Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
22nit	4,054,766	1.92	1.93	922,039	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
22nit	63.2%	1.89	66.9%	88,178	1,849

- **22nit** .. `/ngc/projects2/gm/data/archive/2022/variants/snv/22nithadf-103880783645-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220615_A00559_BHKC2YDSX3-RHGM_LABKA_WGSA`
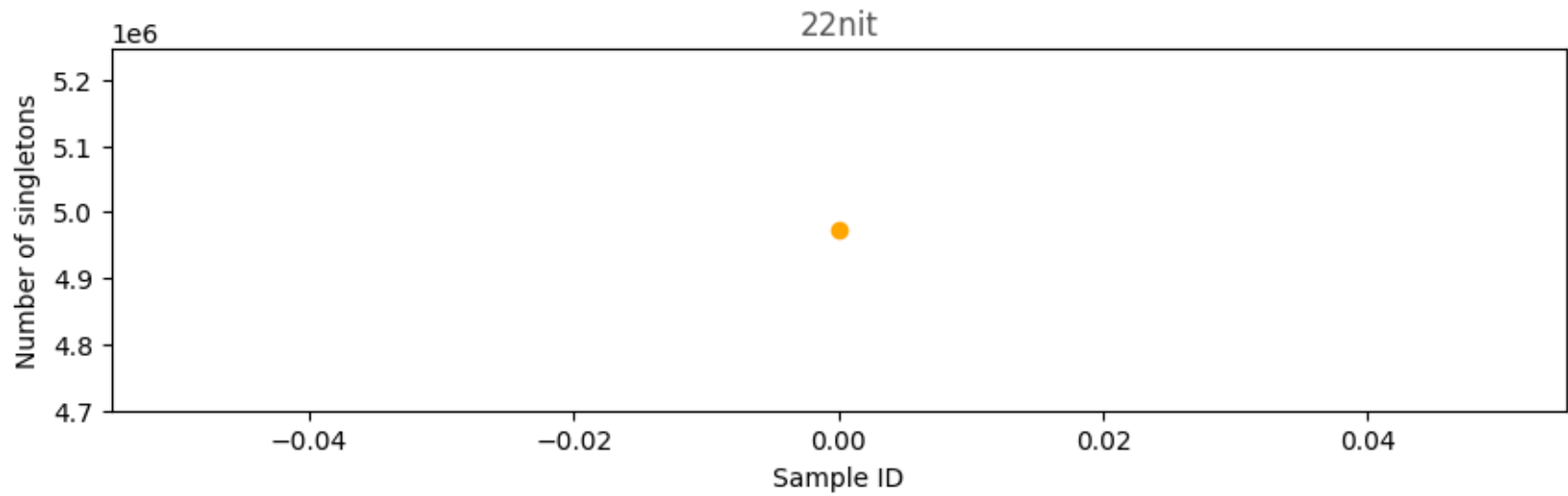 `KUT-WGSAKUT04473_22RKG014060x01_snv_germline_raw.haplotype_caller.vcf.gz`
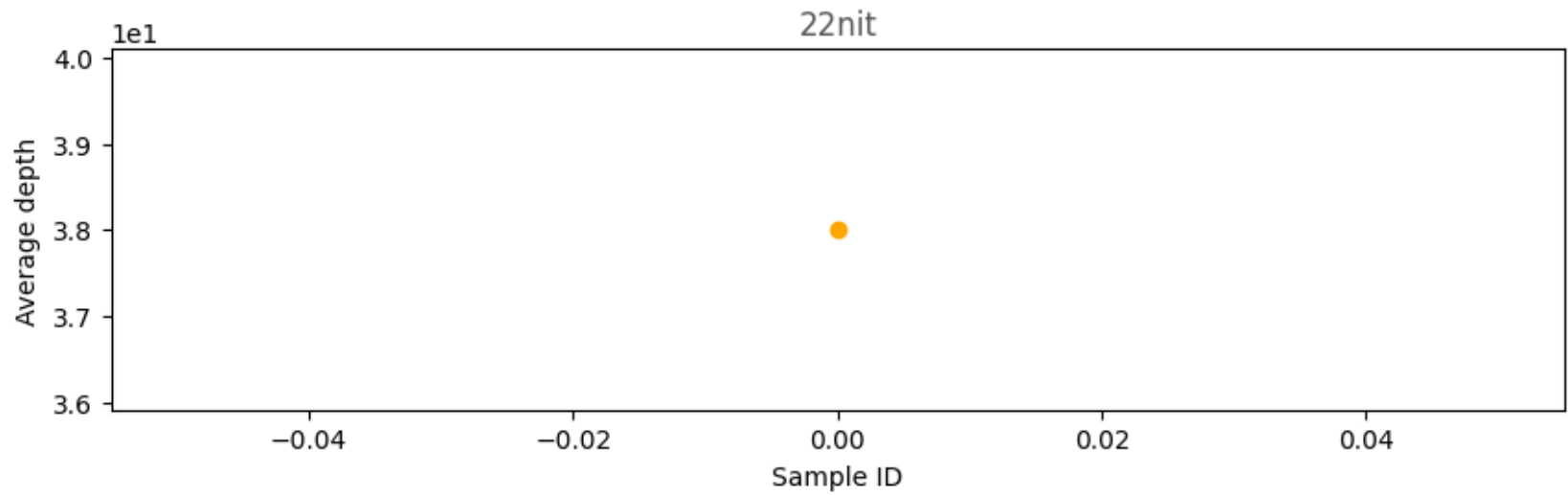
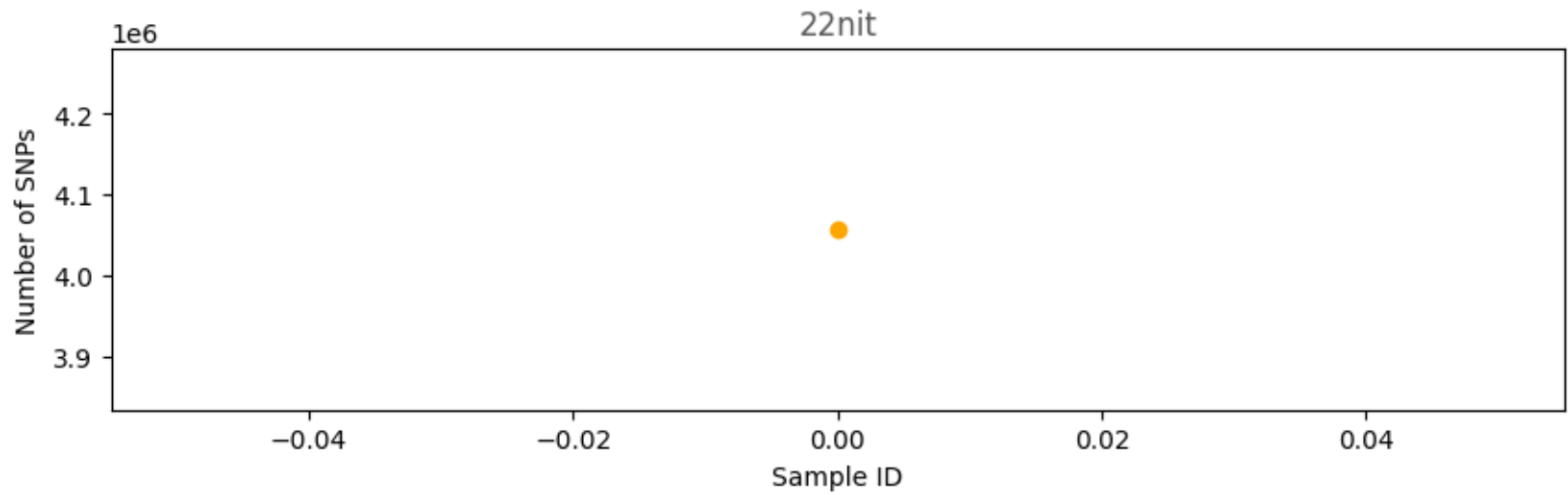# Ts/Tv by sample
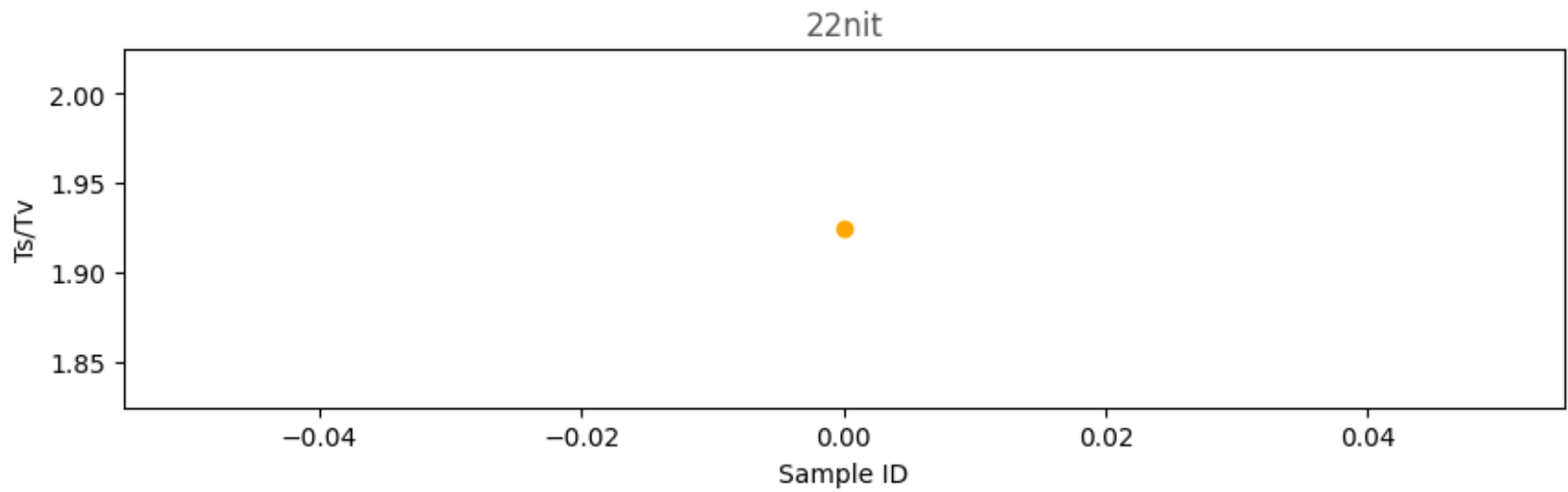
Hets vs non-ref Homs by sample
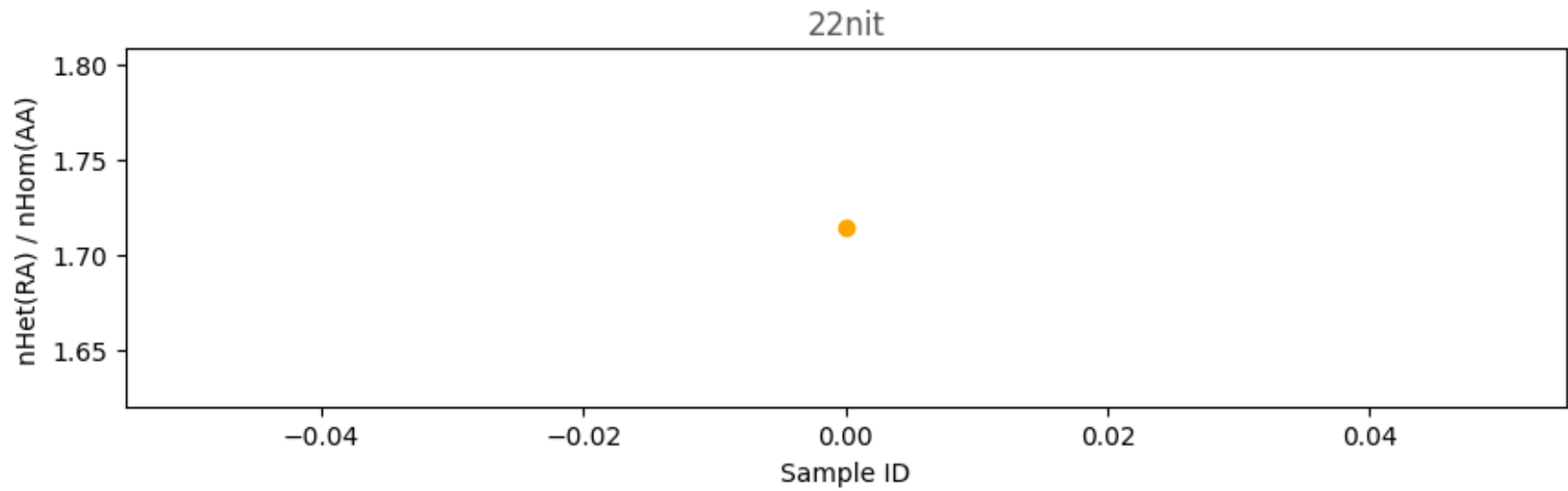
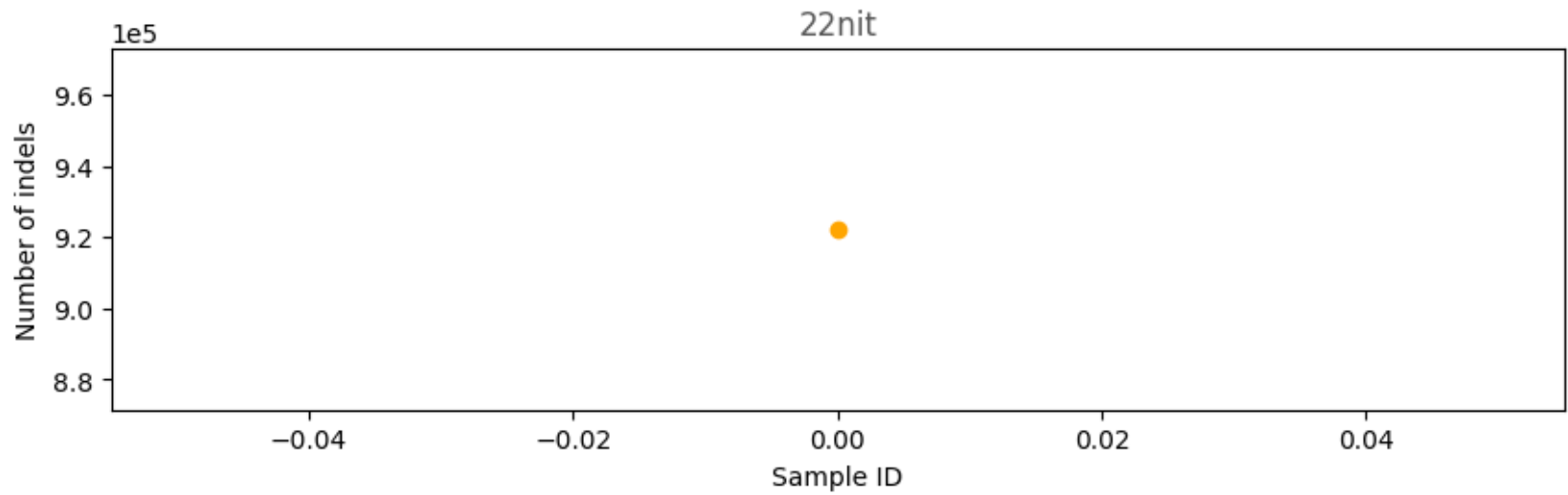# Singletons by sample (hets and homs)
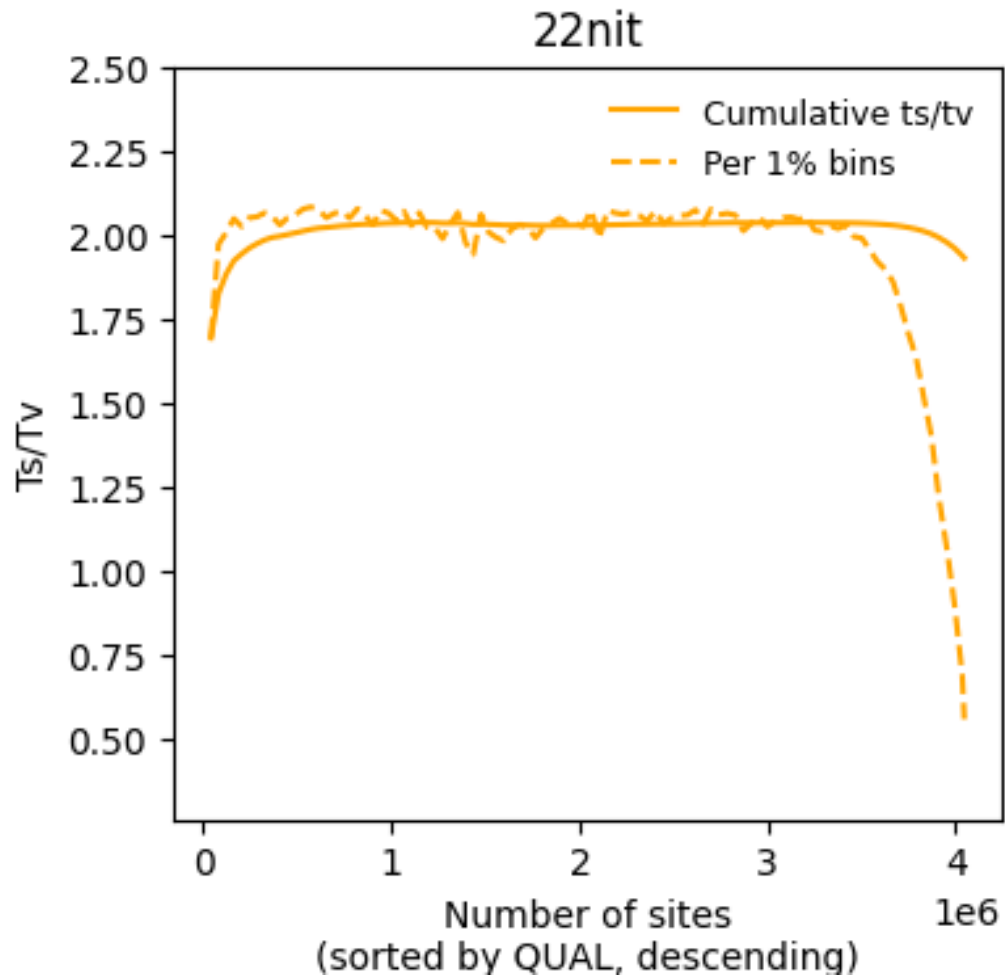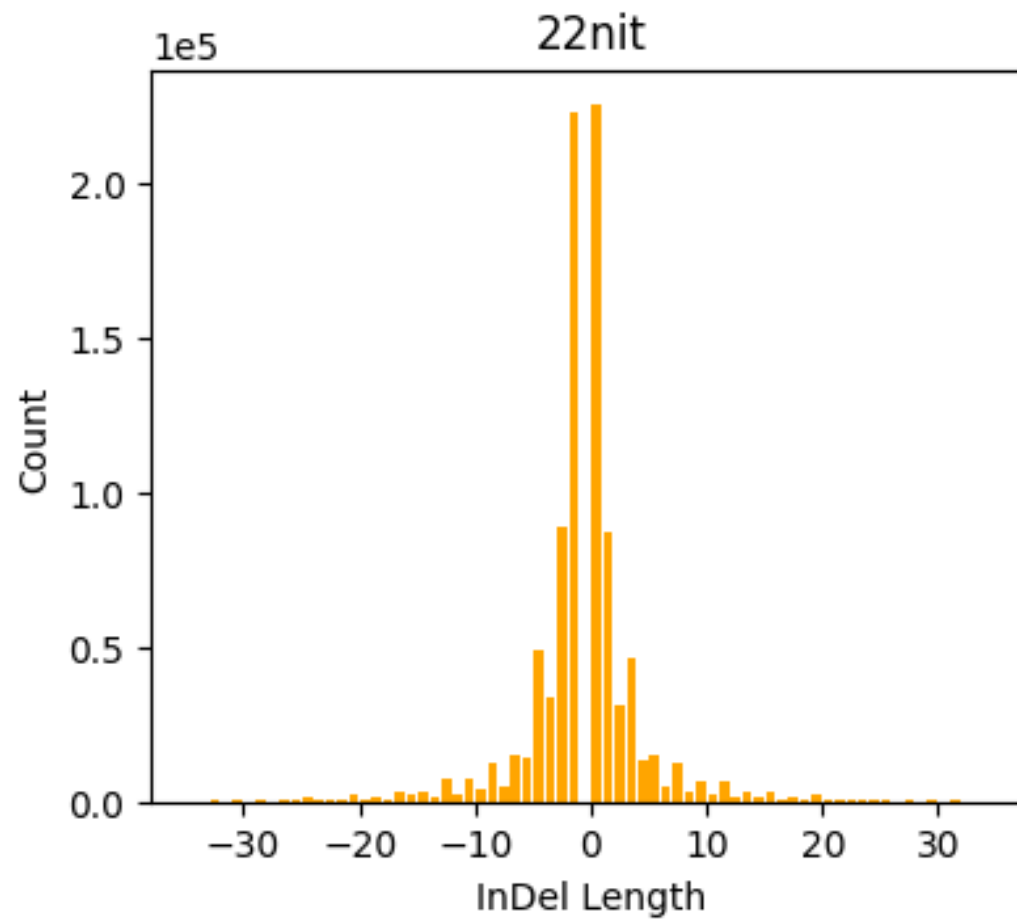
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

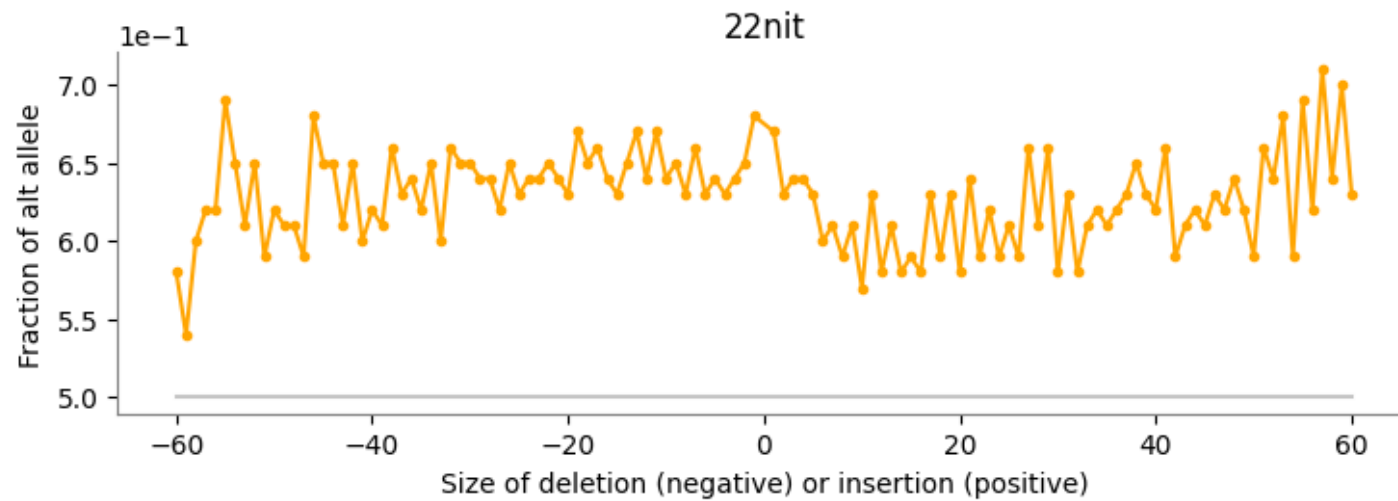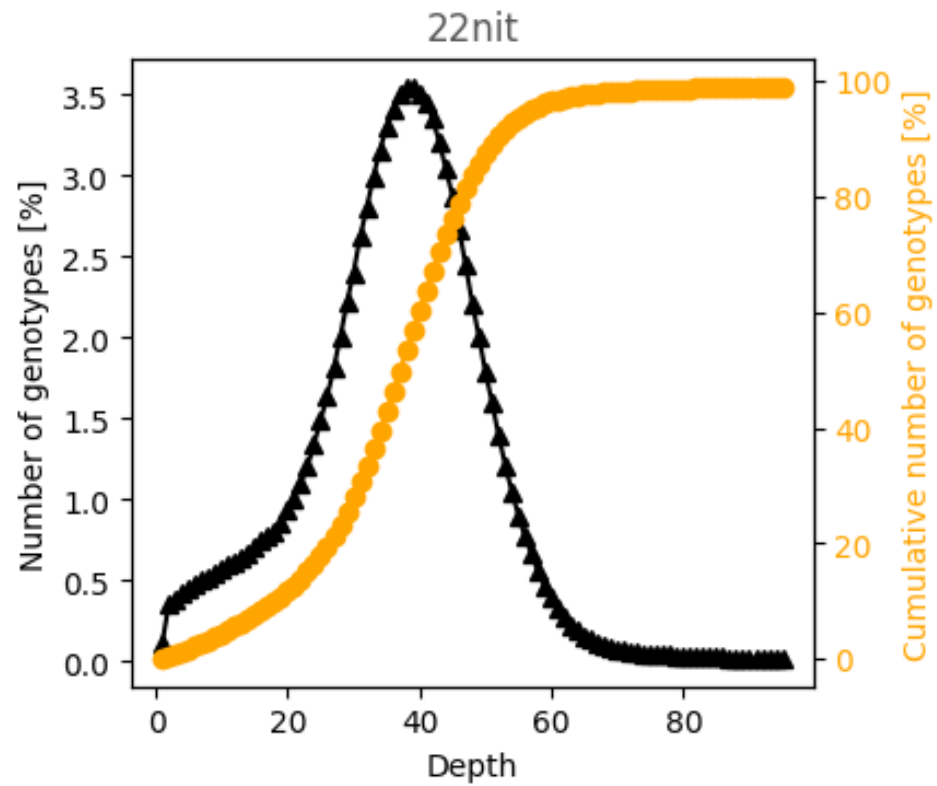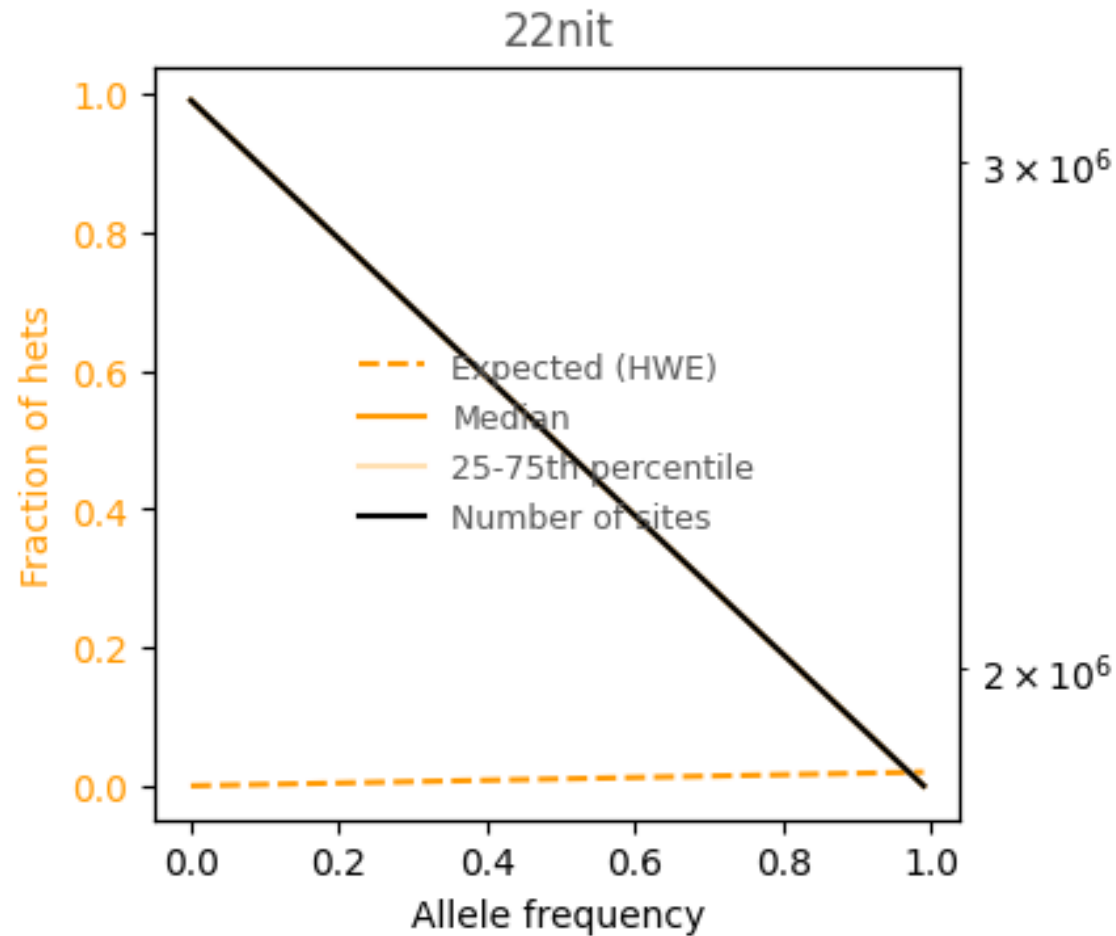# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types