Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
84ide	4,032,924	1.94	1.95	944,105	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
84ide	61.9%	1.91	66.7%	97,157	1,798

- 84ide .. `/ngc/projects2/gm/data/archive/2022/variants/snv/84idesssm-103907628986-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01248-220907_A01176_AHYN73DSX3-EXT_LABKA_NGCWGS-NGCWGS05020_22RKG011087x01_snv_germline_raw.haplotype_caller.vcf.gz`
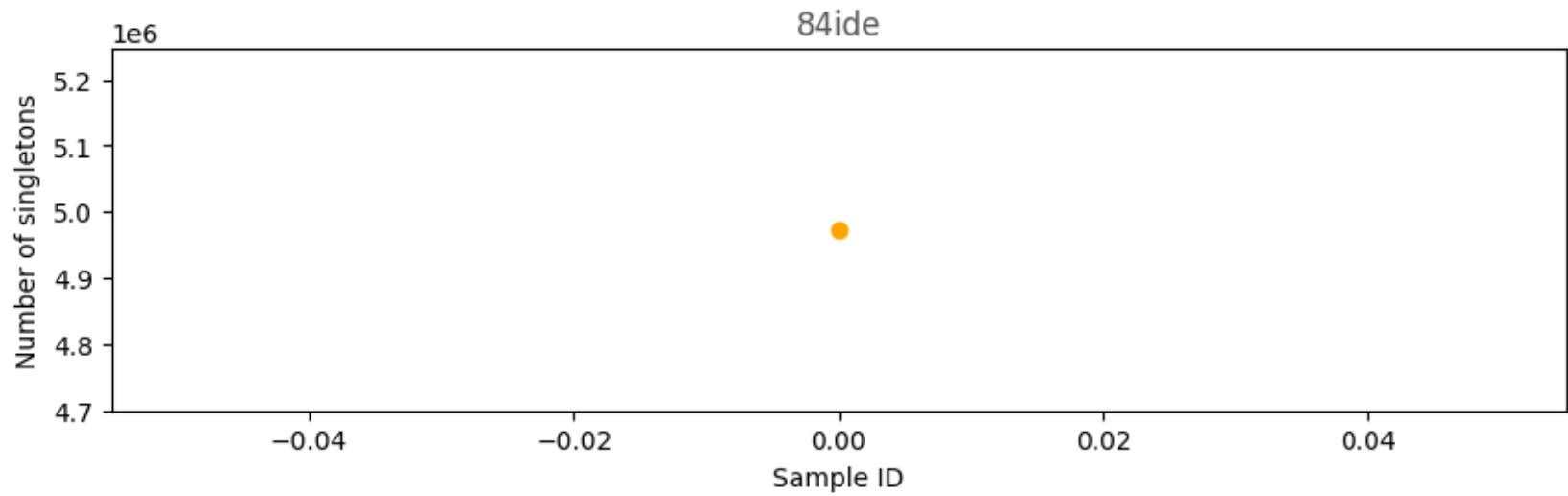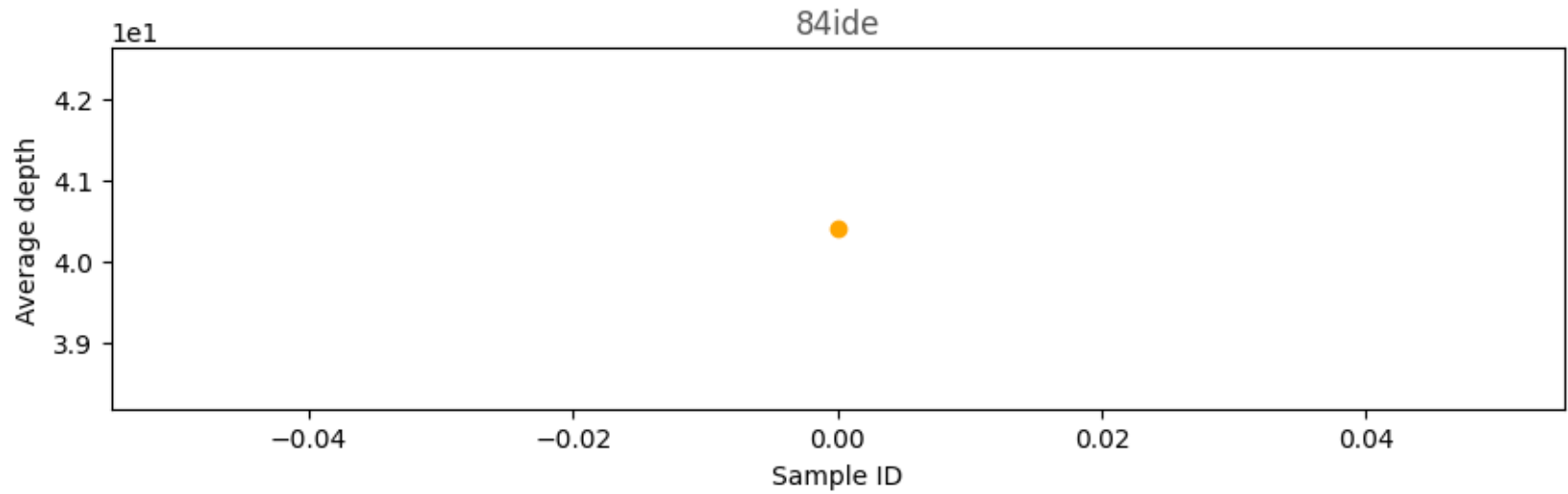
Ts/Tv by sample
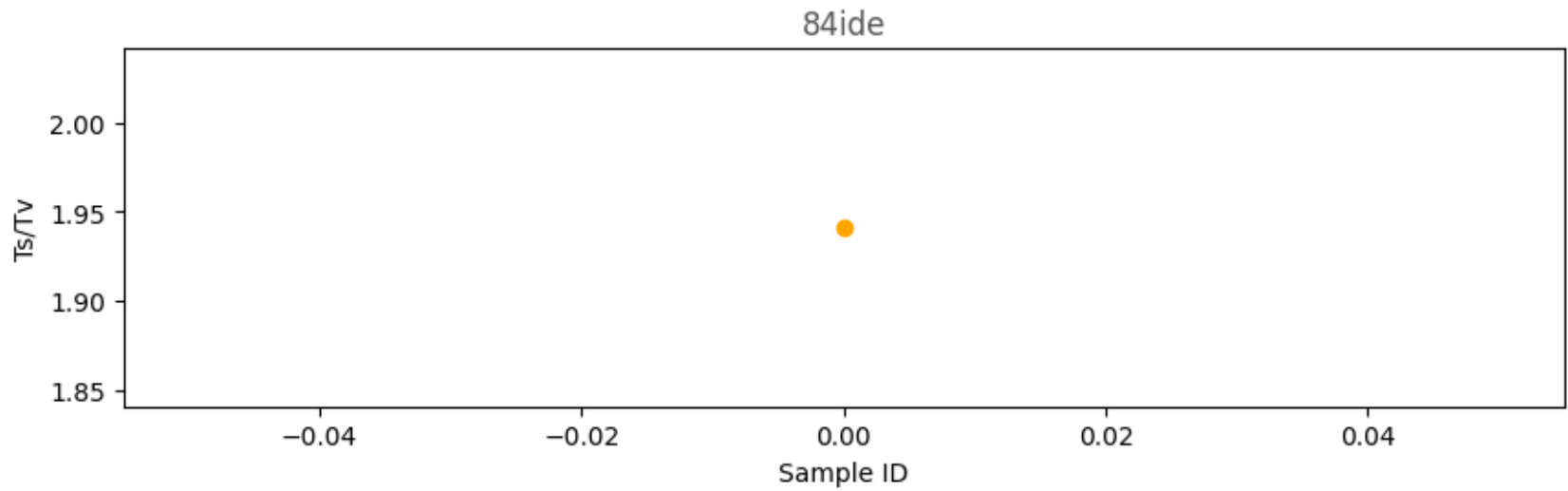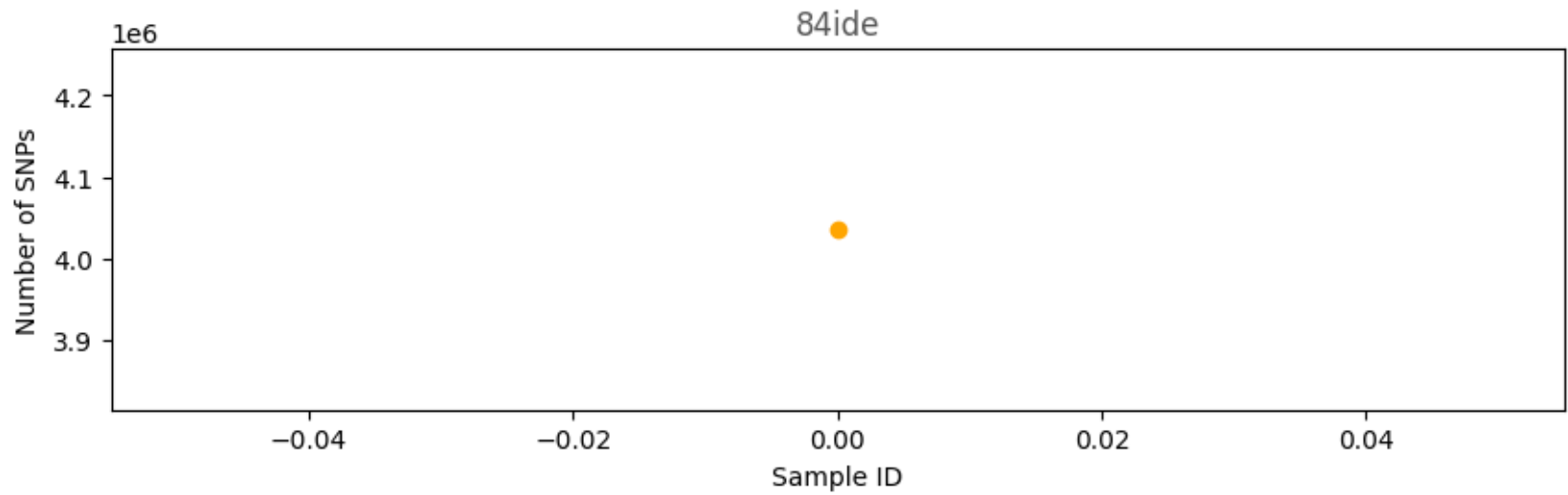
Hets vs non-ref Homs by sample
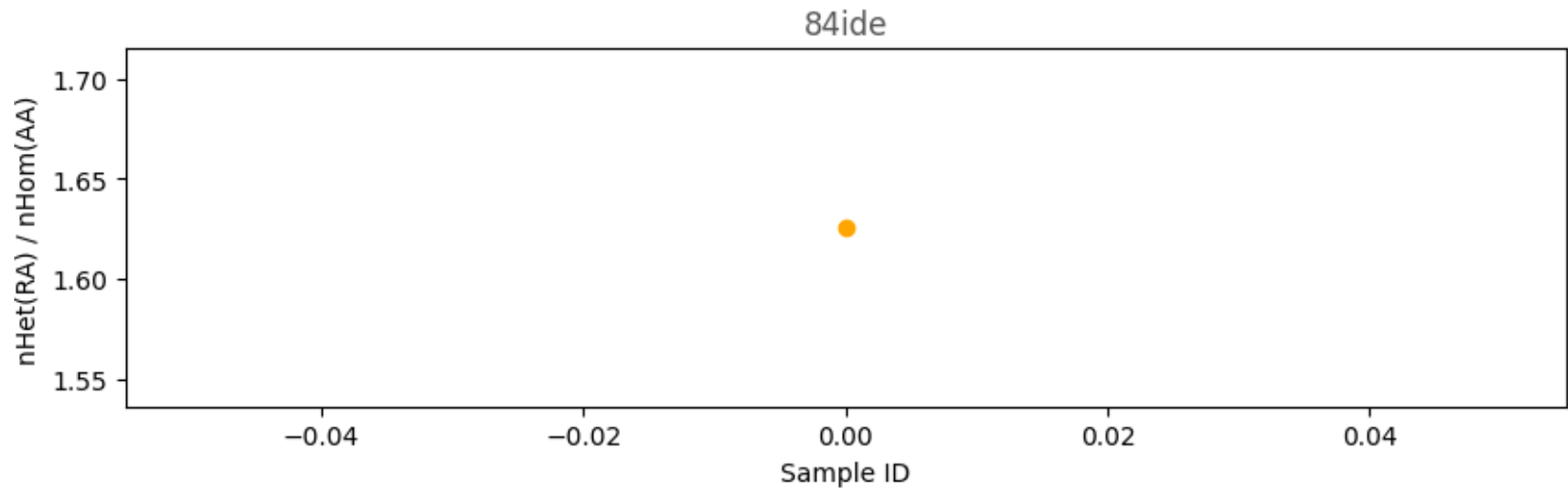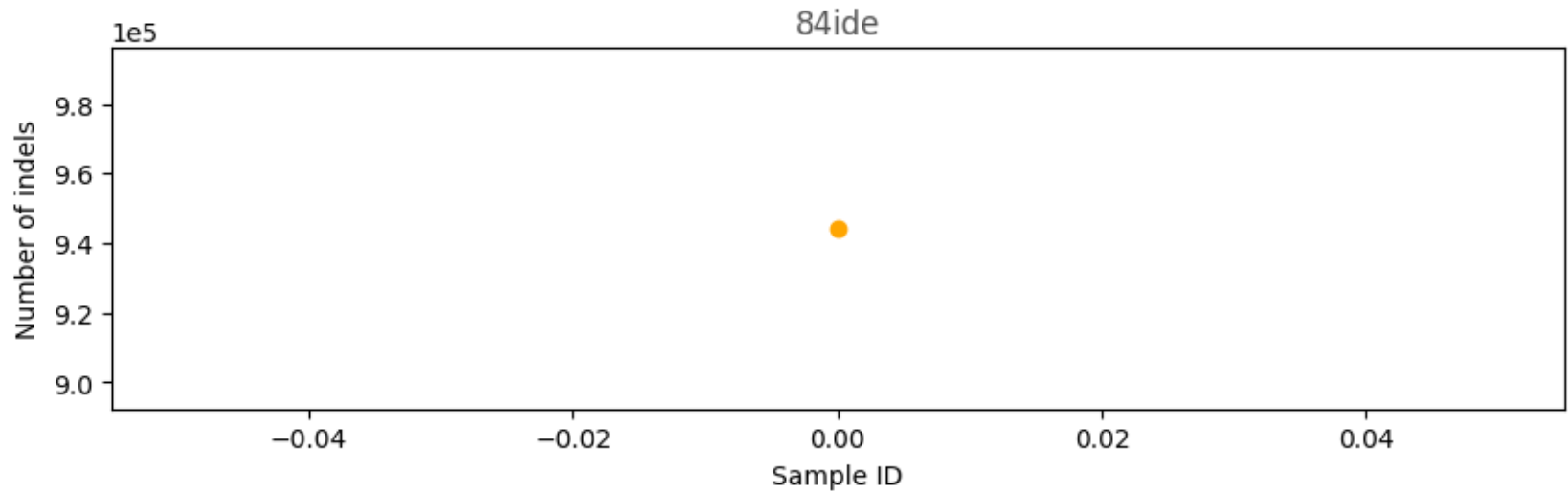
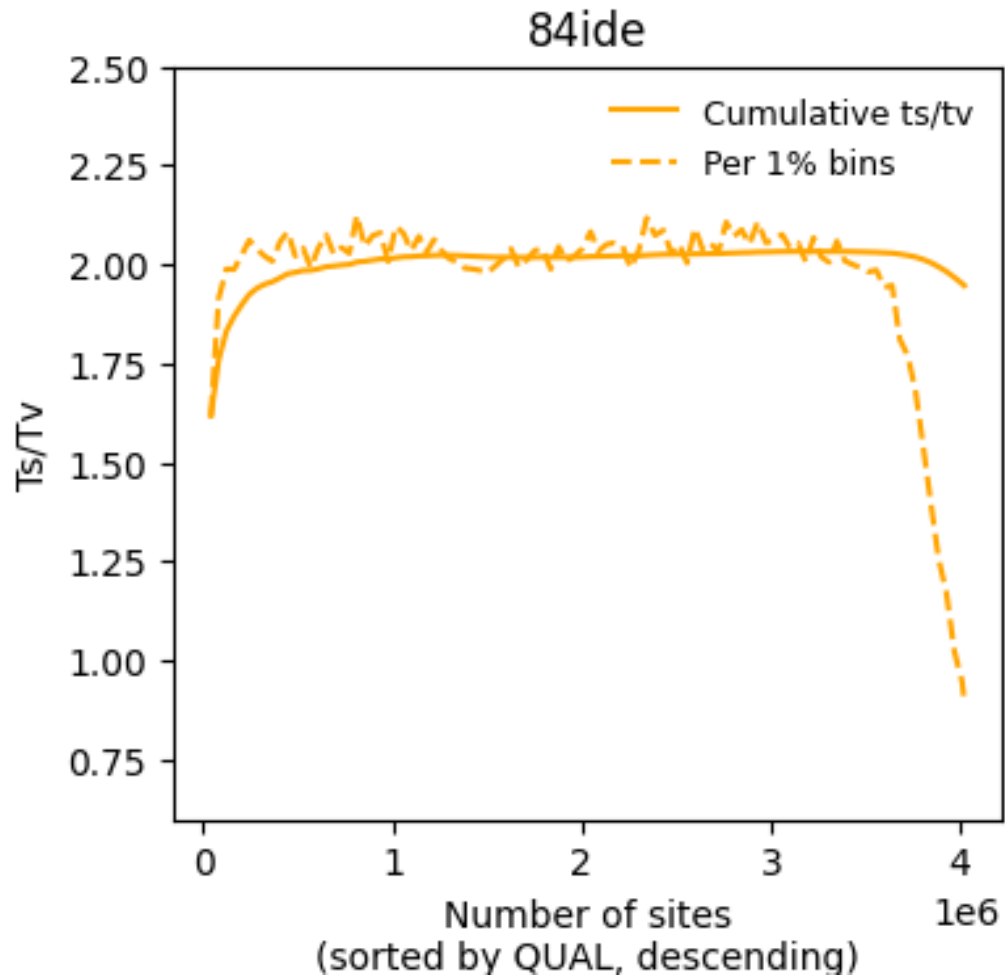# Singletons by sample (hets and homs)
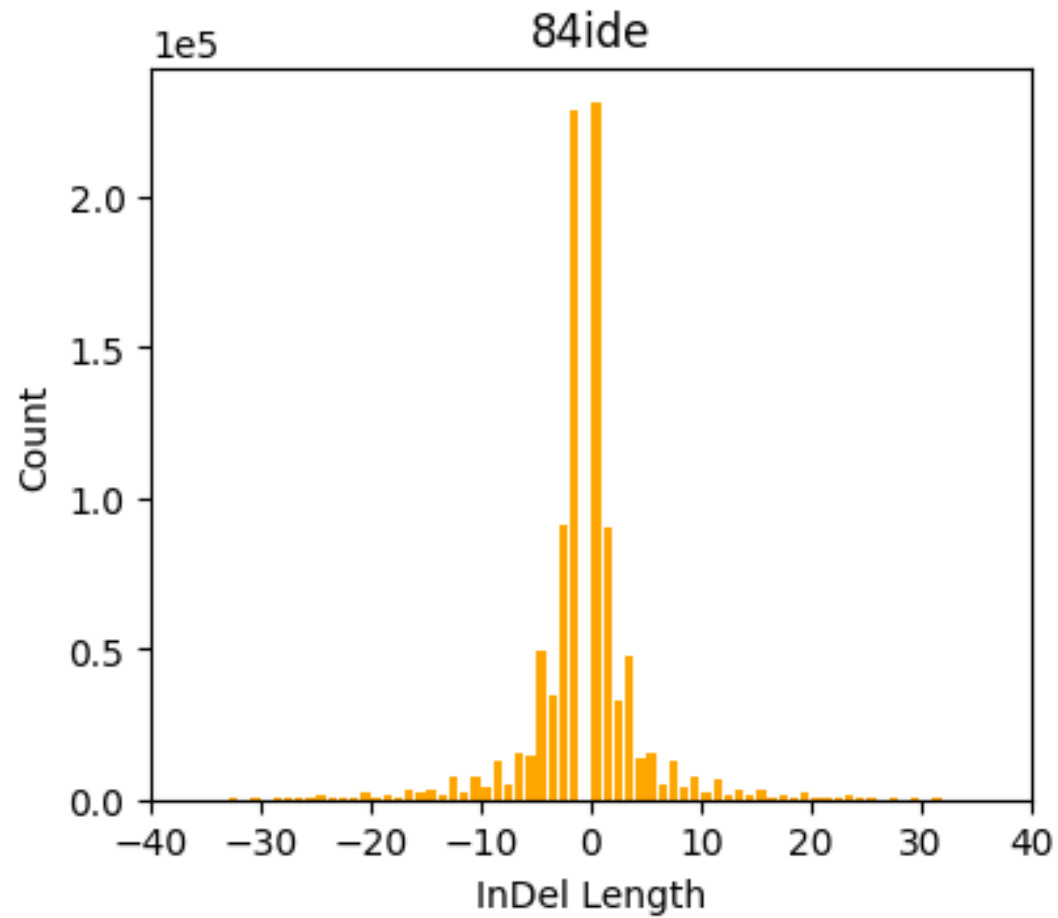
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

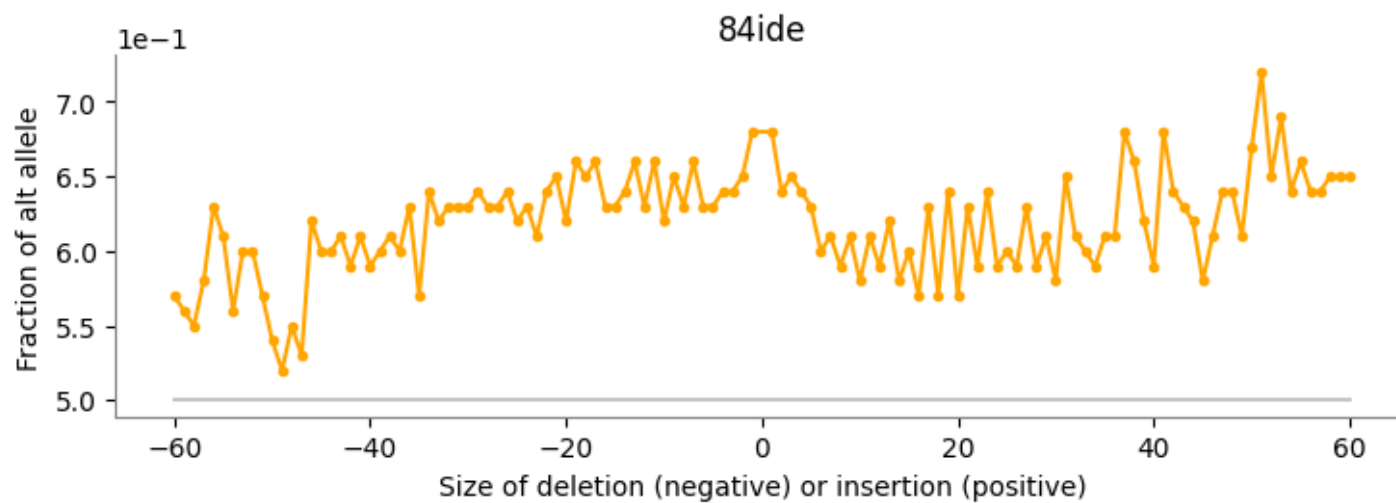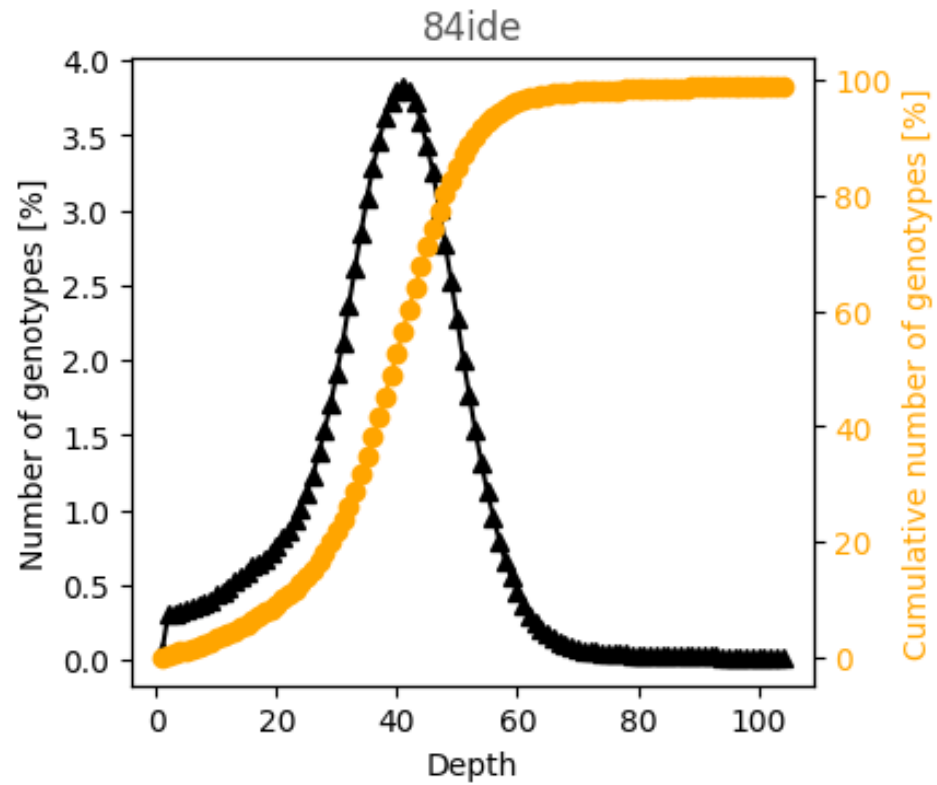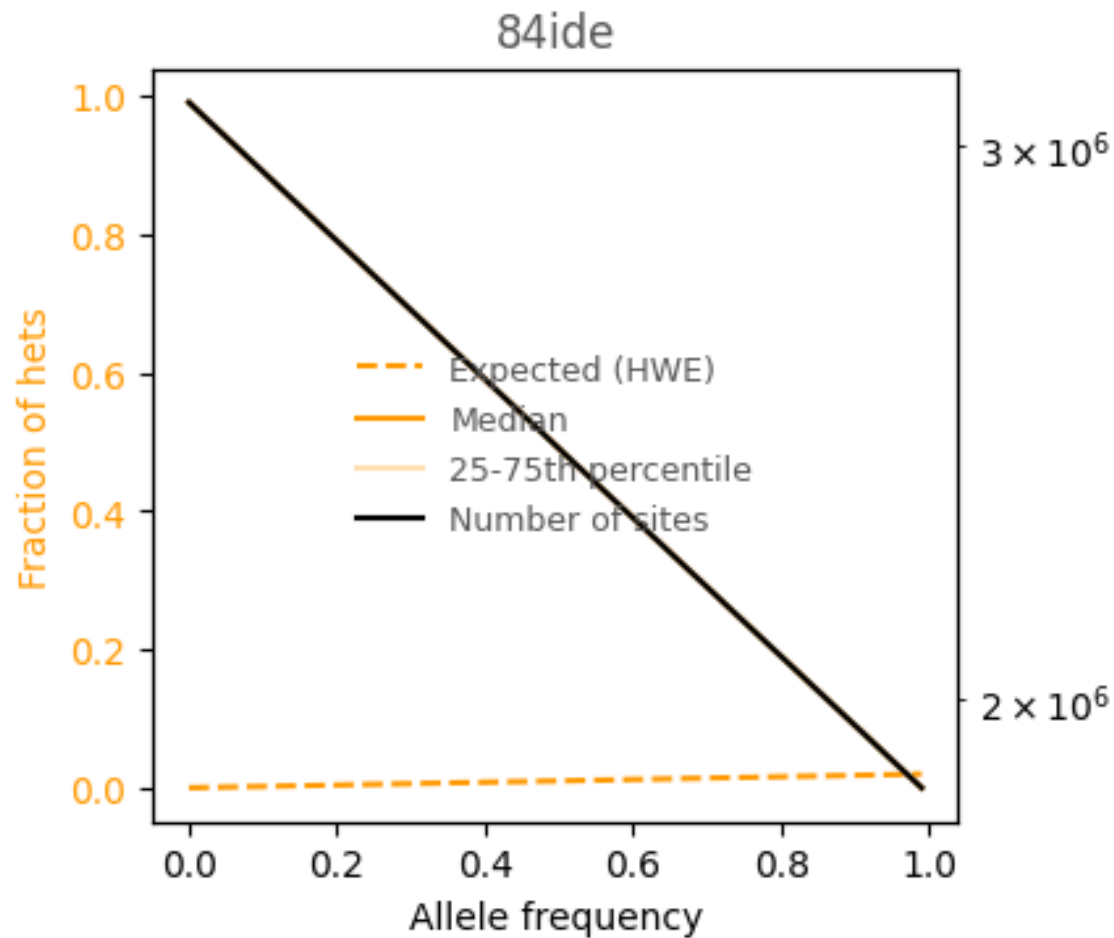# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types