Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
68iar	4,006,073	1.95	1.95	915,585	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
68iar	61.8%	1.92	65.1%	86,503	1,906

- 68iar .. `/ngc/projects2/gm/data/archive/2022/variants/snv/68iarowhm-103755214502-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00216-210629_A00559_BH2GFCDMXY-EXT_LAB`
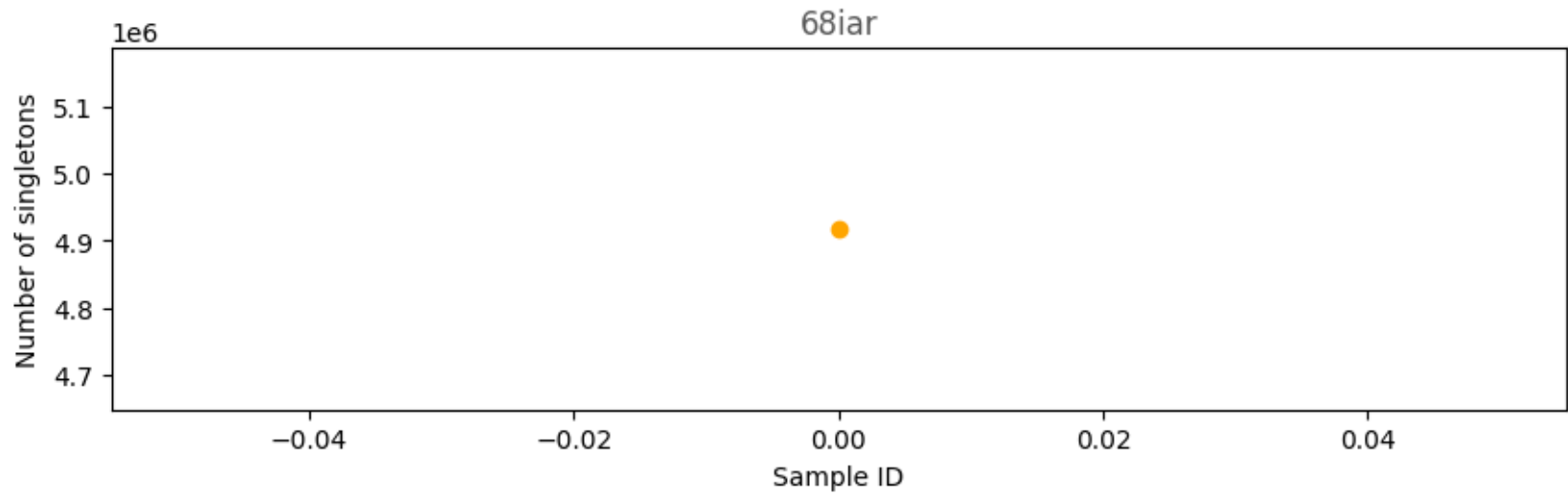 `KA_NGCWGS-NGCWGS00132_68d6phs89_snv_germline_raw.haplotype_caller.vcf.gz`
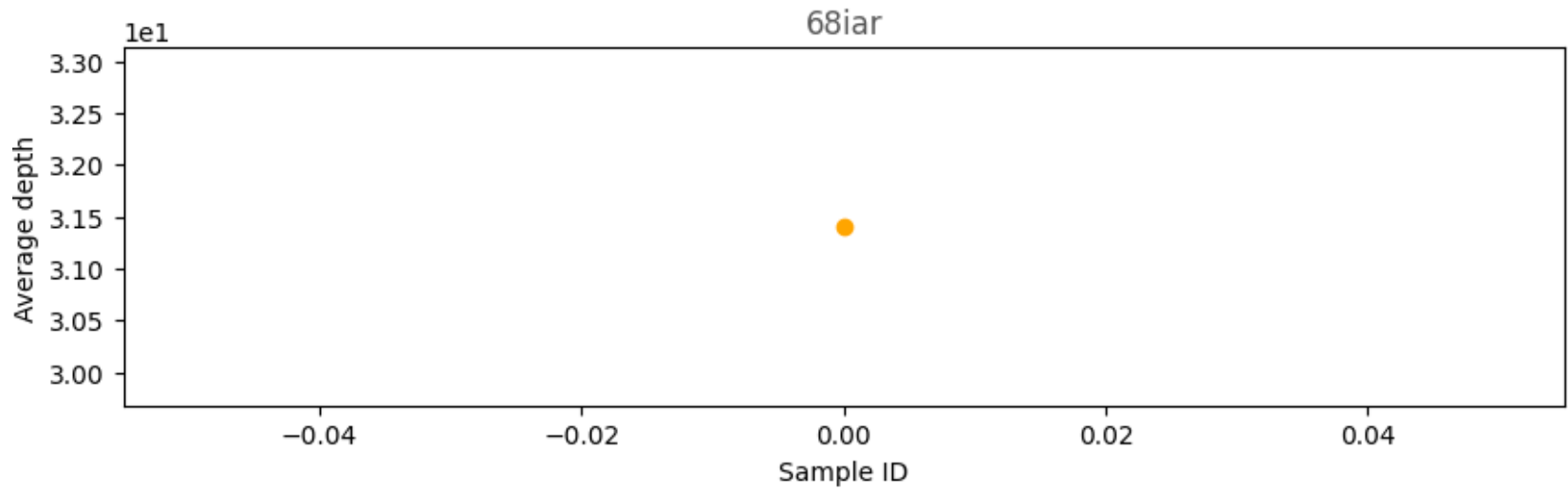
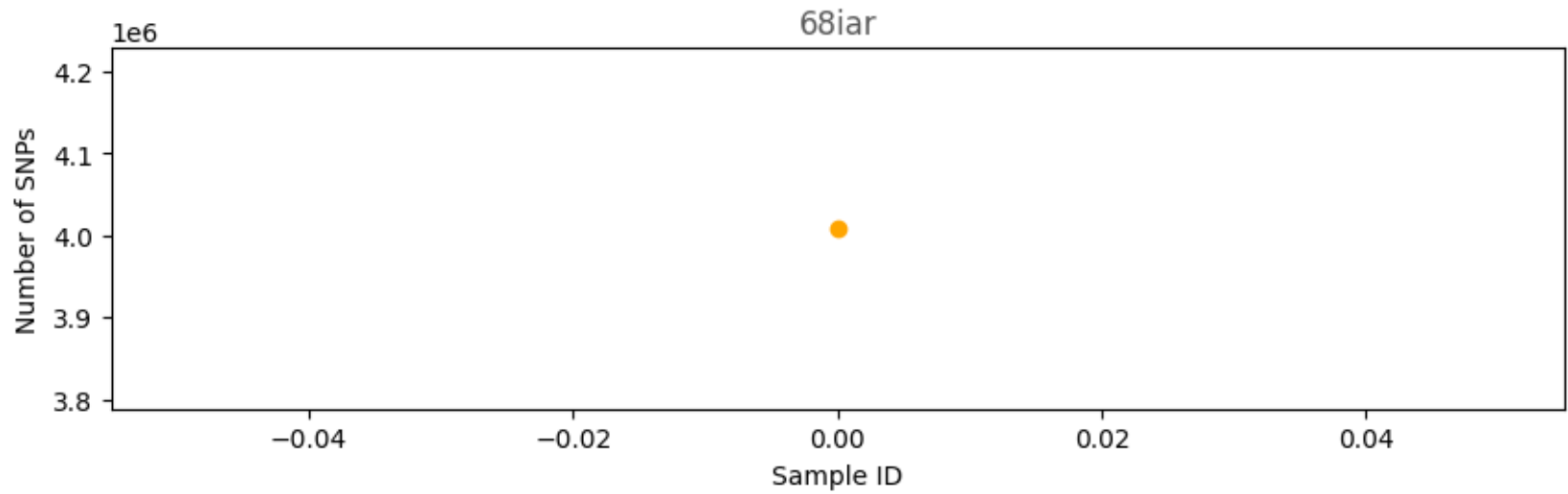# Ts/Tv by sample
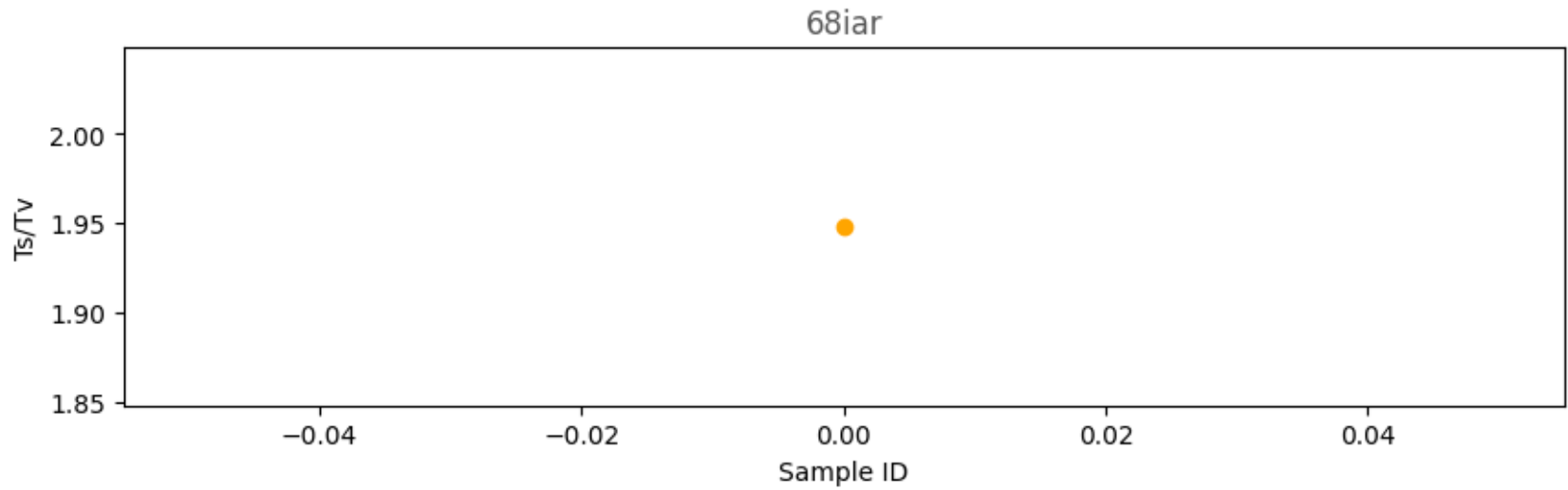
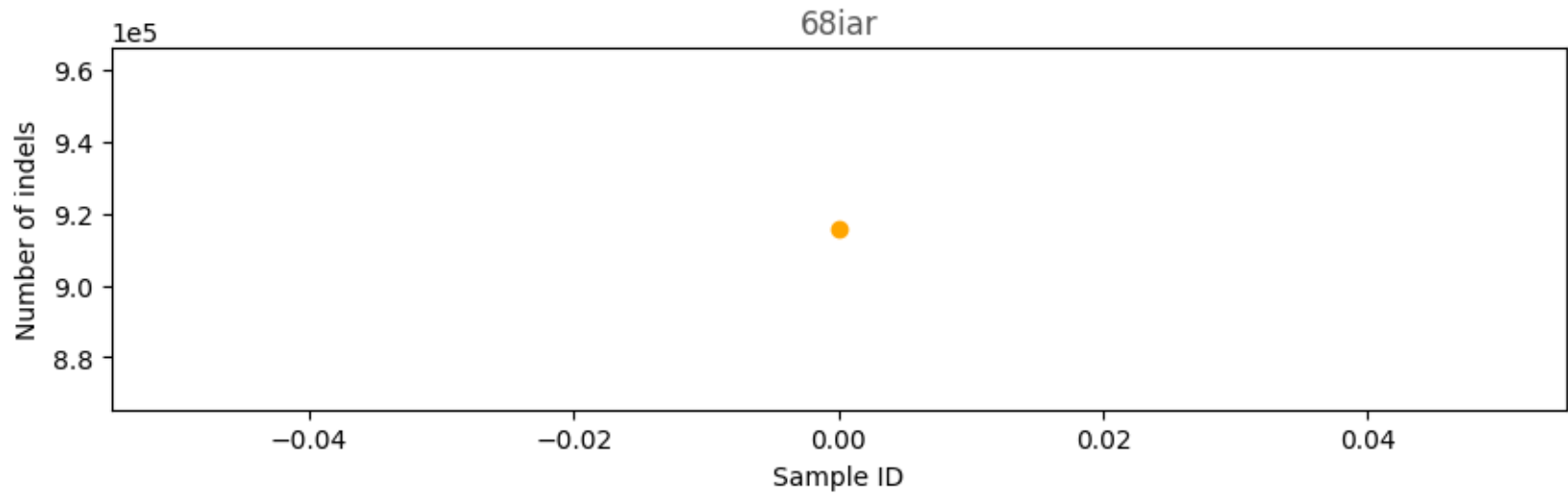# Hets vs non-ref Homs by sample
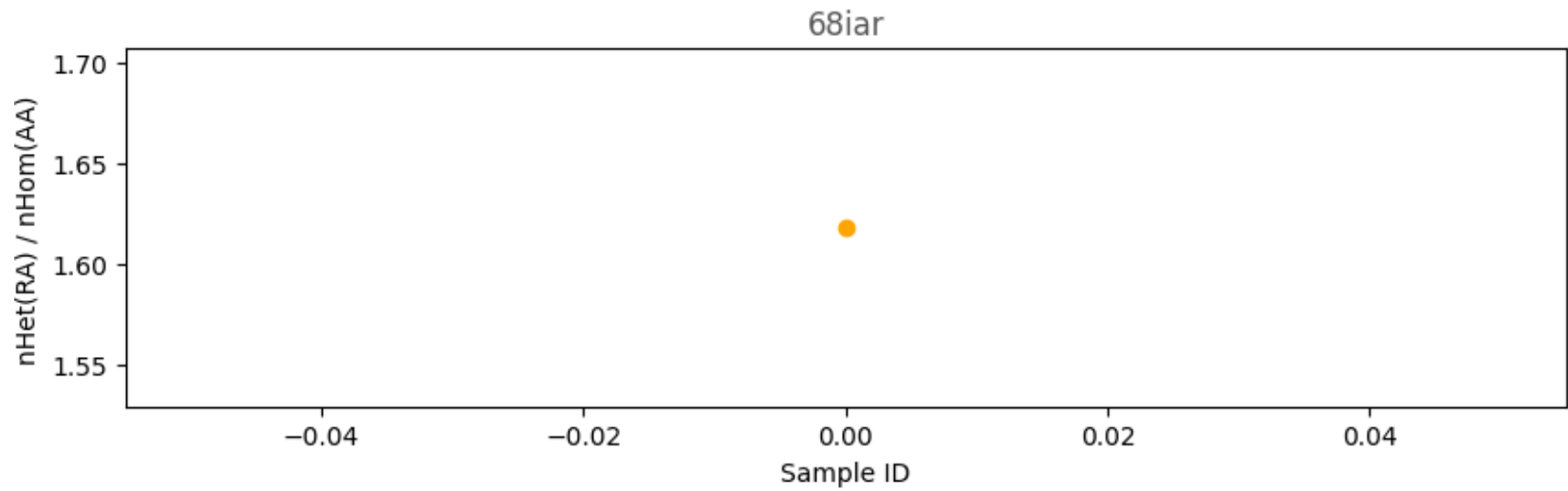
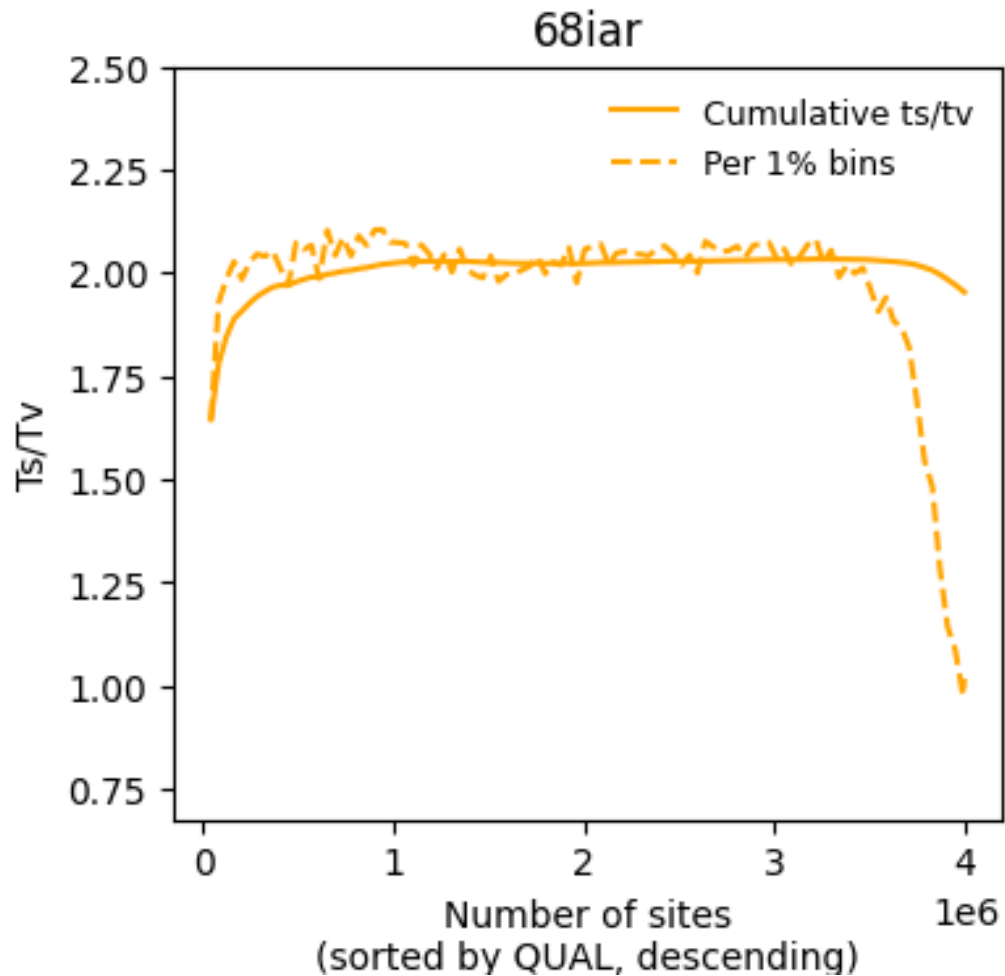# Singletons by sample (hets and homs)
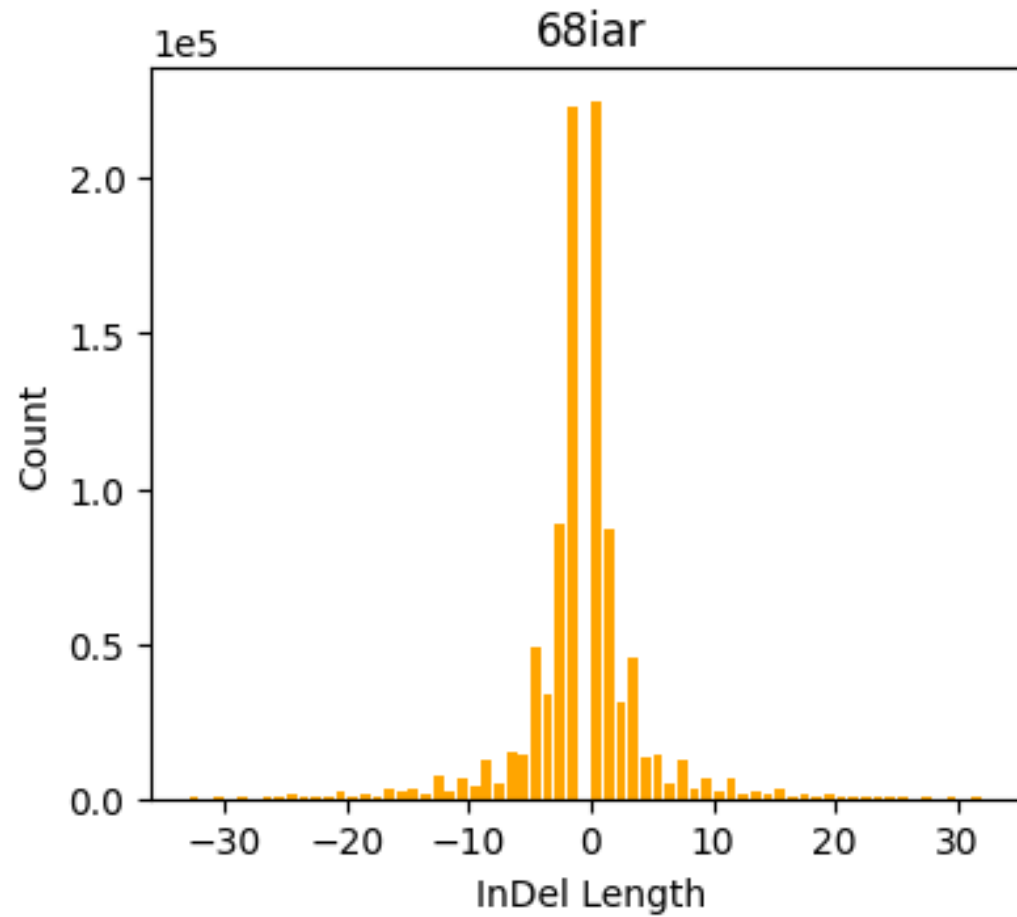
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

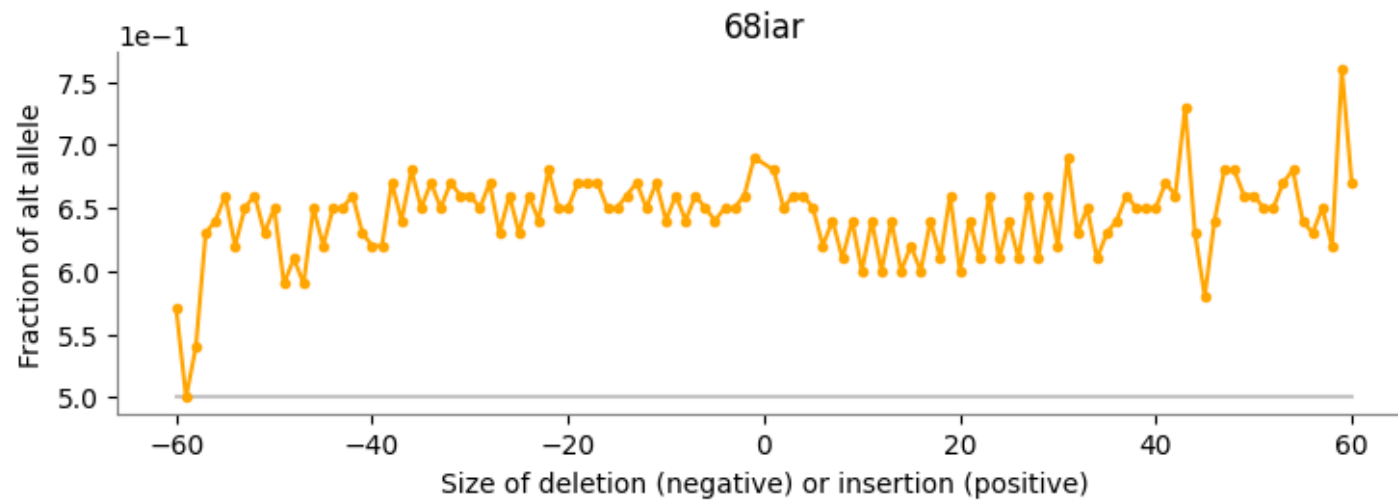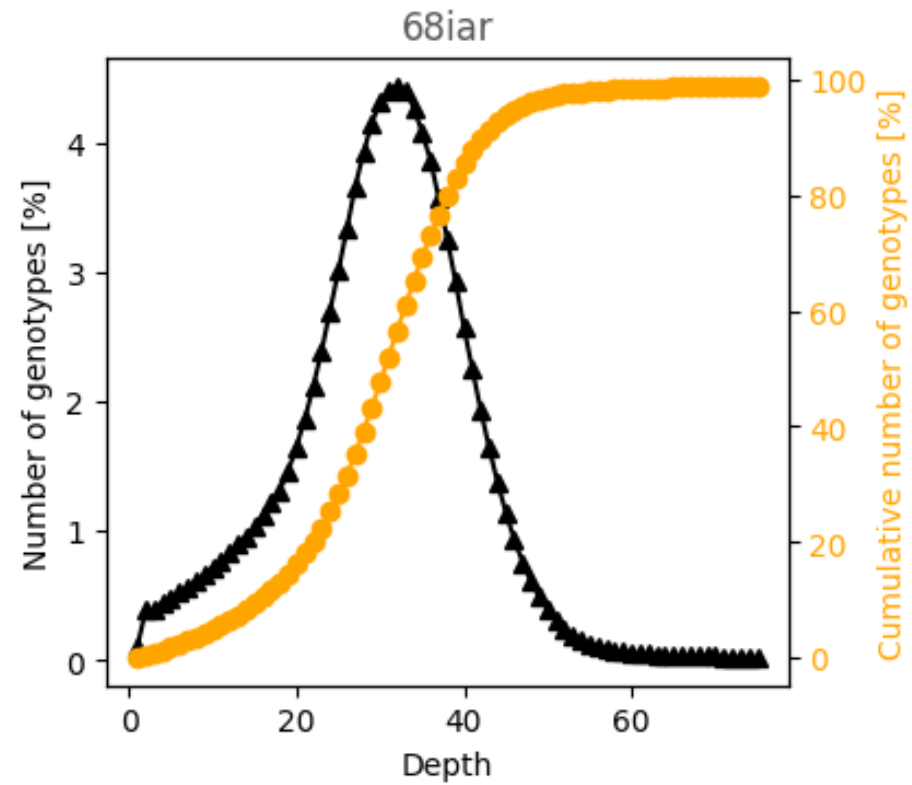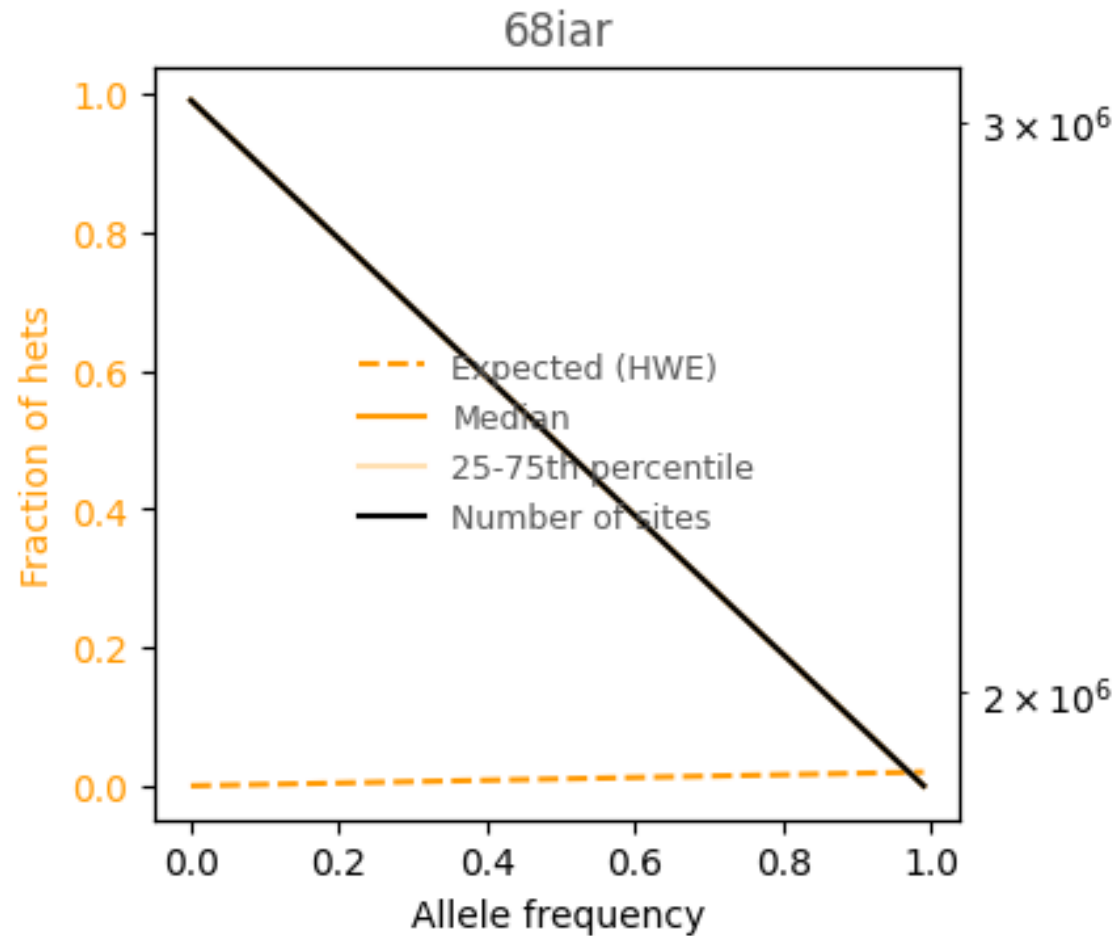# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types