Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
90swh	4,044,540	1.93	1.94	932,550	–	0	0
							* frameshift ratio: out/(out+in)

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
90swh	61.8%	1.90	65.9%	92,914	1,966

- 90swh .. `/ngc/projects2/gm/data/archive/2022/variants/snv/90swhenkm-103849700440-Normal_B` `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-220318_A01411_BHCGF5DSX3-RHGM_LABKA_WGSA` `KUT-WGSAKUT04005_22RKG006278x01_snv_germline_raw.haplotype_caller.vcf.gz`
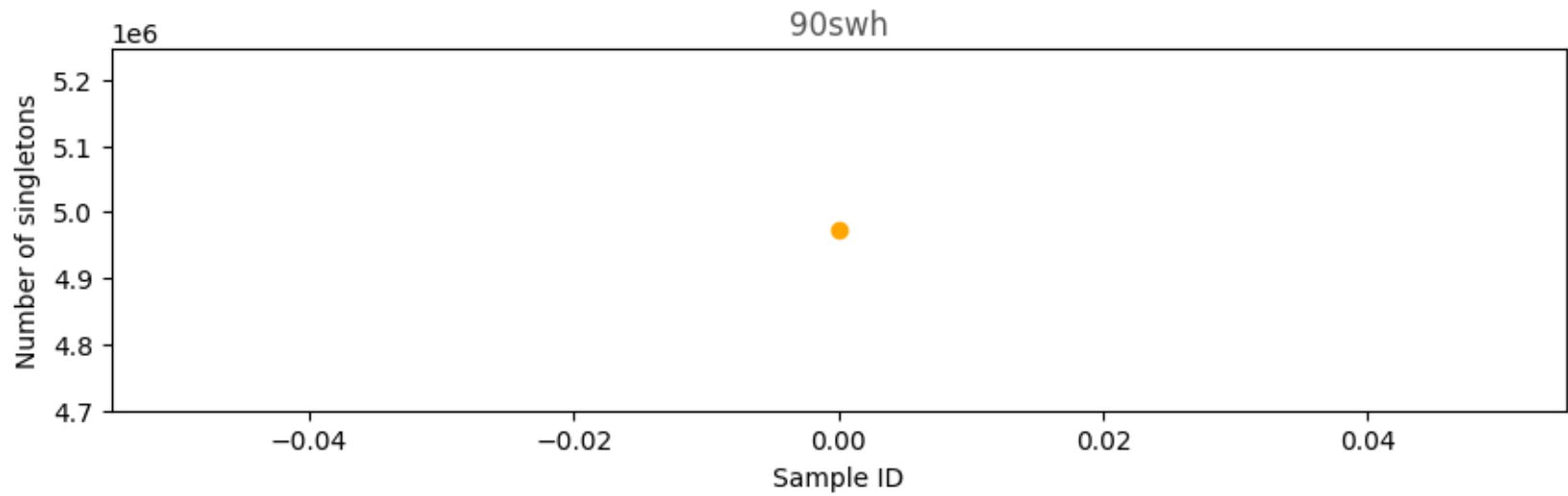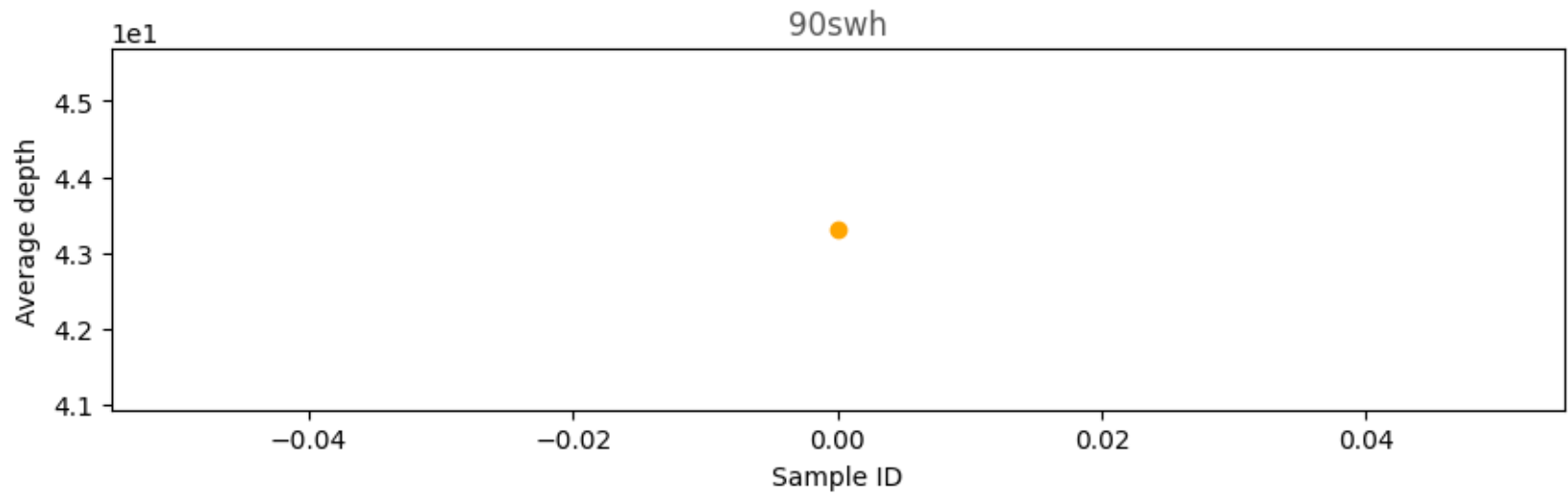
Ts/Tv by sample
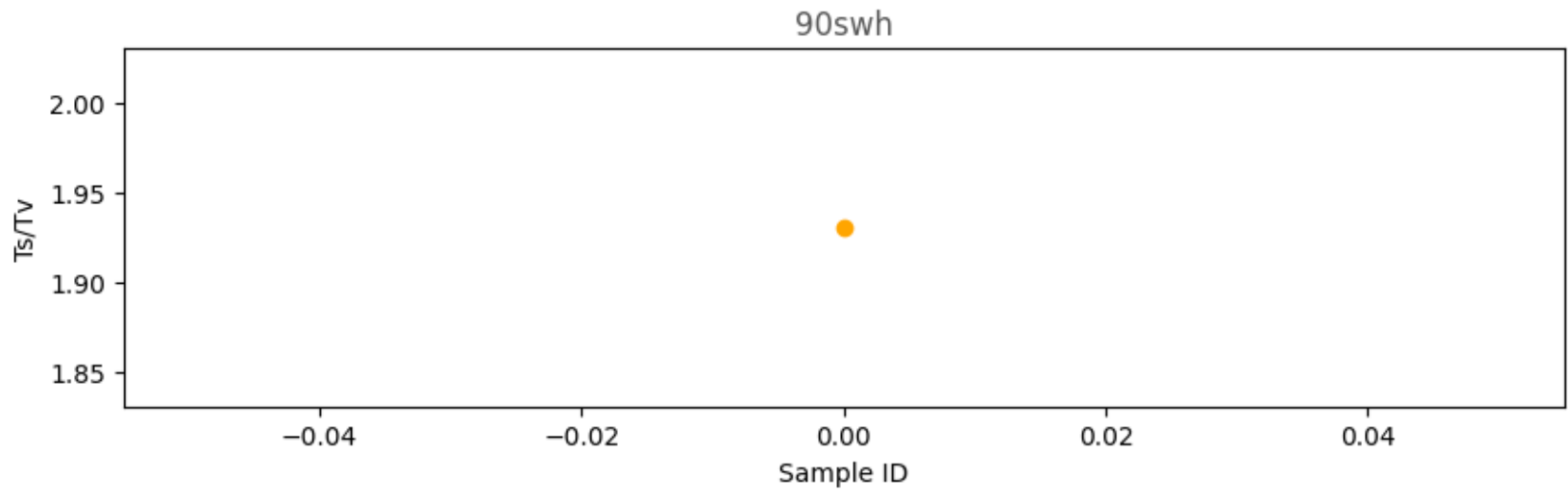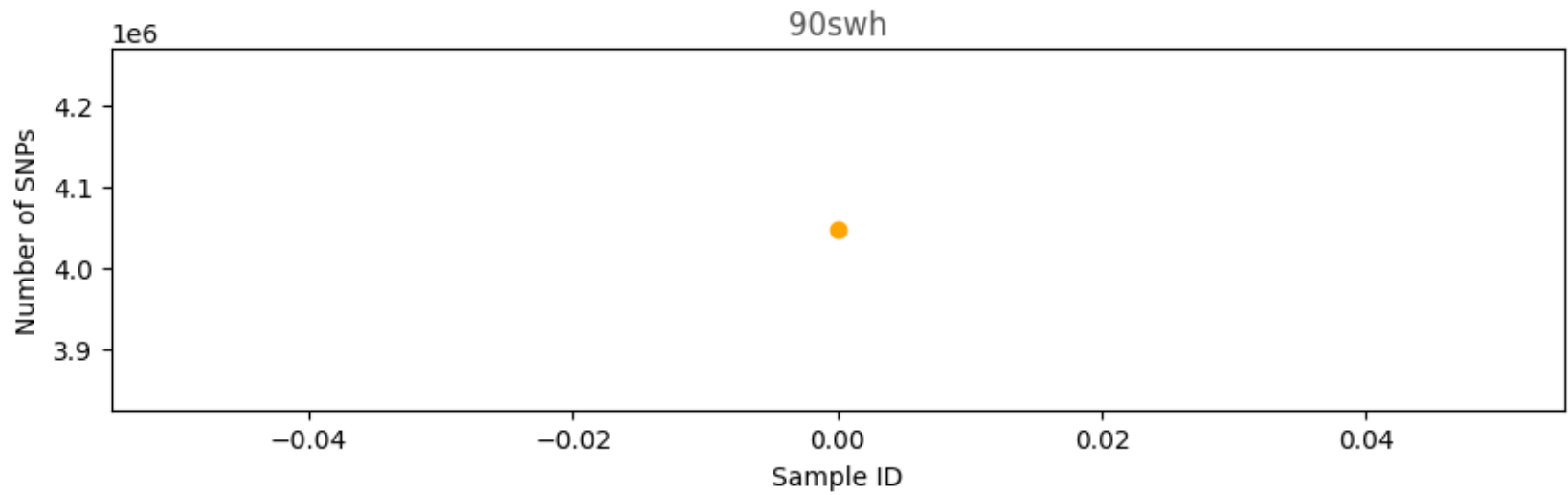
Hets vs non-ref Homs by sample
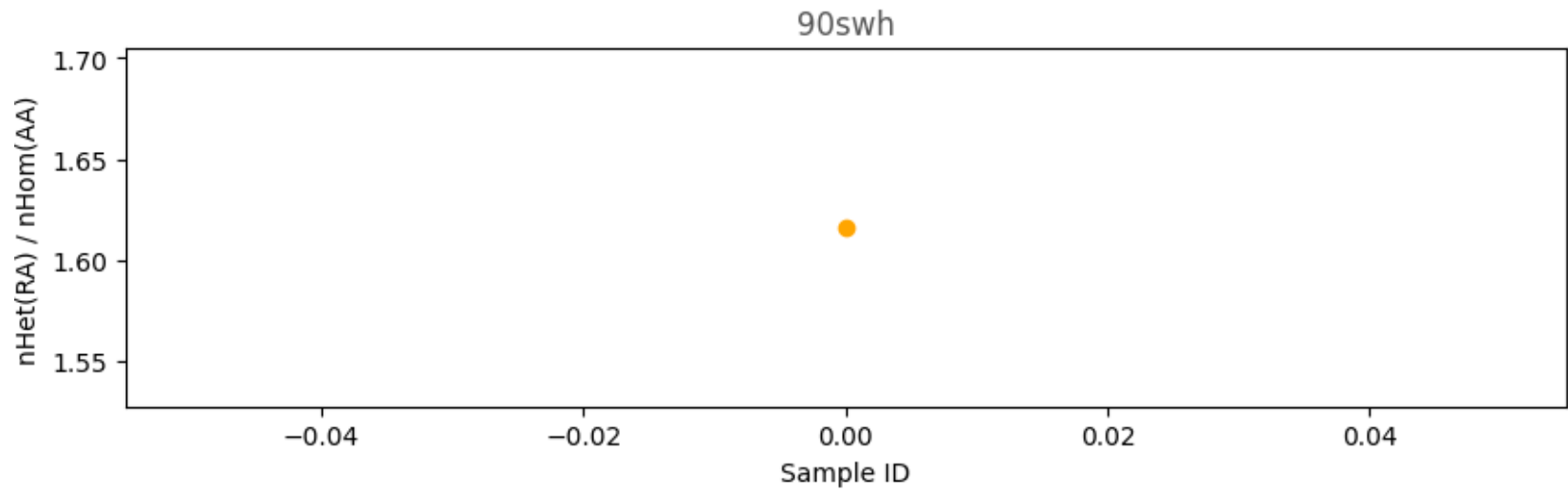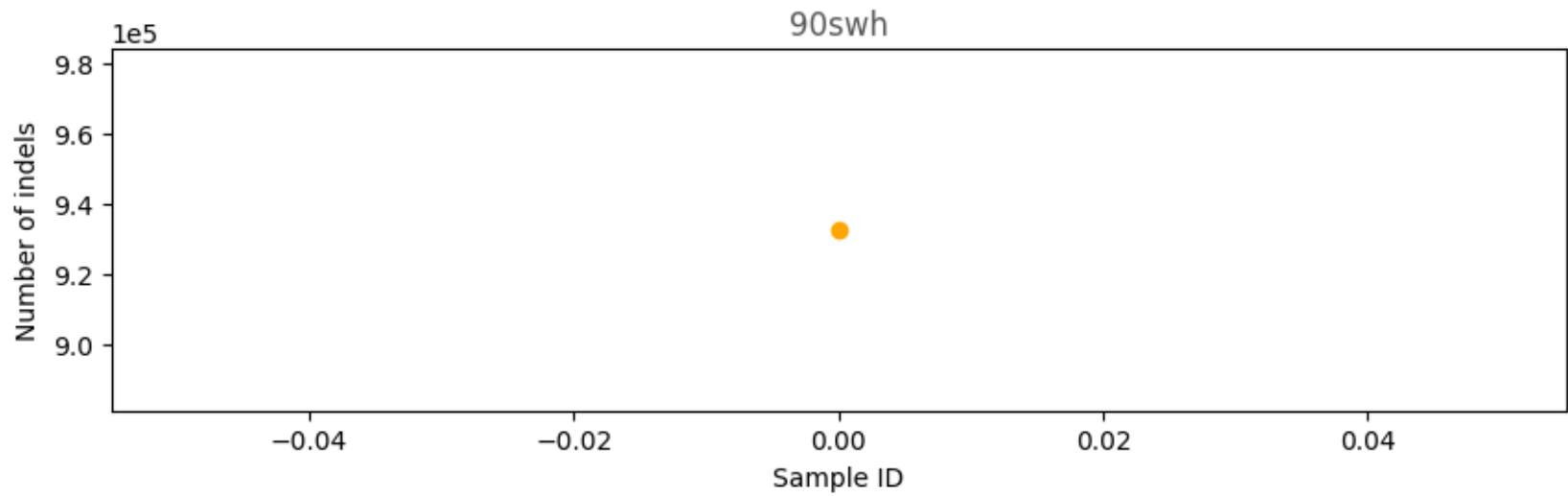
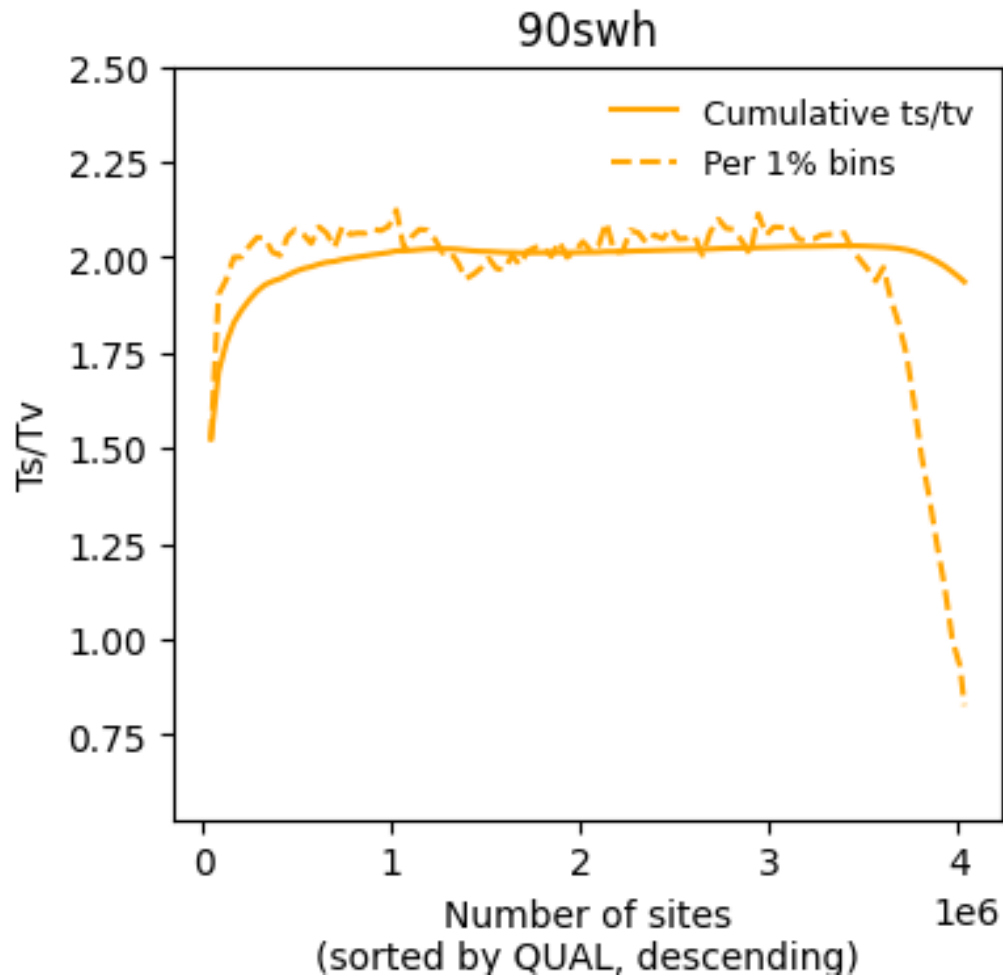# Singletons by sample (hets and homs)

Average depth by sample
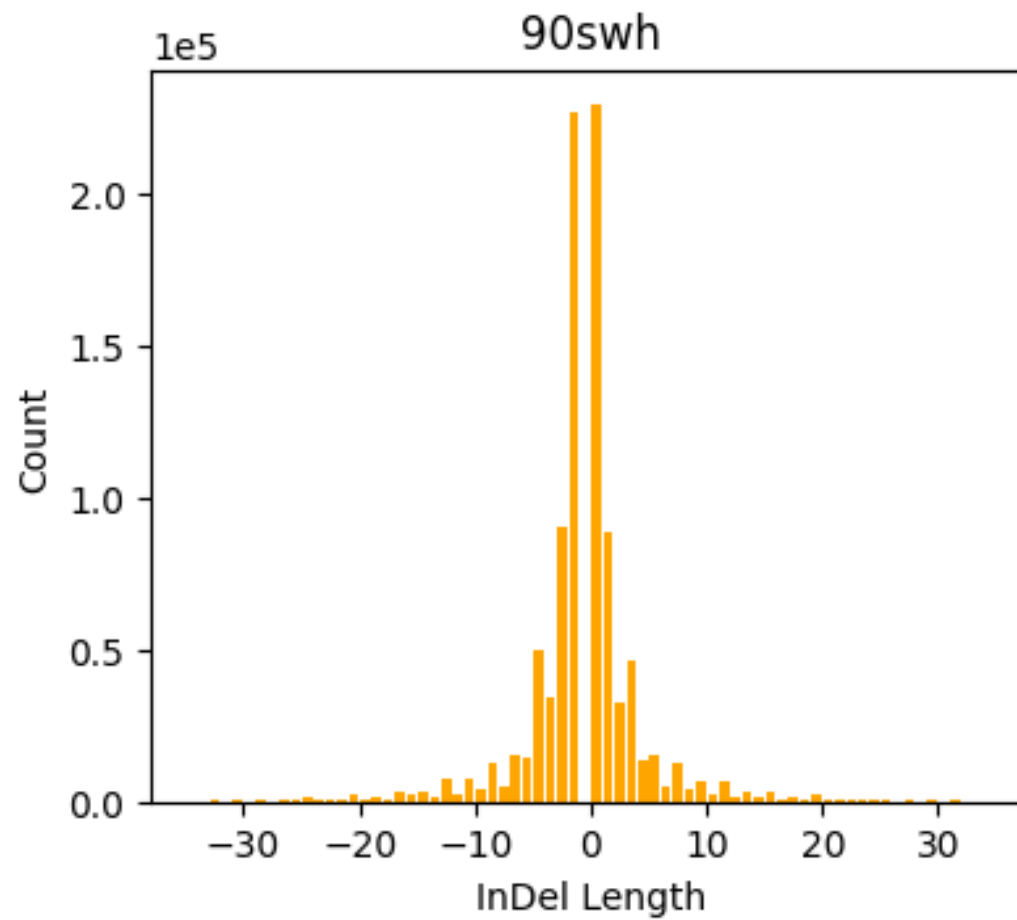
Number of SNPs by sample

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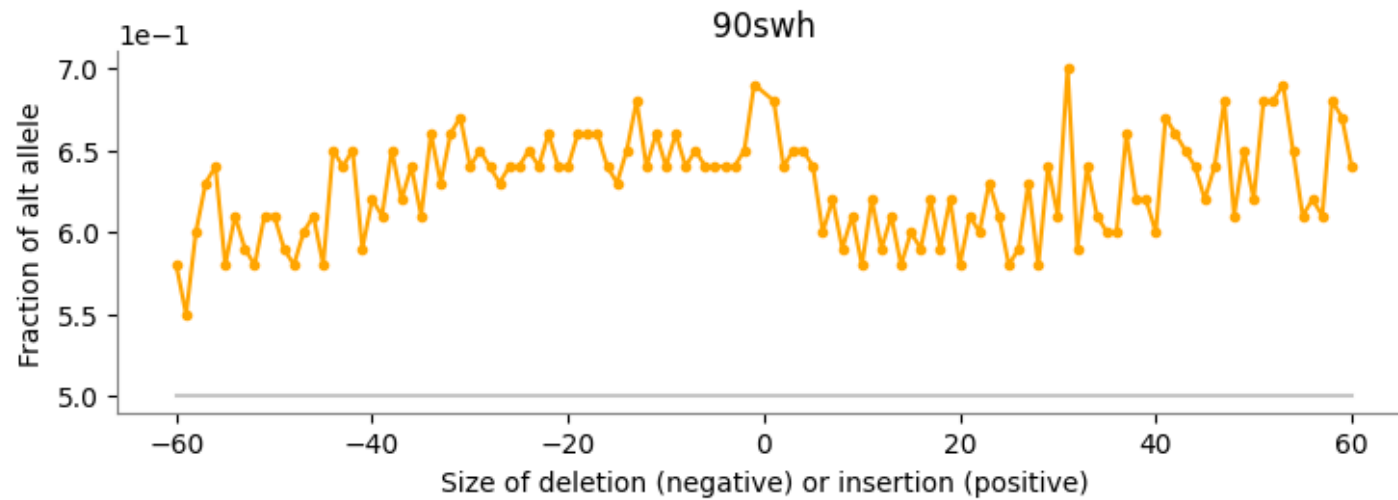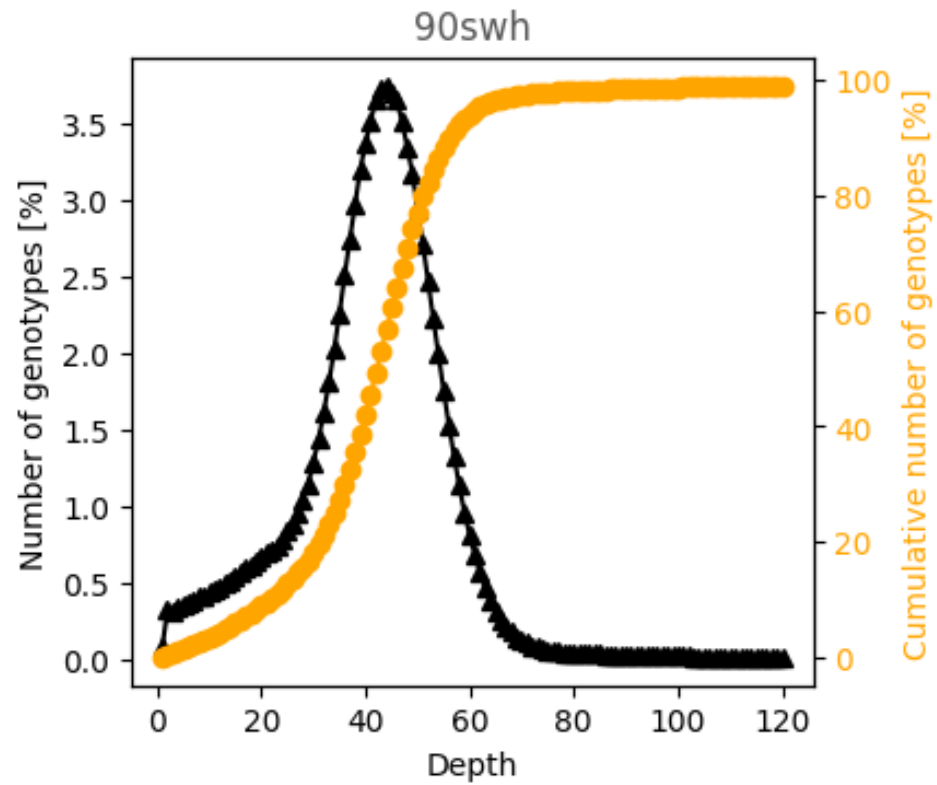
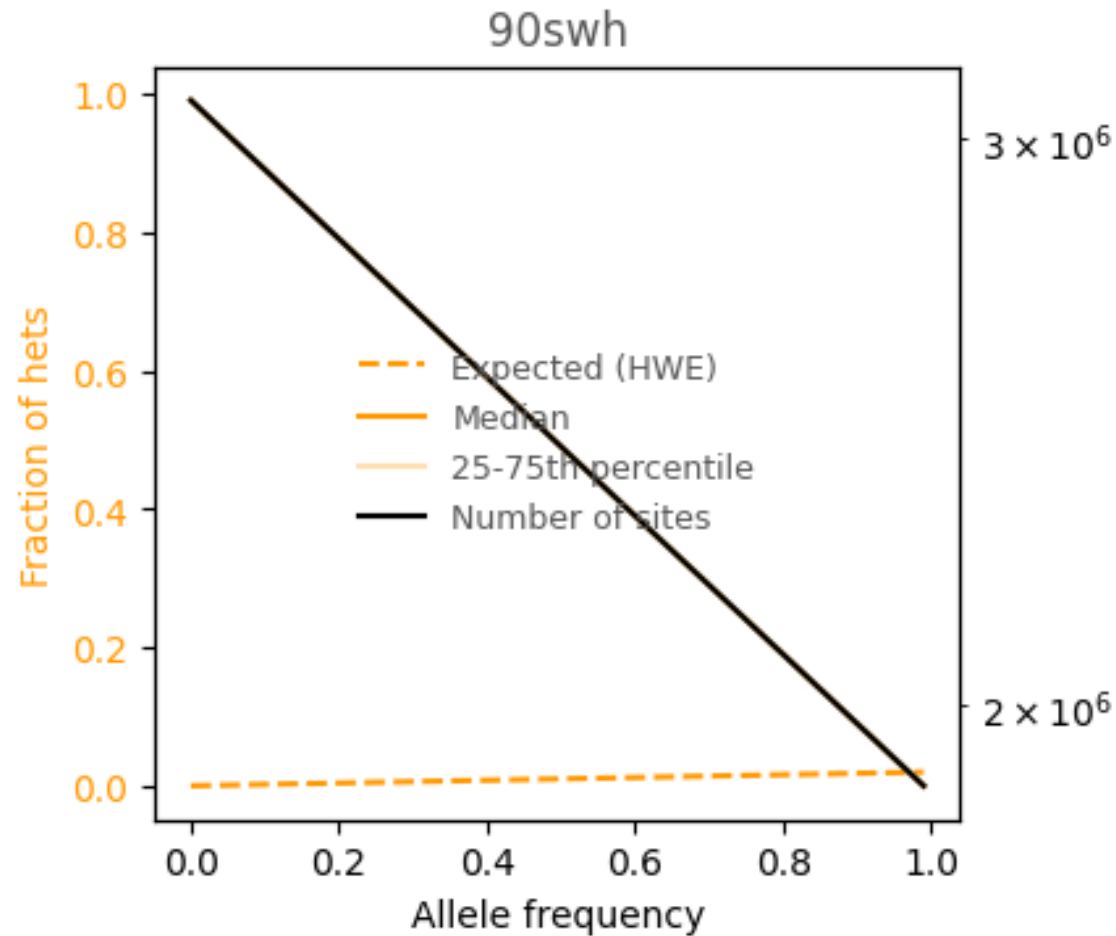# Ts/Tv stratified by QUAL

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types