Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
51tid	4,131,442	1.93	1.93	963,648	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
51tid	63.5%	1.90	68.6%	103,582	2,334

- 51tid .. `/ngc/projects2/gm/data/archive/2022/variants/snv/51tidigrf-103766260489-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211201_A01411_AHVN2GDSX2-RHGM_LABKA_WGS-` `WGS03375_snv_germline_raw.haplotype_caller.vcf.gz`
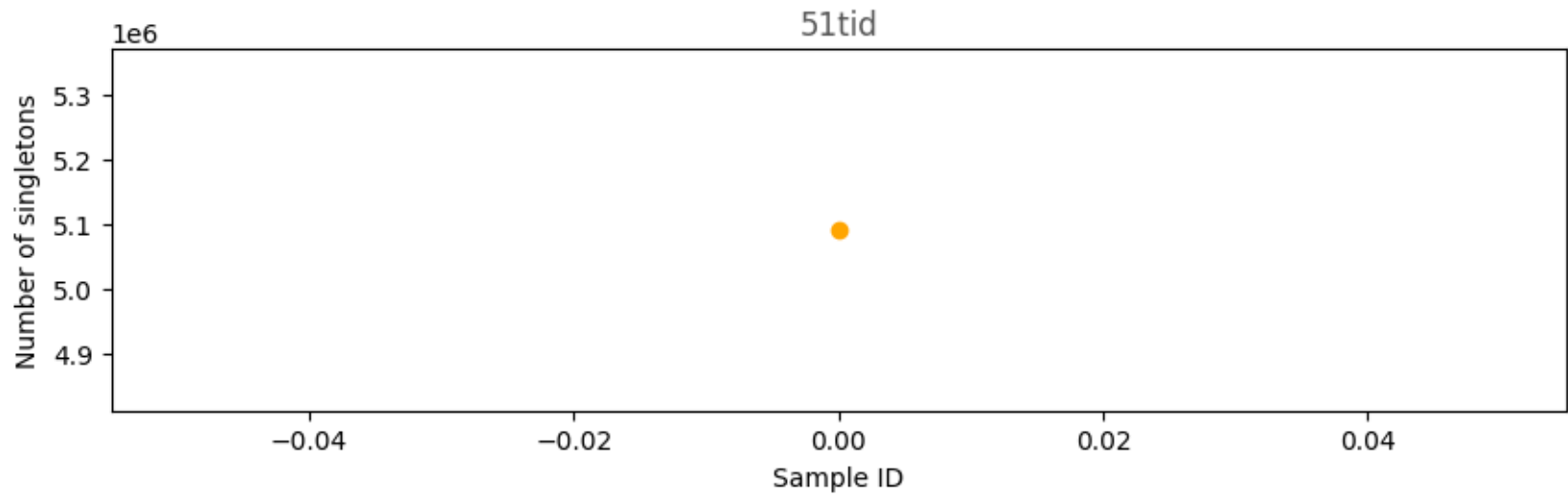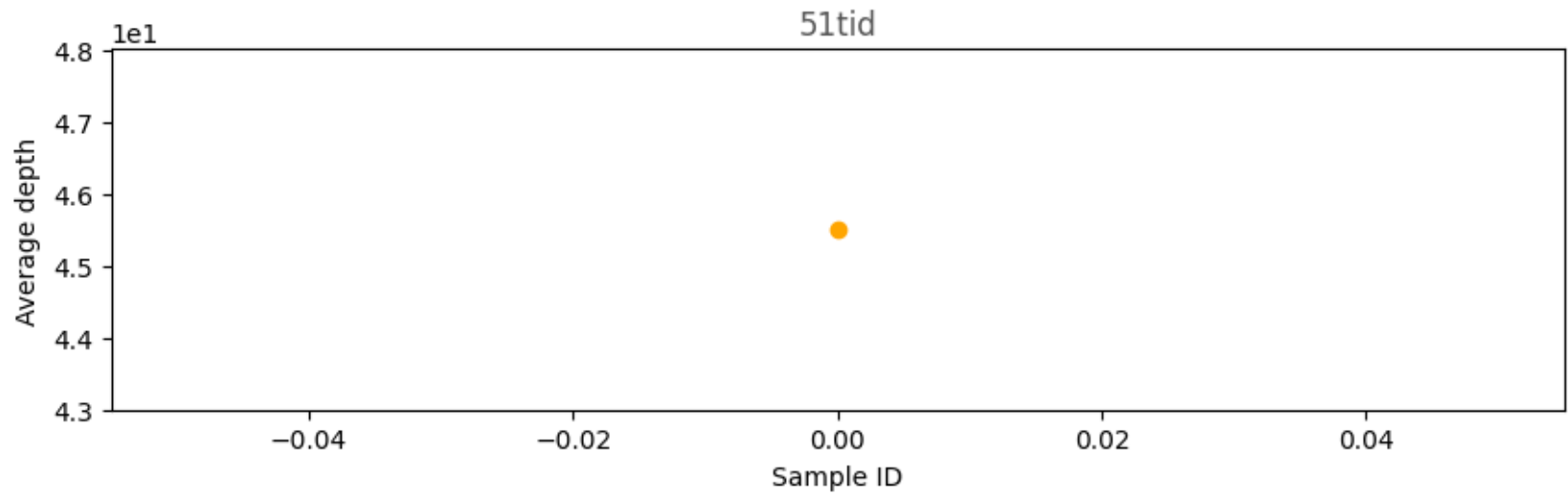
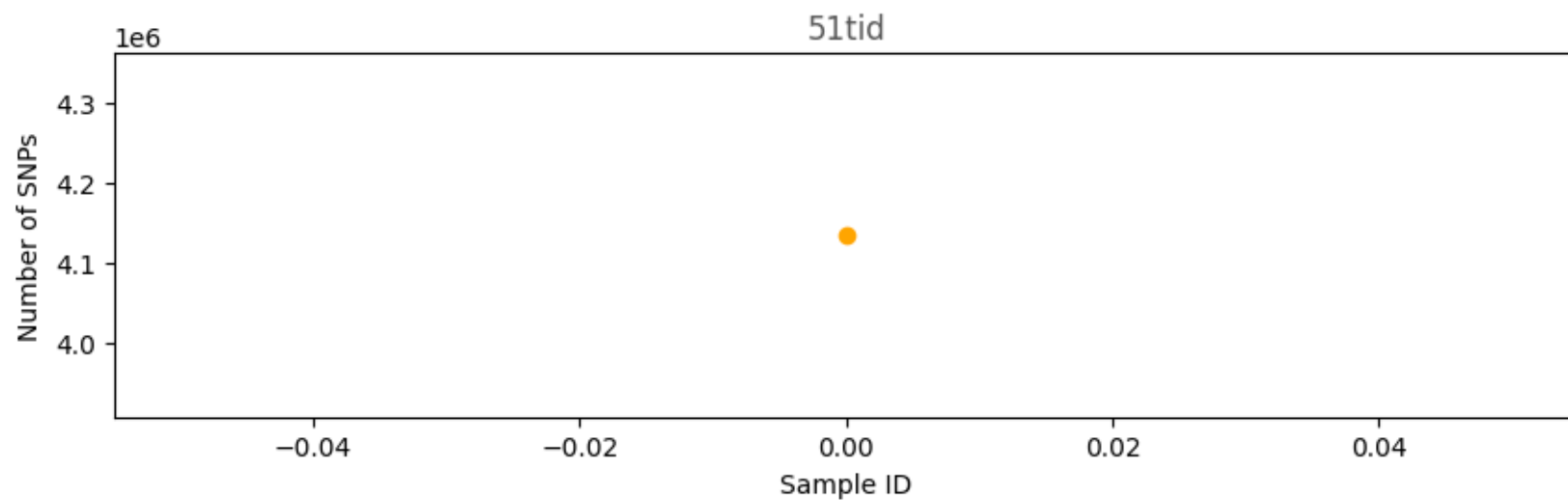# Ts/Tv by sample
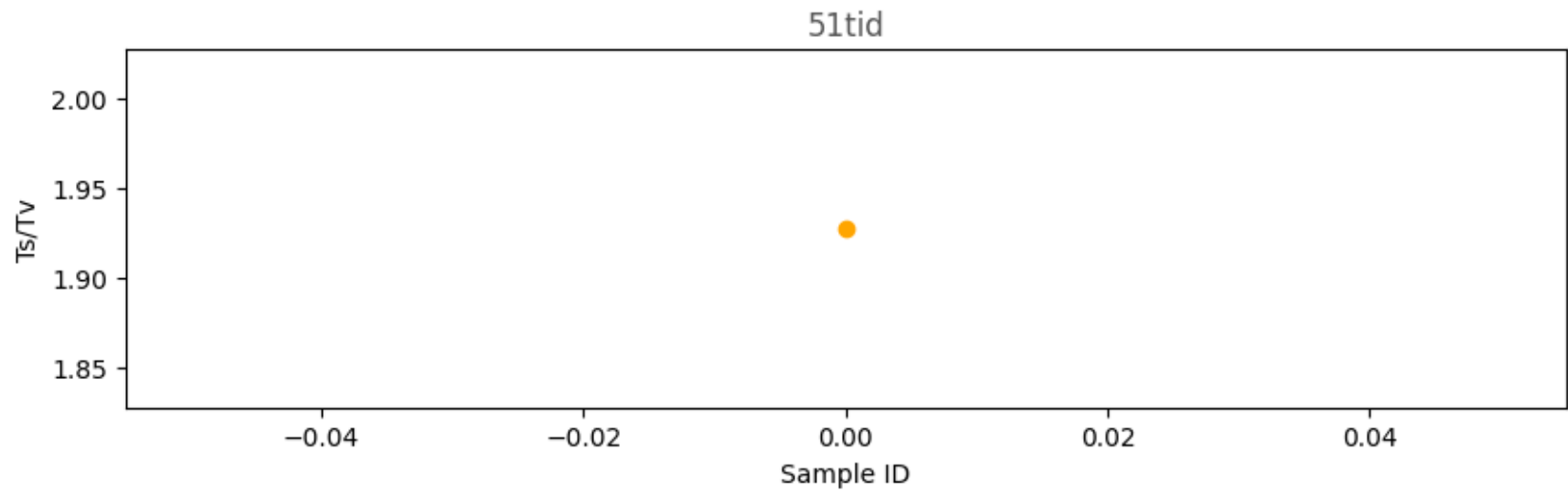
Hets vs non-ref Homs by sample
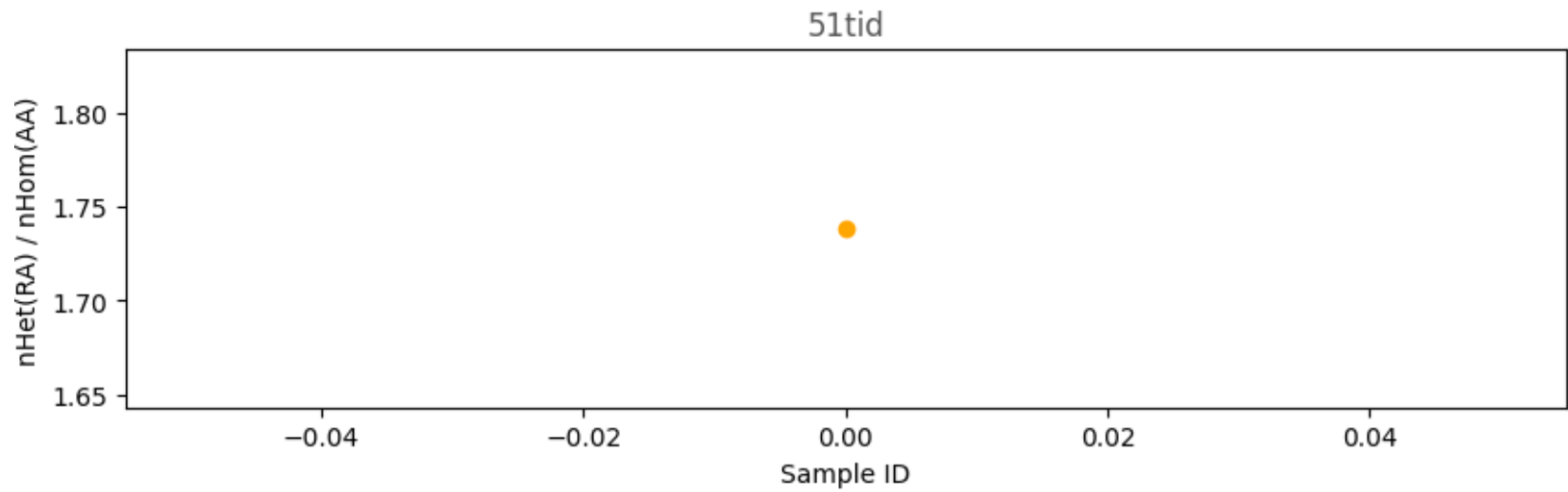
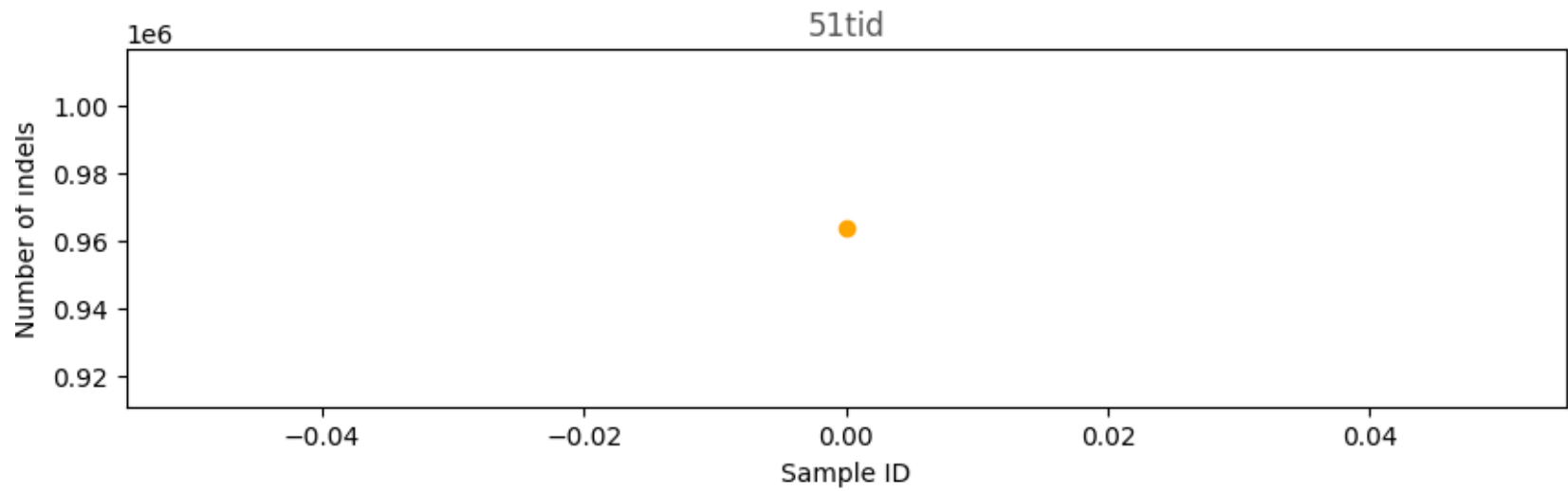# Singletons by sample (hets and homs)
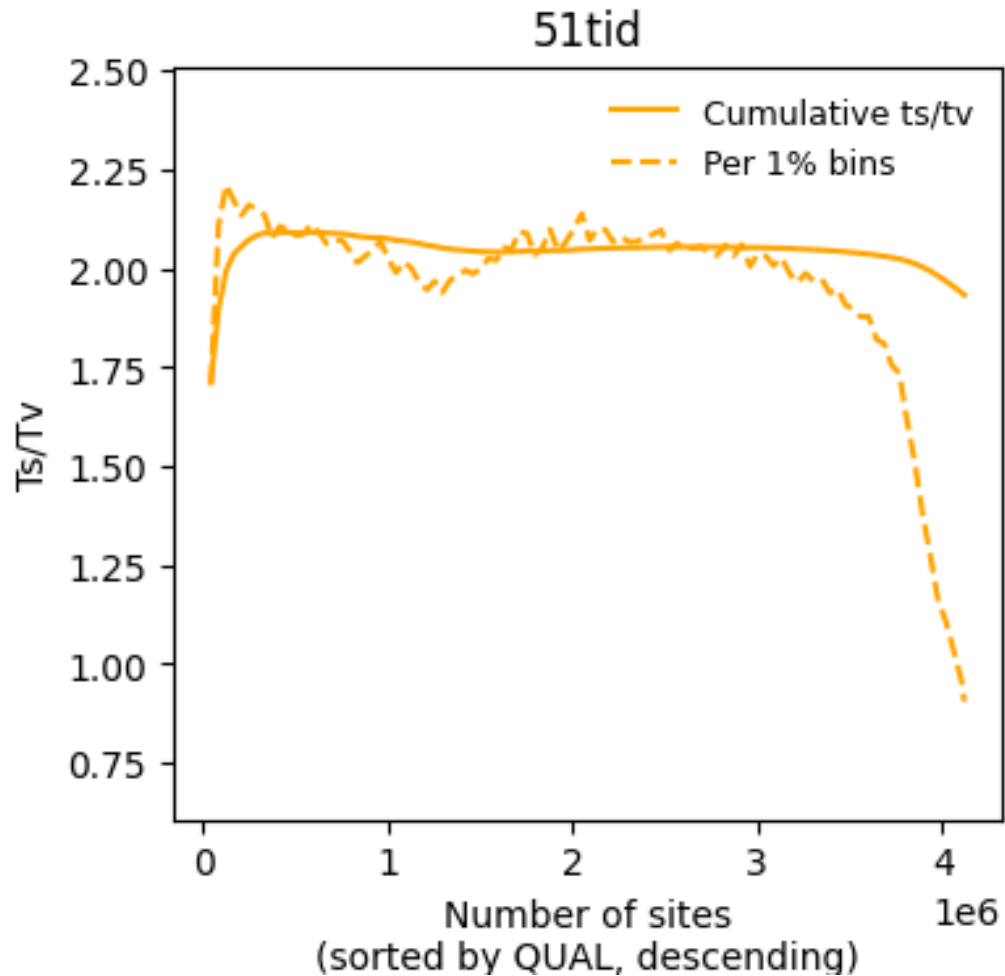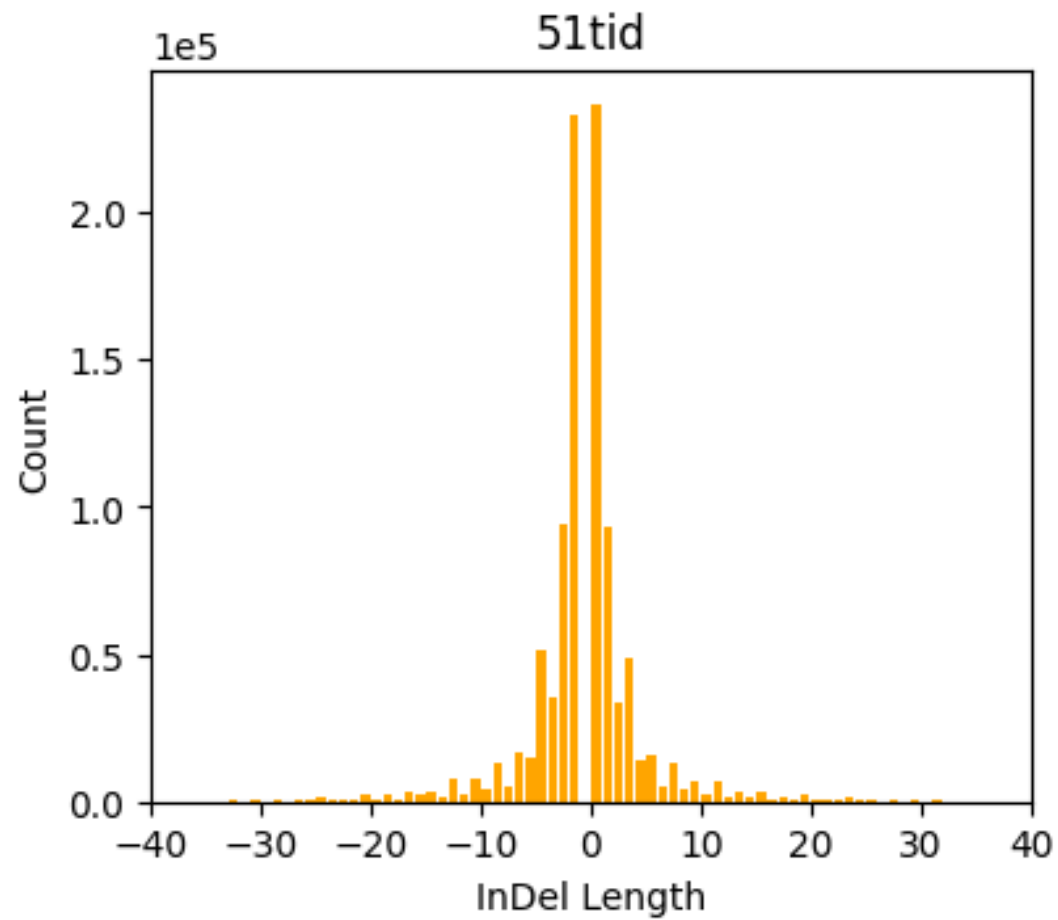
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

51tid

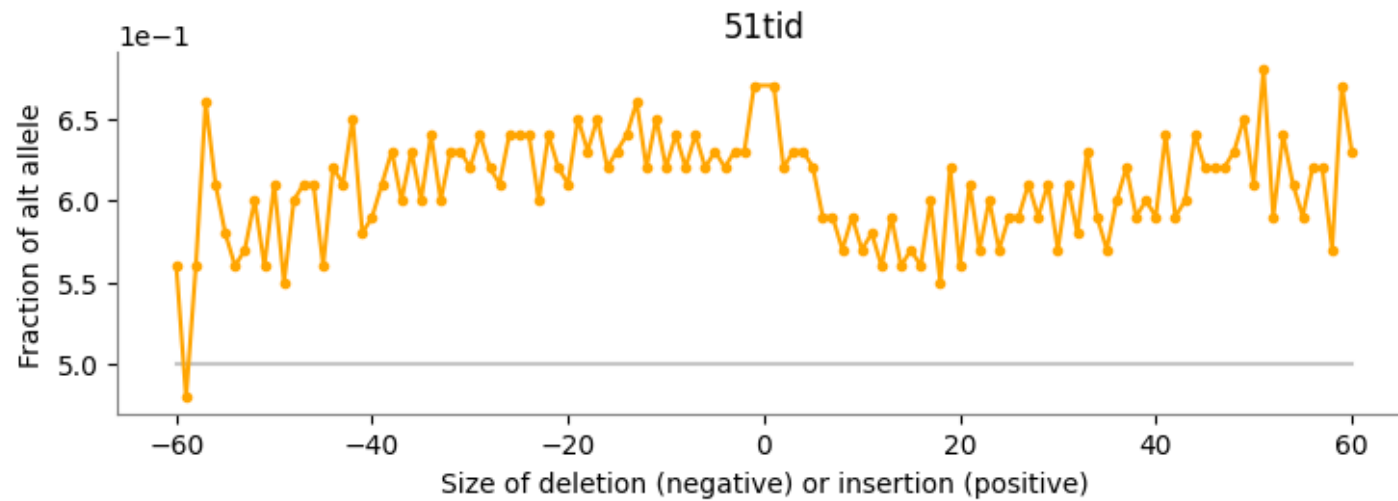# Indel distribution

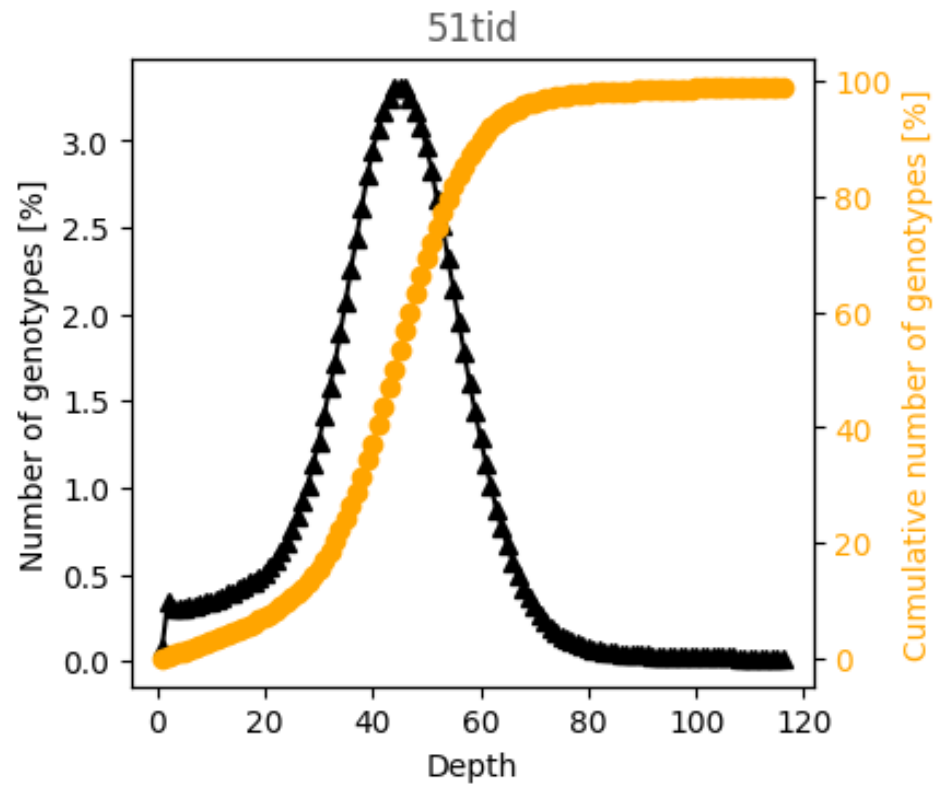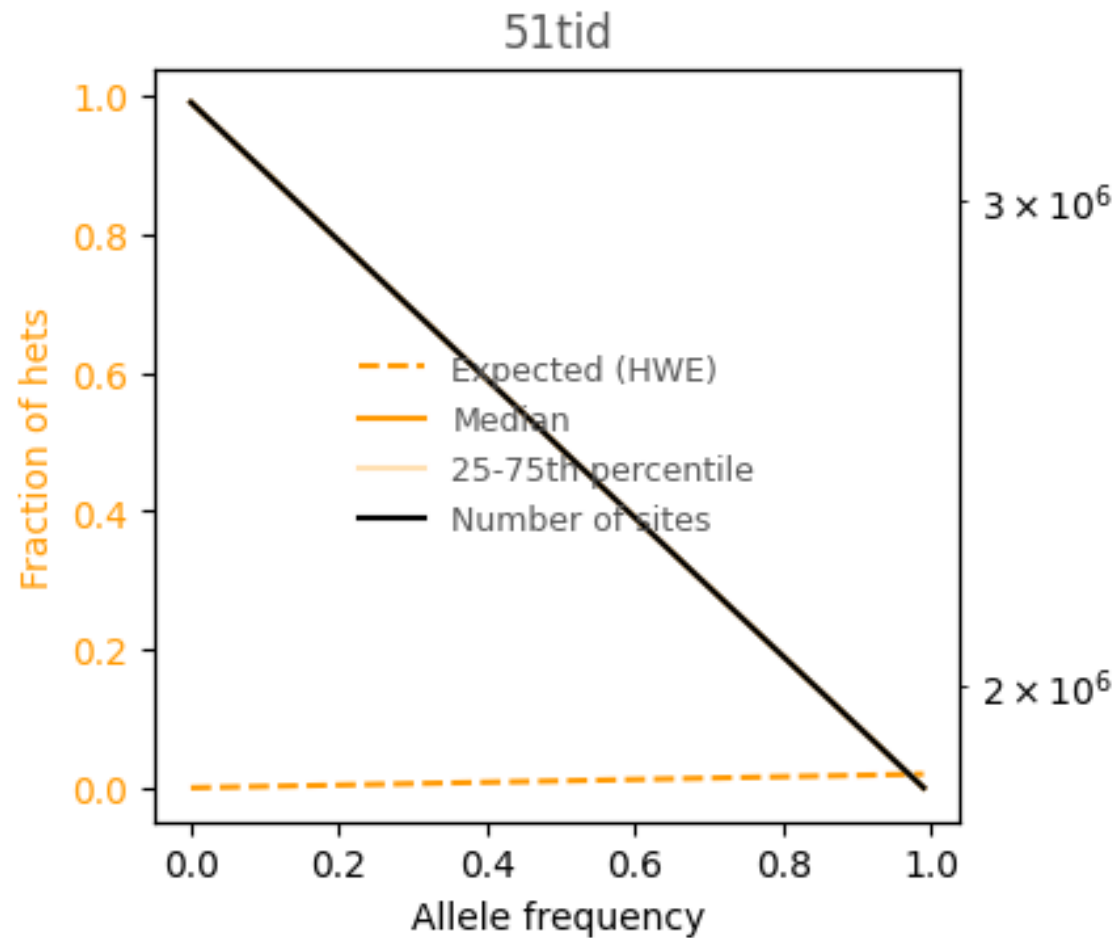# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types