Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76anh	4,060,272	1.93	1.94	945,626	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76anh	62.0%	1.90	66.6%	97,553	2,062

- 76anh .. `/ngc/projects2/gm/data/archive/2022/variants/snv/76anhergm-103903131785-Normal_B`
    `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01496-220914_A01176_BHTNVLDSX3-EXT_LAB`
    `KA_NGCWGS-NGCWGS05270_22RKG020012_snv_germline_raw.haplotype_caller.vcf.gz`
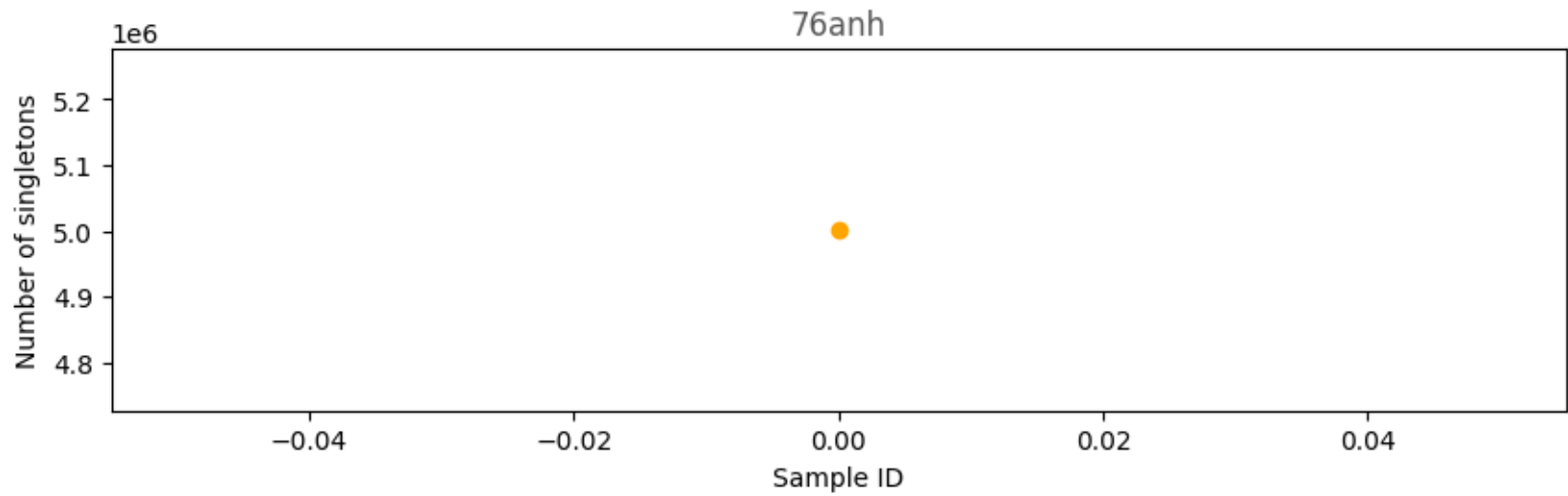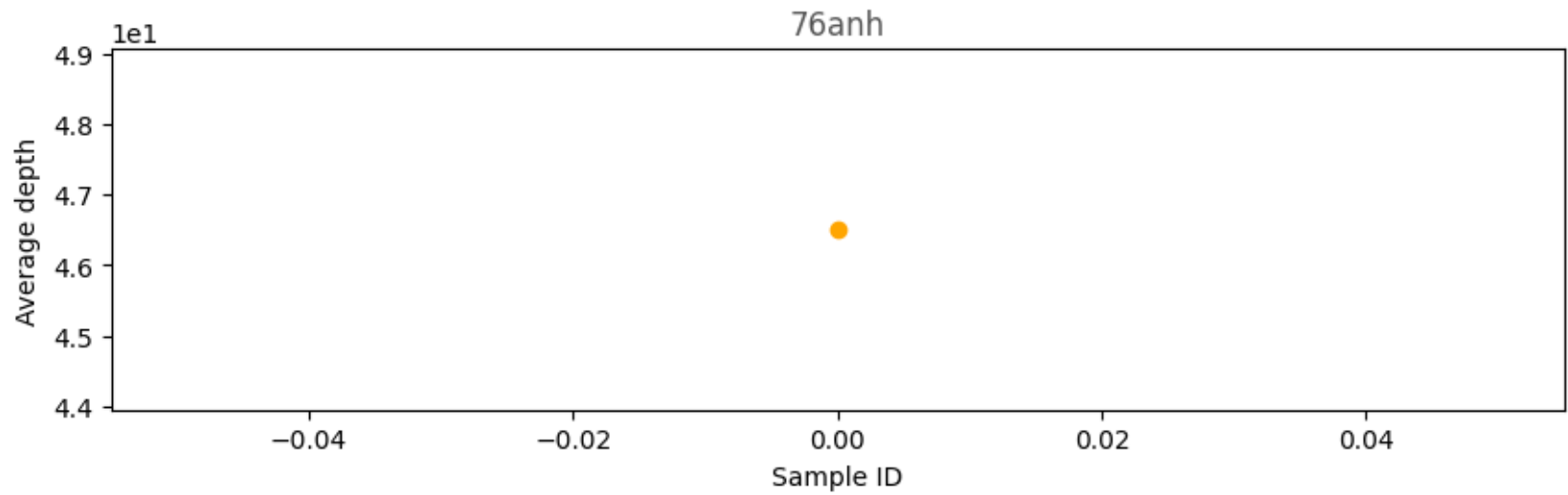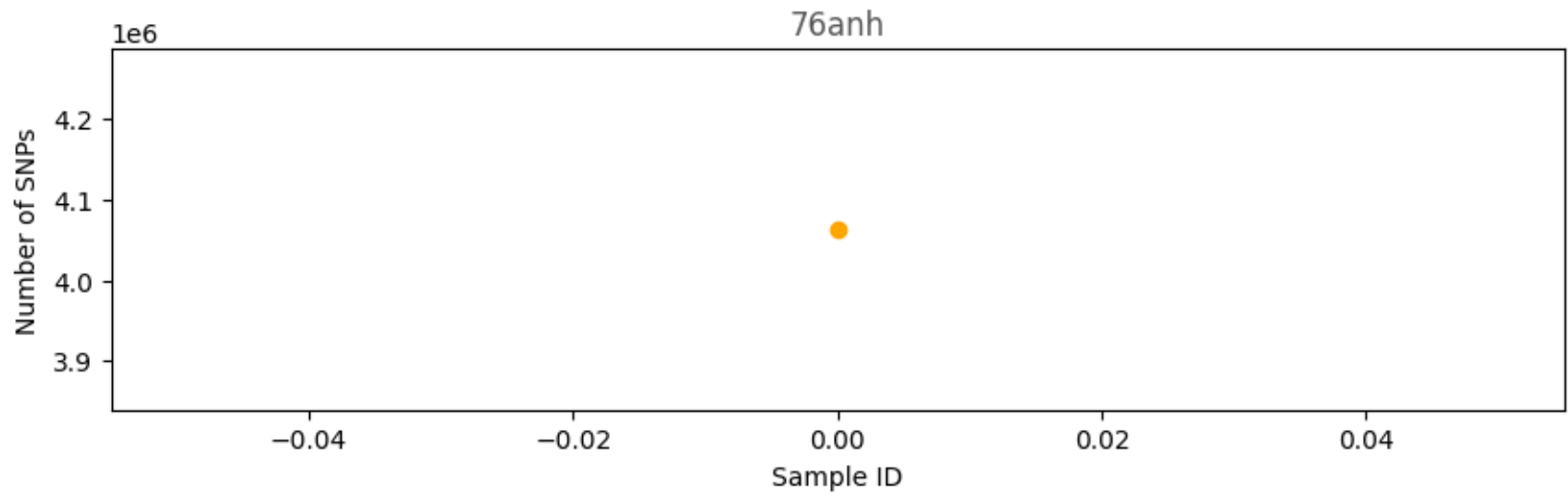
Ts/Tv by sample
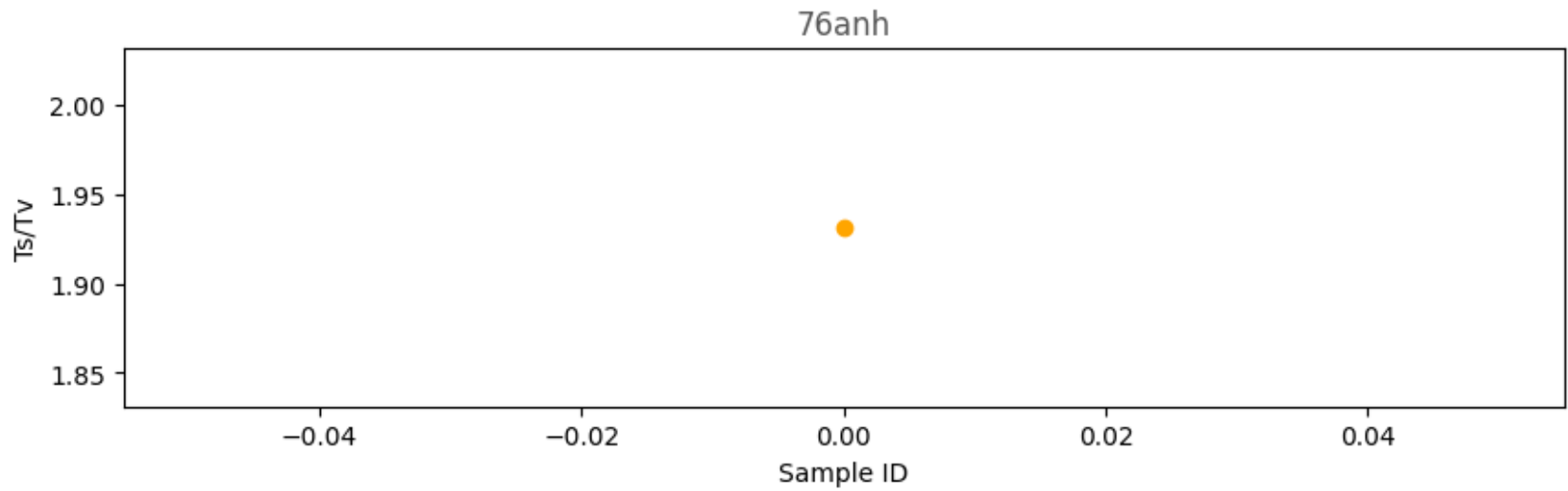
Hets vs non-ref Homs by sample
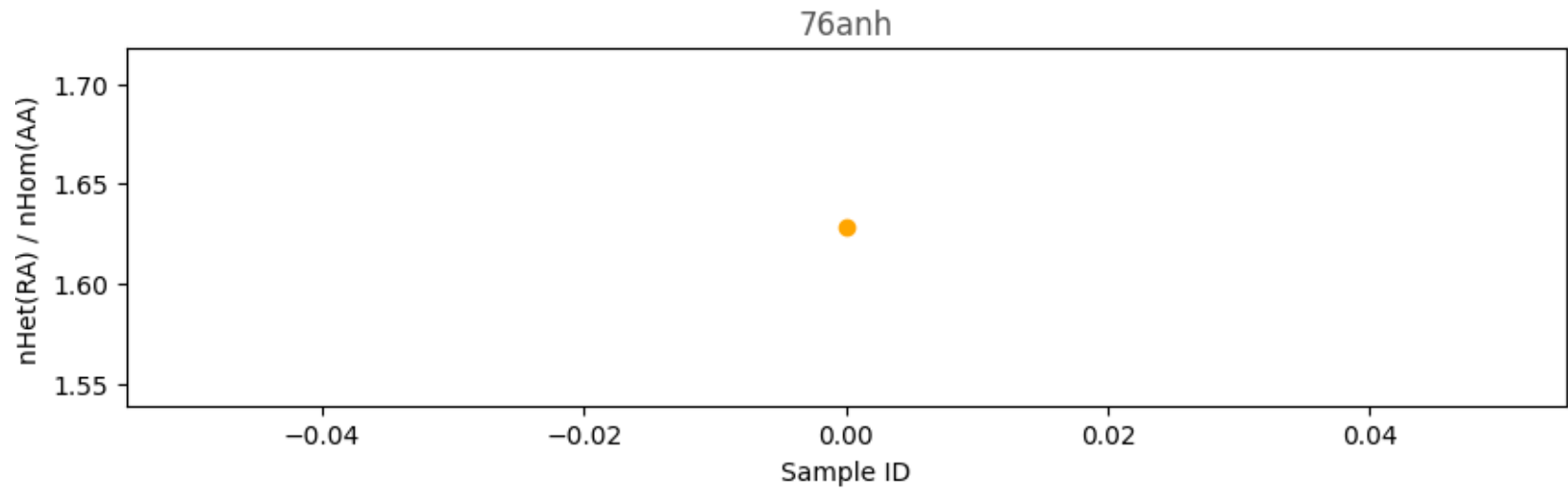
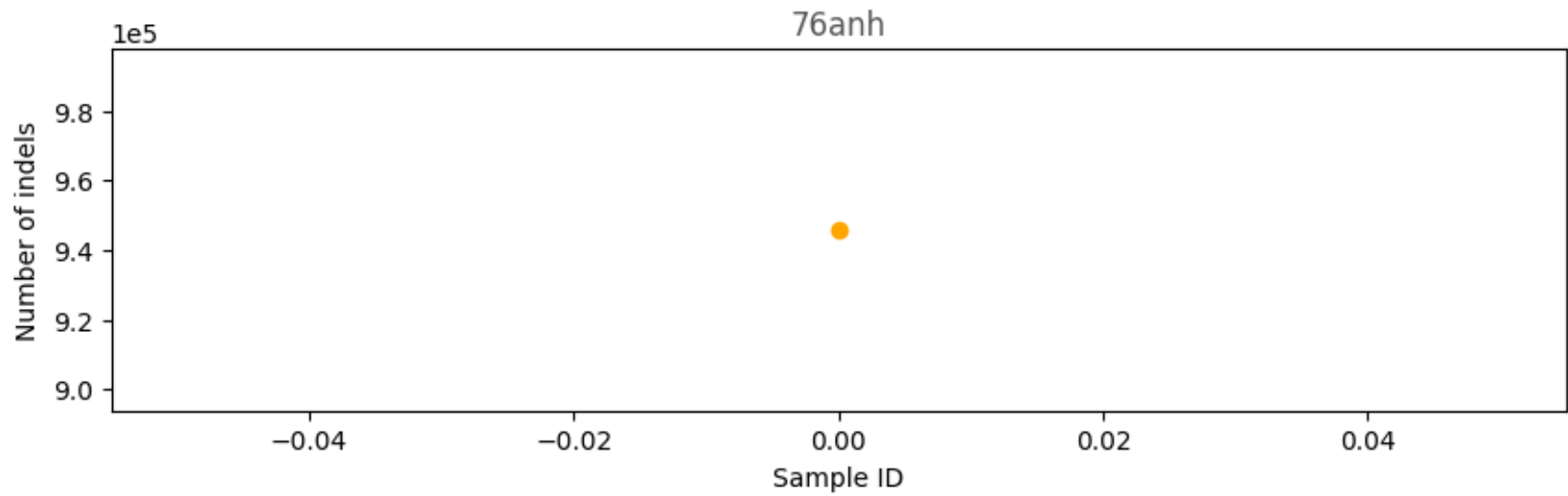# Singletons by sample (hets and homs)
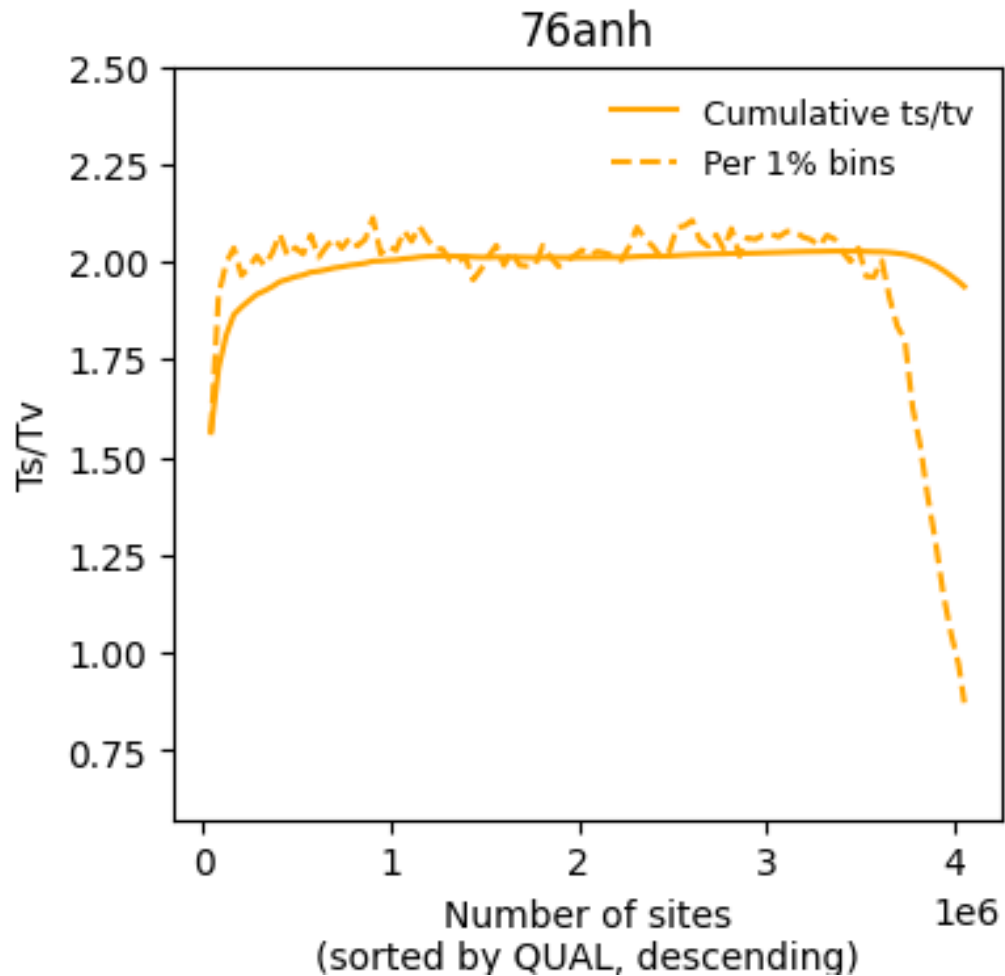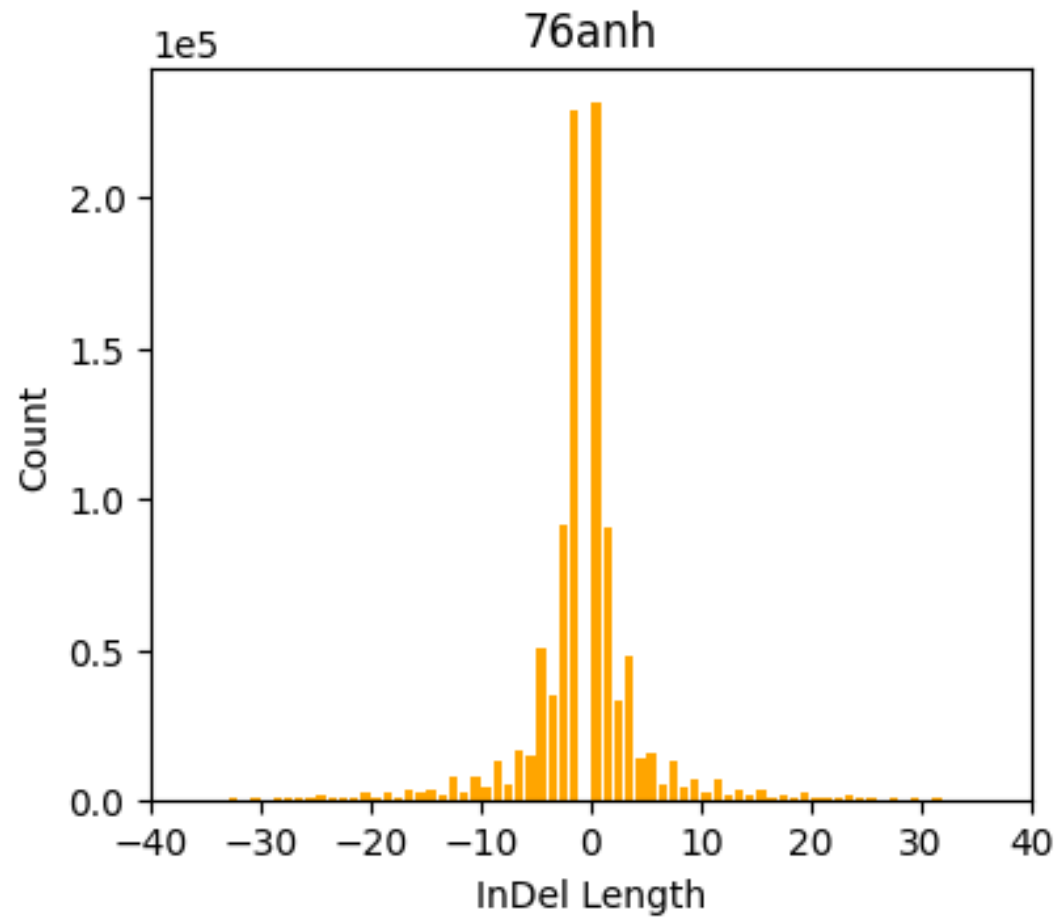
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

8

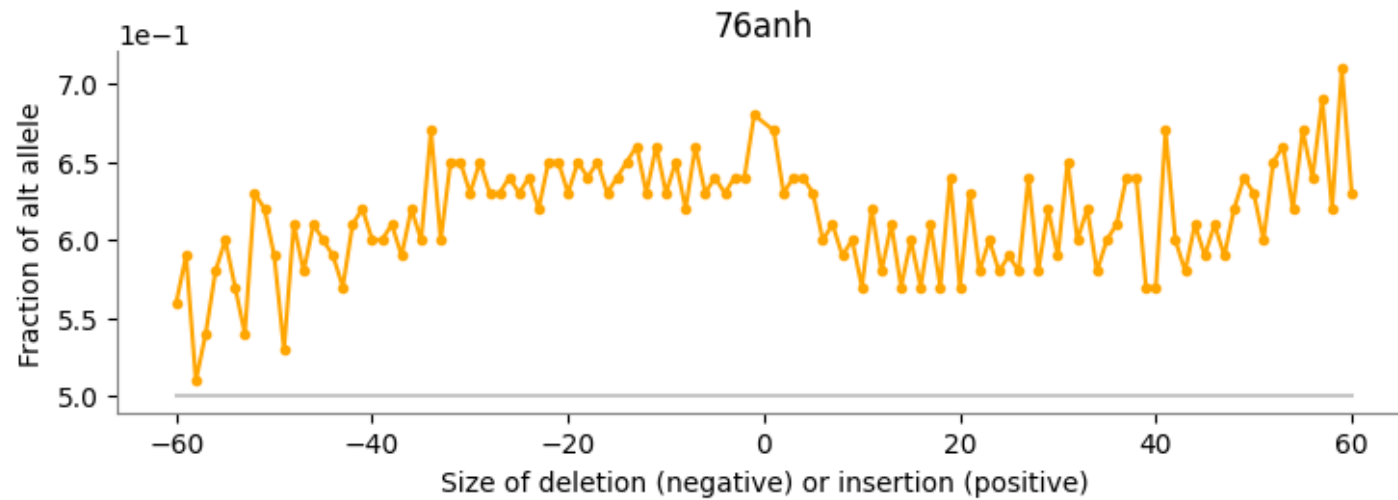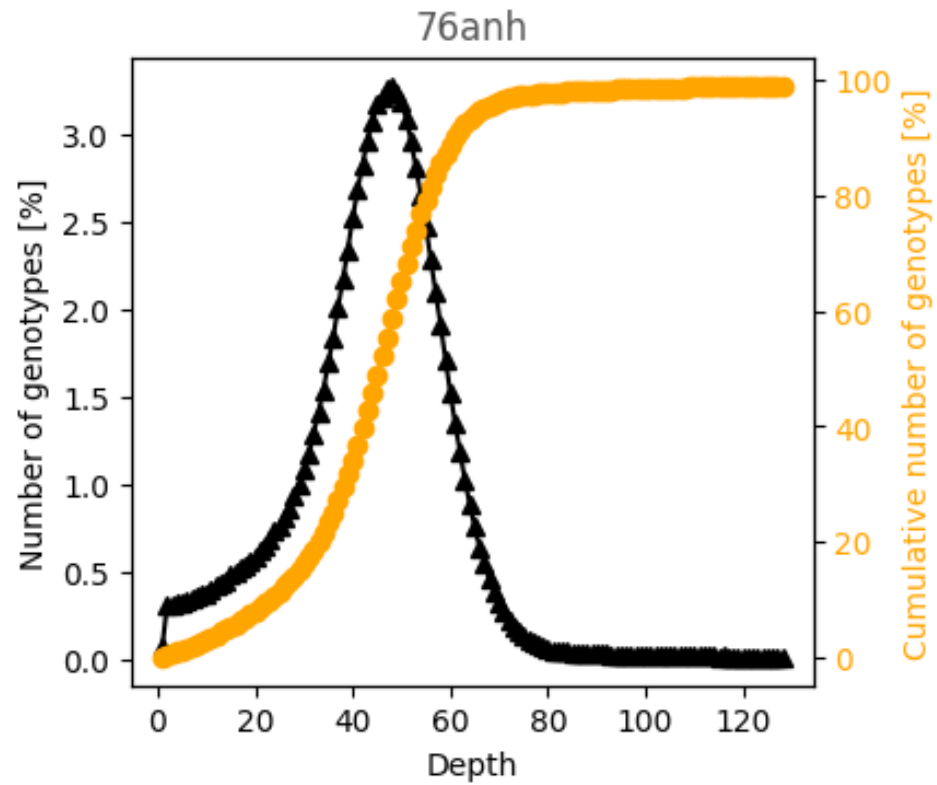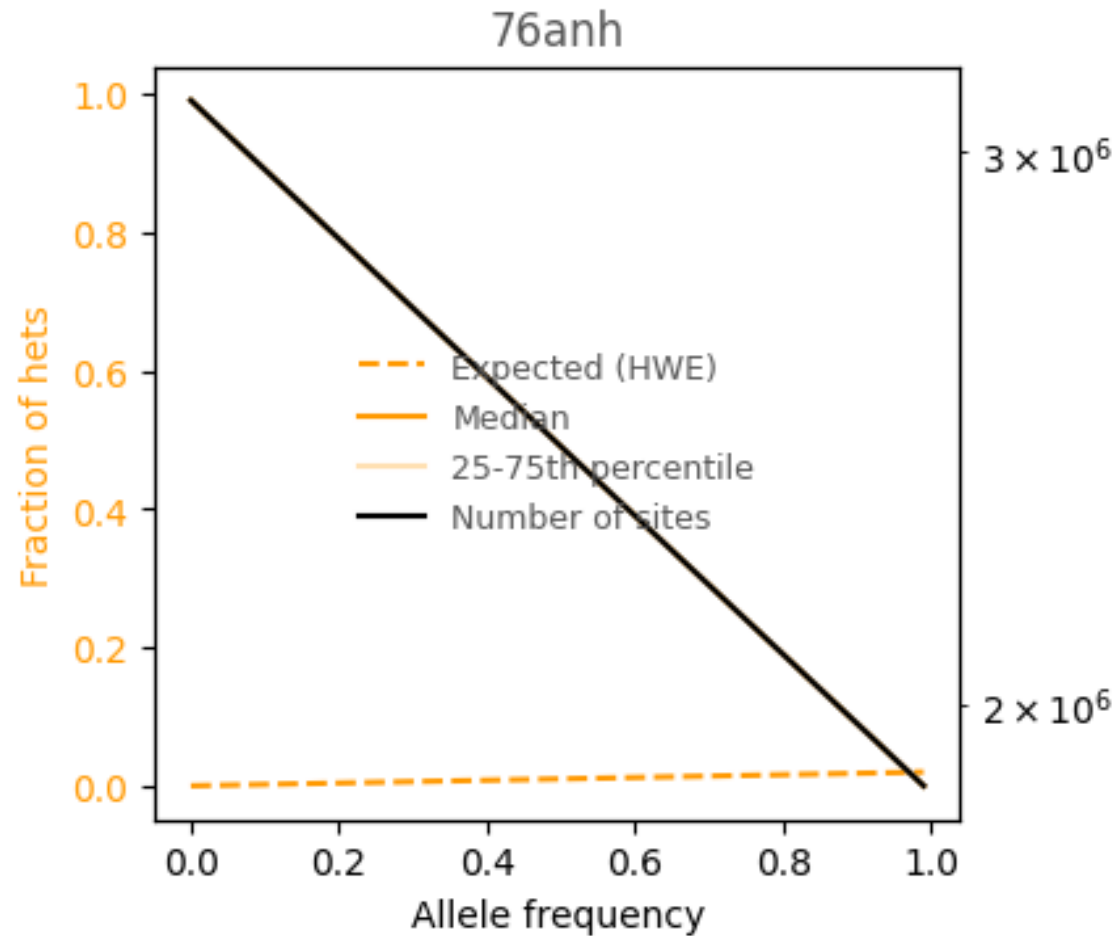# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types