Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
86ior	4,116,713	1.93	1.93	962,235	–	0	0
						* frameshift ratio: out/(out+in)	

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
86ior	63.1%	1.90	68.5%	105,983	2,282

- 86ior .. `/ngc/projects2/gm/data/archive/2022/variants/snv/86ioreinf-110296384686-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-221012_A01176_AH2CWMDSX5-RHGM_LABKA_WGS-`
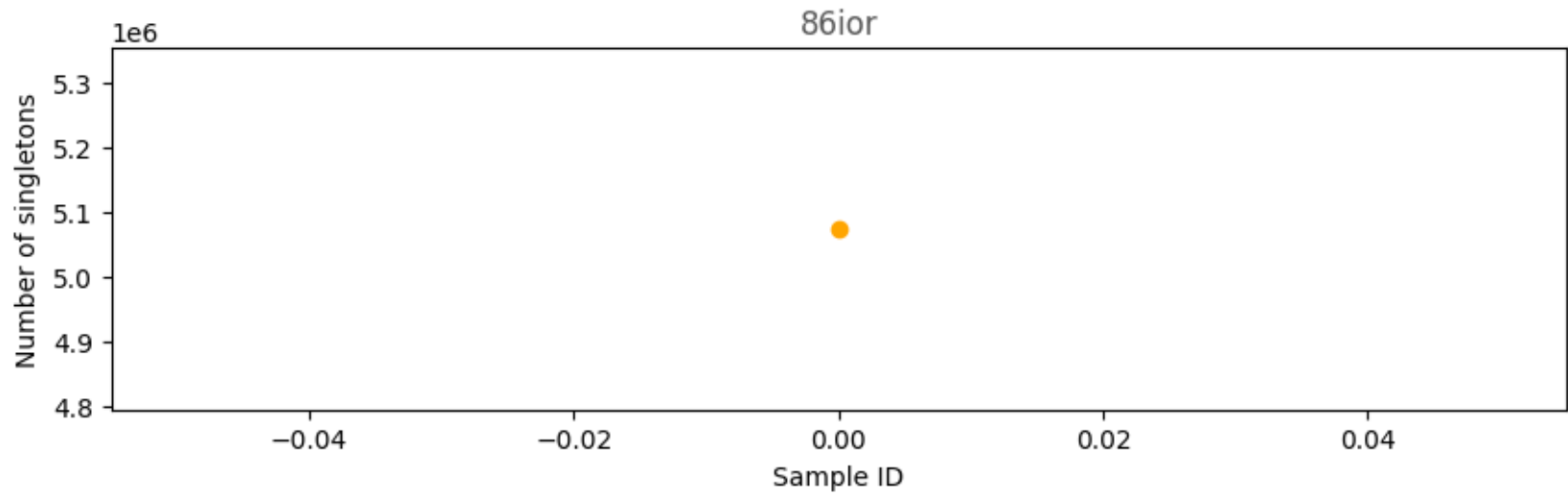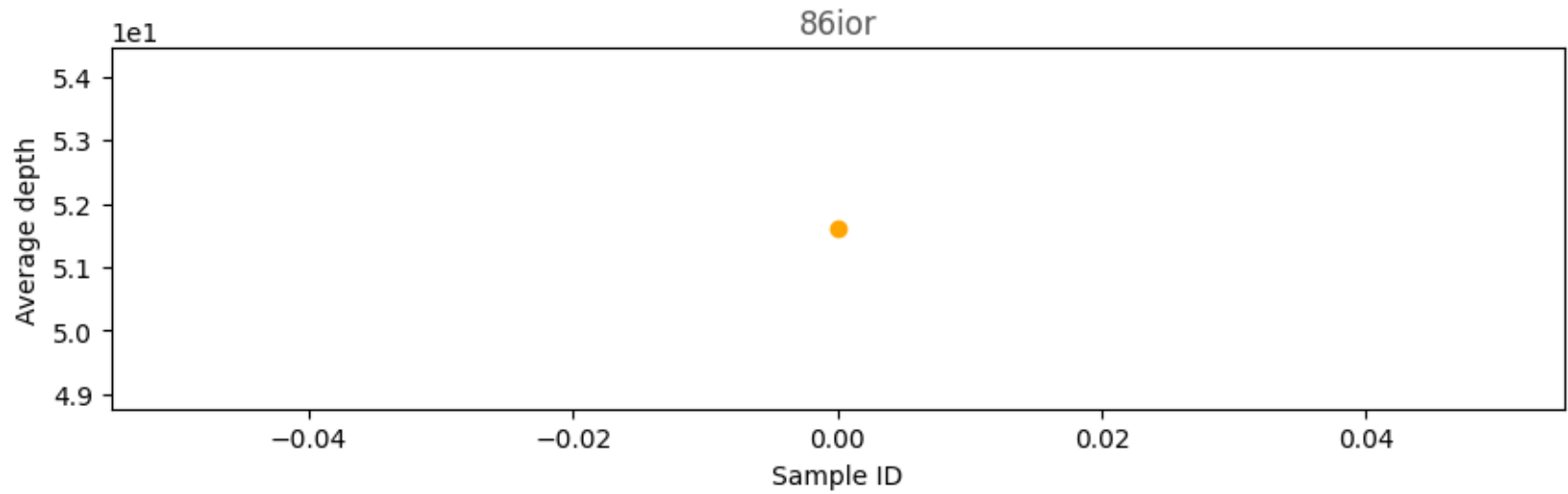 `WGS05819_snv_germline_raw.haplotype_caller.vcf.gz`
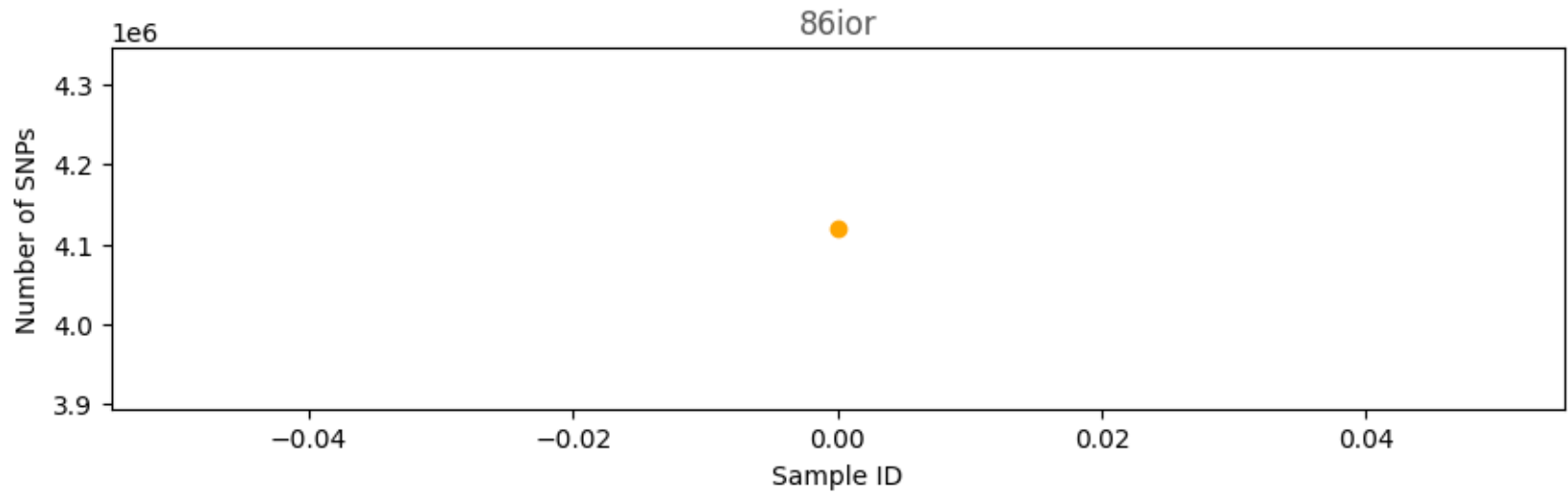
Ts/Tv by sample
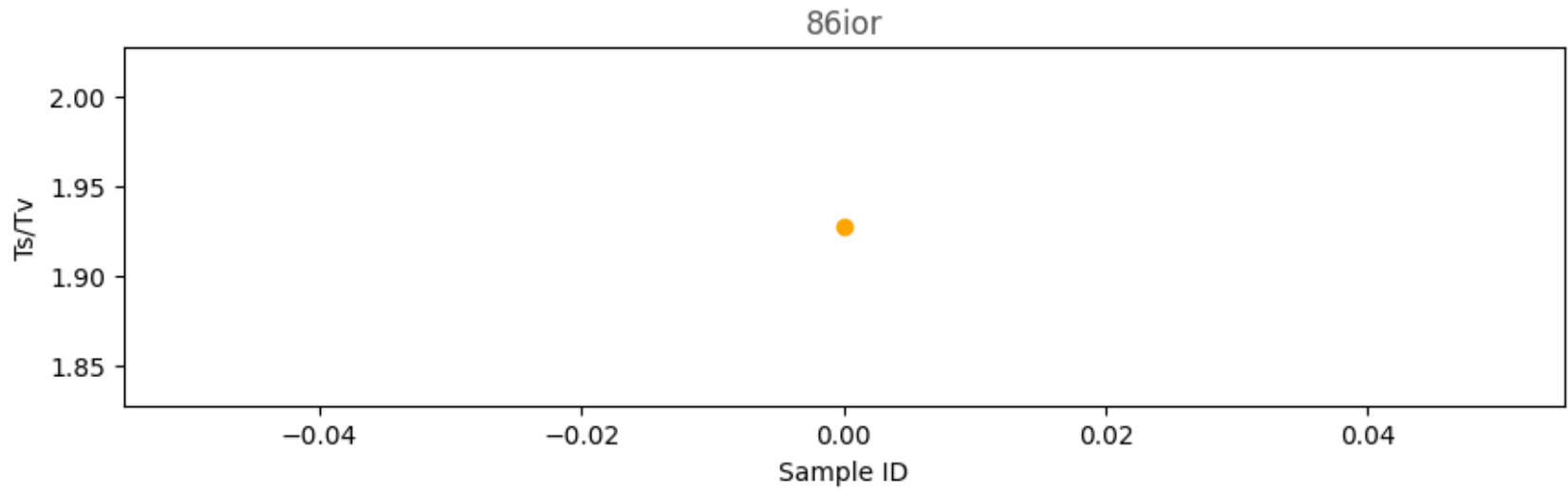
Hets vs non-ref Homs by sample
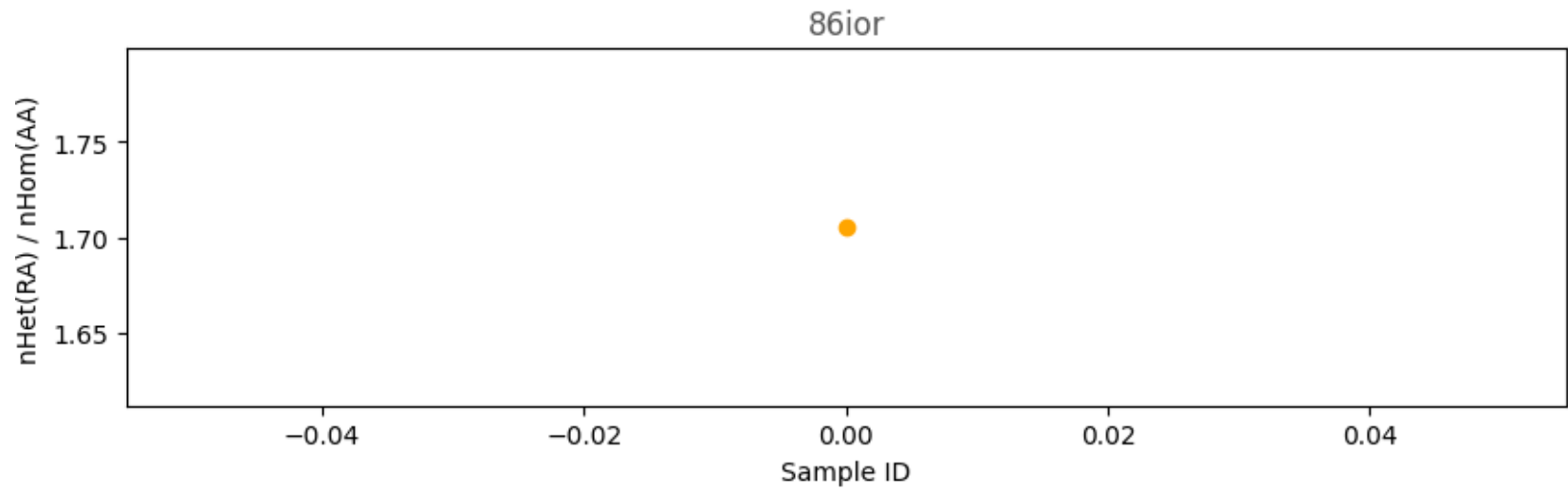
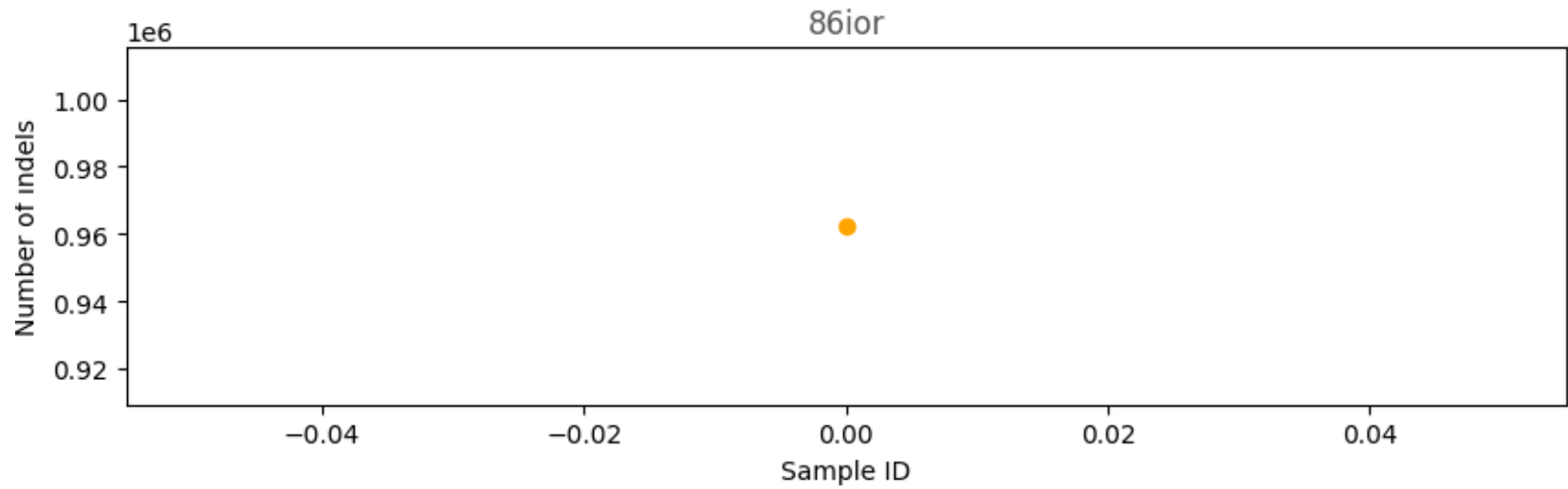# Singletons by sample (hets and homs)
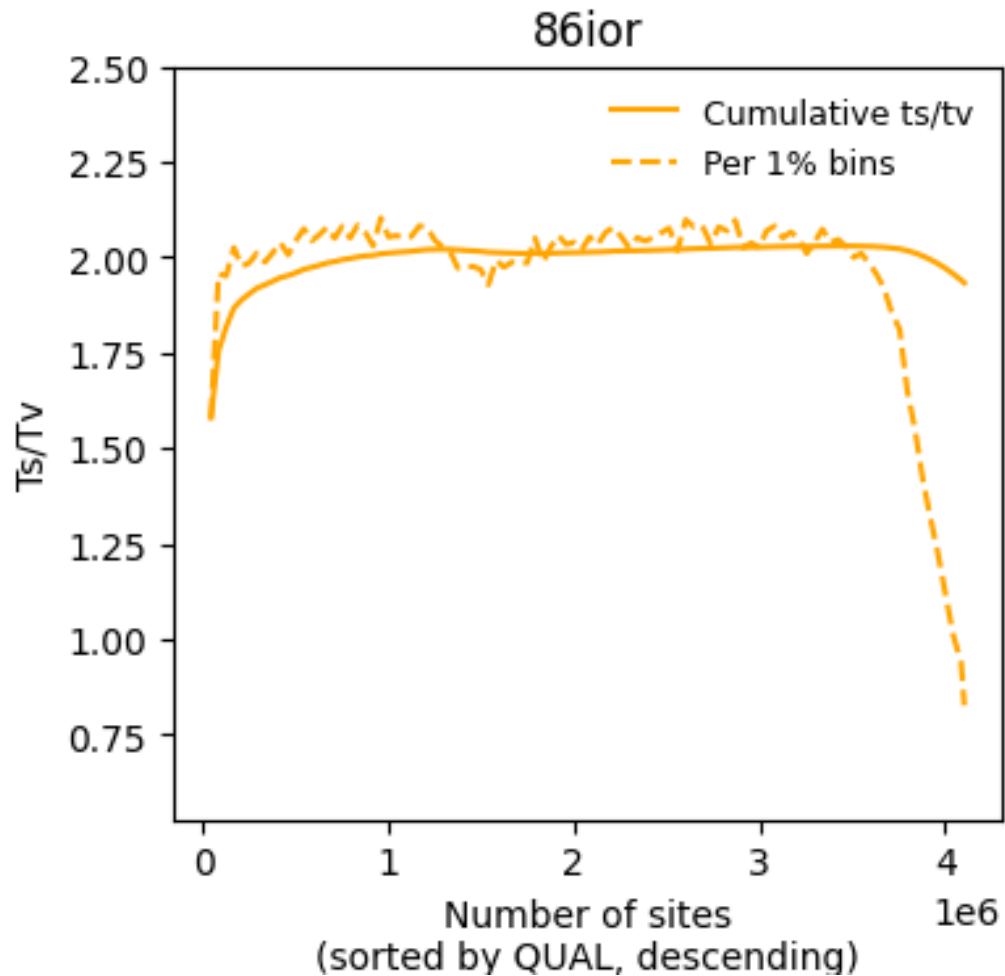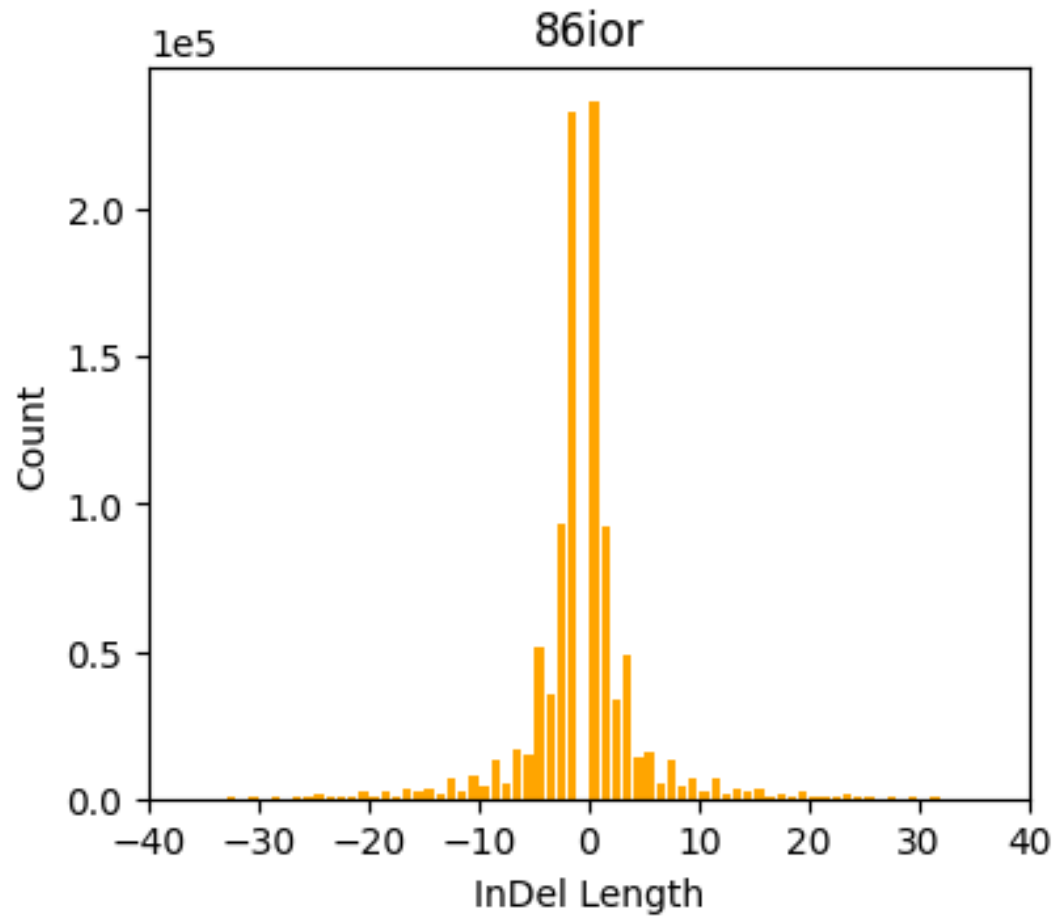
Average depth by sample

Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

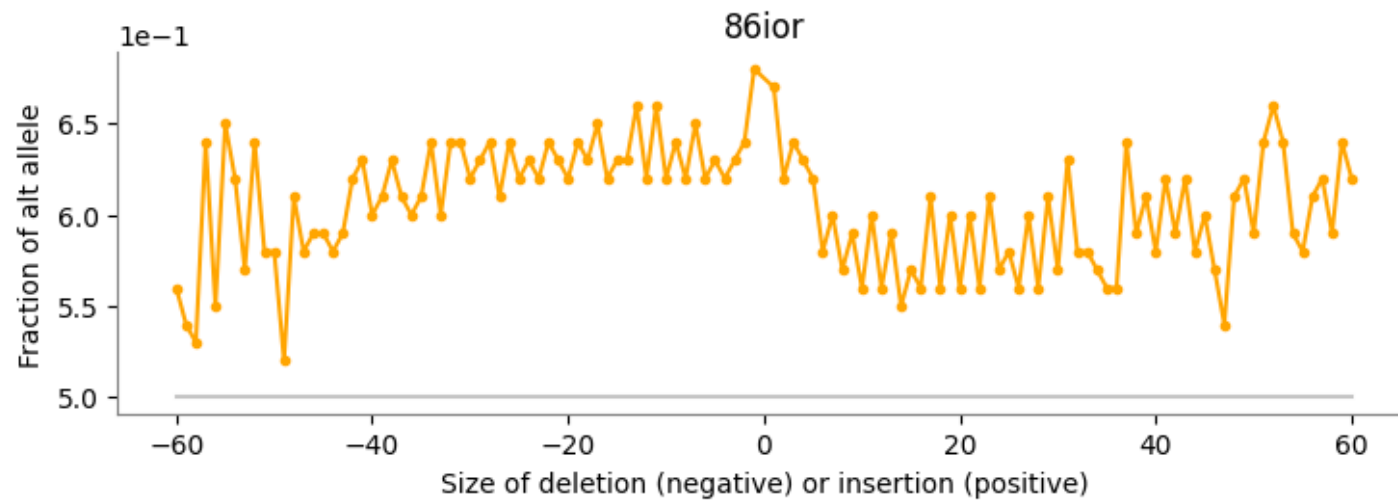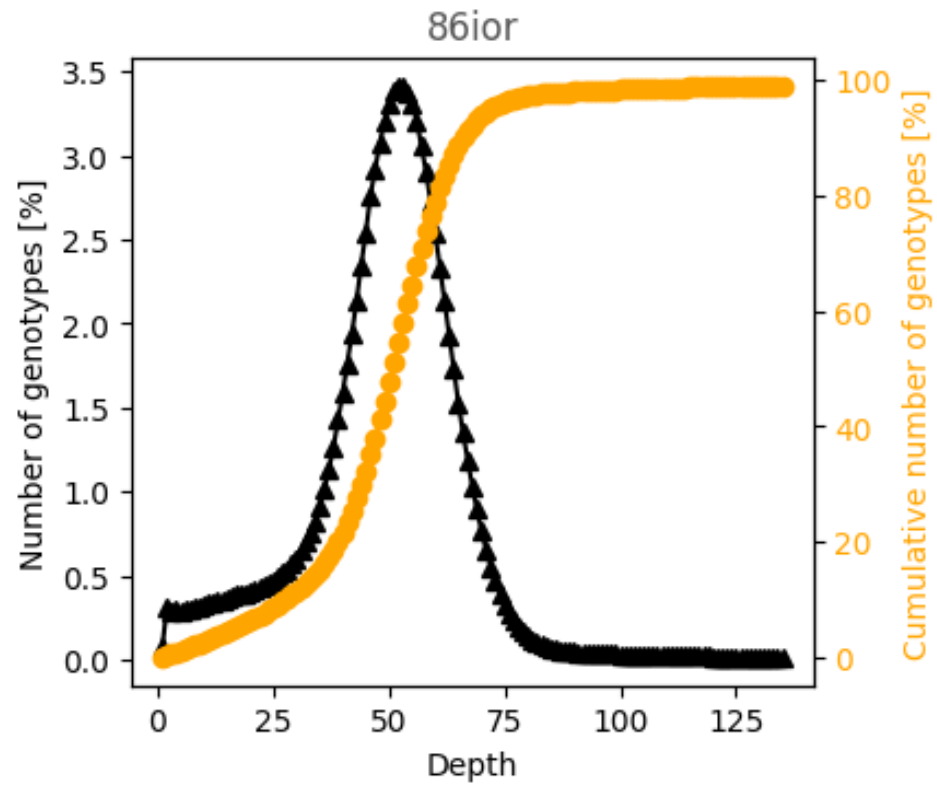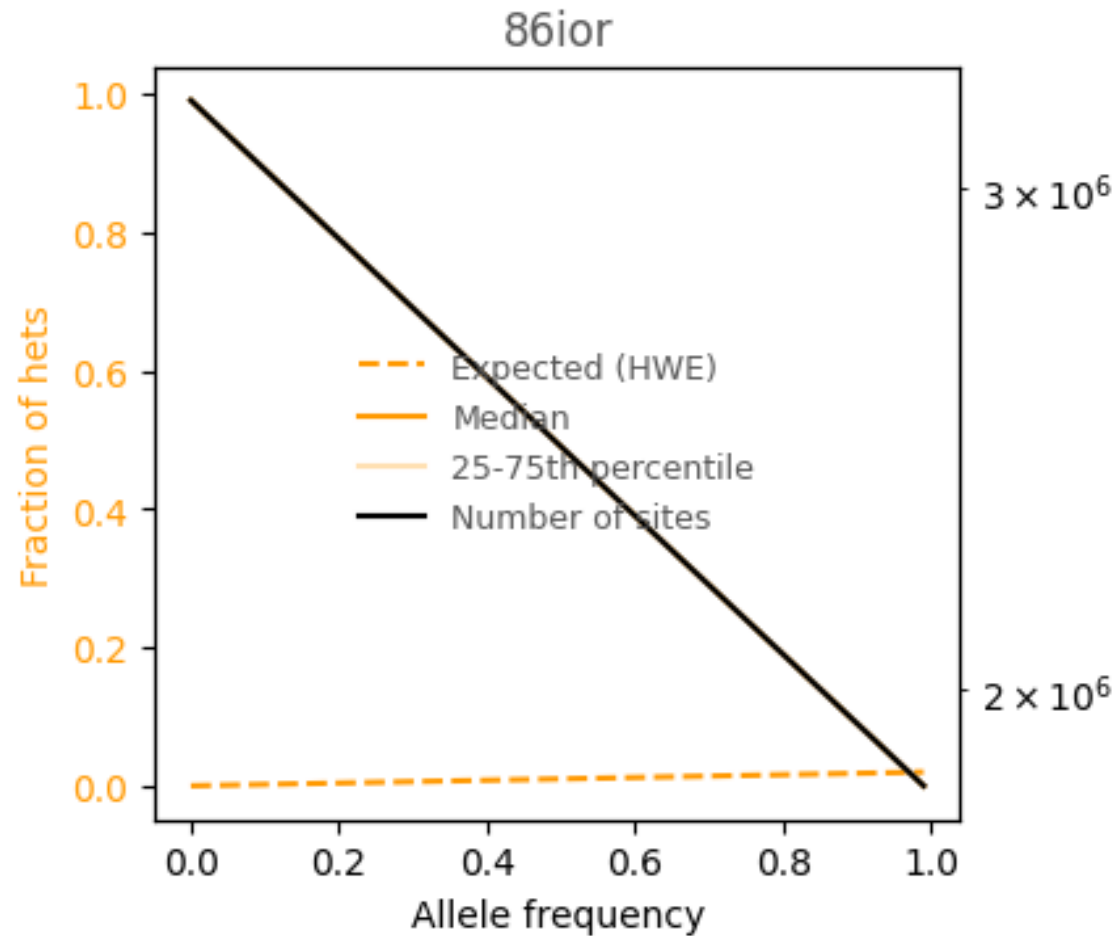# Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types