Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
81psh	4,088,937	1.93	1.94	953,589	–	0	0
					* frameshift ratio: out/(out+in)		

Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
81psh	63.3%	1.91	68.2%	101,015	2,053

- 81psh .. `/ngc/projects2/gm/data/archive/2022/variants/snv/81pshausf-103818402617-Normal_B`
   `lood_noinfo-WGS_v1_IlluminaDNAPCRFree_X-211222_A01466_AH2CJCDSX3-RHGM_LABKA_WGSA`
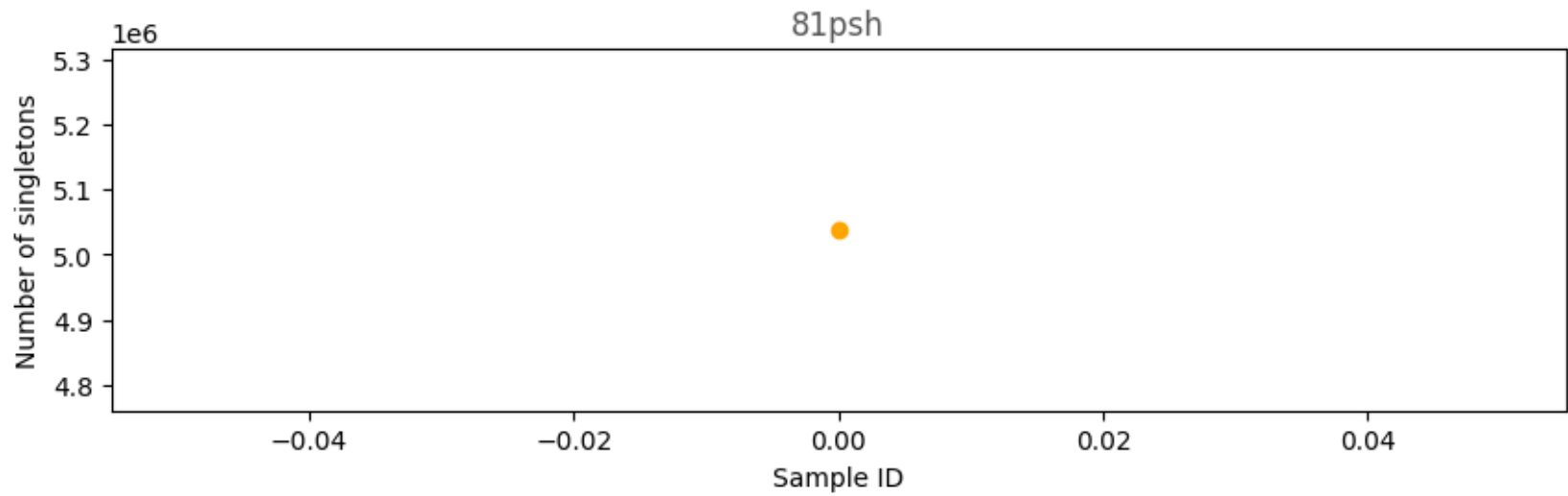   `KUT-WGSAKUT03550_21RKG029750x01_snv_germline_raw.haplotype_caller.vcf.gz`
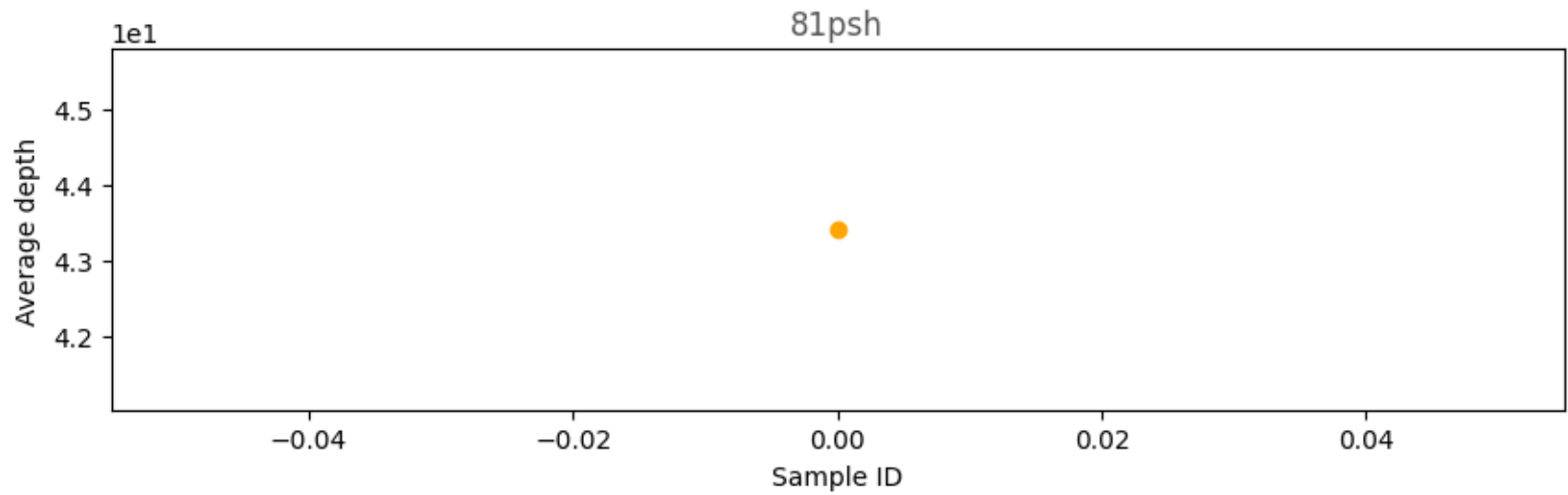
Ts/Tv by sample

81psh
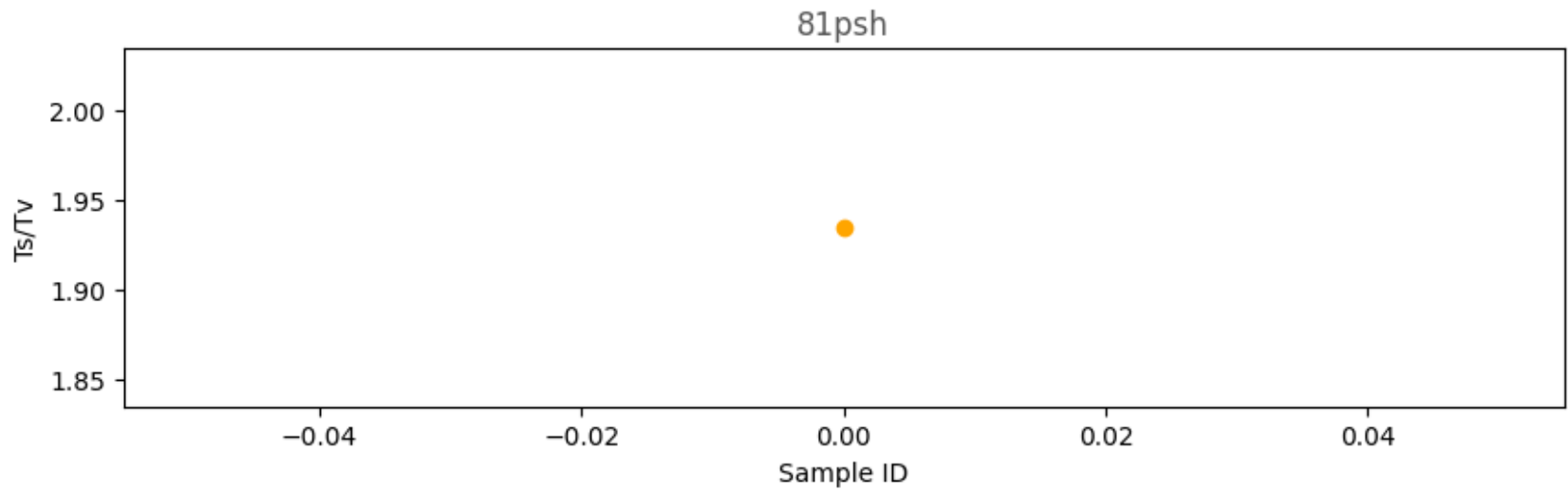
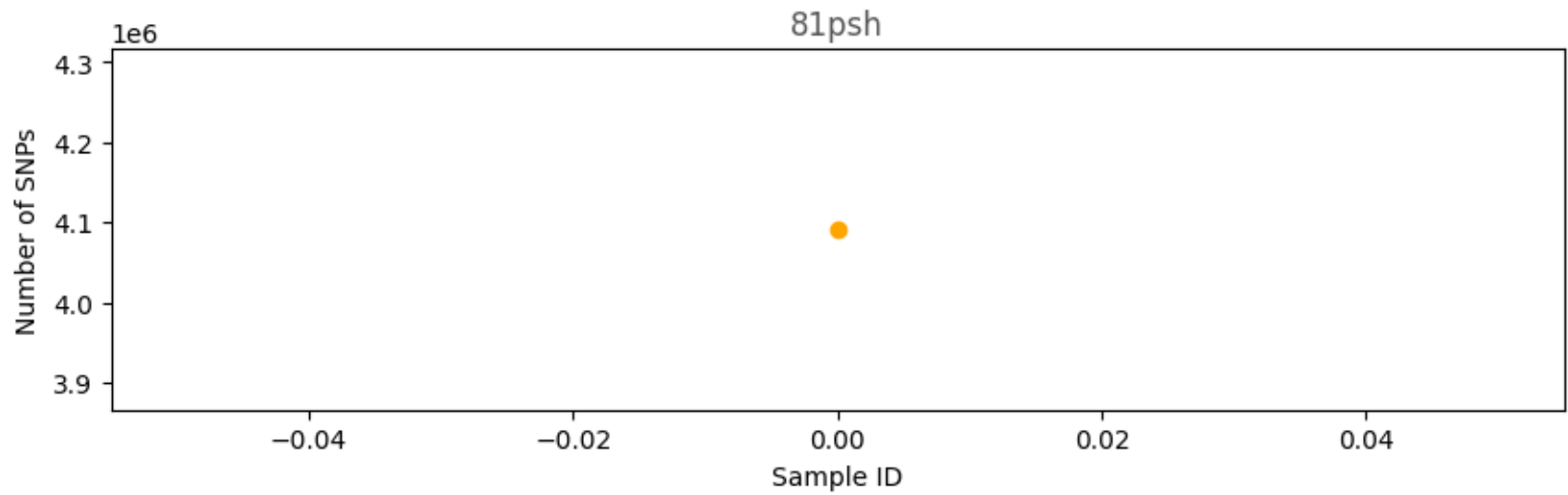# Hets vs non-ref Homs by sample
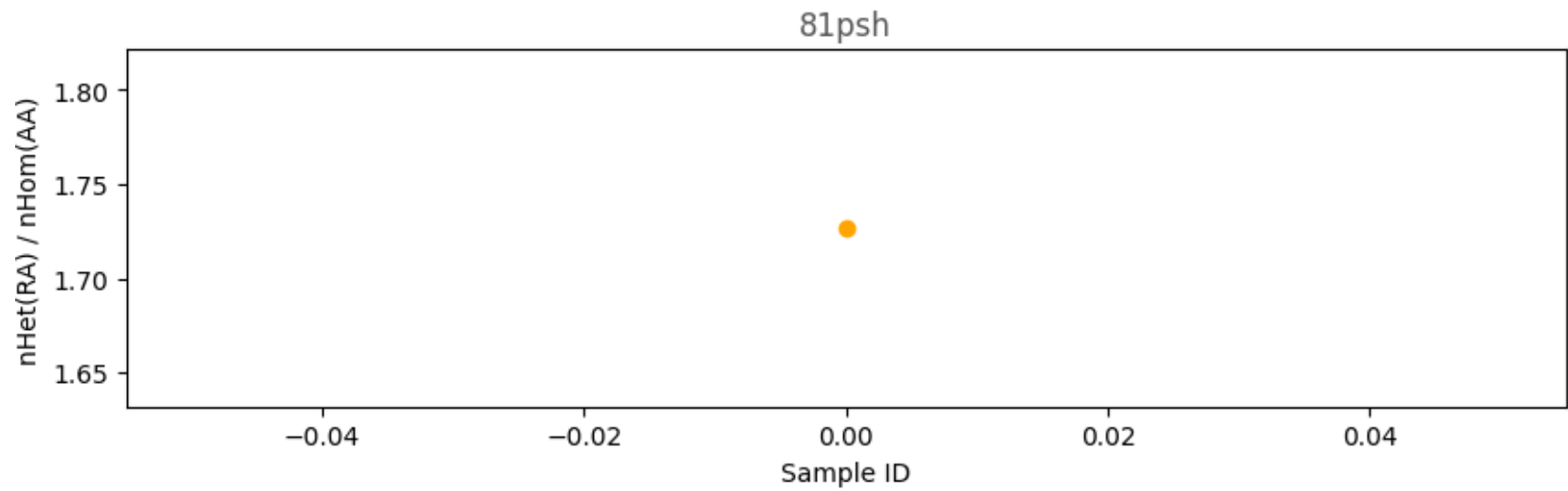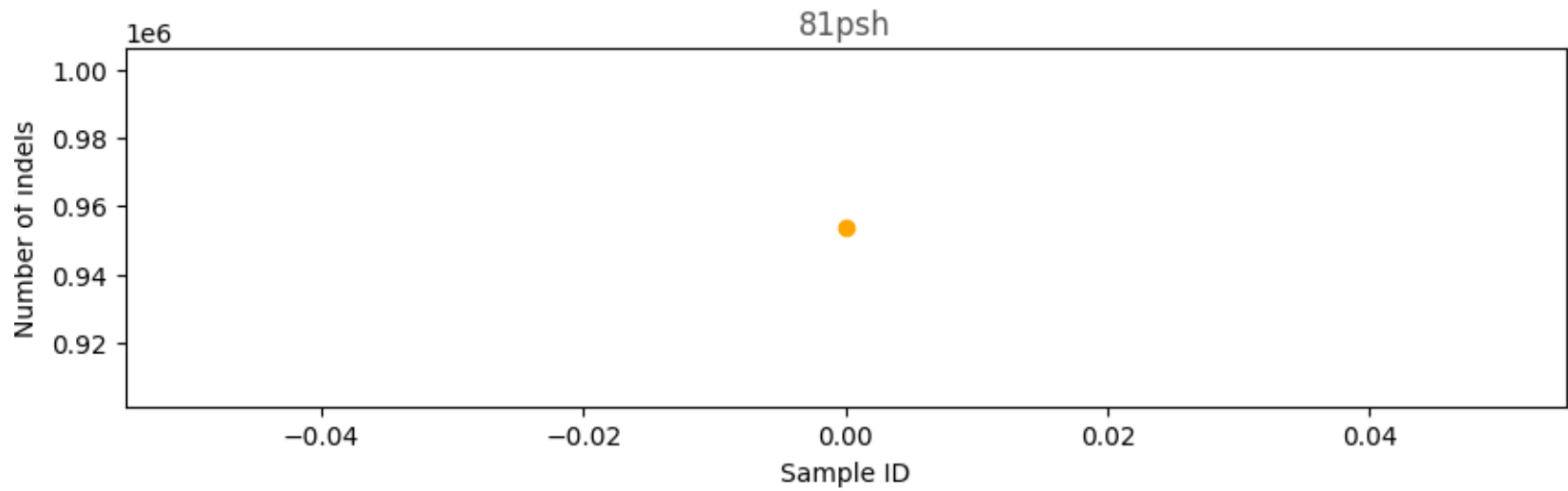
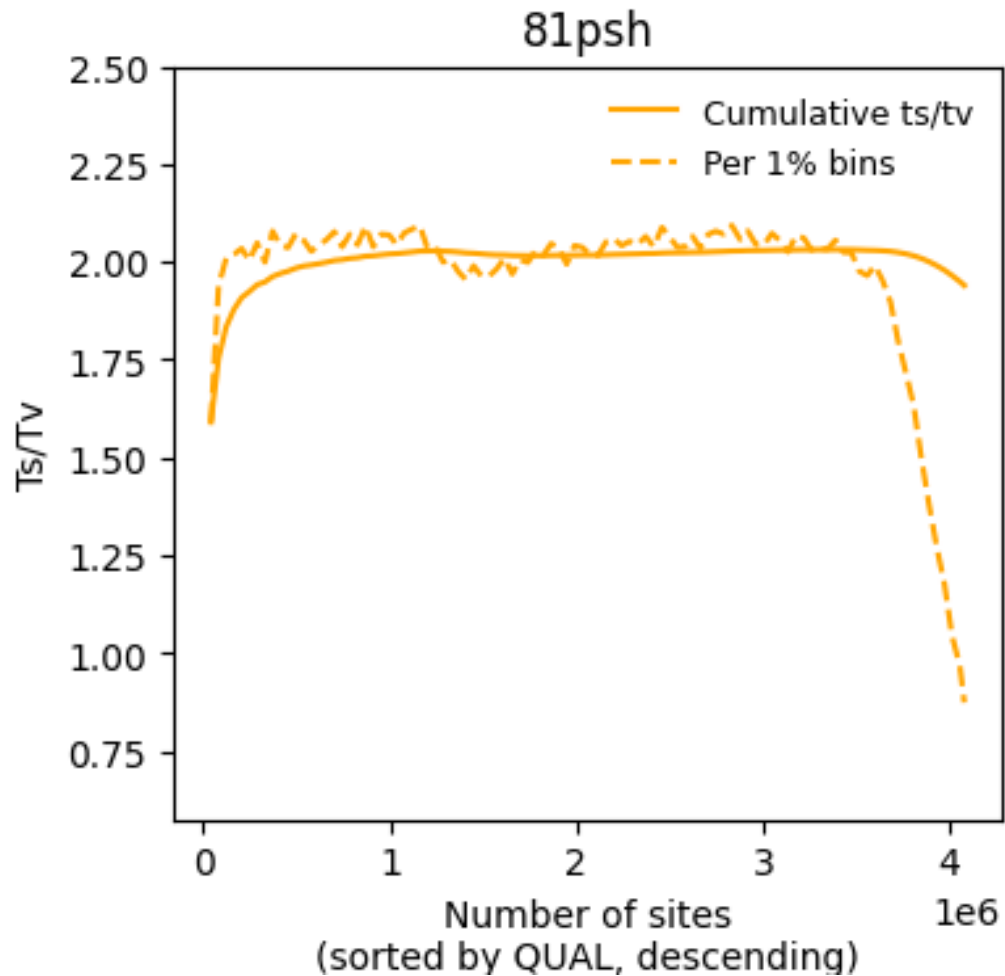# Singletons by sample (hets and homs)

Average depth by sample

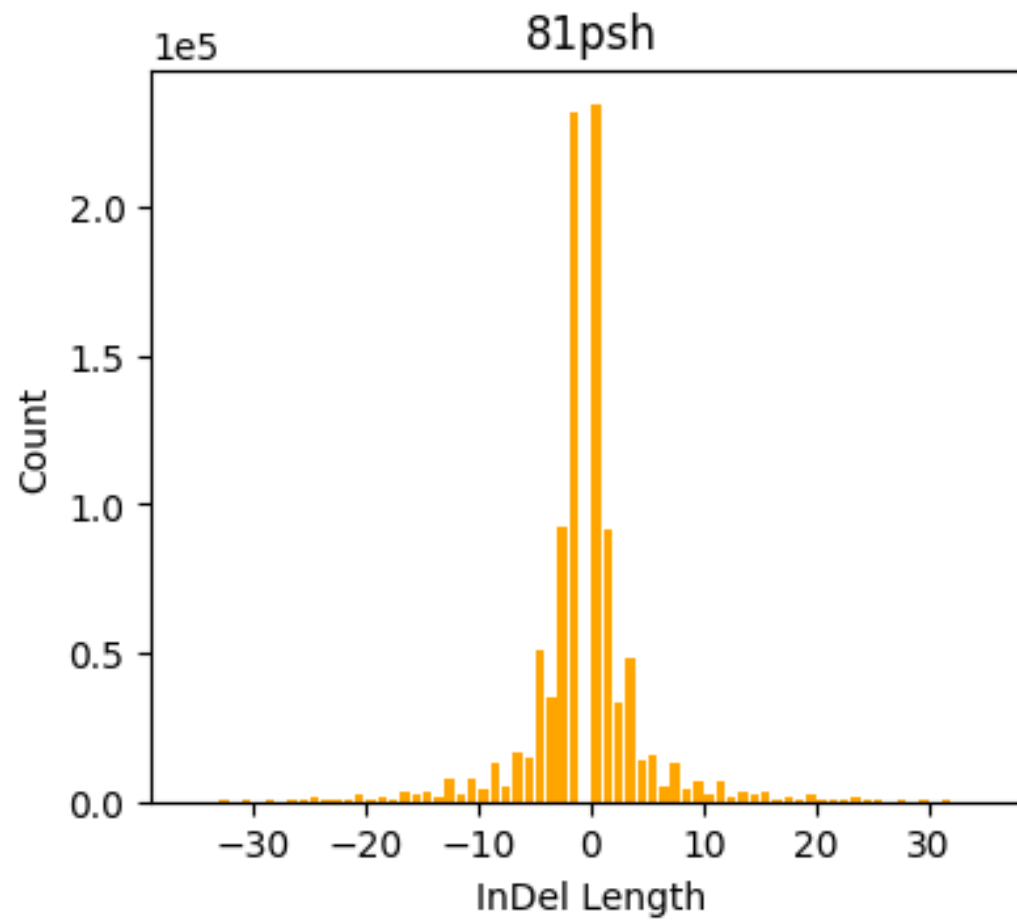# Number of SNPs by sample

Number of indels by sample
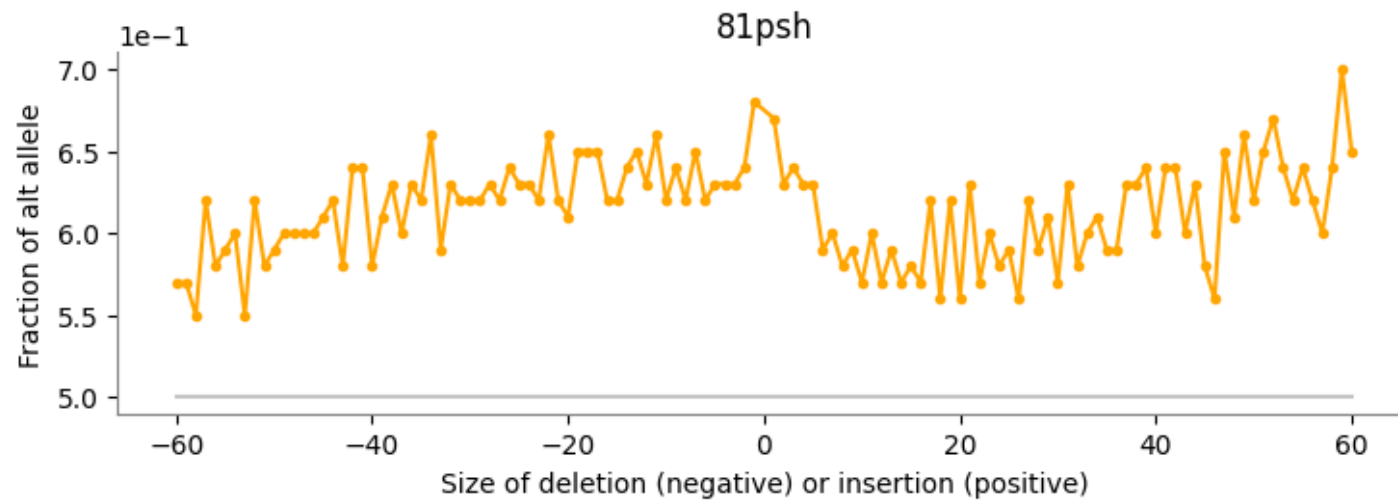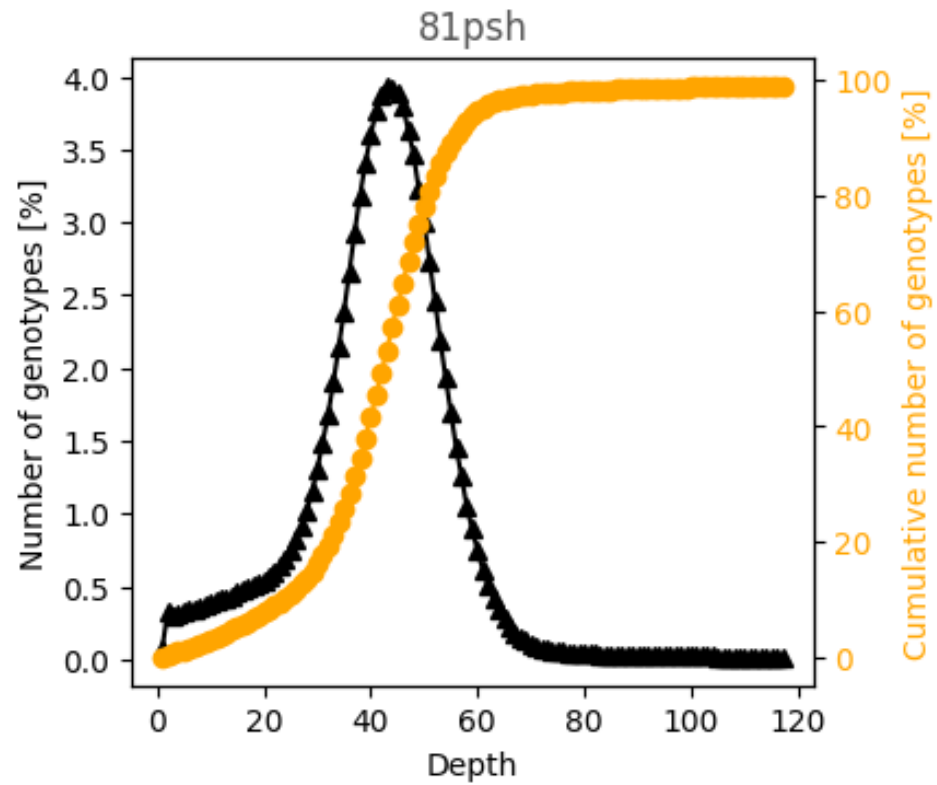
Ts/Tv stratified by QUAL

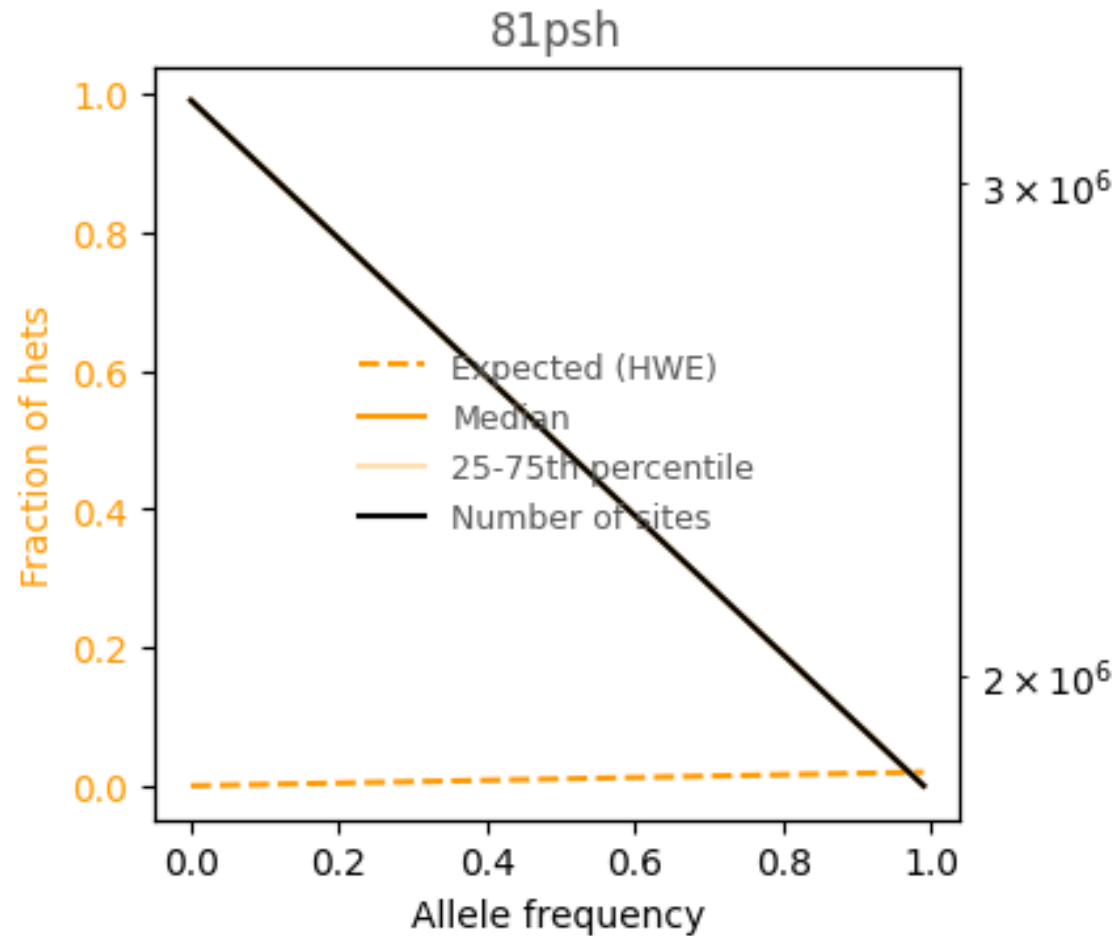

81psh

Indel distribution

Fraction of alternate indel allele

Depth distribution

Number of HETs by AF

Substitution types