Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
89ofo	4,079,128	1.93	1.94	943,448	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
89ofo	62.2%	1.90	66.6%	96,105	2,003

- 89ofo .. `/ngc/projects2/gm/data/archive/2022/variants/snv/89oforecm-110302560289-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM02259-221109_A01411_BH3NLCDSX5-EXT_LAB` `KA_NGCWGS-NGCWGS06144_19RKG020136_snv_germline_raw.haplotype_caller.vcf.gz`
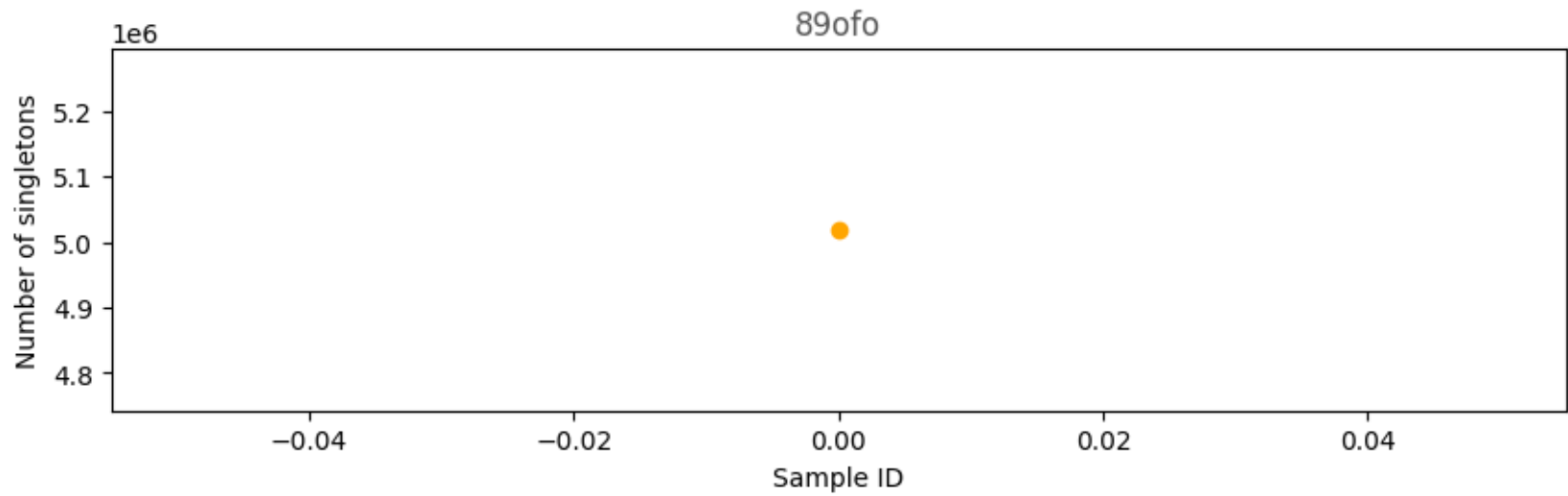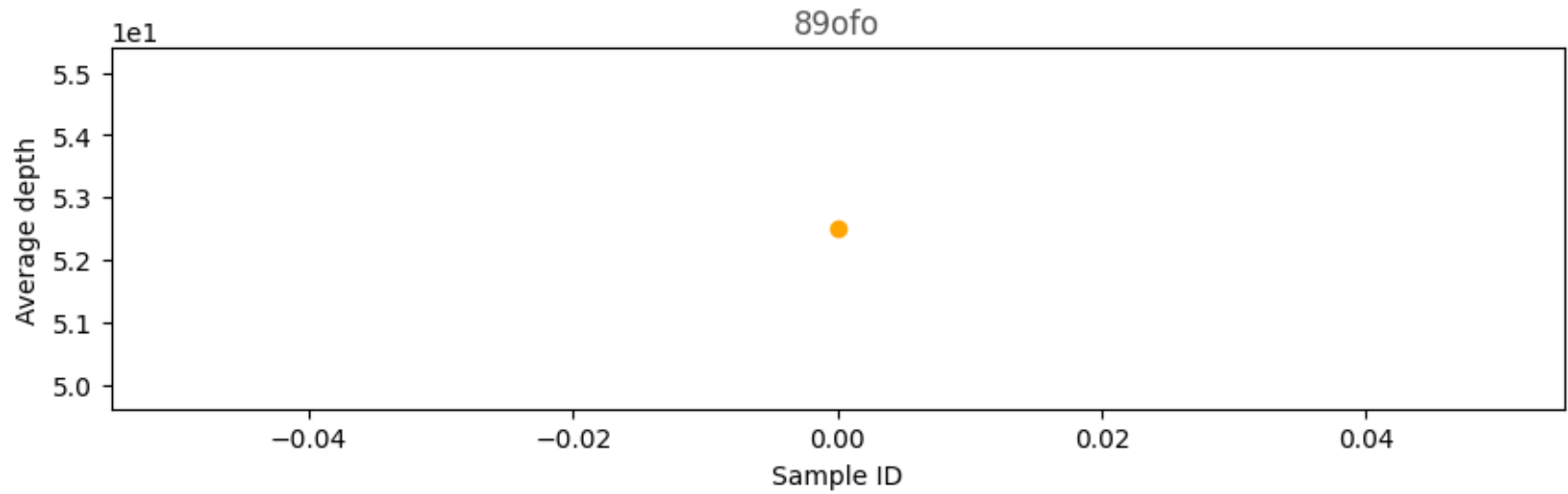
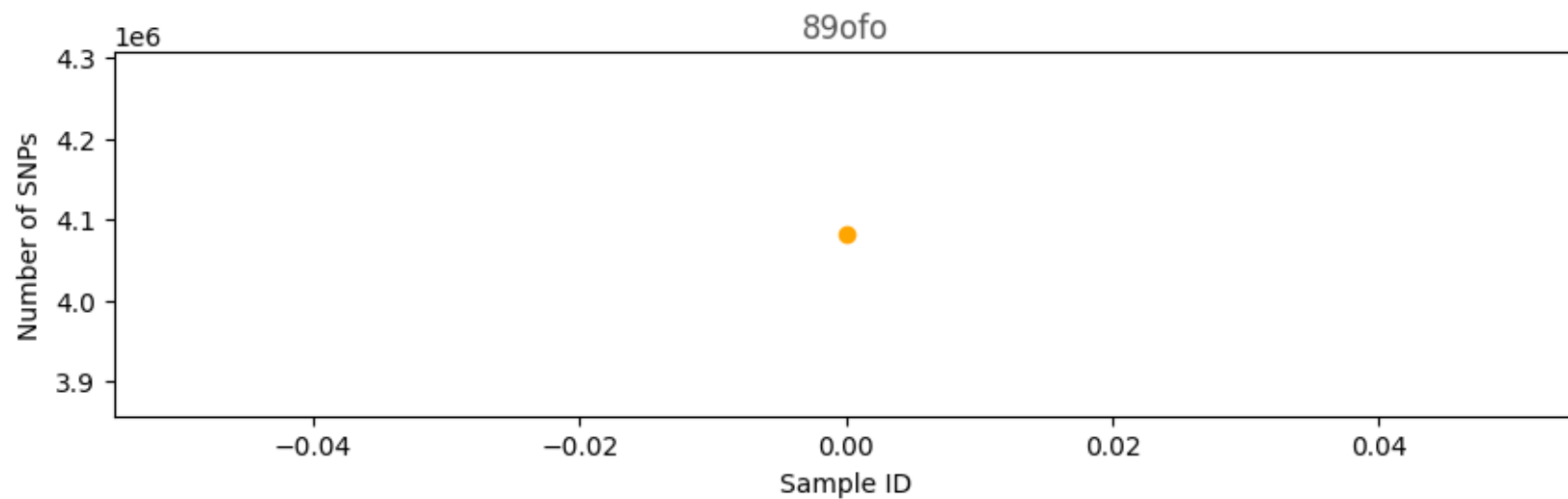# Ts/Tv by sample
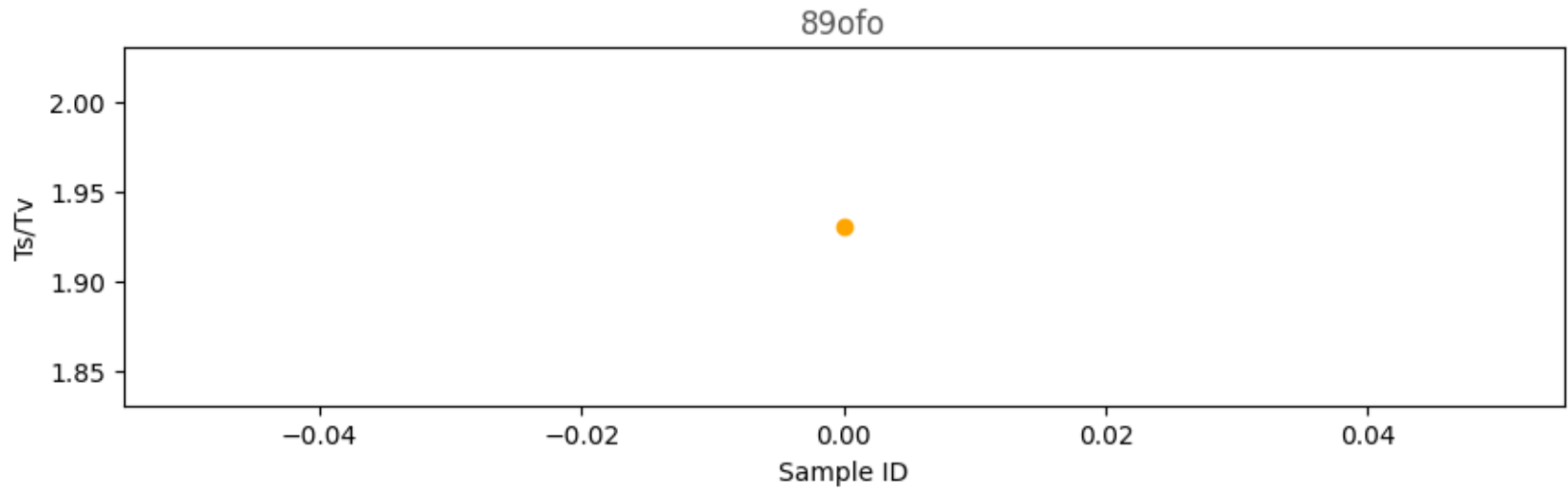
Hets vs non-ref Homs by sample
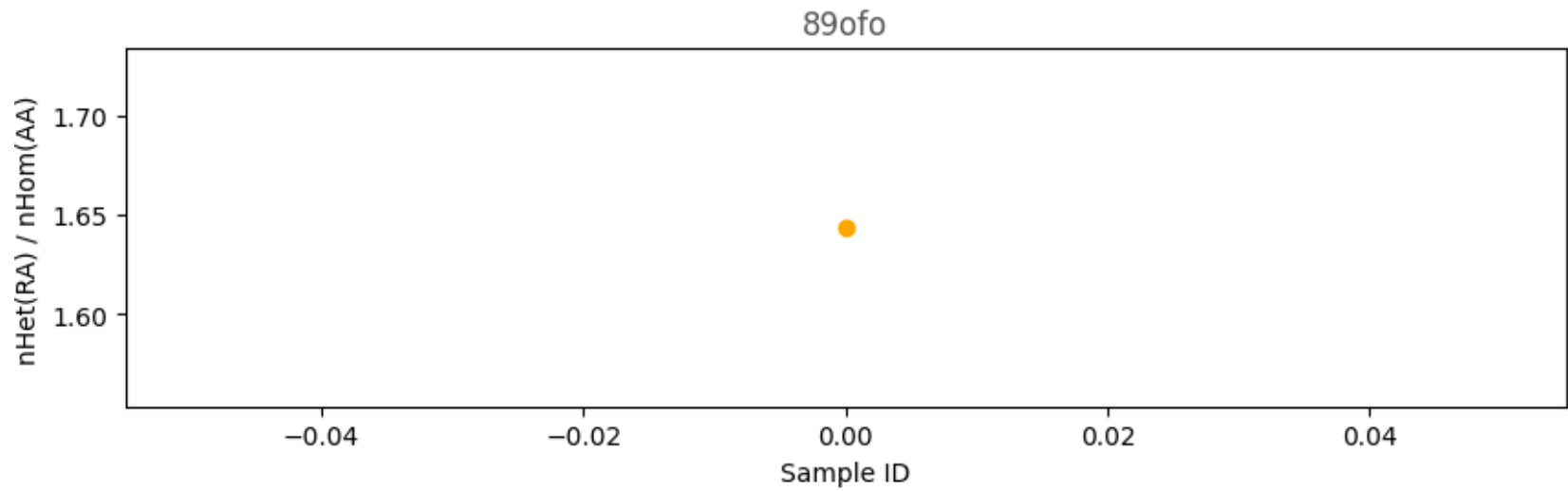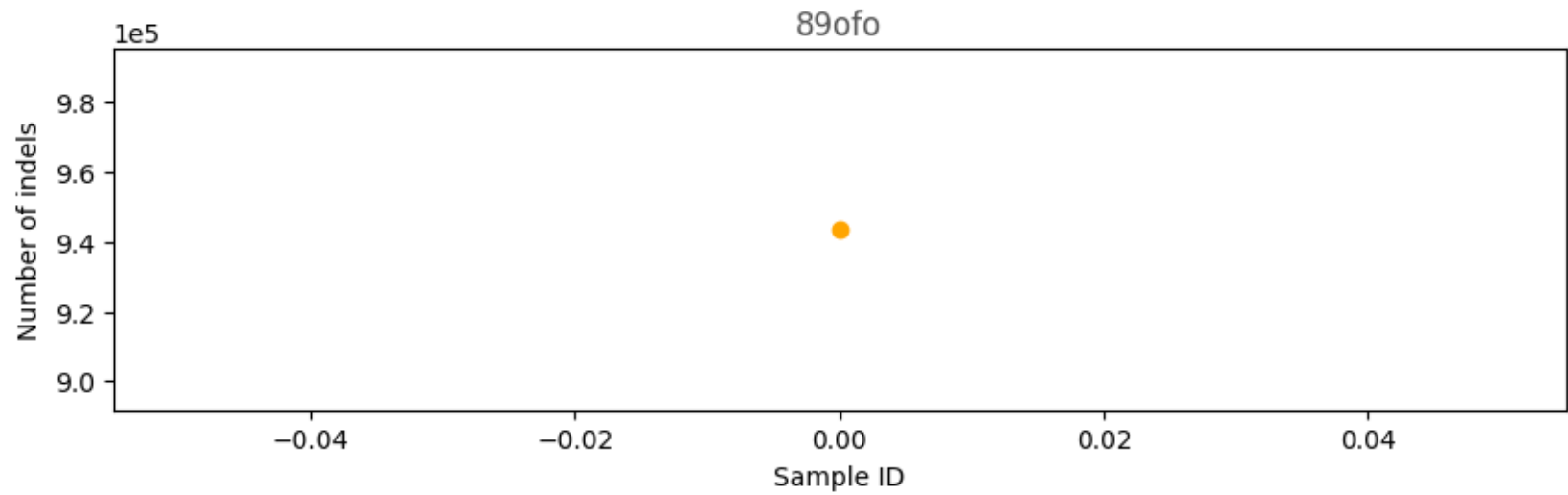
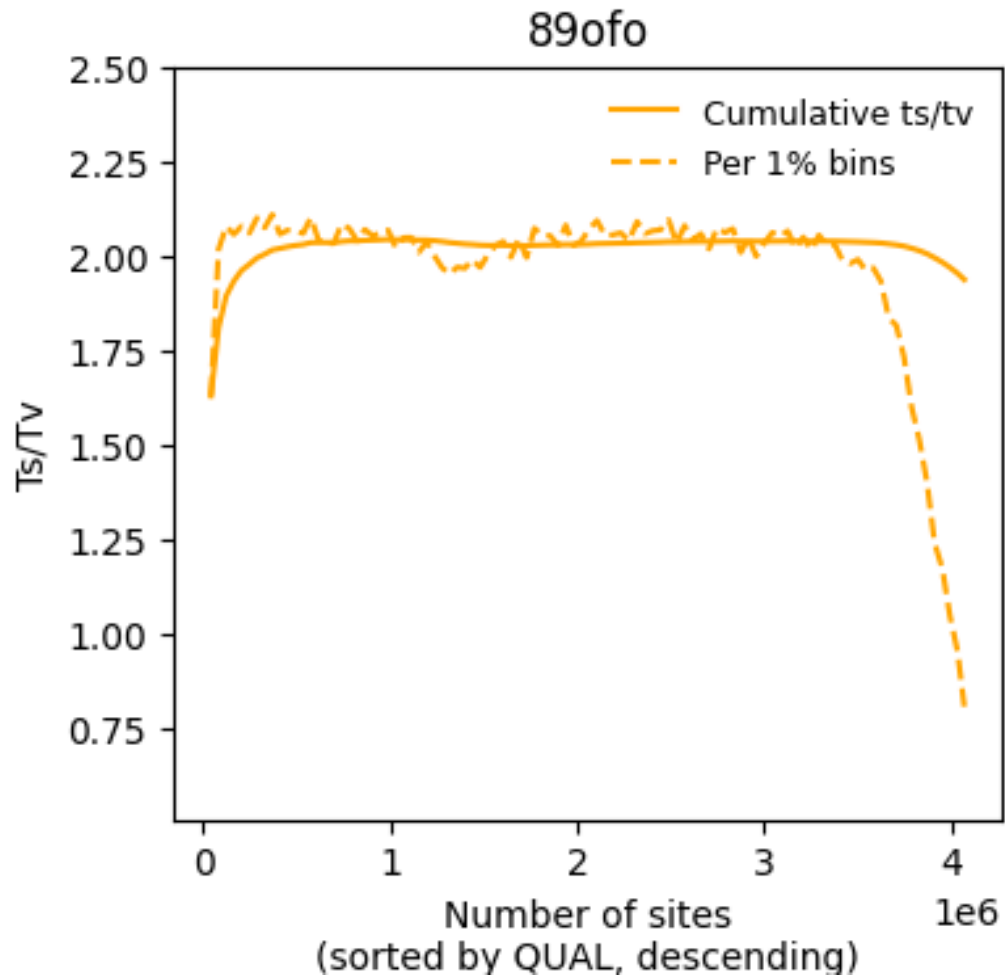# Singletons by sample (hets and homs)
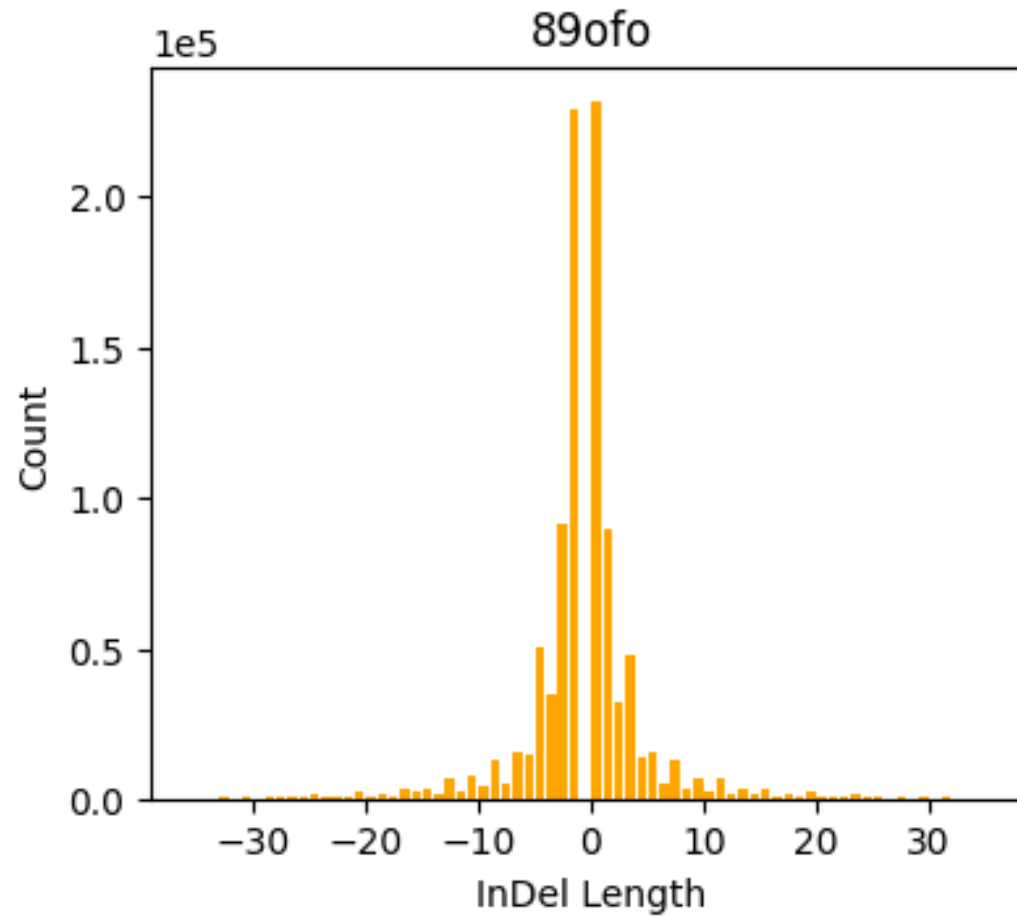
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

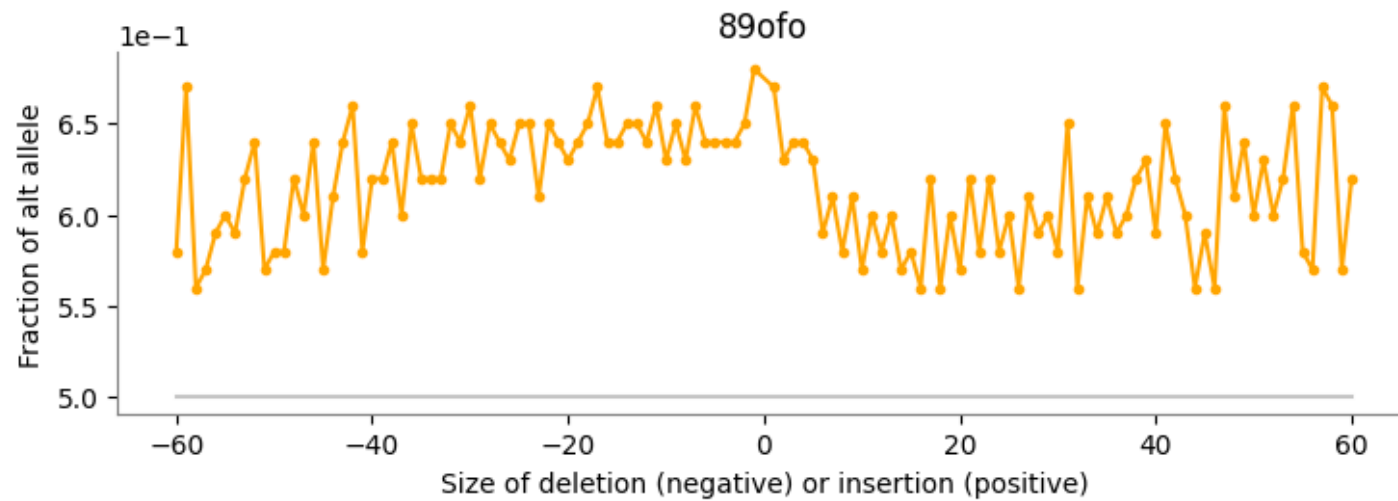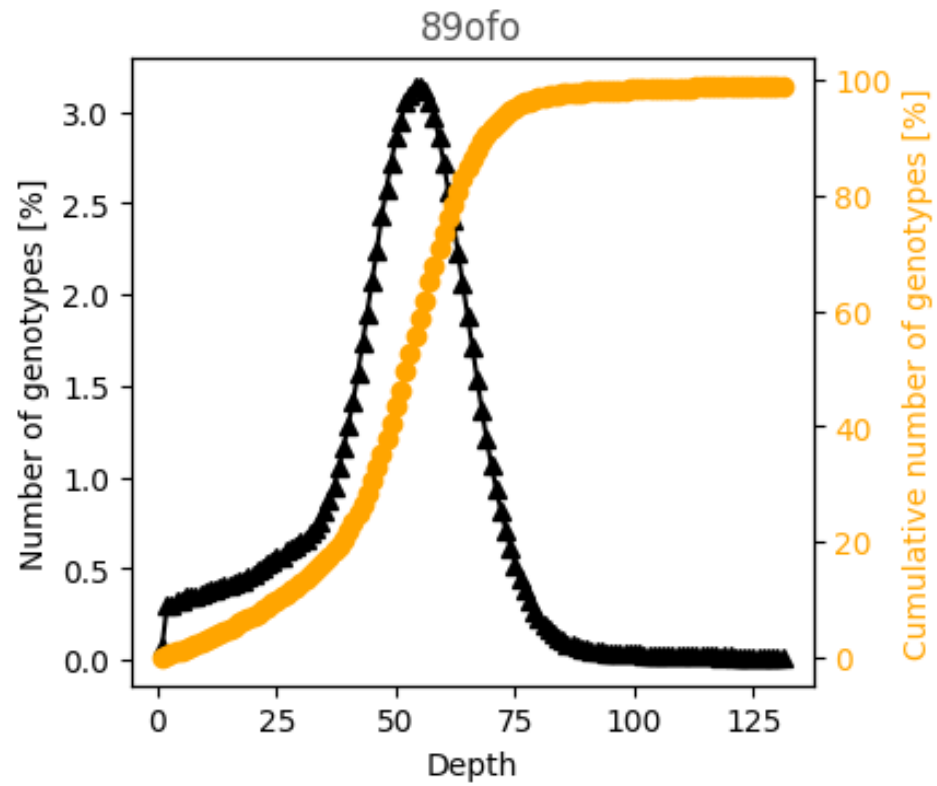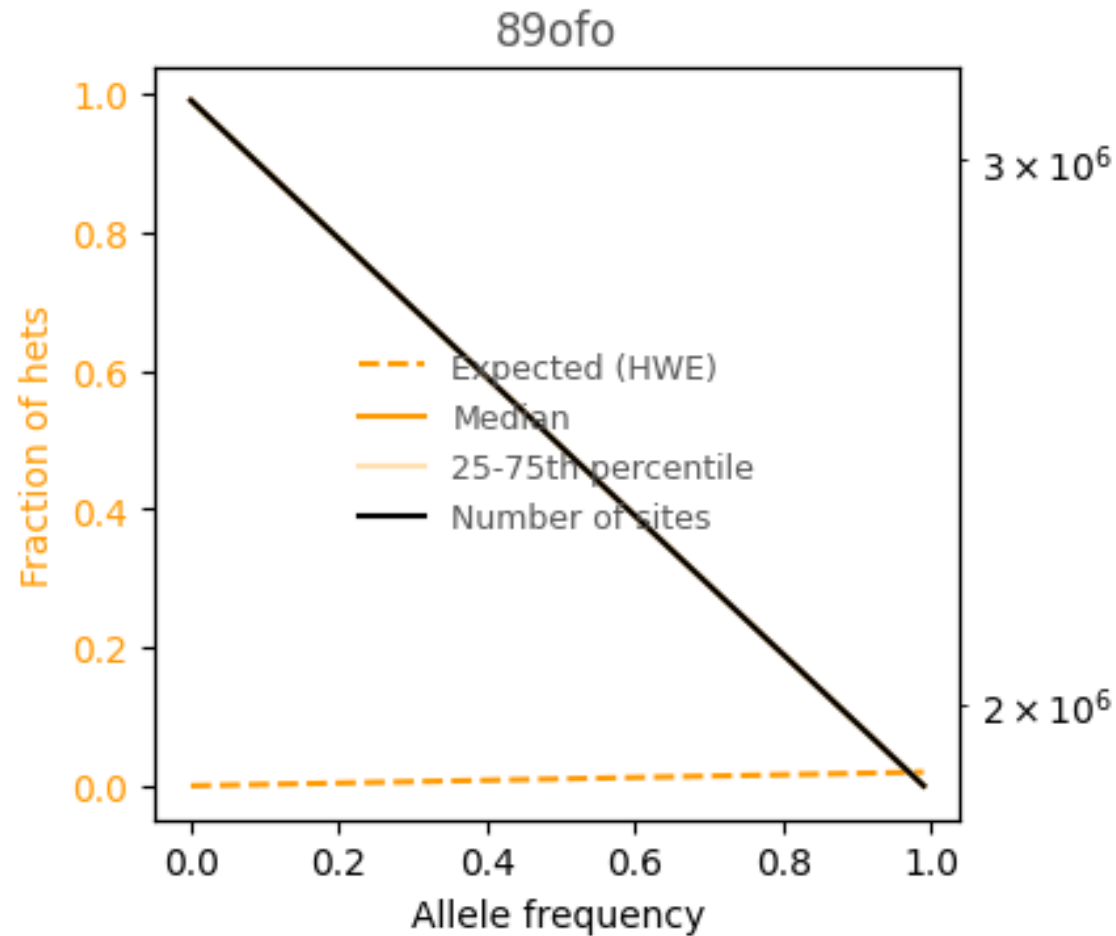# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types