Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
88mit	4,001,020	1.96	1.97	932,307	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
88mit	61.7%	1.94	65.7%	88,600	1,813

- 88mit .. `/ngc/projects2/gm/data/archive/2022/variants/snv/88mitoftm-103776919629-Normal_B lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00294-210907_A01176_AHJVNLDSX2-EXT_LAB KA_NGCWGS-NGCWGS00188_21RKG020650x01_snv_germline_raw.haplotype_caller.vcf.gz`
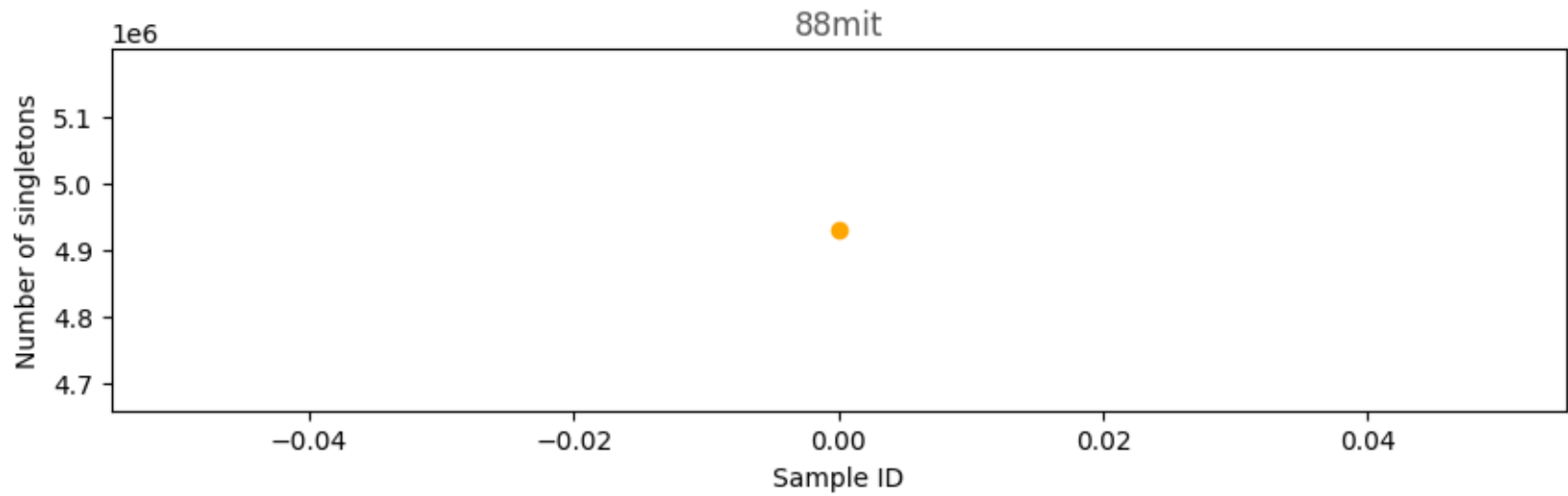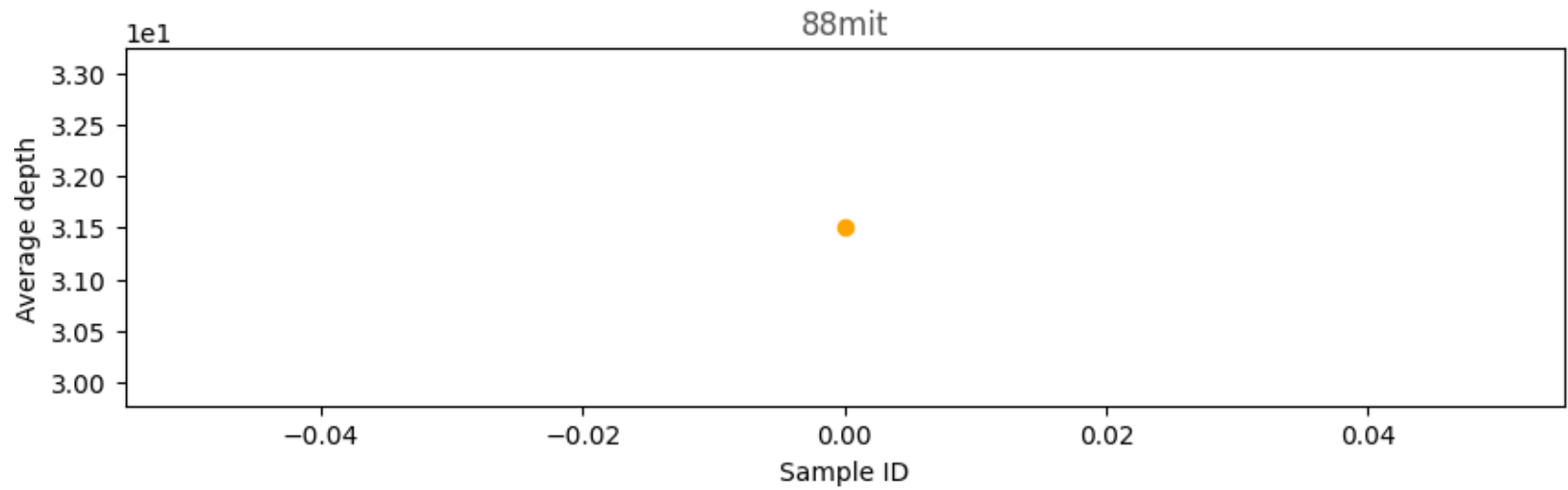
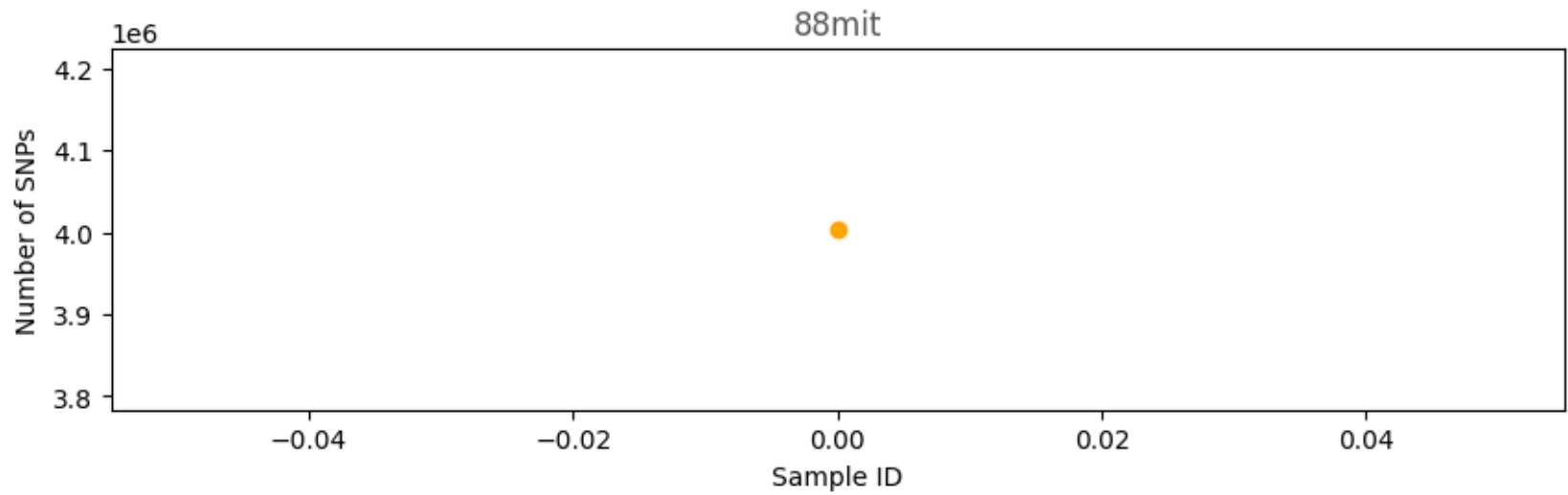# Ts/Tv by sample
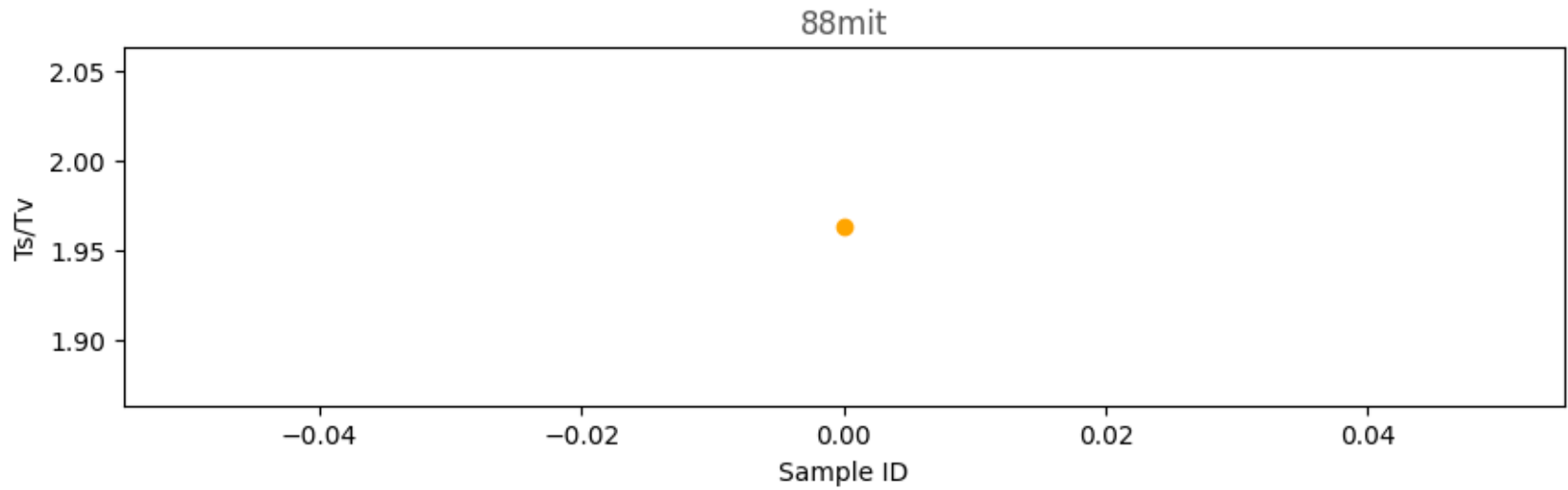
Hets vs non-ref Homs by sample
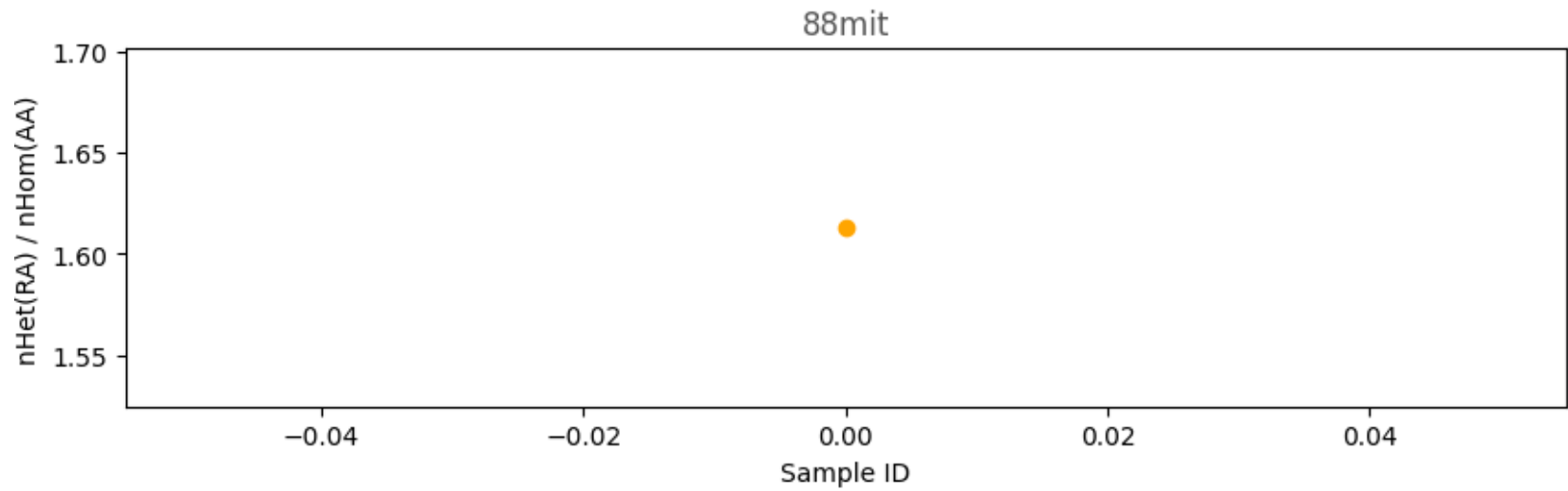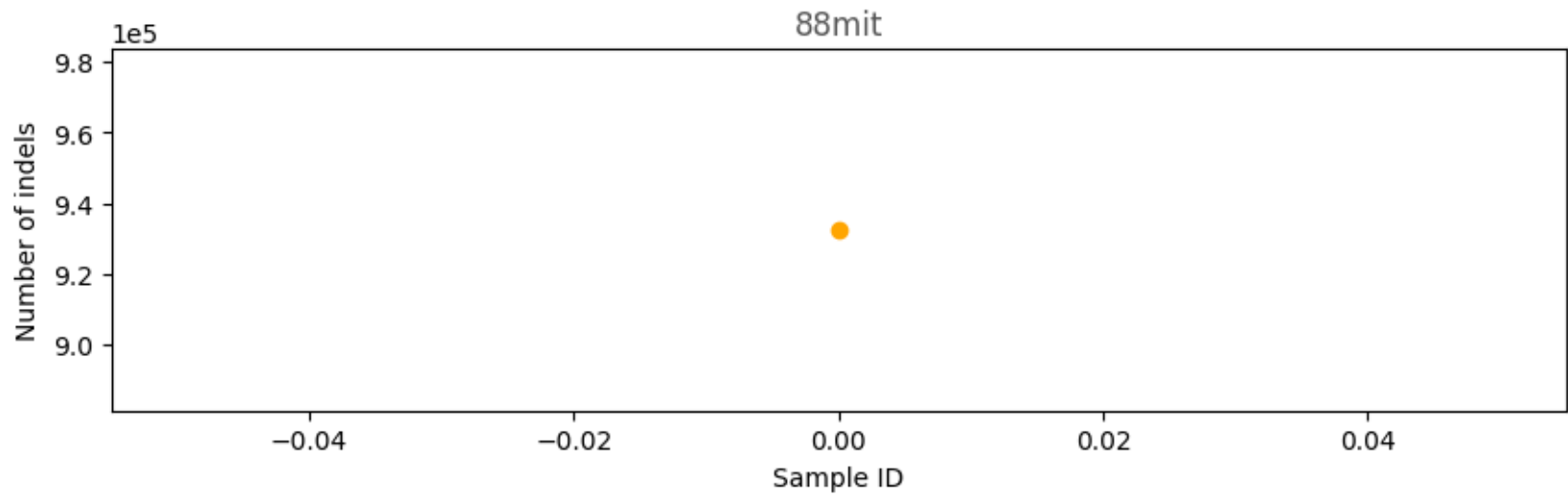
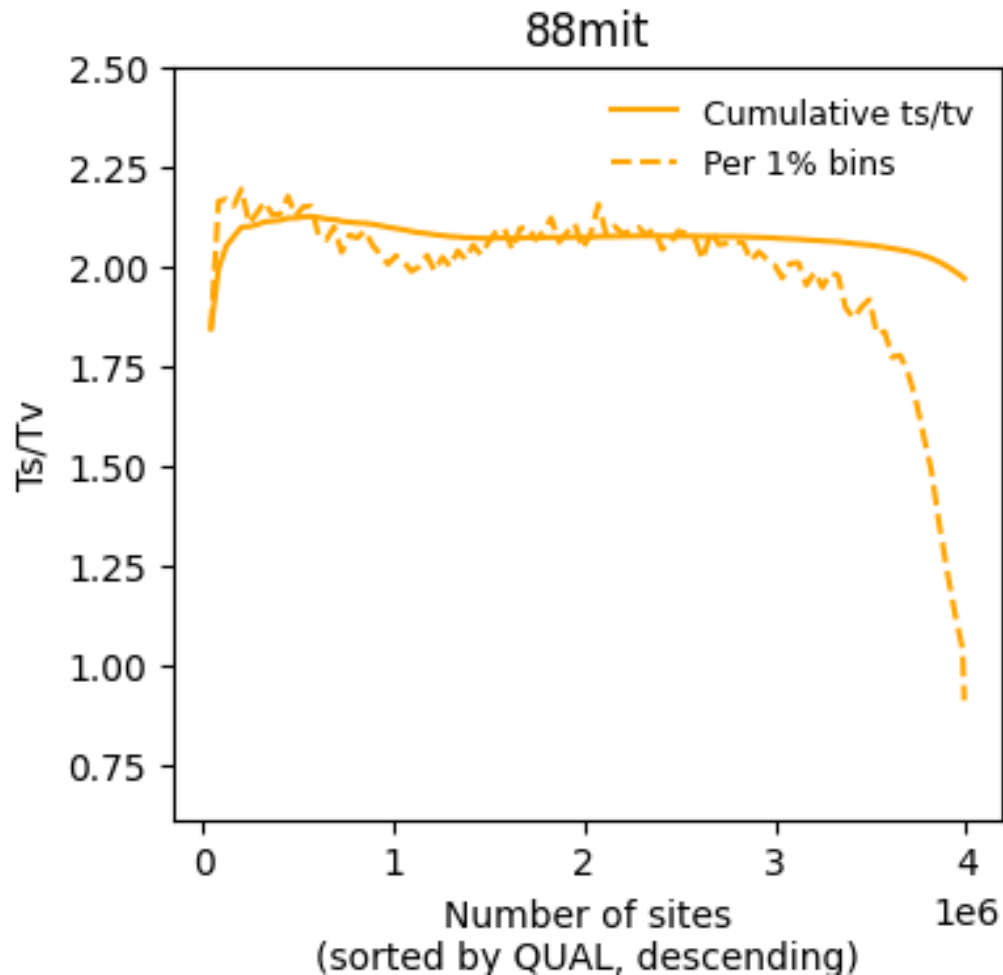# Singletons by sample (hets and homs)
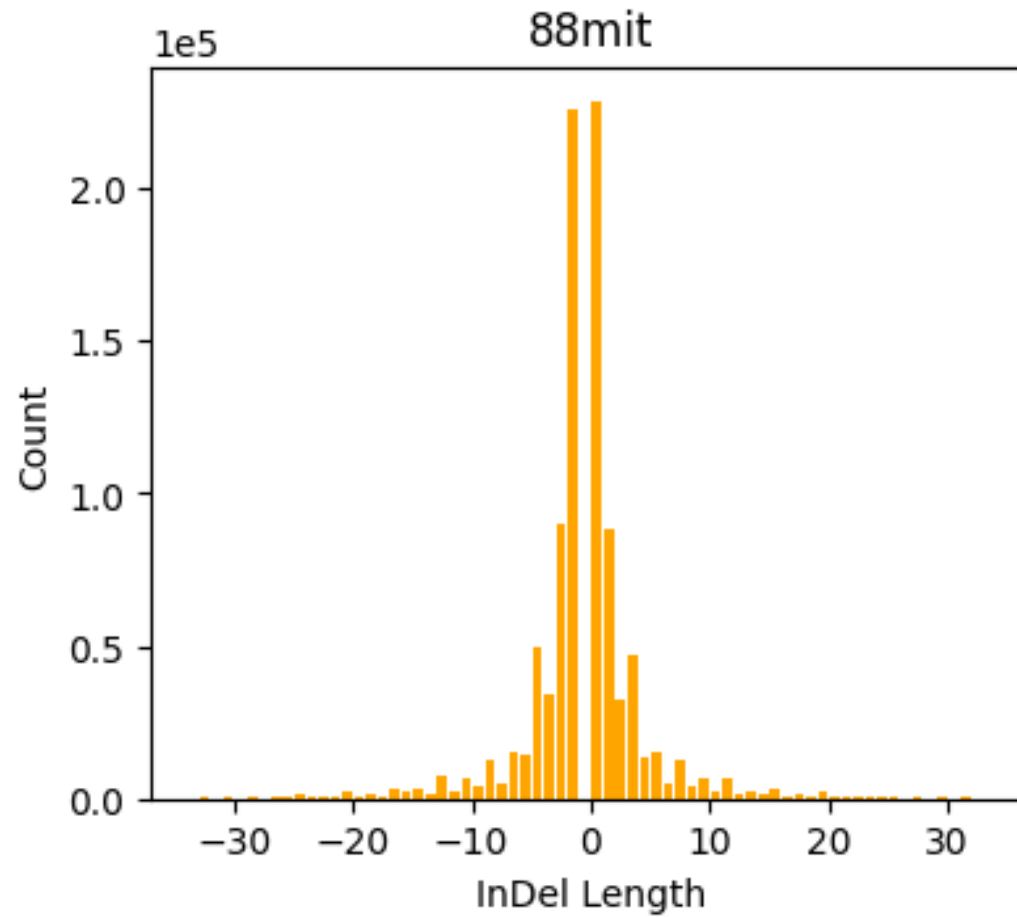
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

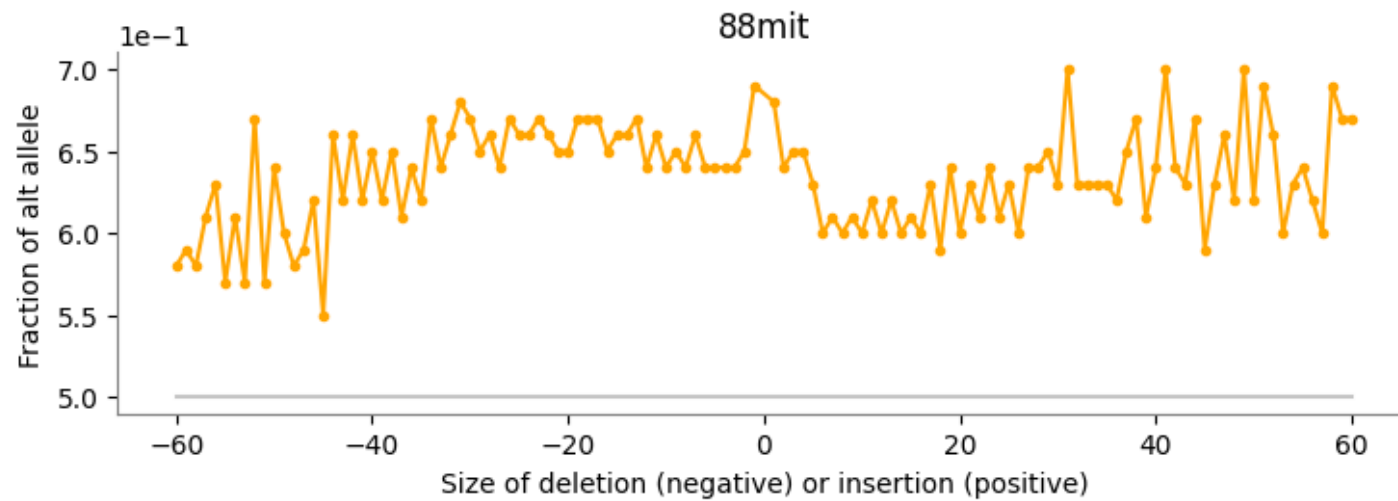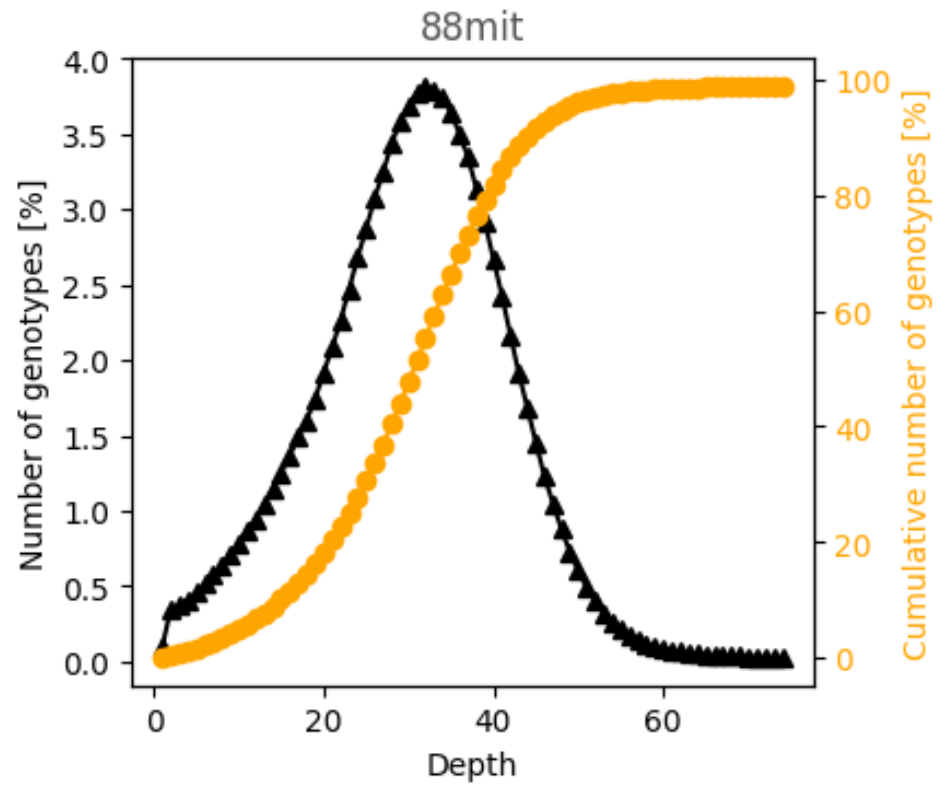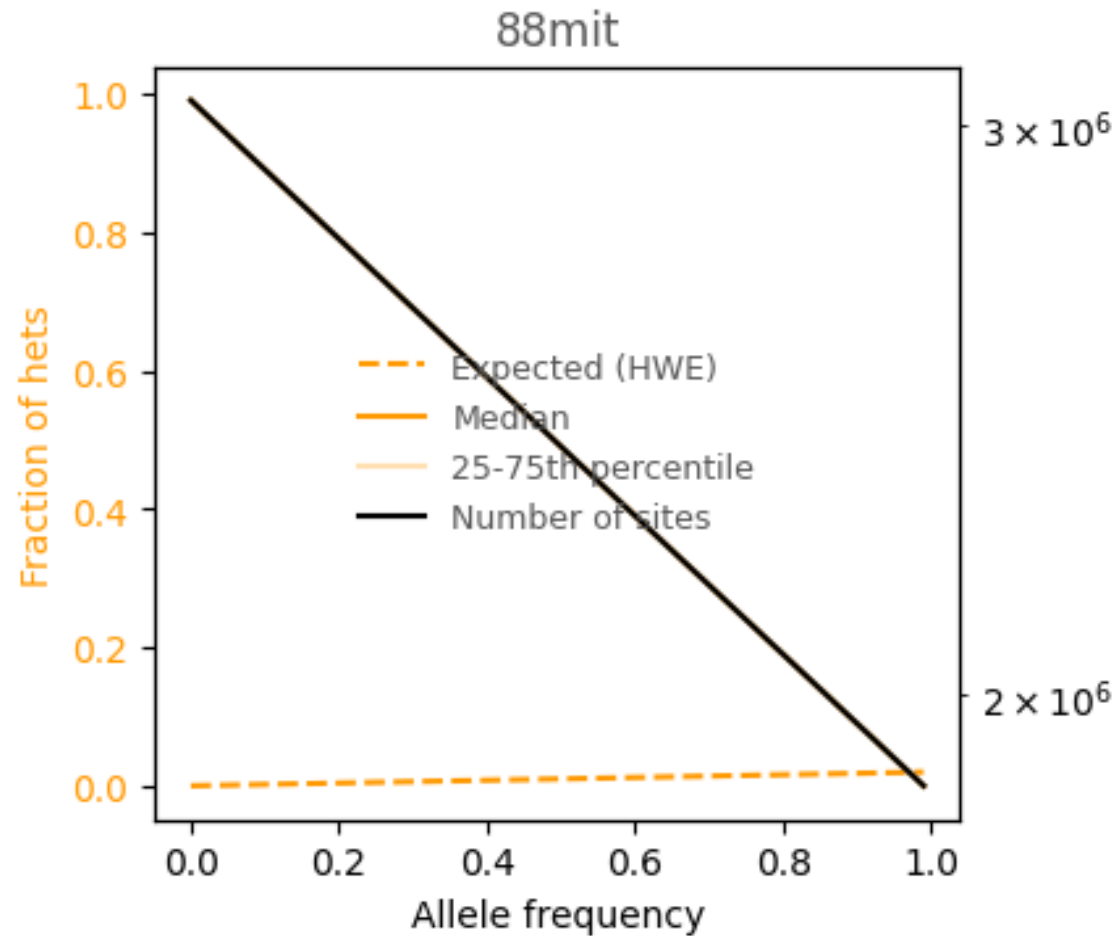# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types