Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
55tit	4,166,628	1.91	1.92	974,476	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
55tit	63.7%	1.88	69.5%	111,079	2,246

- 55tit .. `/ngc/projects2/gm/data/archive/2022/variants/snv/55titoutf-110297742306-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02113-221026_A00559_BH3MJVDSX5-EXT_LAB`
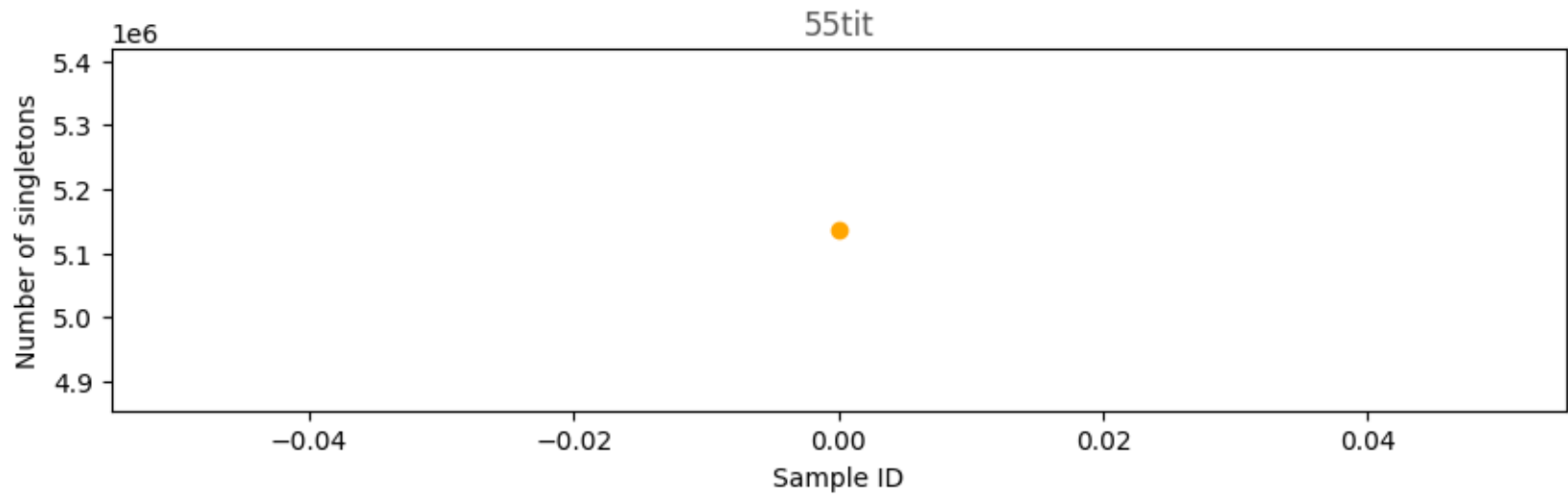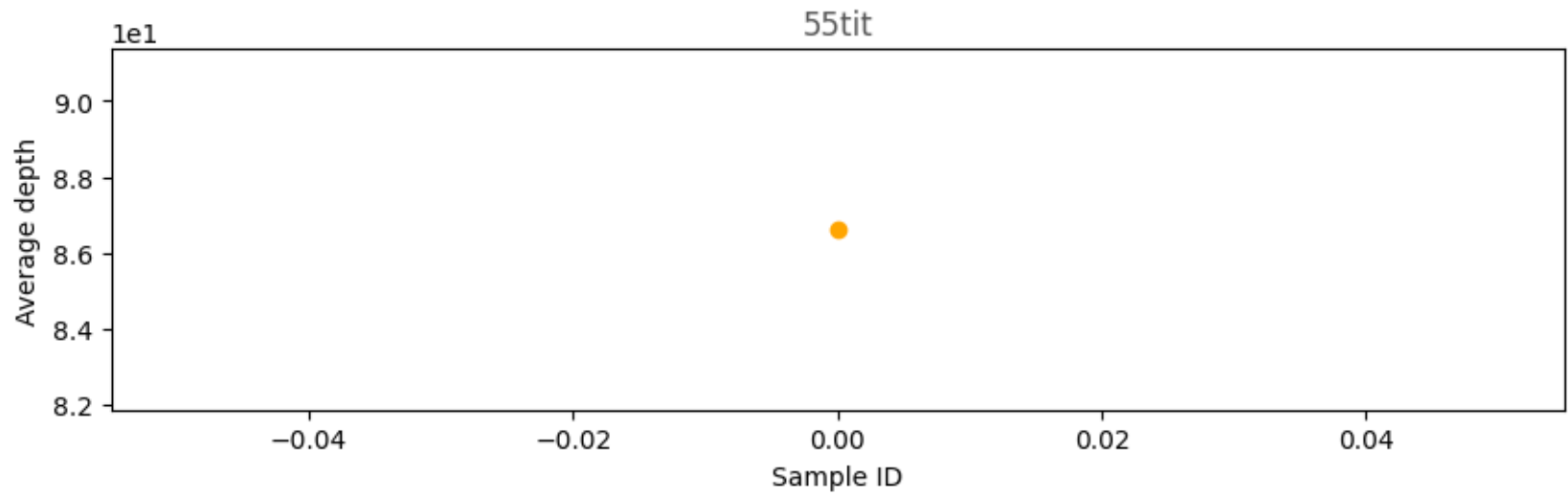 `KA_NGCWGS-NGCWGS05987_snv_germline_raw.haplotype_caller.vcf.gz`

Ts/Tv by sample

55tit
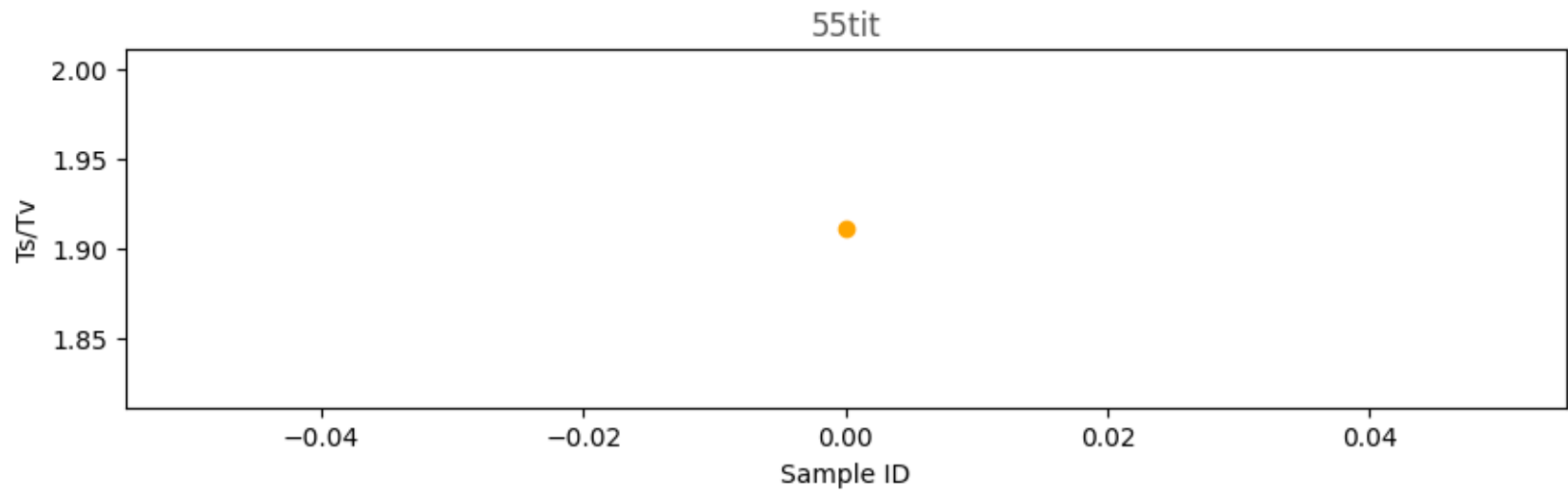
Hets vs non-ref Homs by sample

55tit
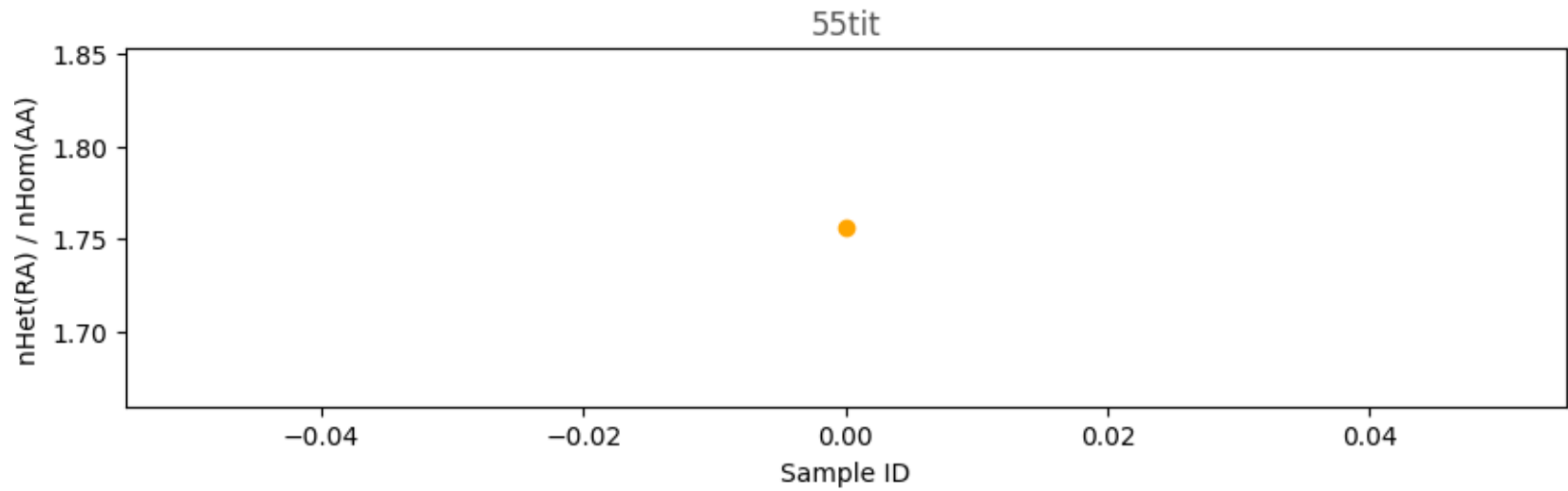
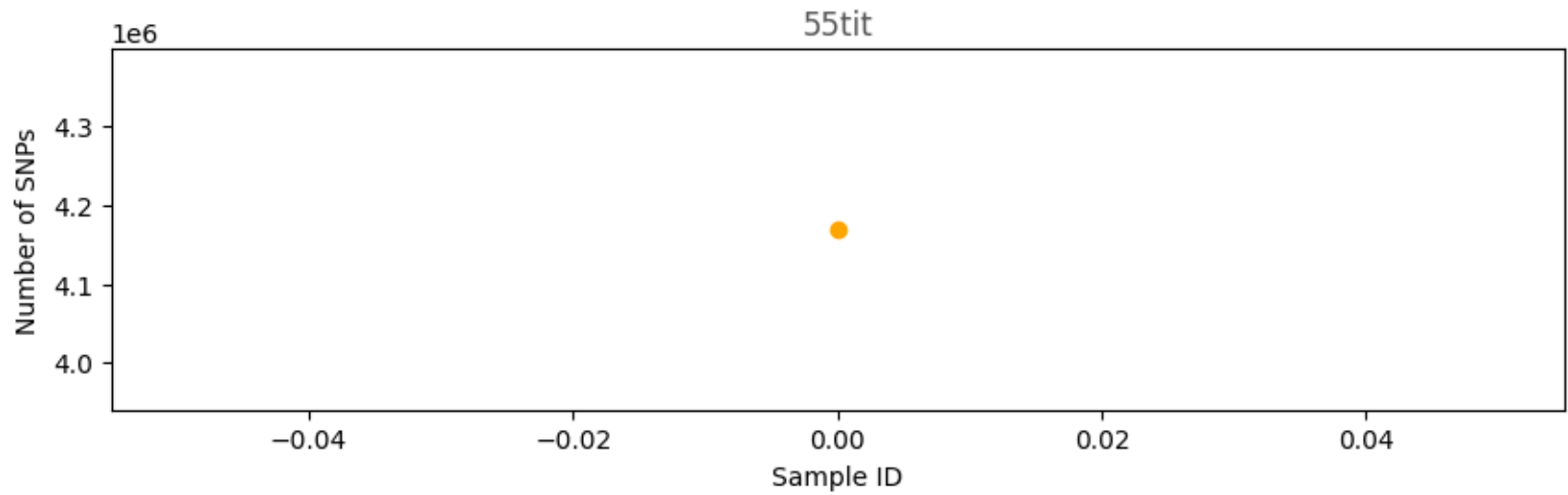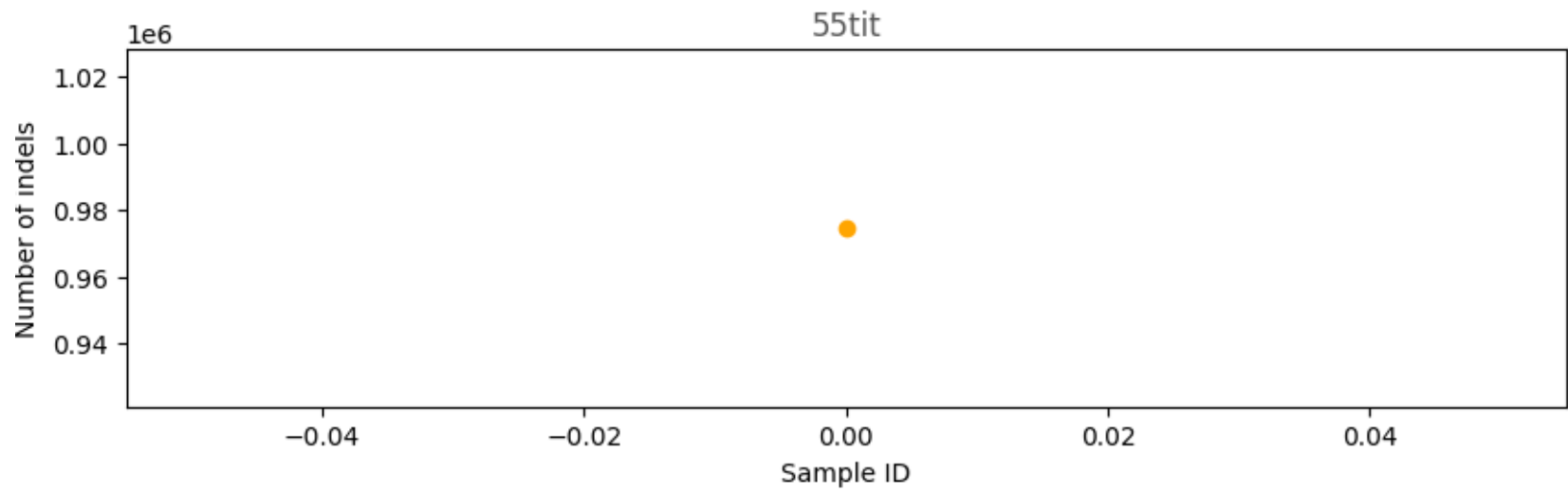# Singletons by sample (hets and homs)
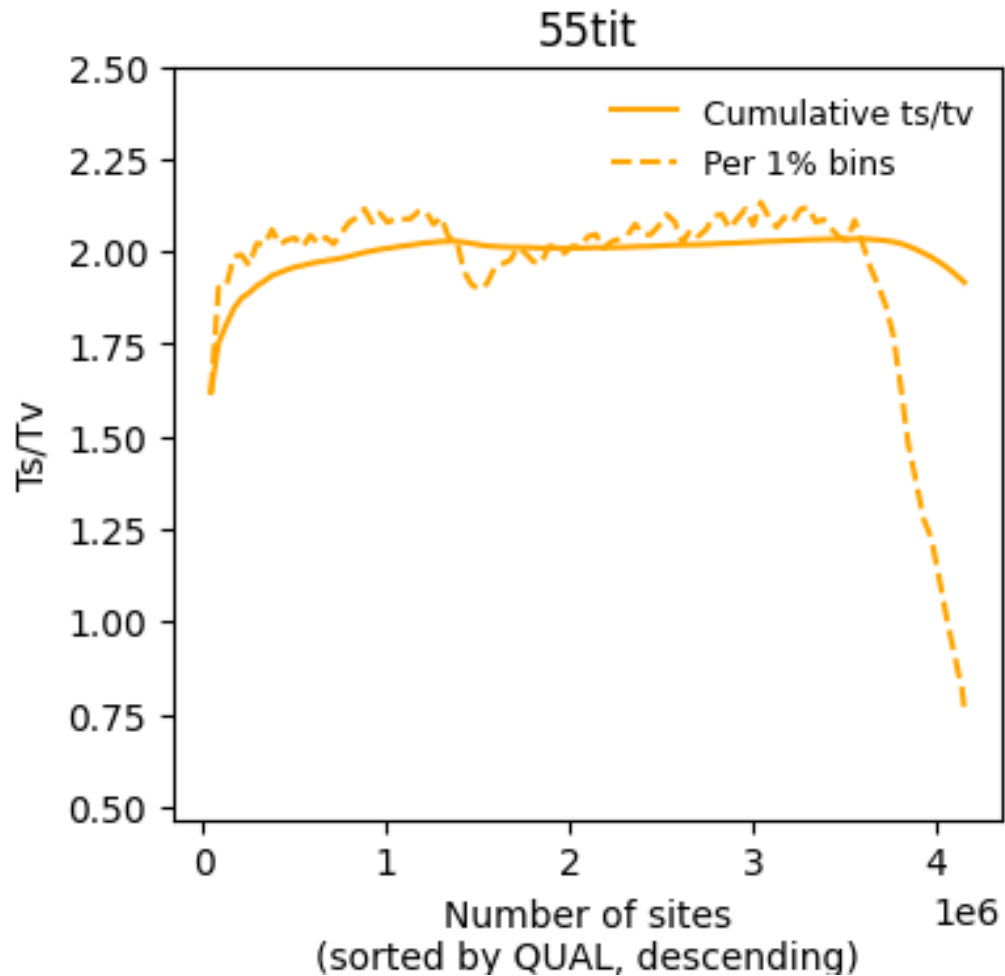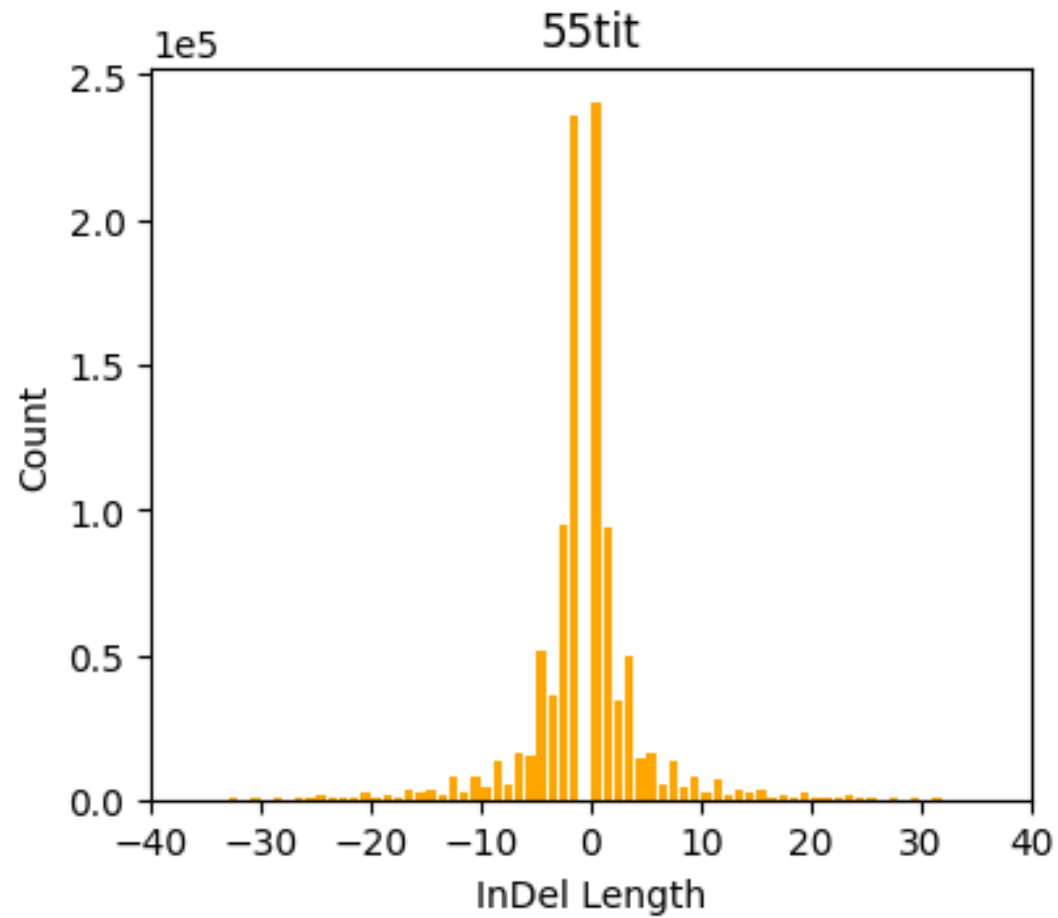
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

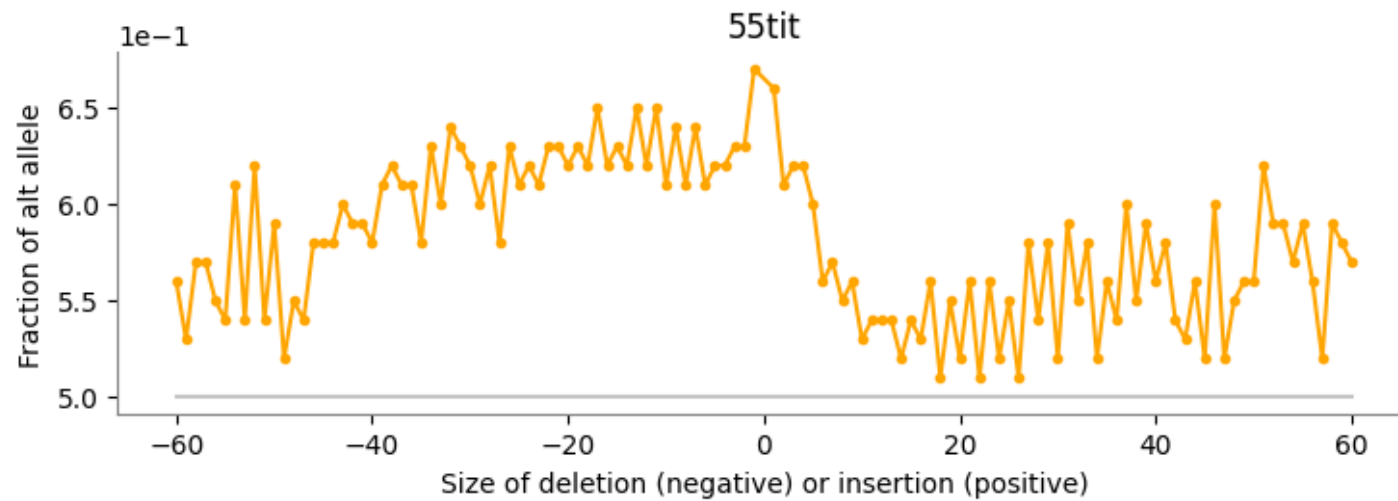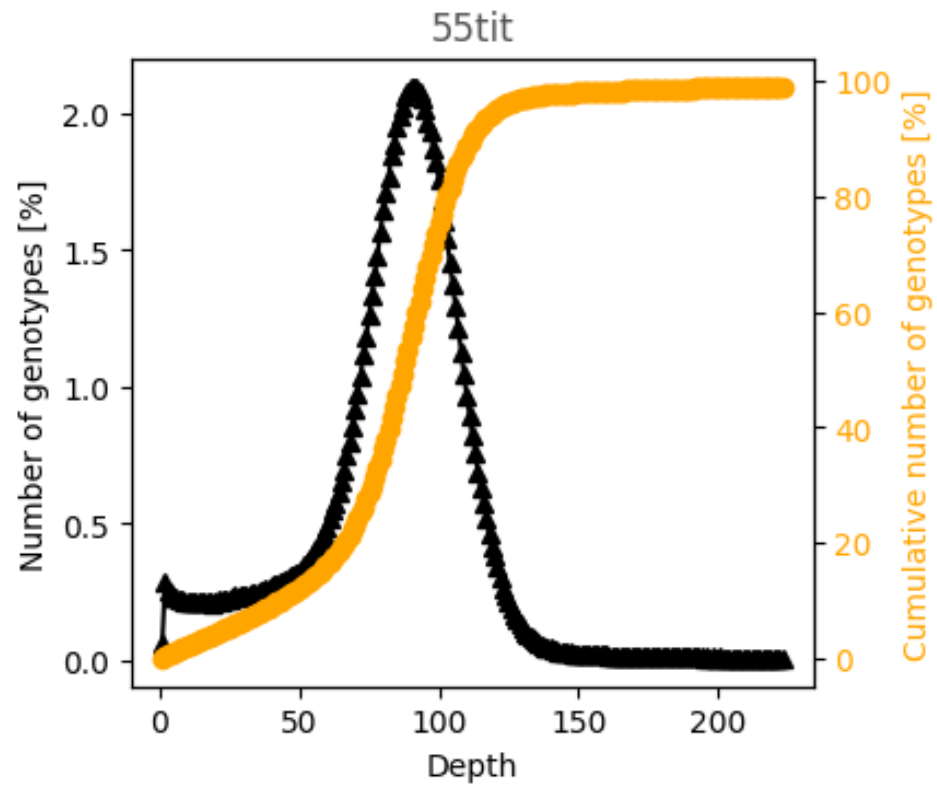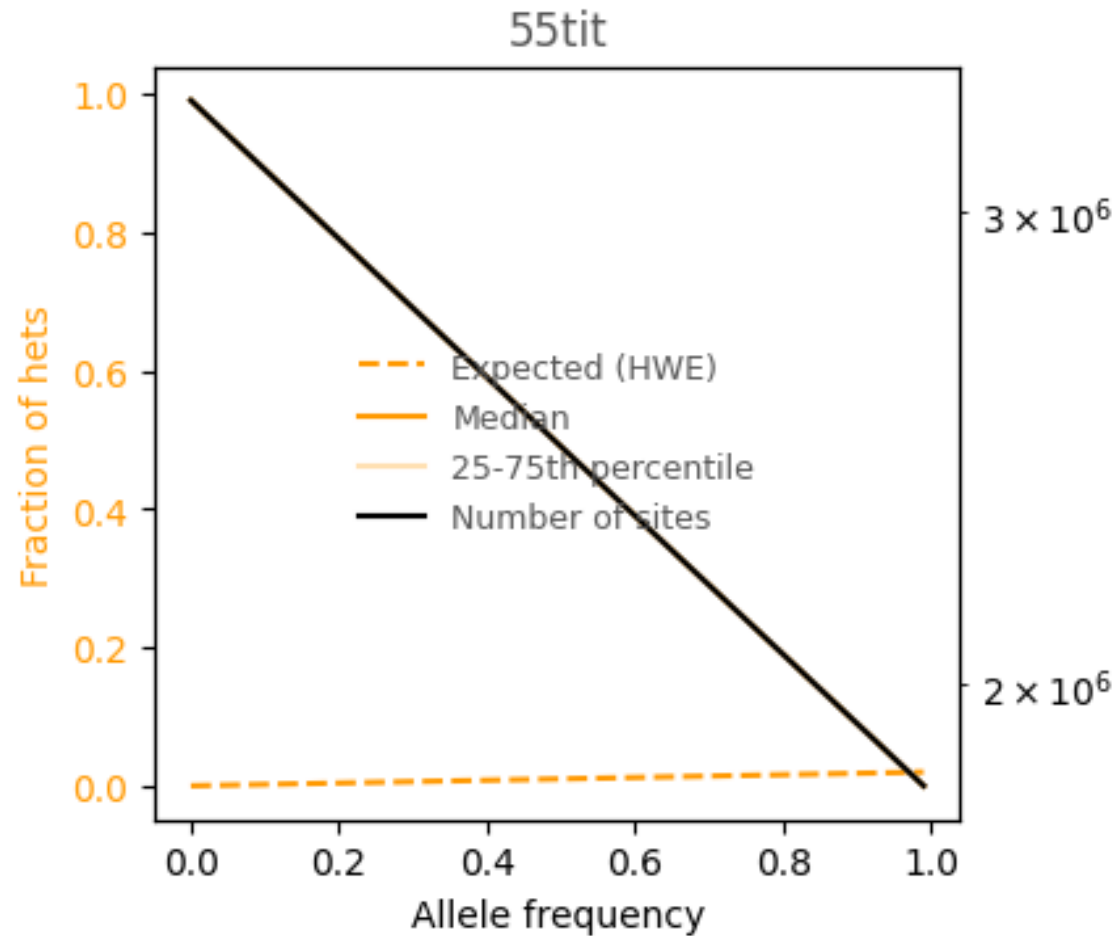# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types