Summary Numbers

	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
92dme	4,083,859	1.93	1.93	951,389	–	0	0
					* frameshift ratio: out/(out+in)		

	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
92dme	62.4%	1.90	67.2%	99,181	2,001

- 92dme .. `/ngc/projects2/gm/data/archive/2022/variants/snv/92dmererm-103903011488-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220819_A00559_AHMW72DSX3-RHGM_LABKA_WGSA`
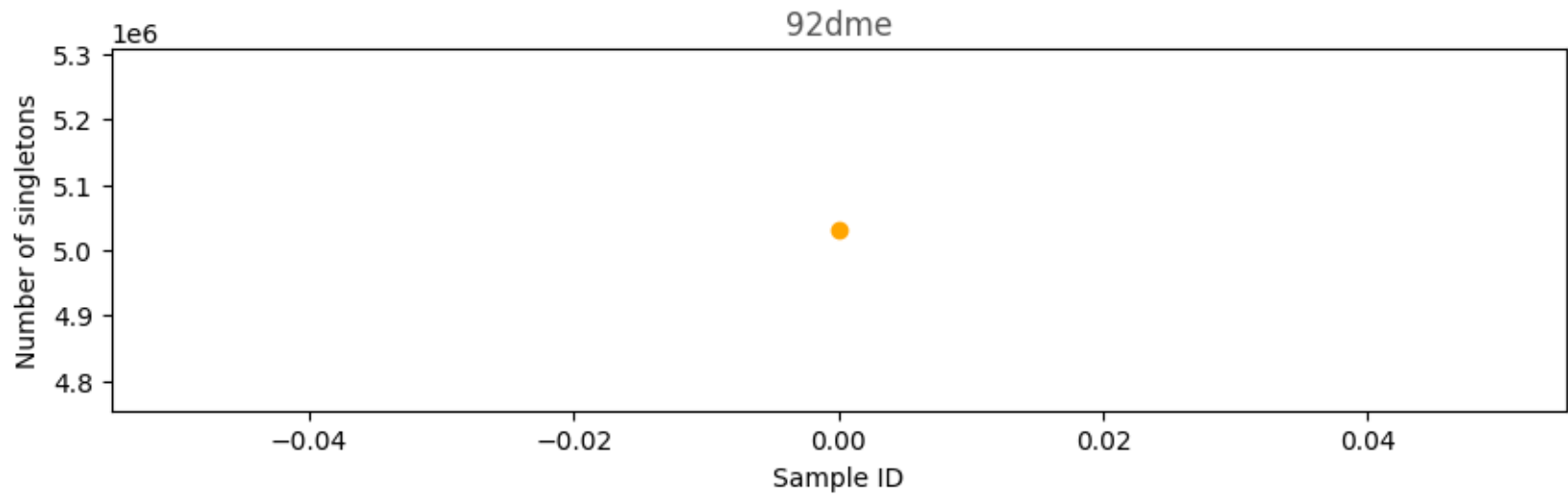 `KUT-WGSAKUT04852_22RKG019488x01_snv_germline_raw.haplotype_caller.vcf.gz`
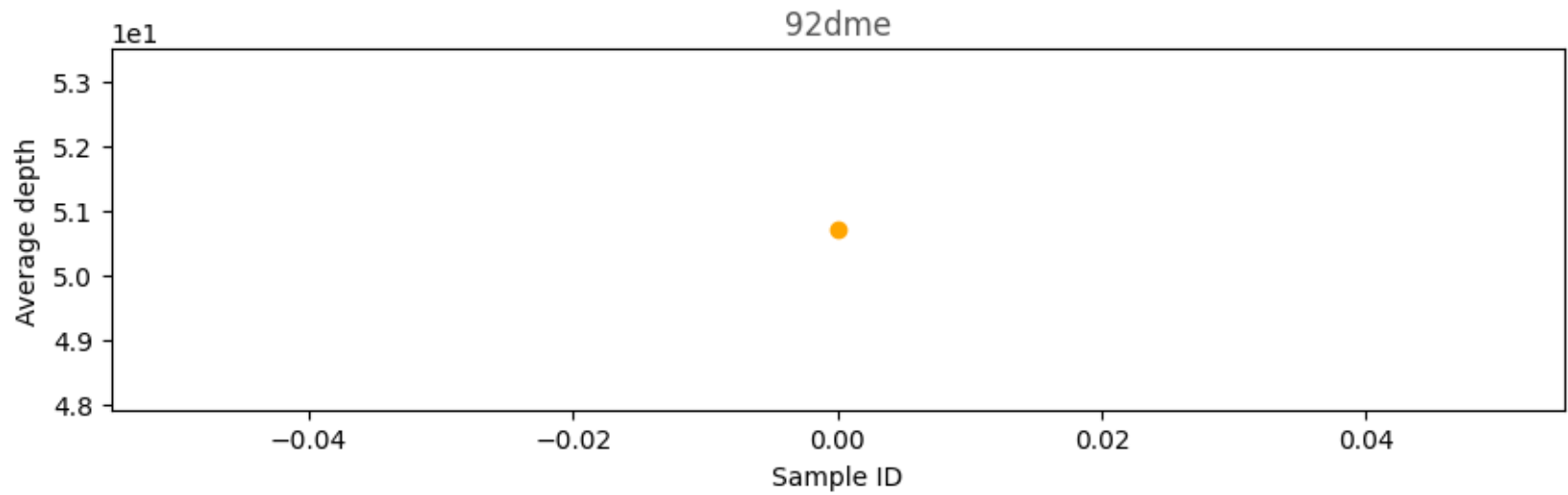
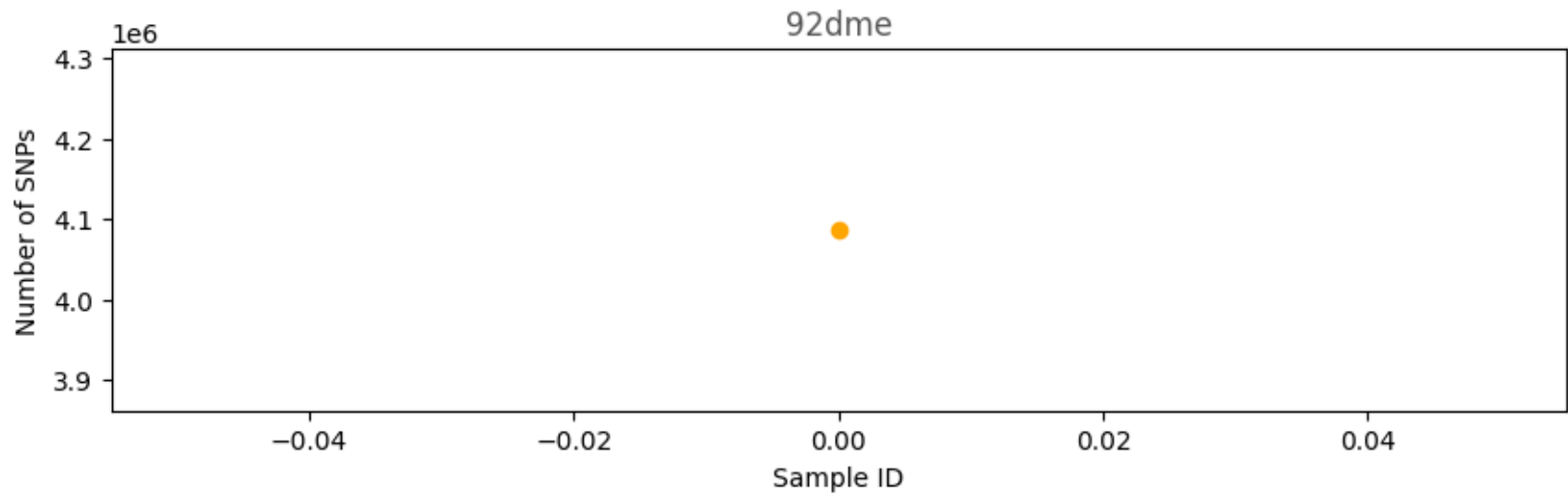# Ts/Tv by sample
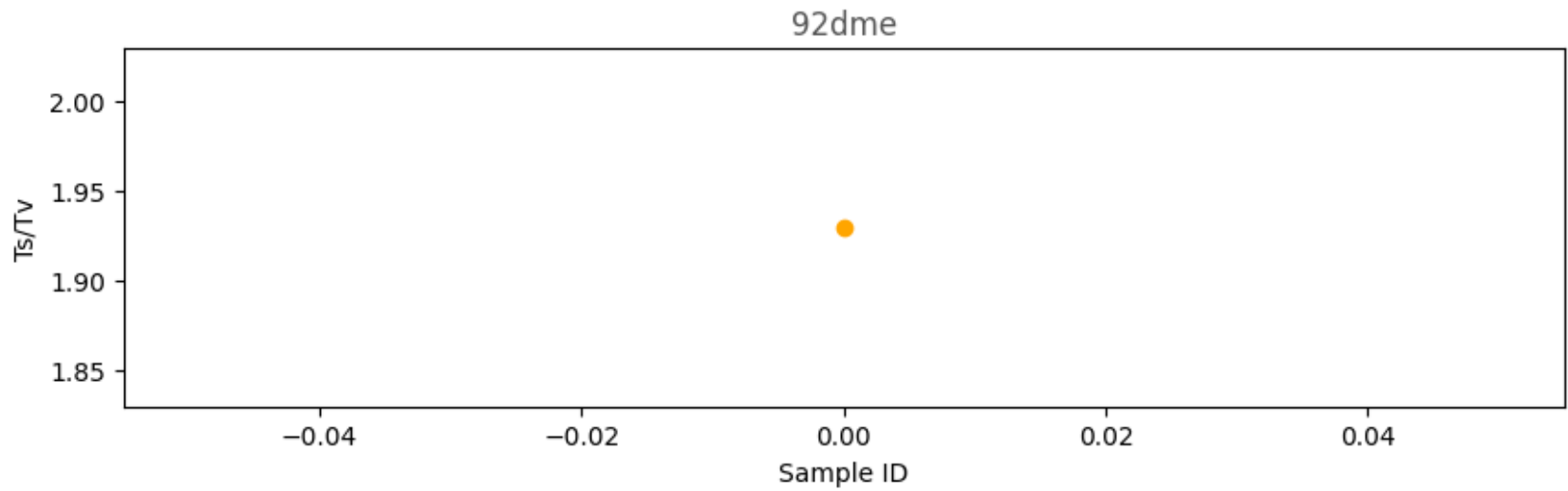
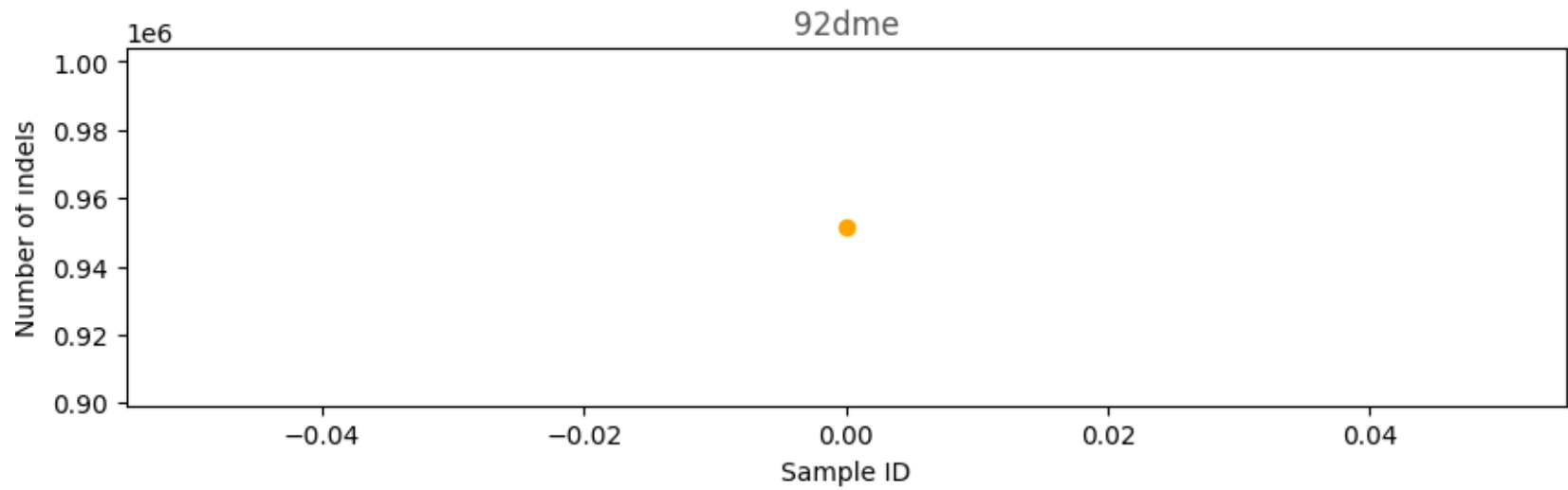# Hets vs non-ref Homs by sample
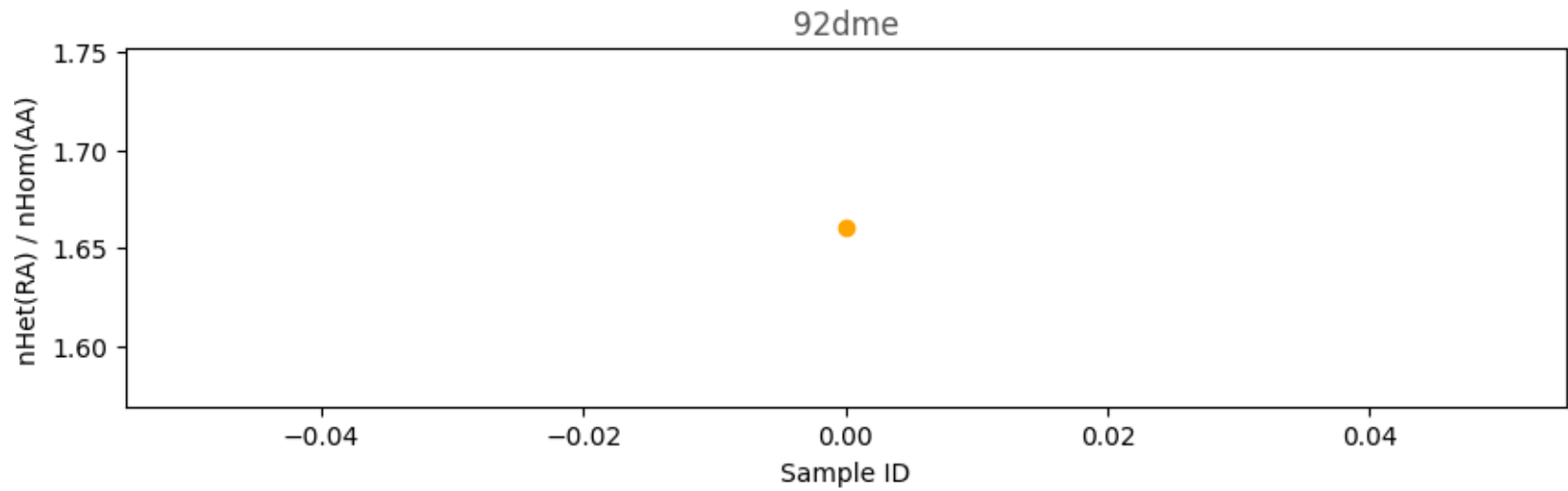
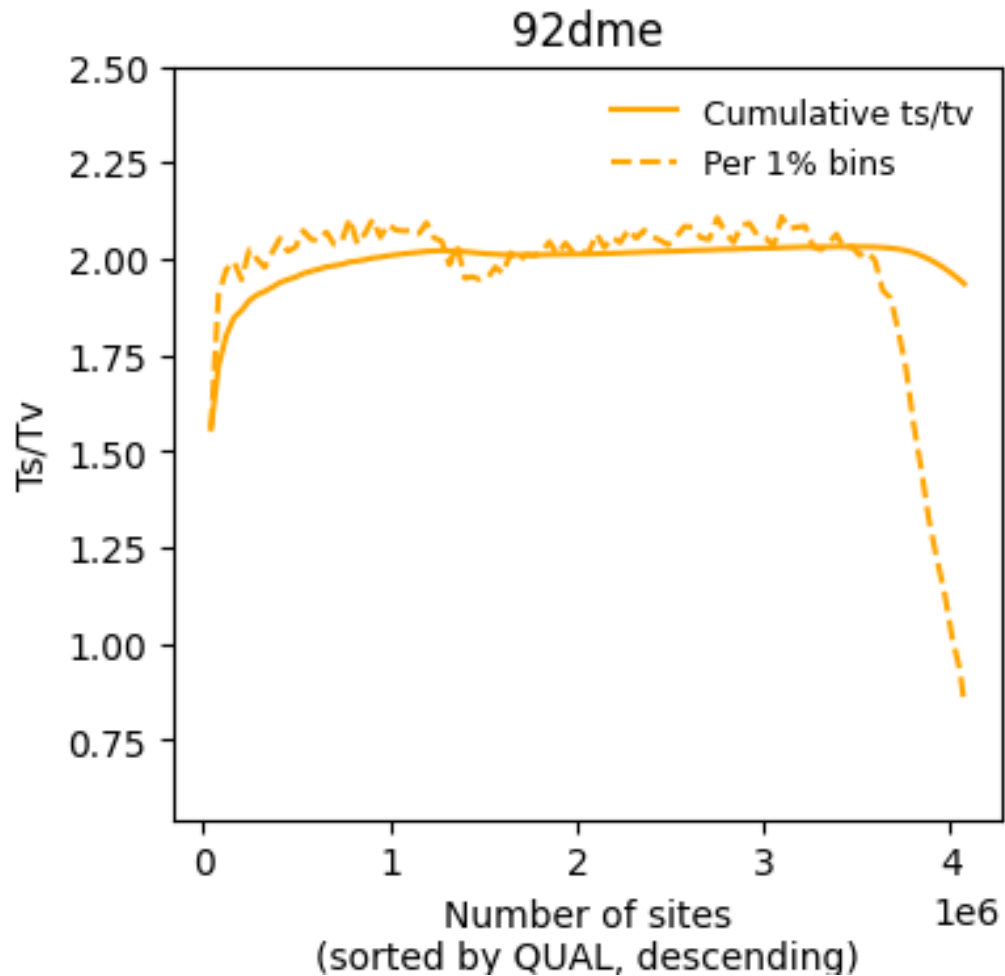# Singletons by sample (hets and homs)
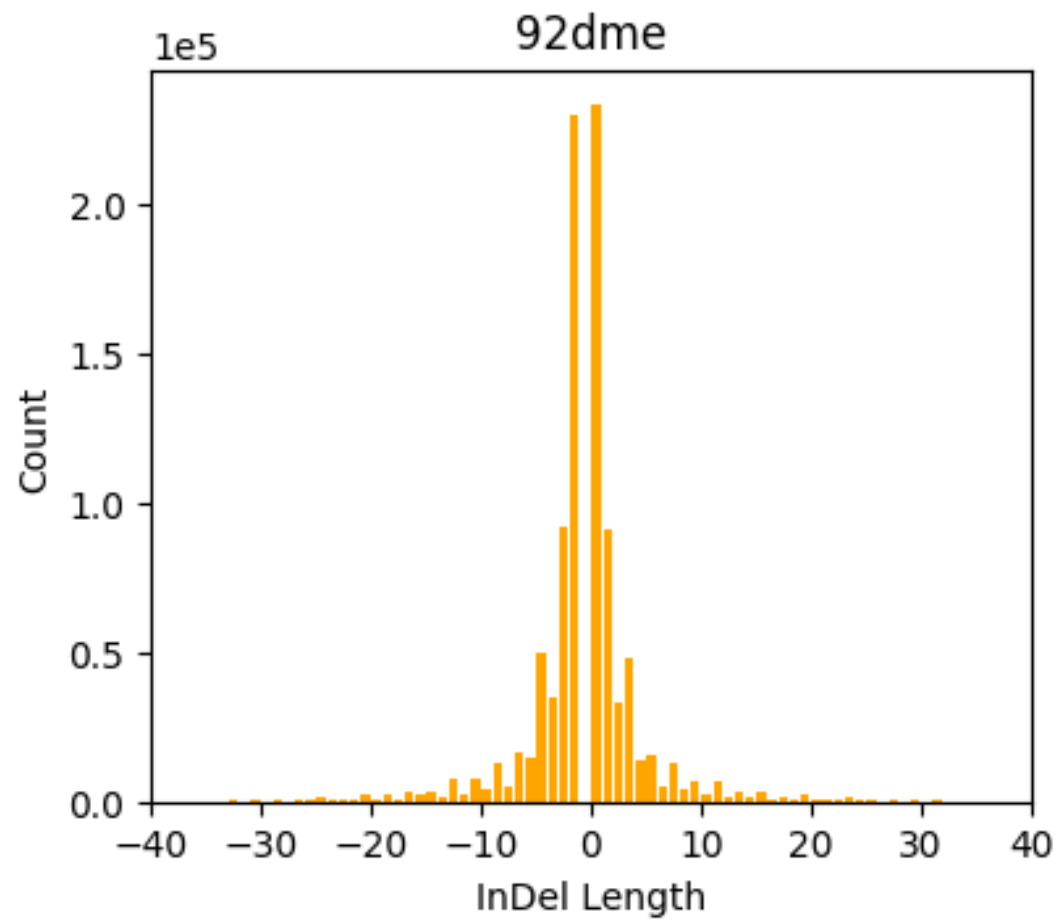
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

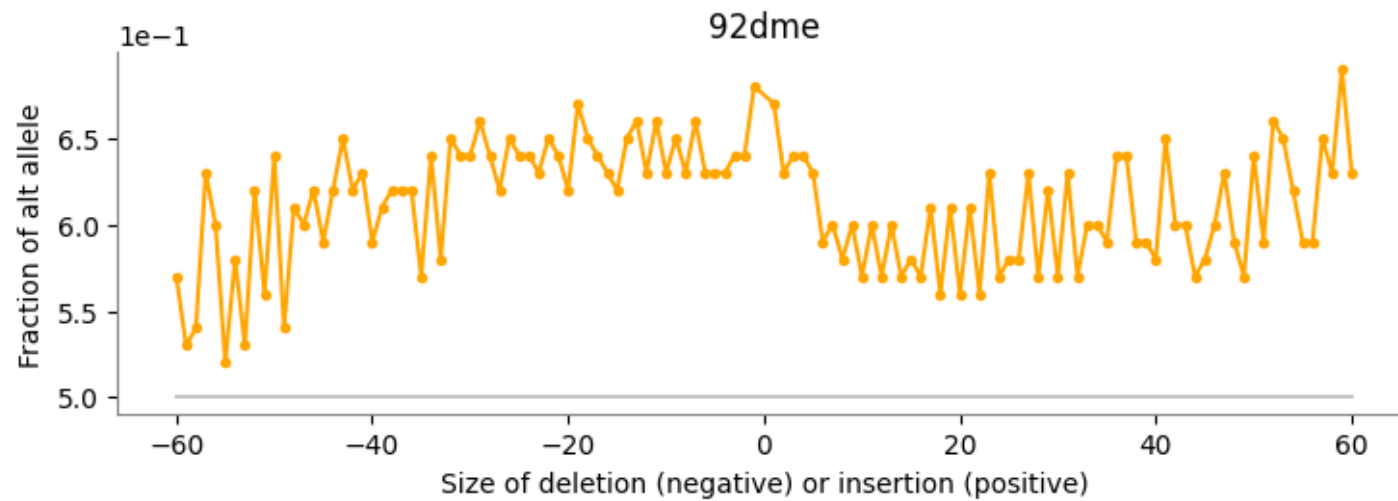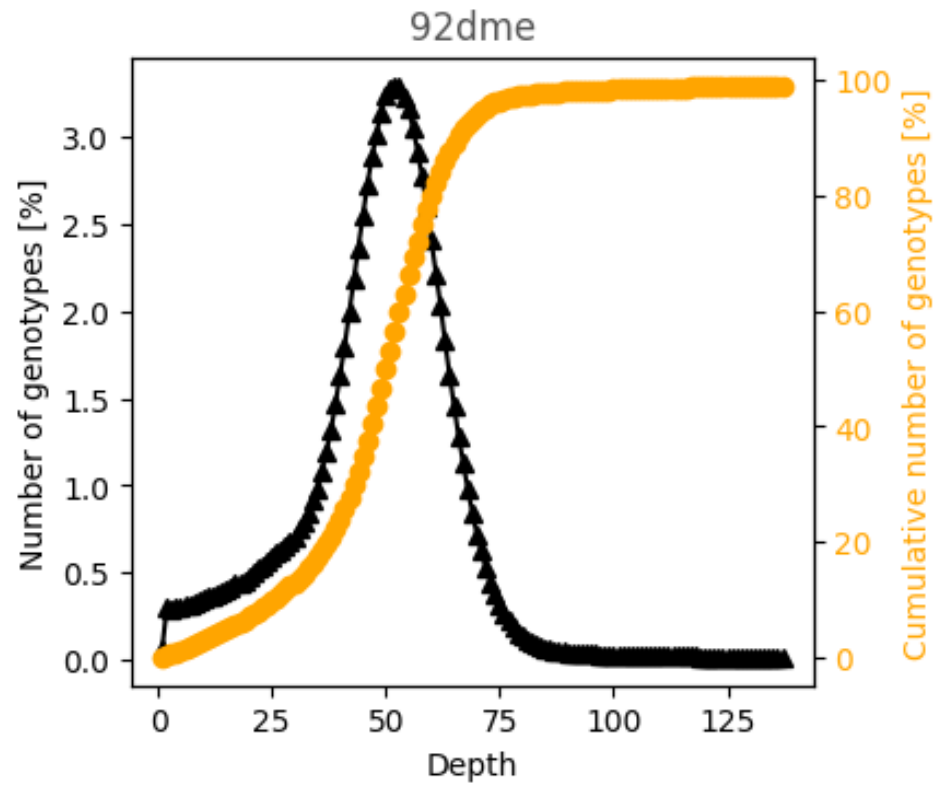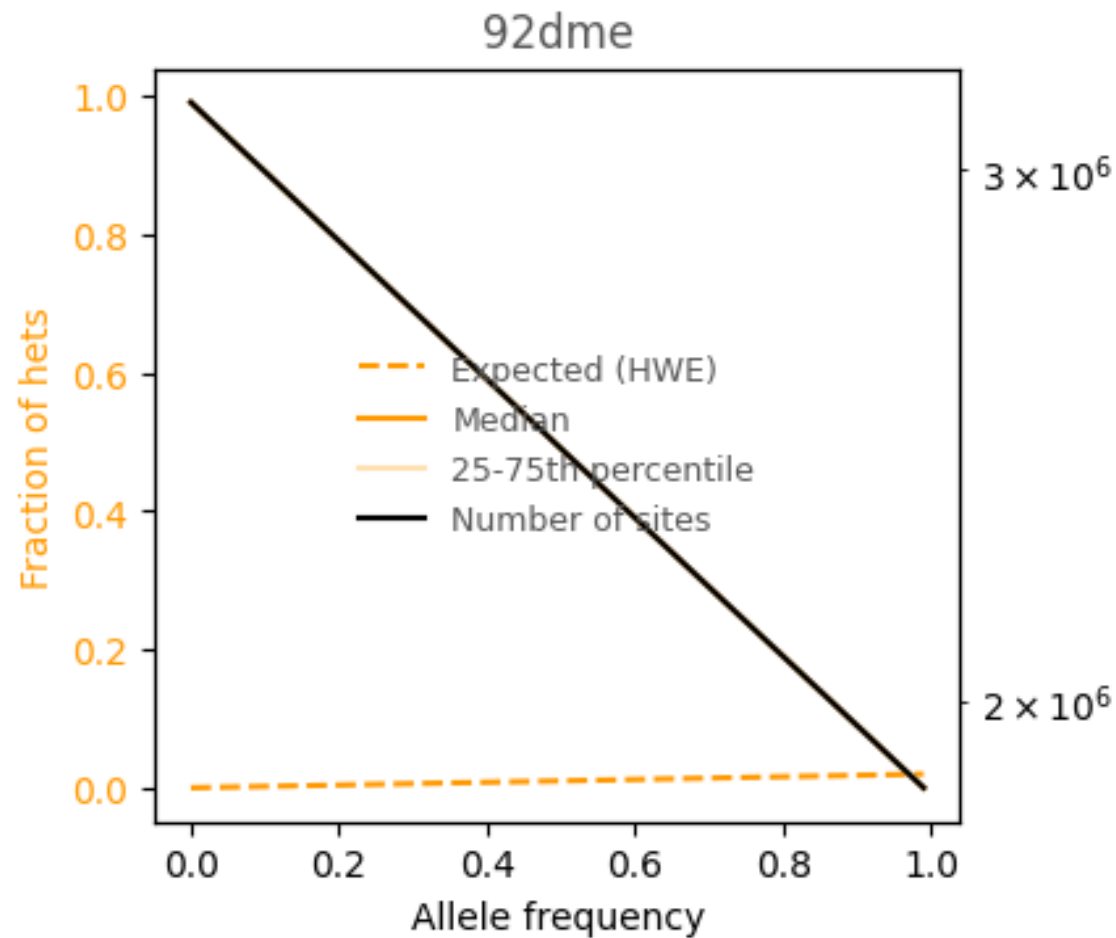# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types