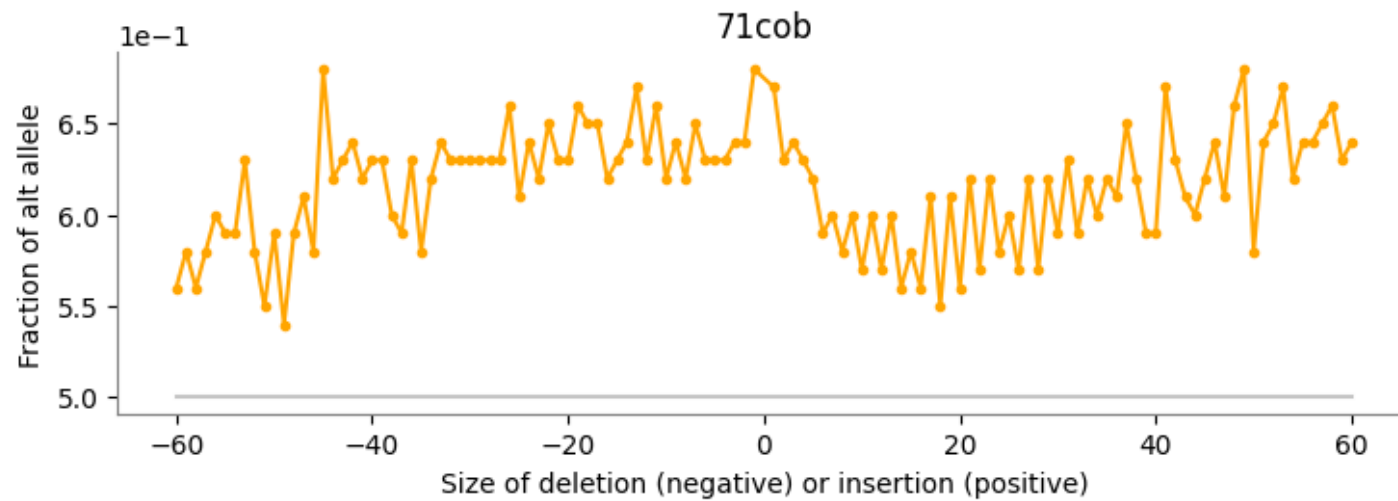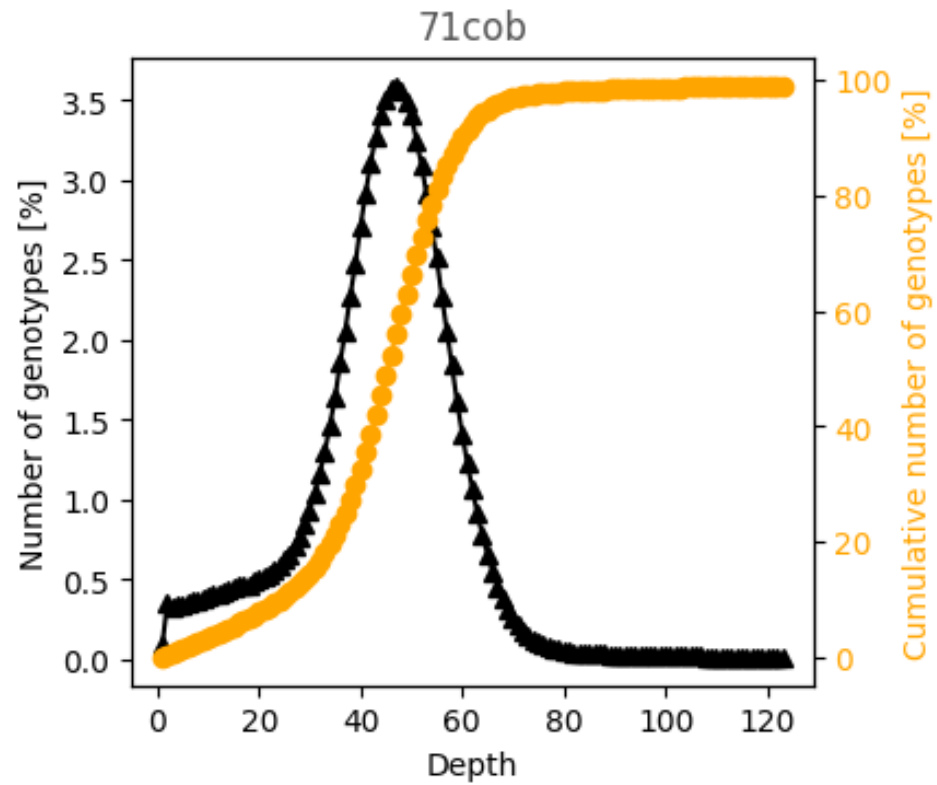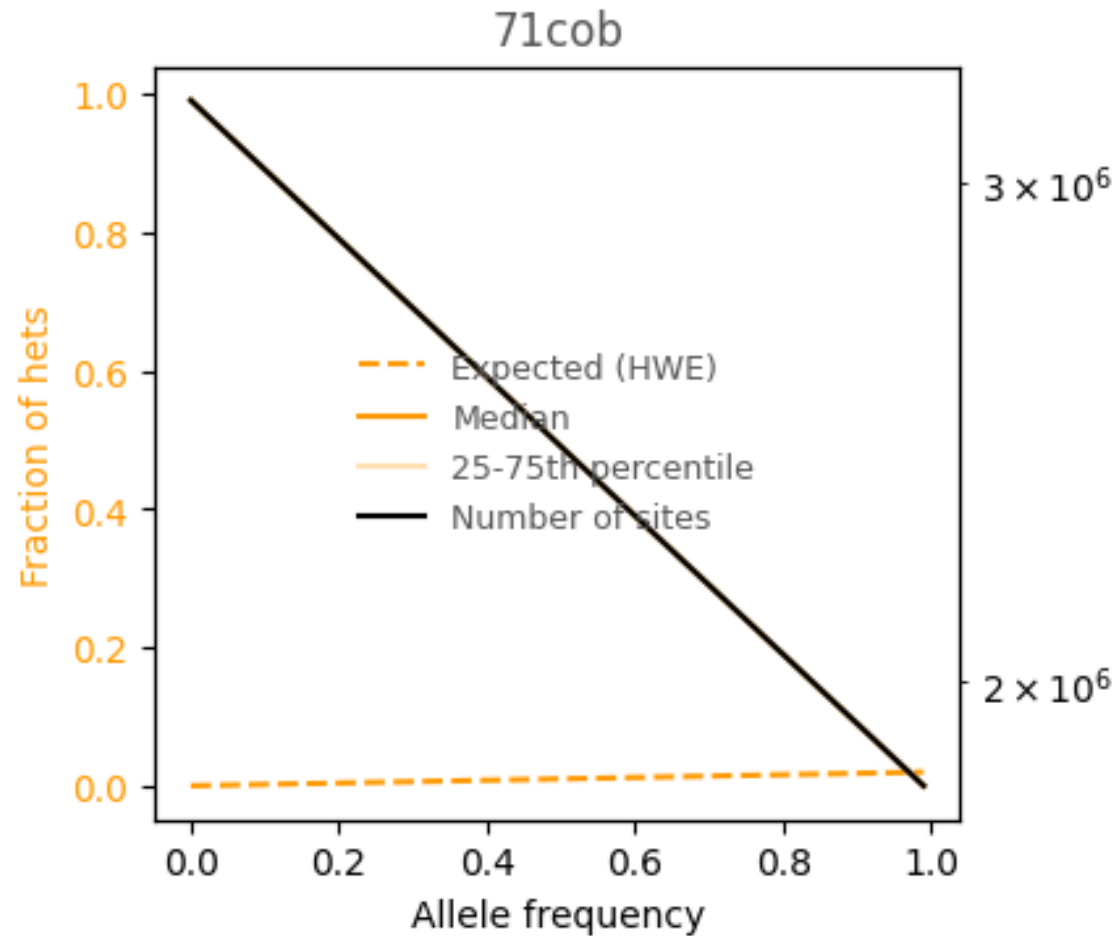# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types