Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
76ore	4,070,272	1.94	1.95	939,133	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
76ore	63.0%	1.91	67.2%	94,629	2,139

- 76ore .. `/ngc/projects2/gm/data/archive/2022/variants/snv/76orenfdf-103745707719-Normal_B`
    `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00164-210608_A01176_AHCYWGDSX2-EXT_LAB`
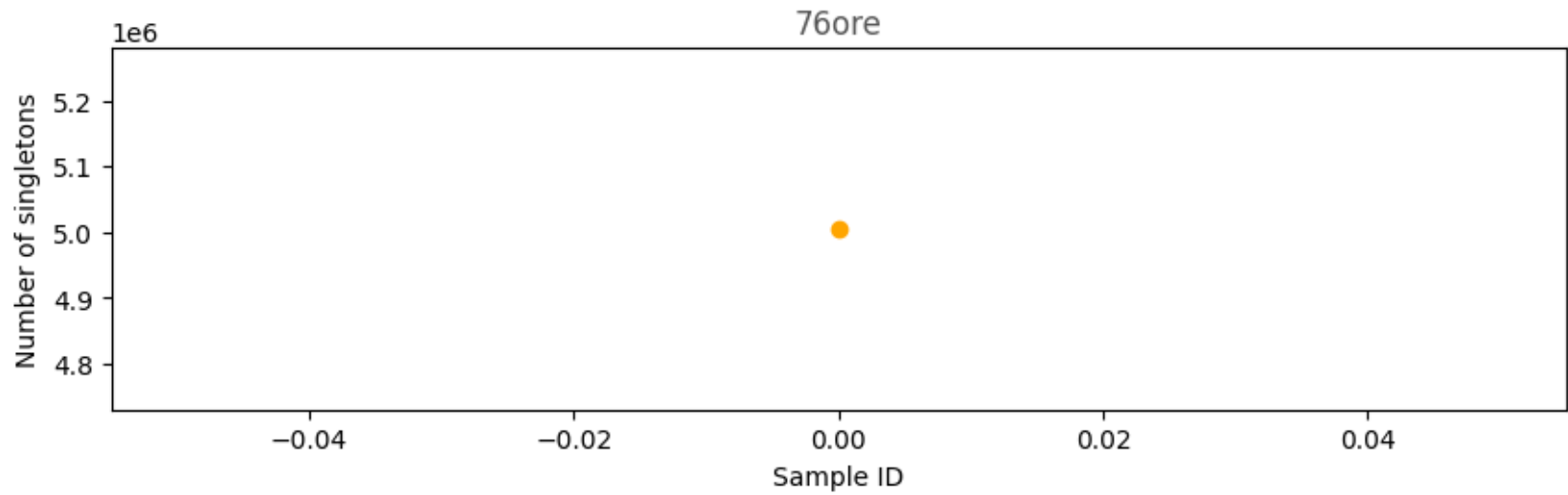   `KA_NGCWGS-NGCWGS00079_7634u2df8_snv_germline_raw.haplotype_caller.vcf.gz`
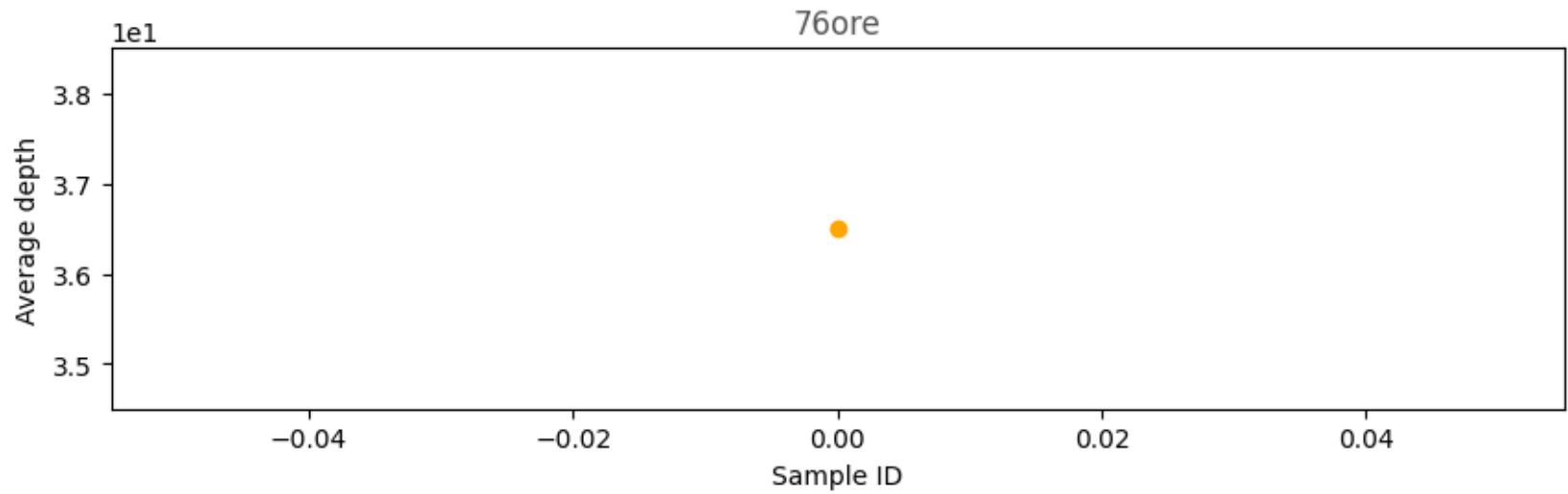
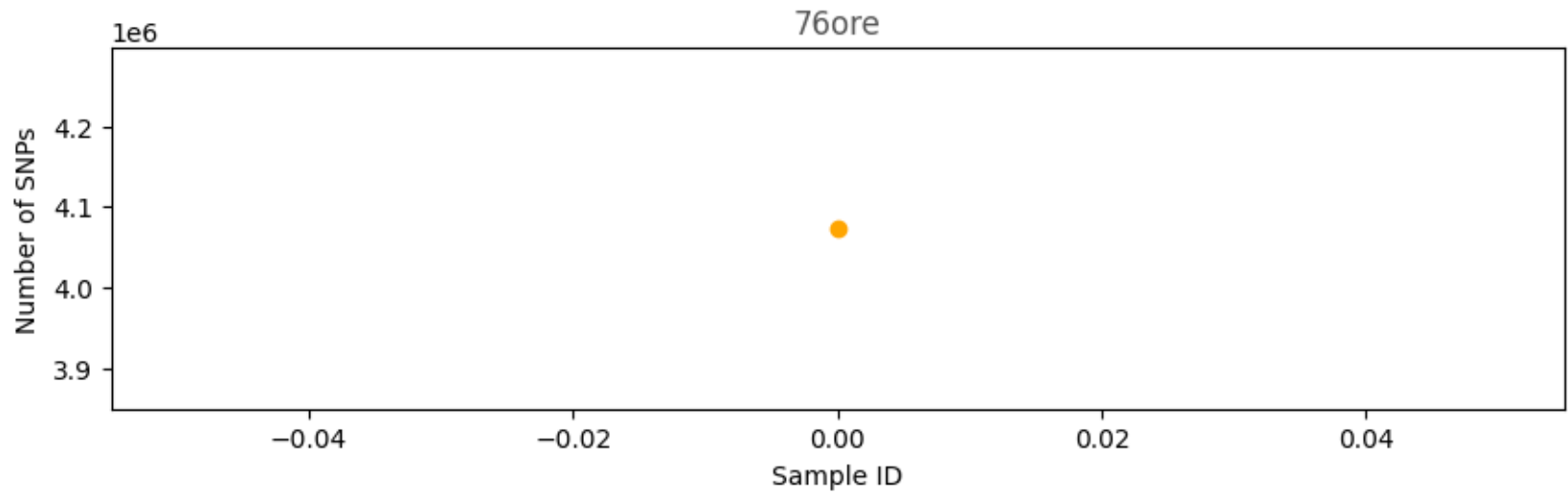# Ts/Tv by sample
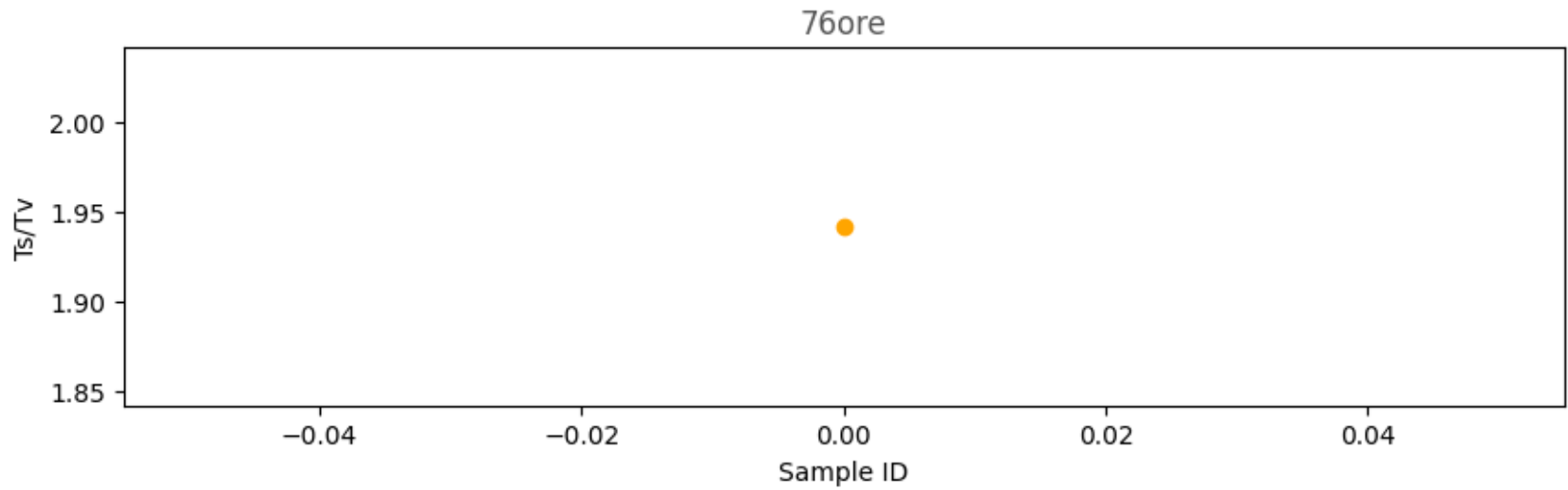
Hets vs non-ref Homs by sample
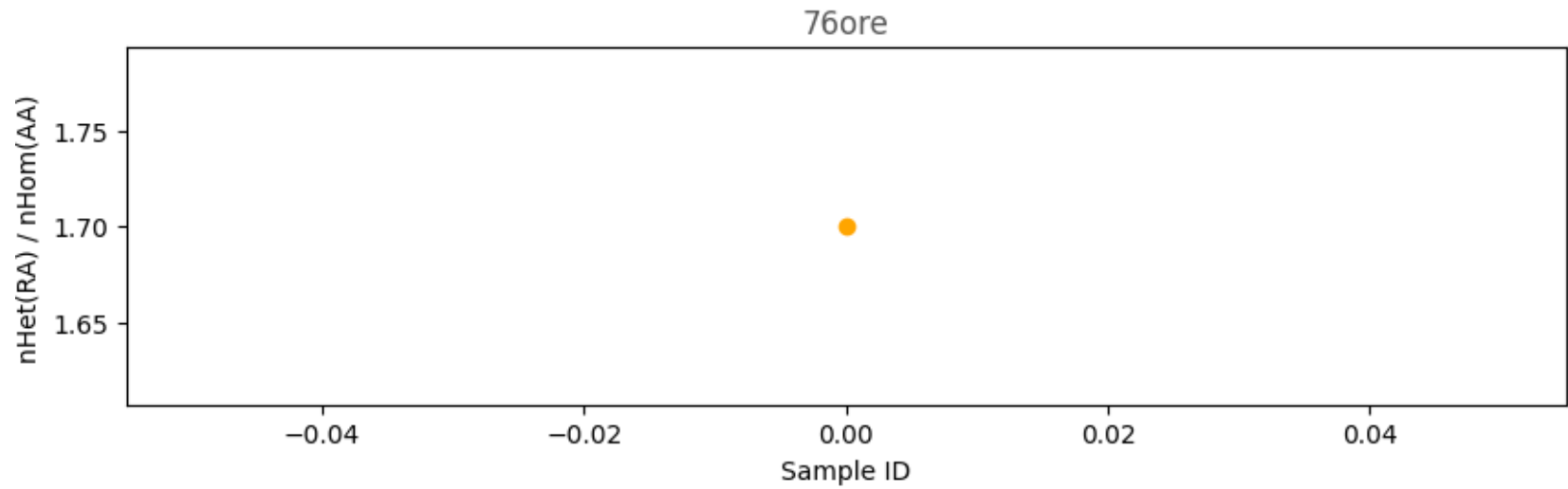
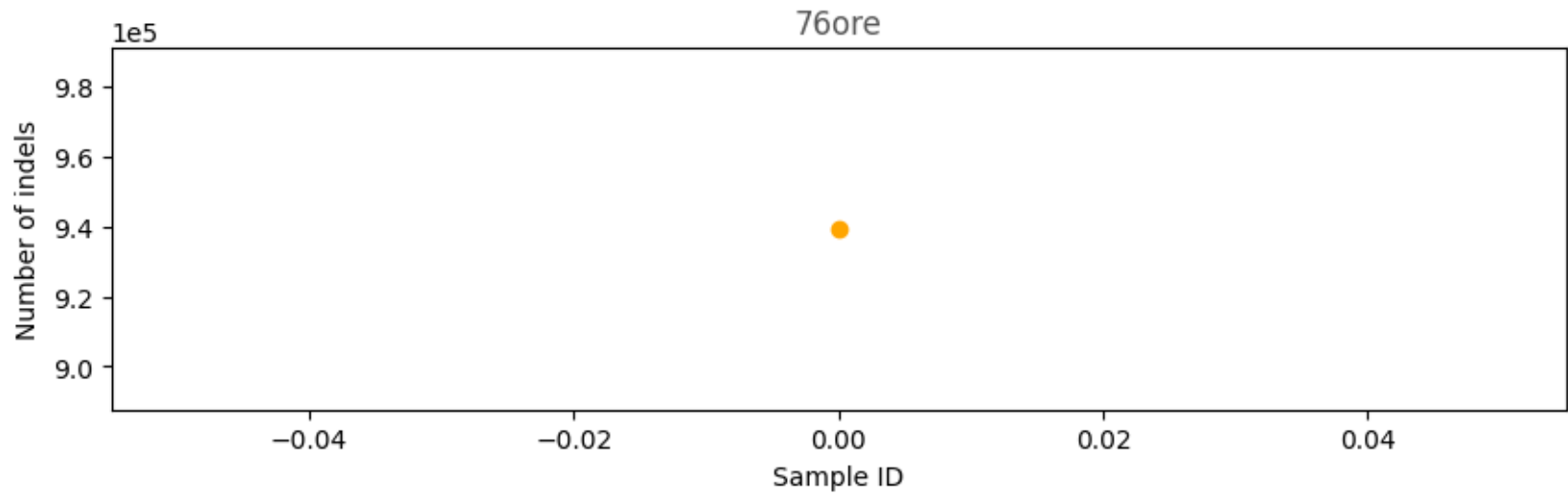# Singletons by sample (hets and homs)
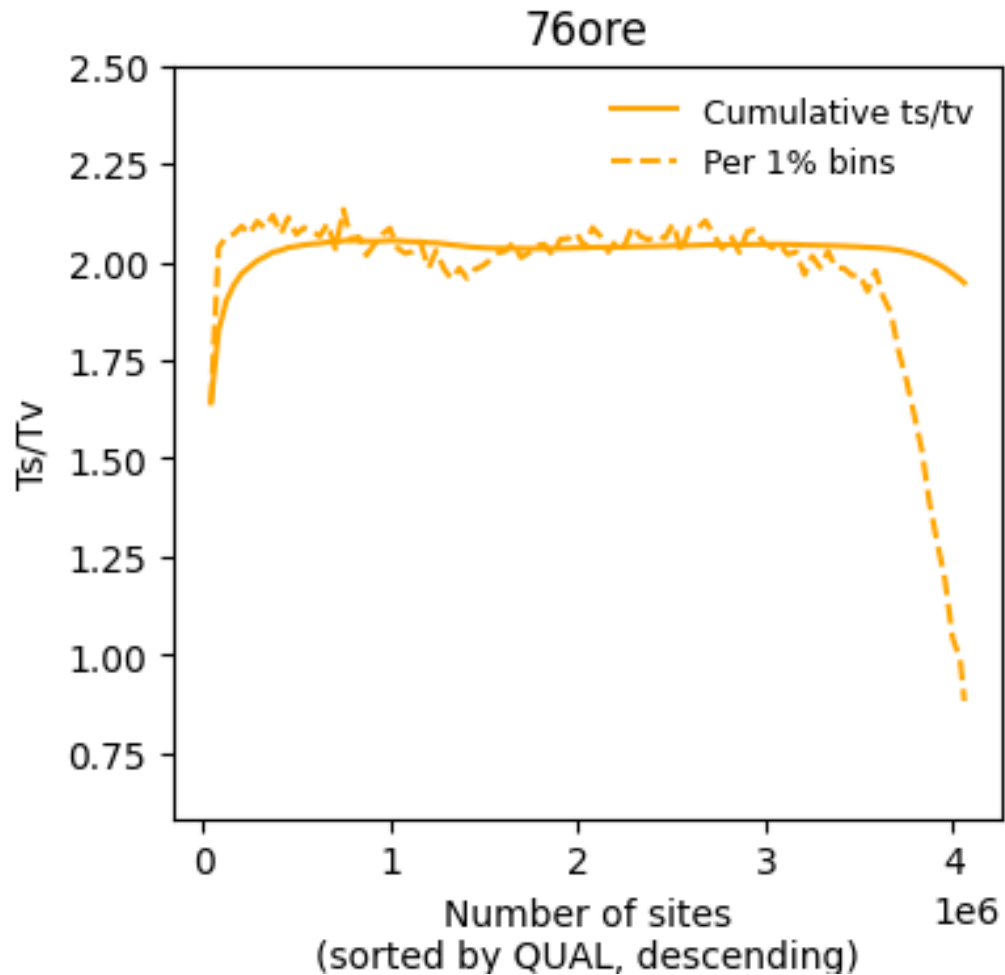

76ore
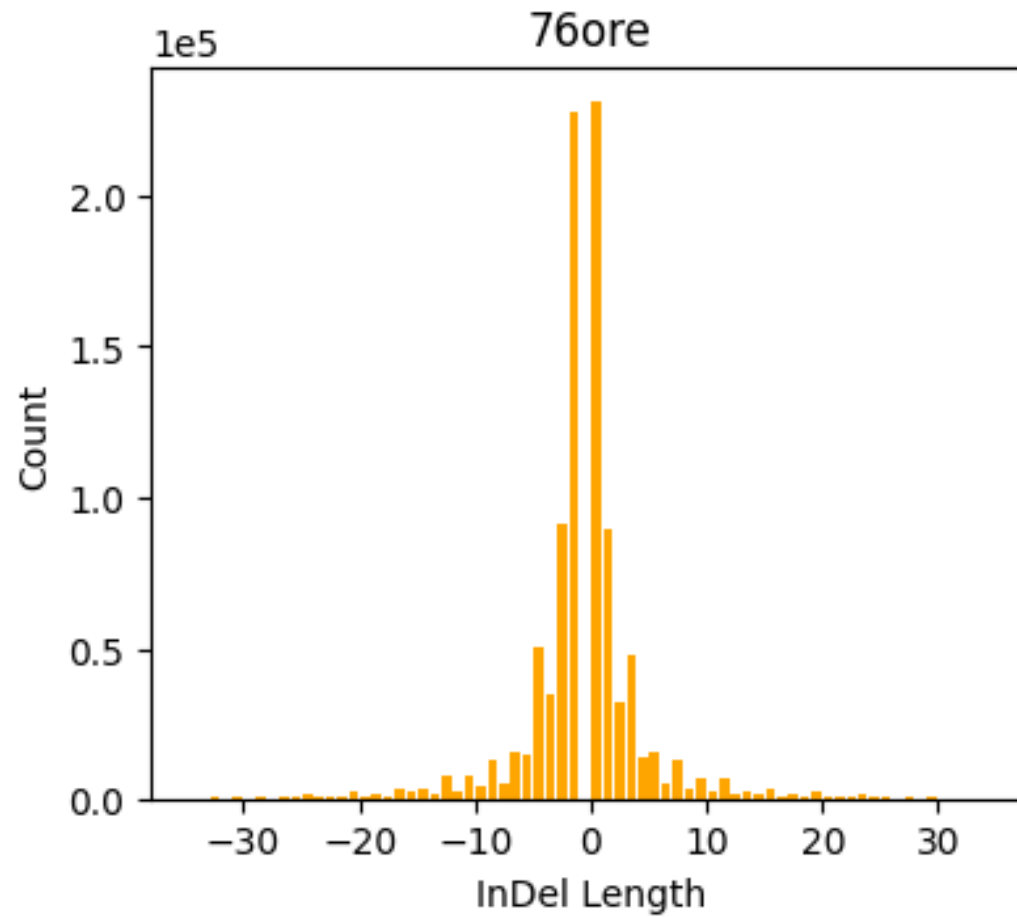
Average depth by sample
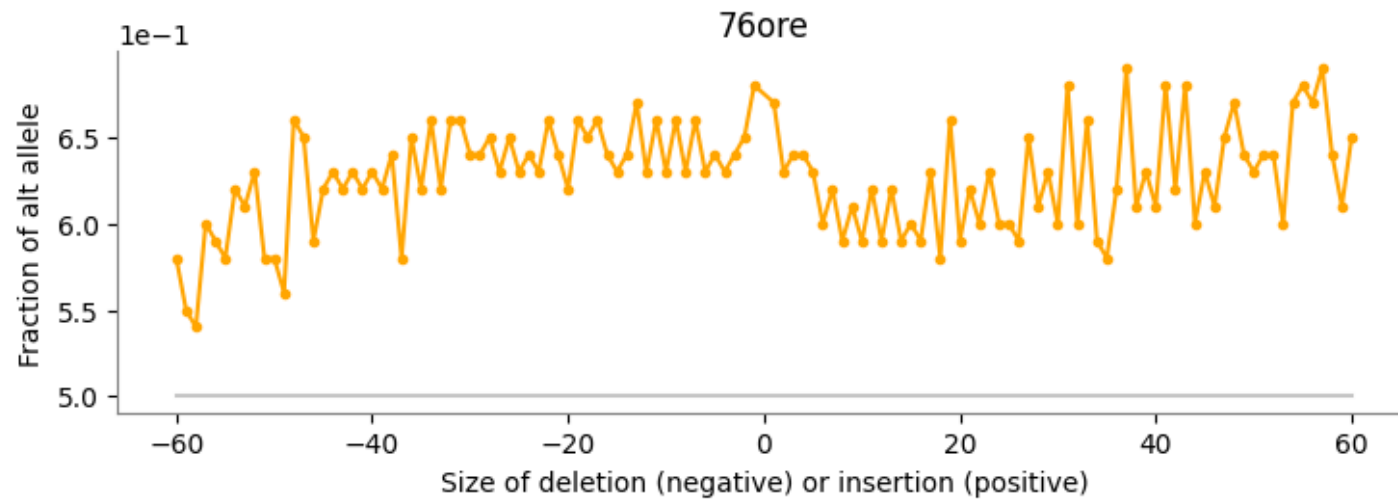
Number of SNPs by sample

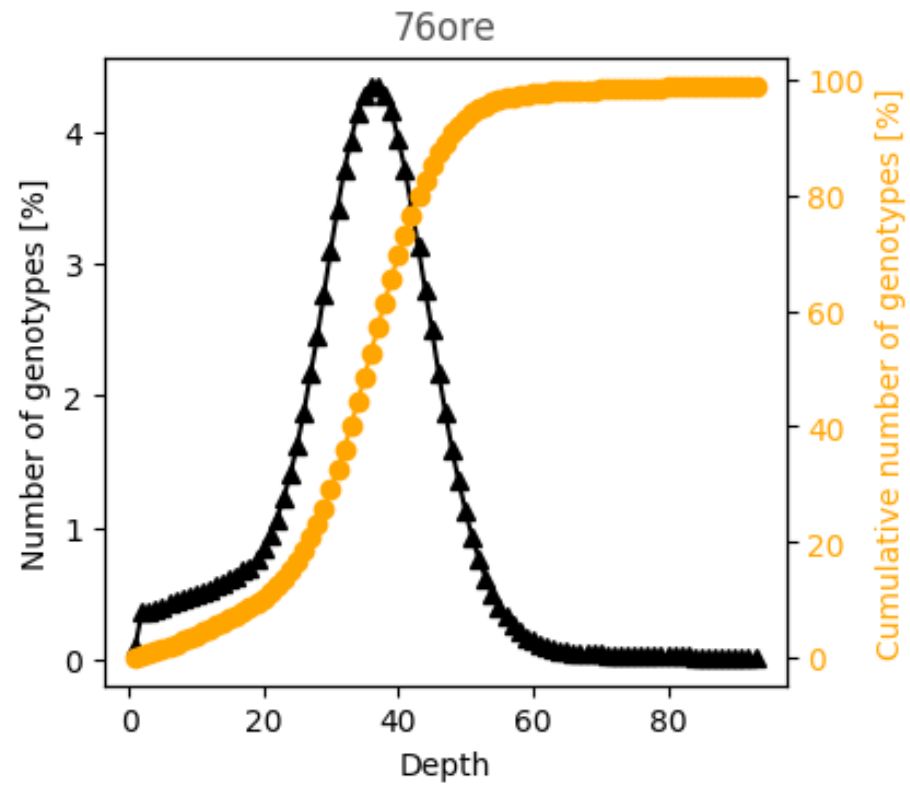# Number of indels by sample
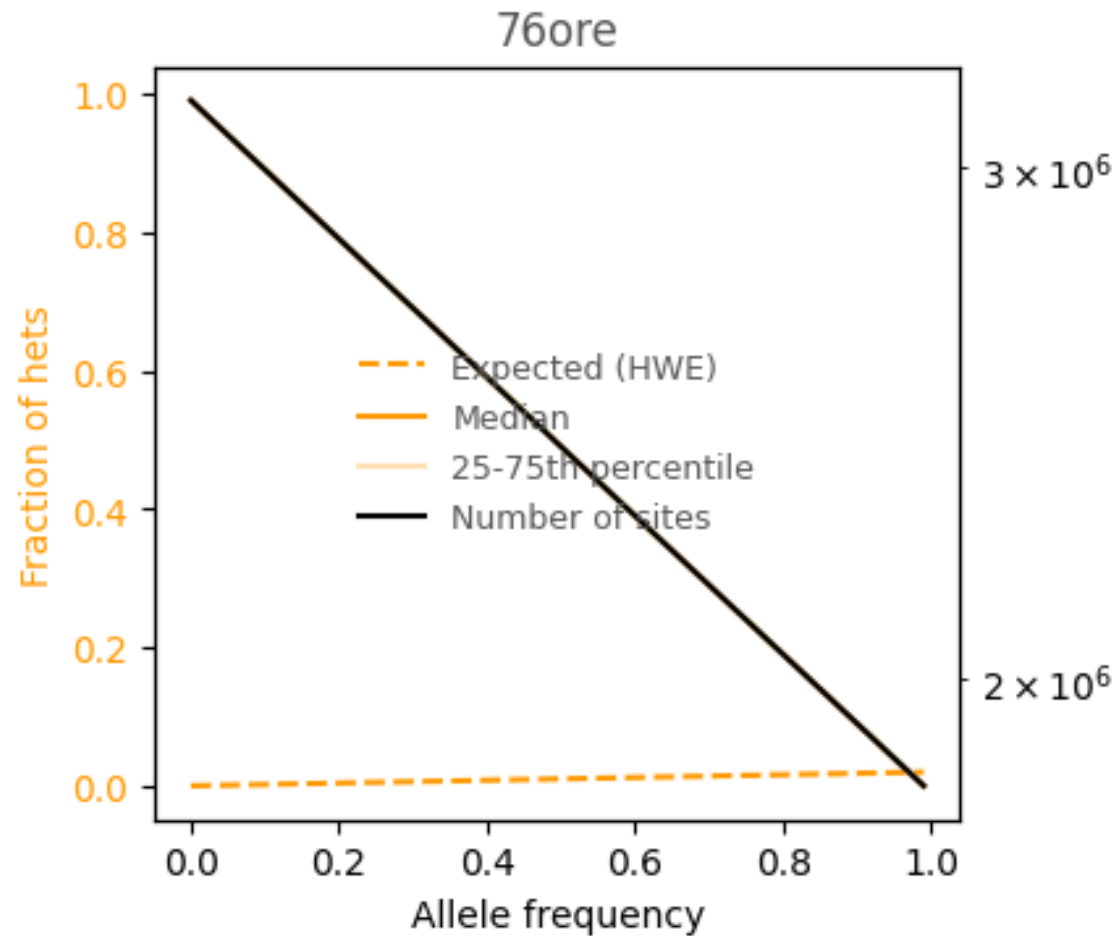
Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types