Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
59ang	4,076,801	1.92	1.93	951,073	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
59ang	62.3%	1.89	67.2%	100,883	2,058

- 59ang .. `/ngc/projects2/gm/data/archive/2022/variants/snv/59angraam-103914577534-Normal_B`
  `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM01782-220928_A01176_BH3FHLDSX5-EXT_LAB`
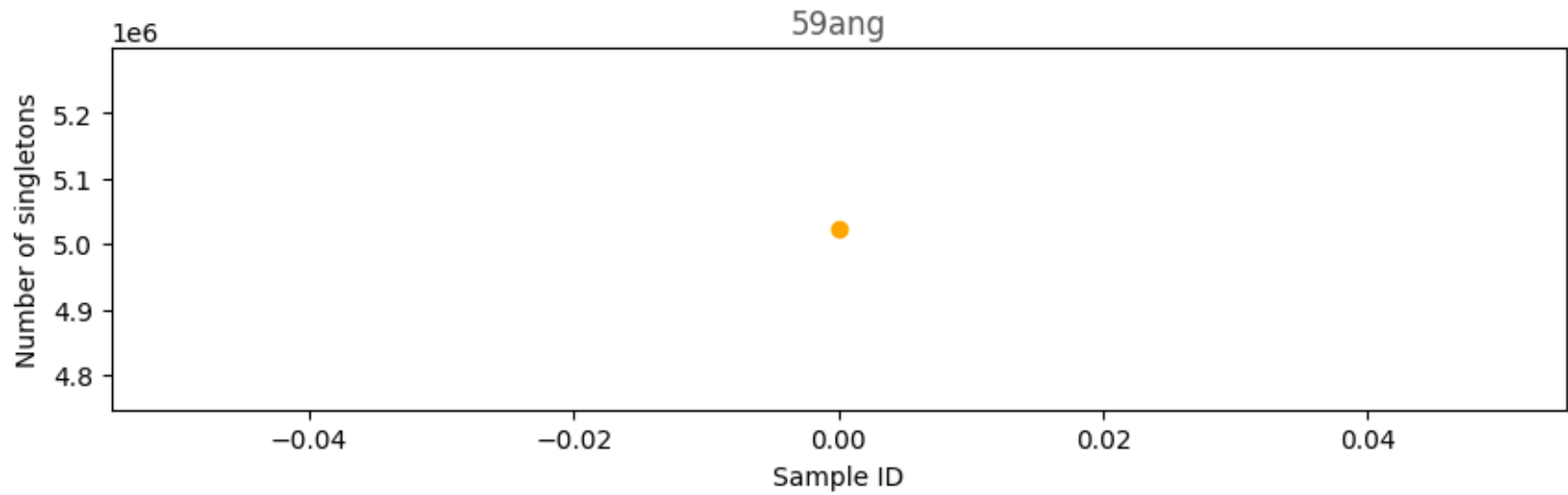 `KA_NGCWGS-NGCWGS05616_22RKG010104_snv_germline_raw.haplotype_caller.vcf.gz`
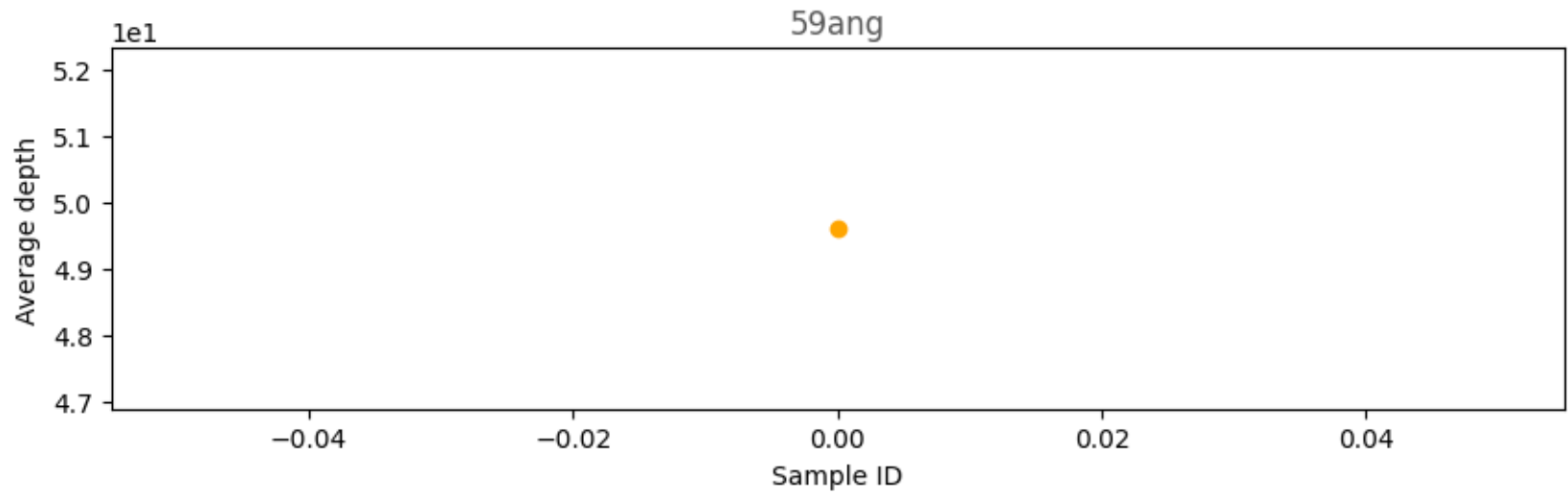
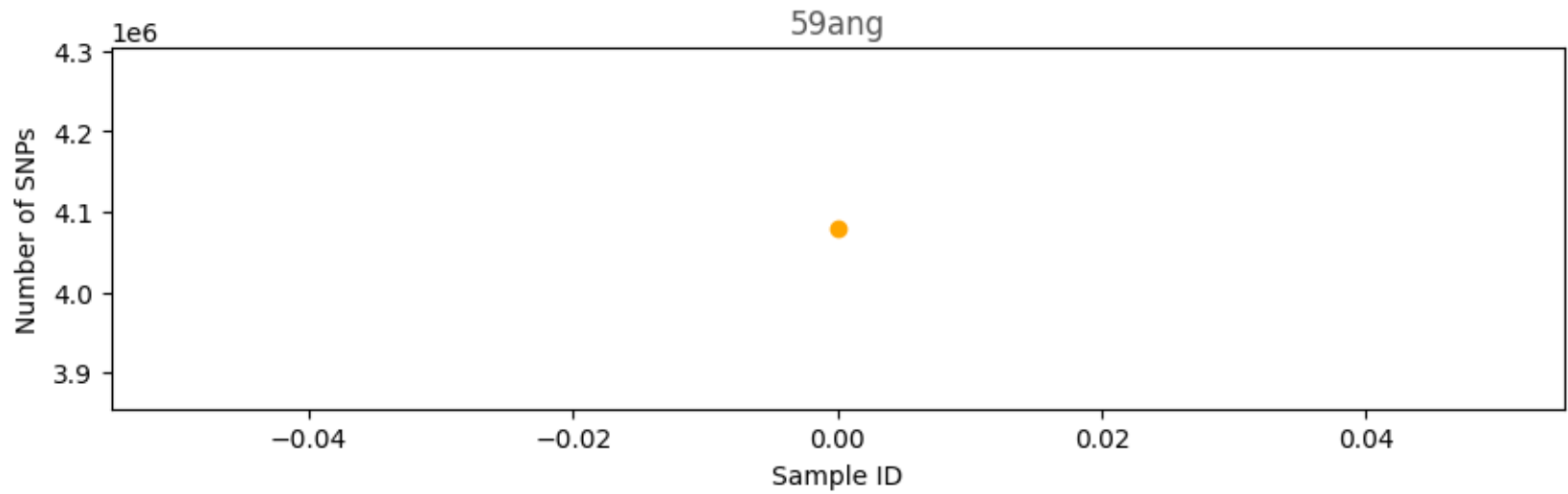# Ts/Tv by sample

59ang
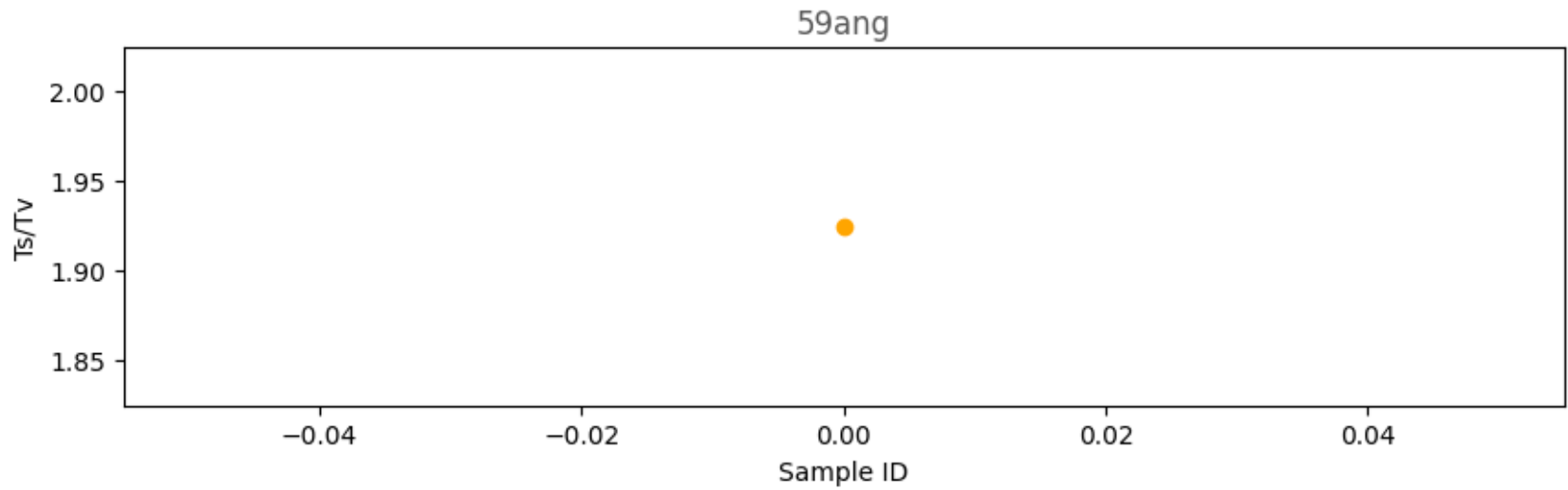
Hets vs non-ref Homs by sample
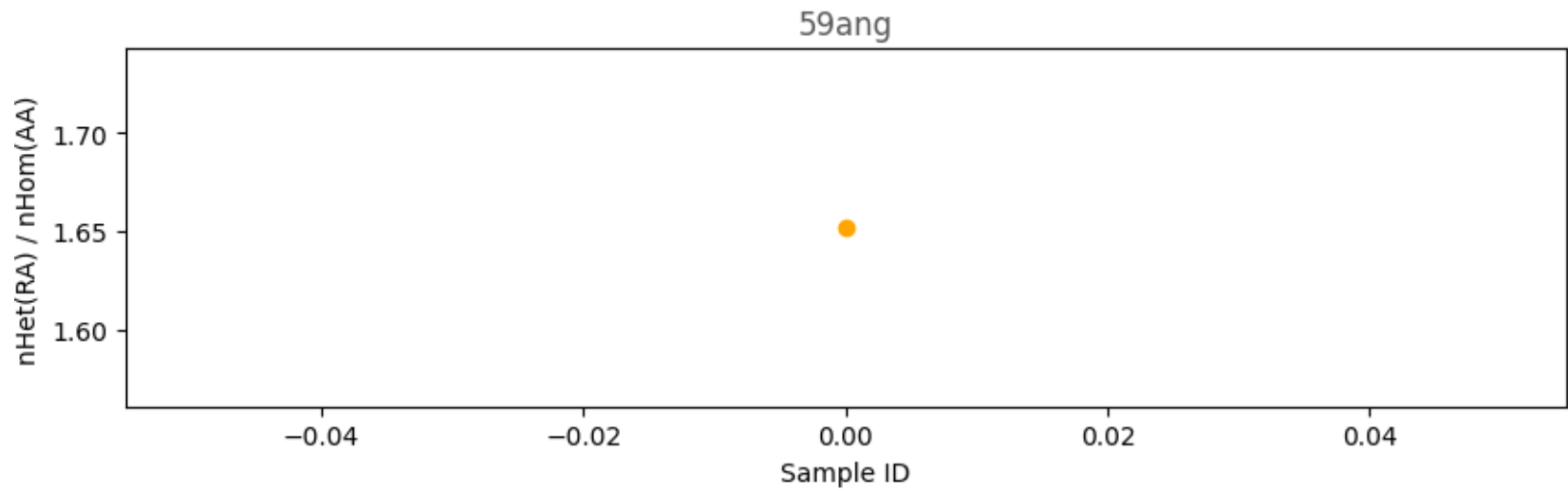
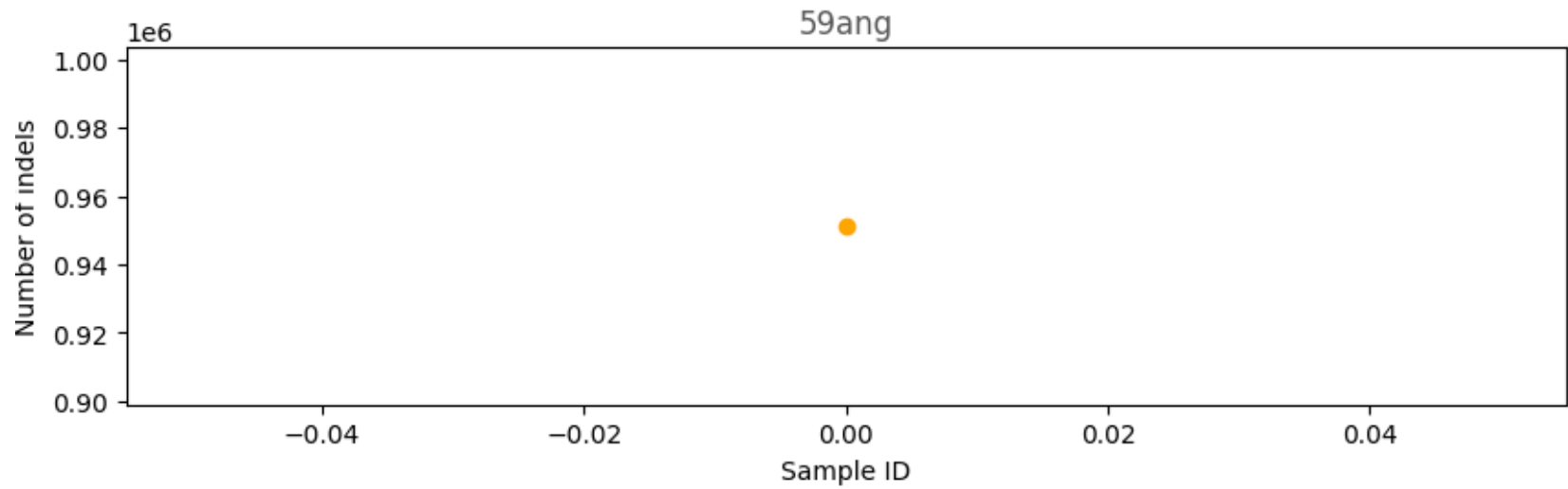# Singletons by sample (hets and homs)
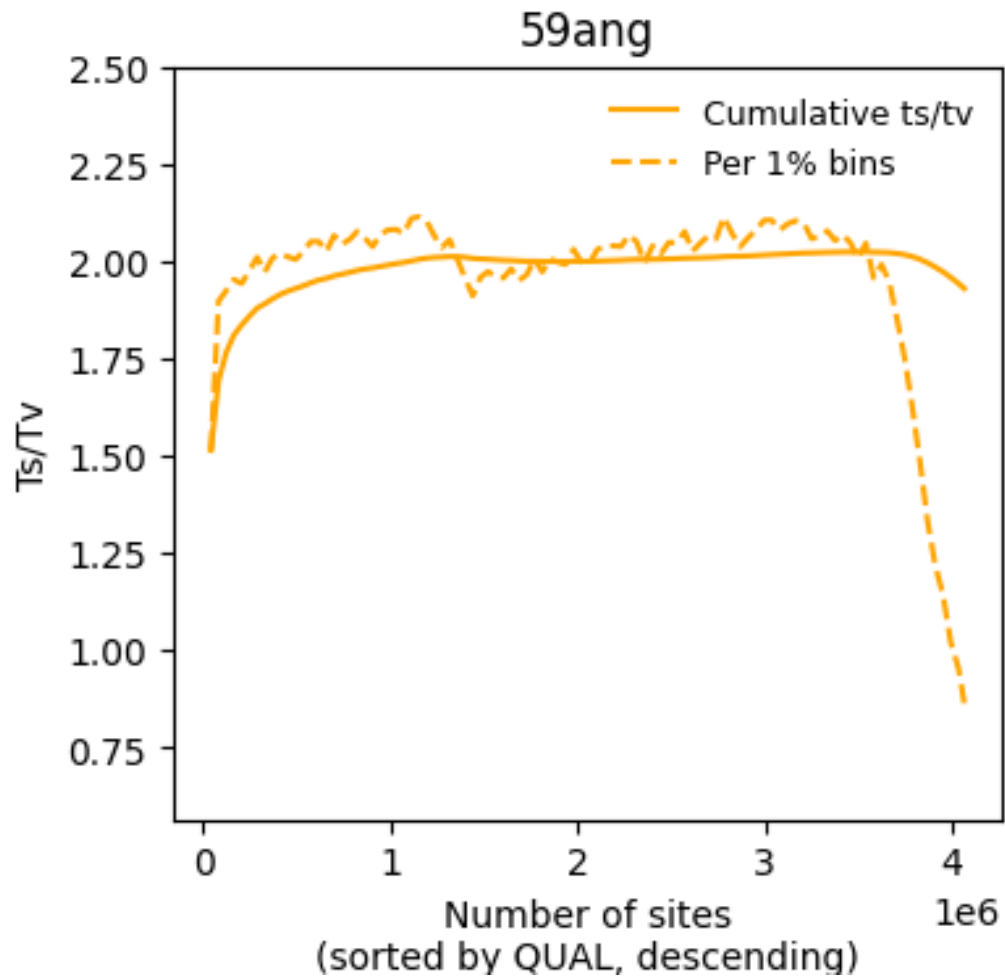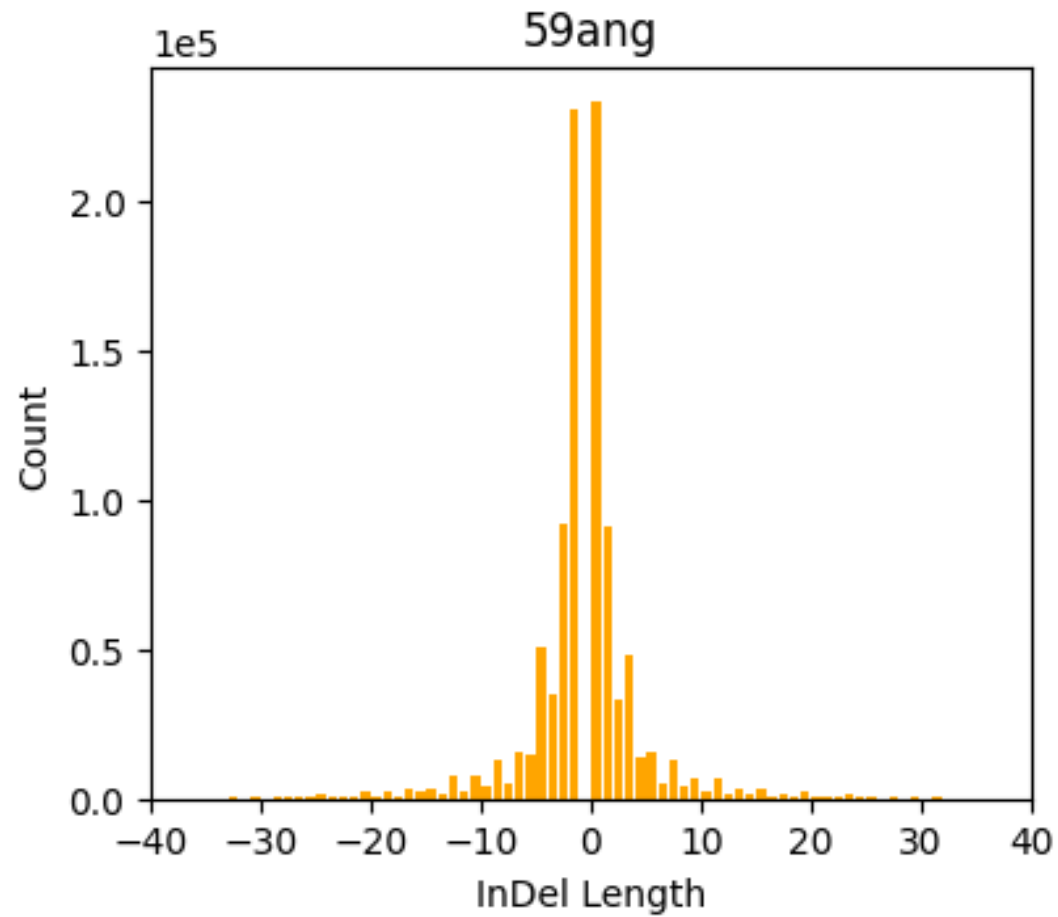
Average depth by sample
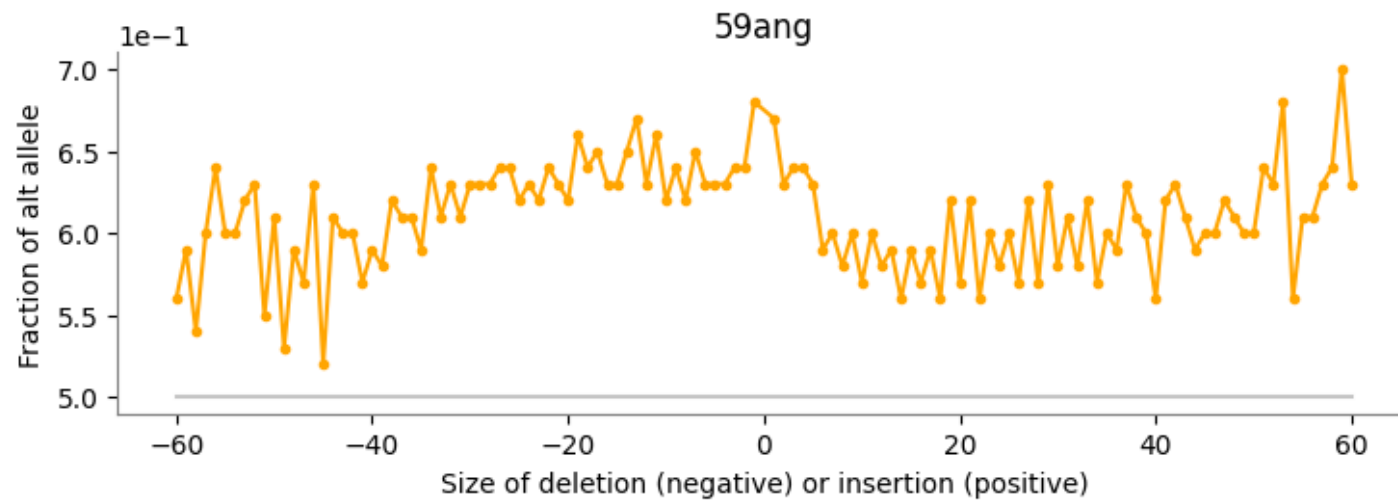
Number of SNPs by sample

Number of indels by sample

Ts/Tv stratified by QUAL

59ang

Indel distribution

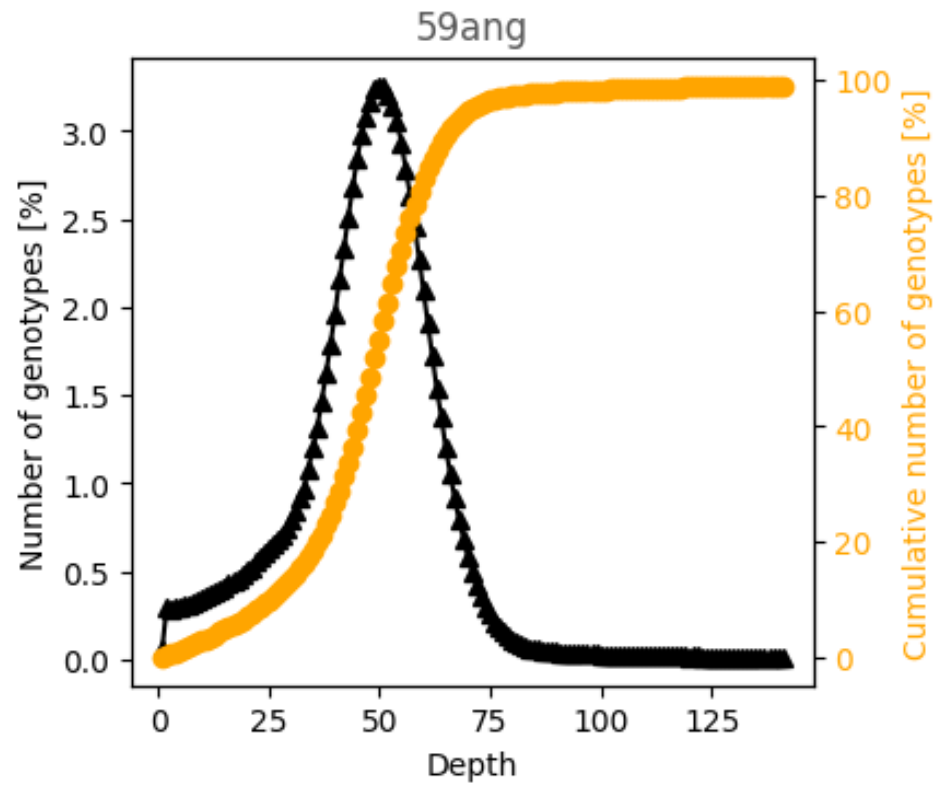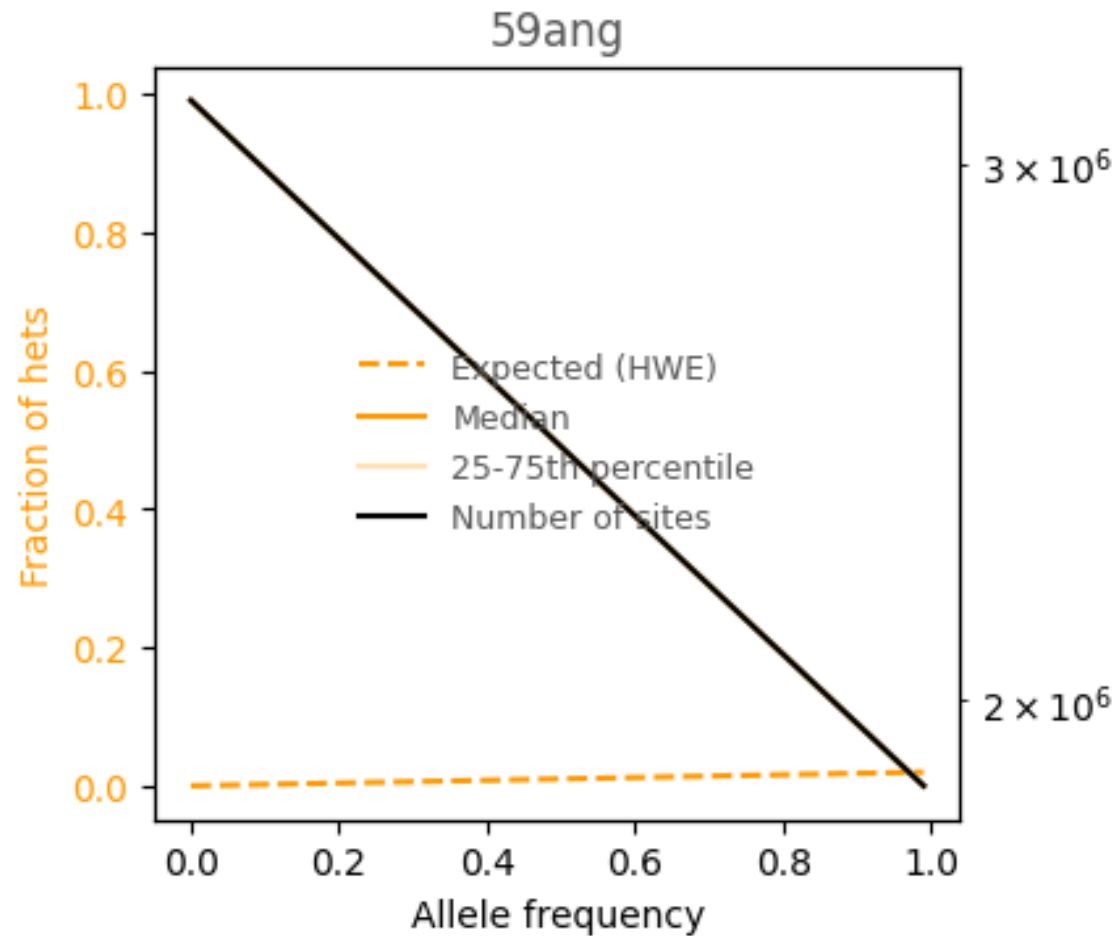# Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types