

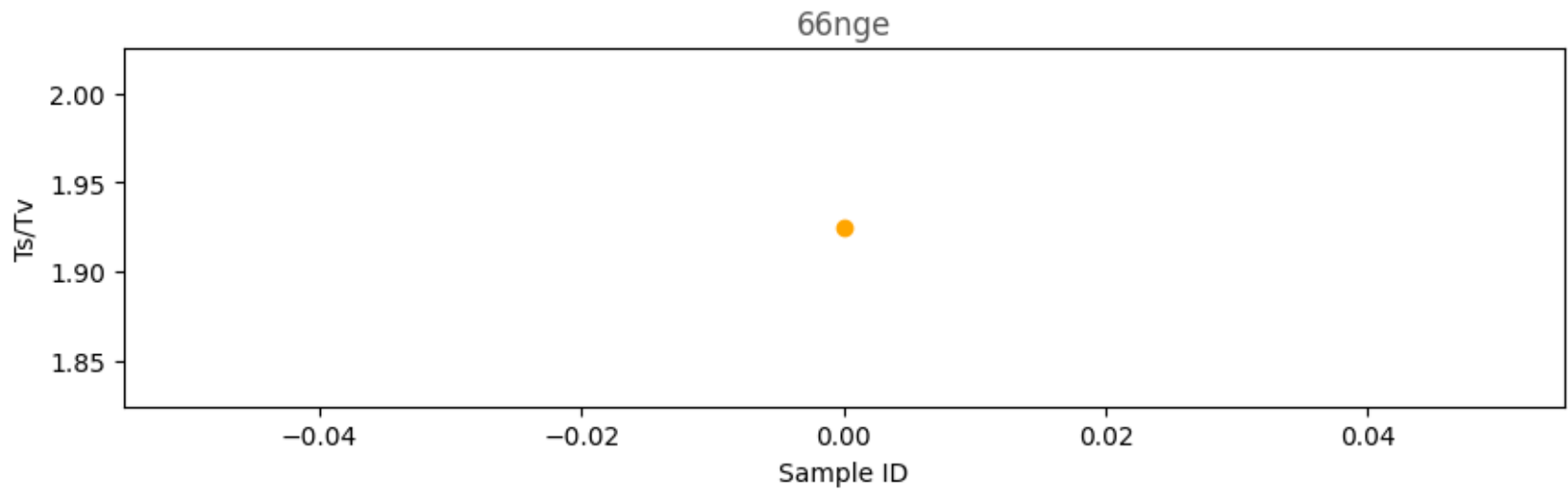
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
66nge	4,099,622	1.92	1.93	957,328	–	0	0
* frameshift ratio: out/(out+in)							

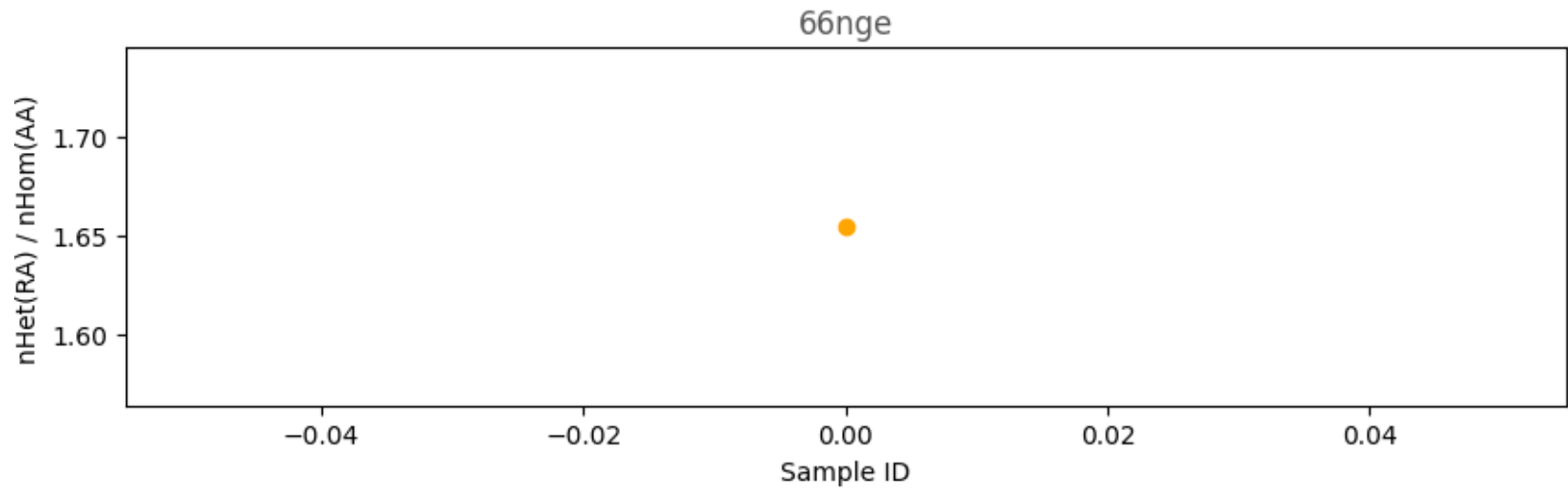
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
66nge	62.3%	1.89	67.2%	101,137	2,061

- 66nge .. /ngc/projects2/gm/data/archive/2022/variants/snv/66nge_twnm-103760977365-Normal_Blood_noinfo-WGS_v1_IlluminaDNAPCRFree_RHGM00258-210720_A01411_AHGW5DSX2-EXT_LAB_KA_NGCWGS-NGCWGS00151-661c71xu5-snv_germline_raw.haplotype_caller.vcf.gz

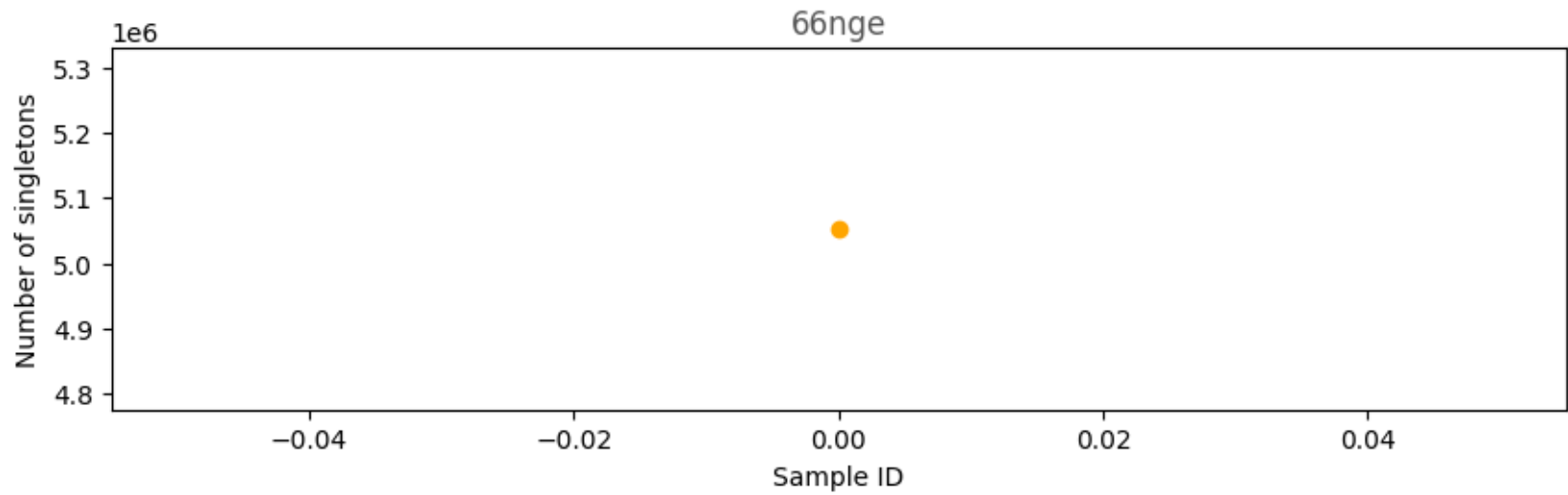
Ts/Tv by sample



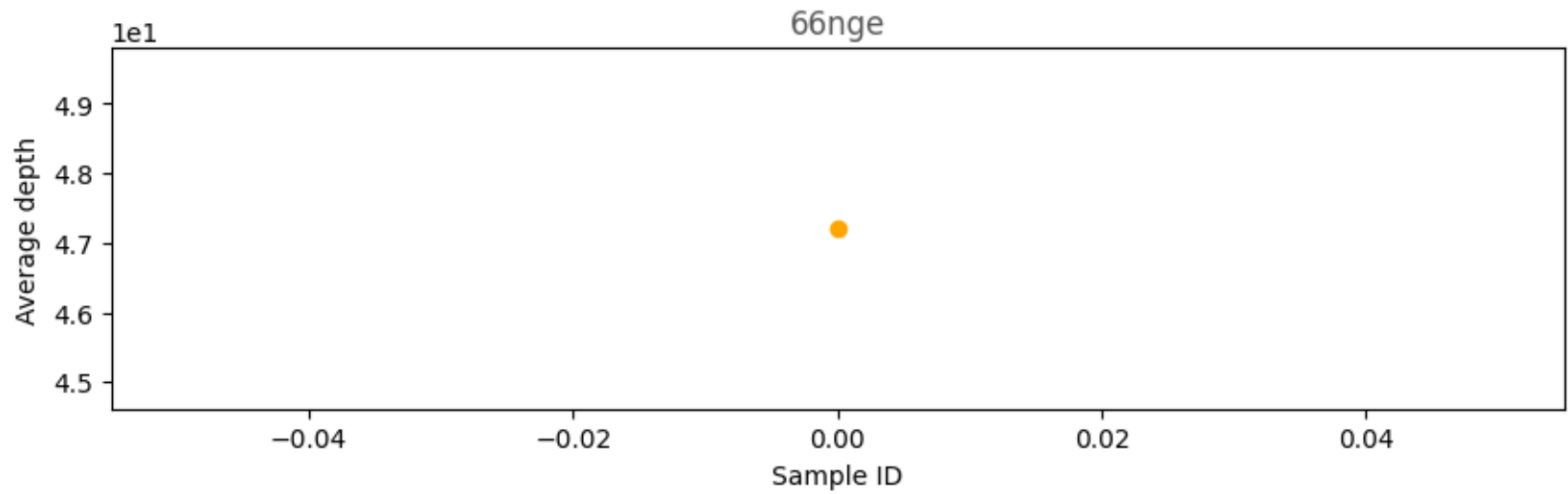
Hets vs non-ref Homs by sample



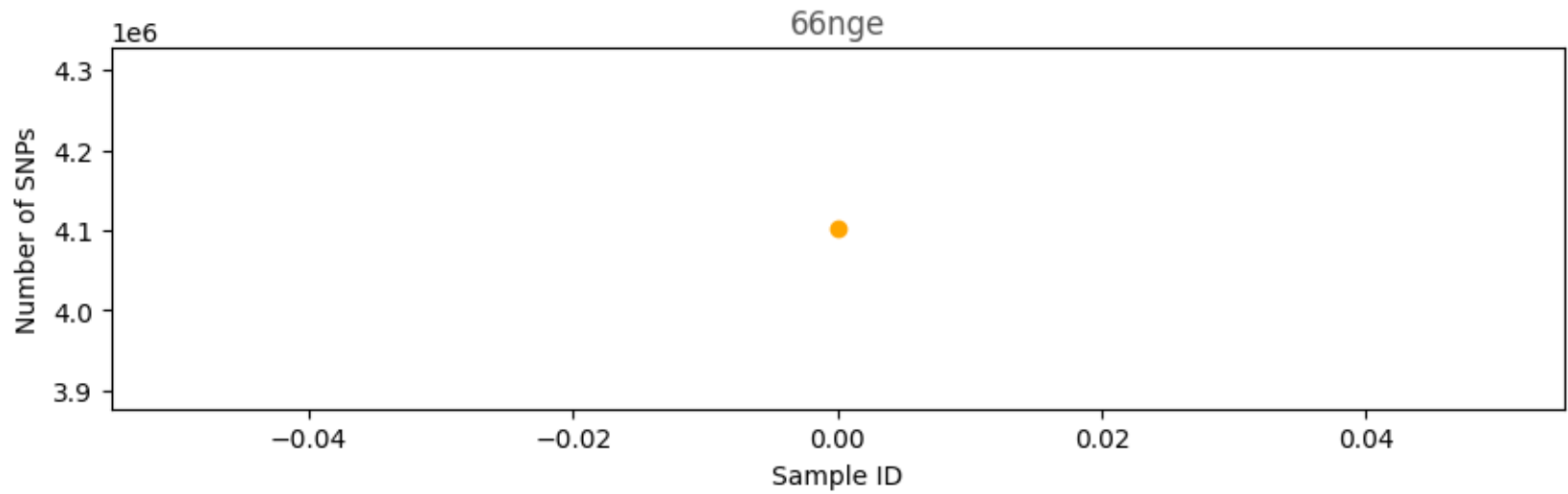
Singletons by sample (hets and homs)



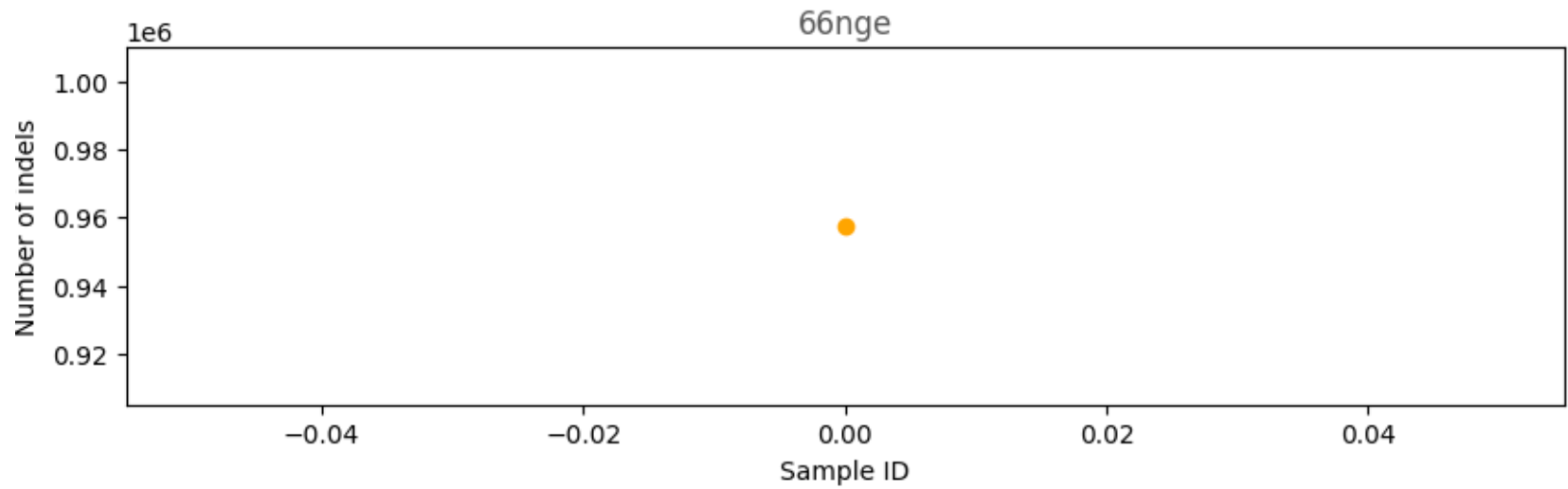
Average depth by sample



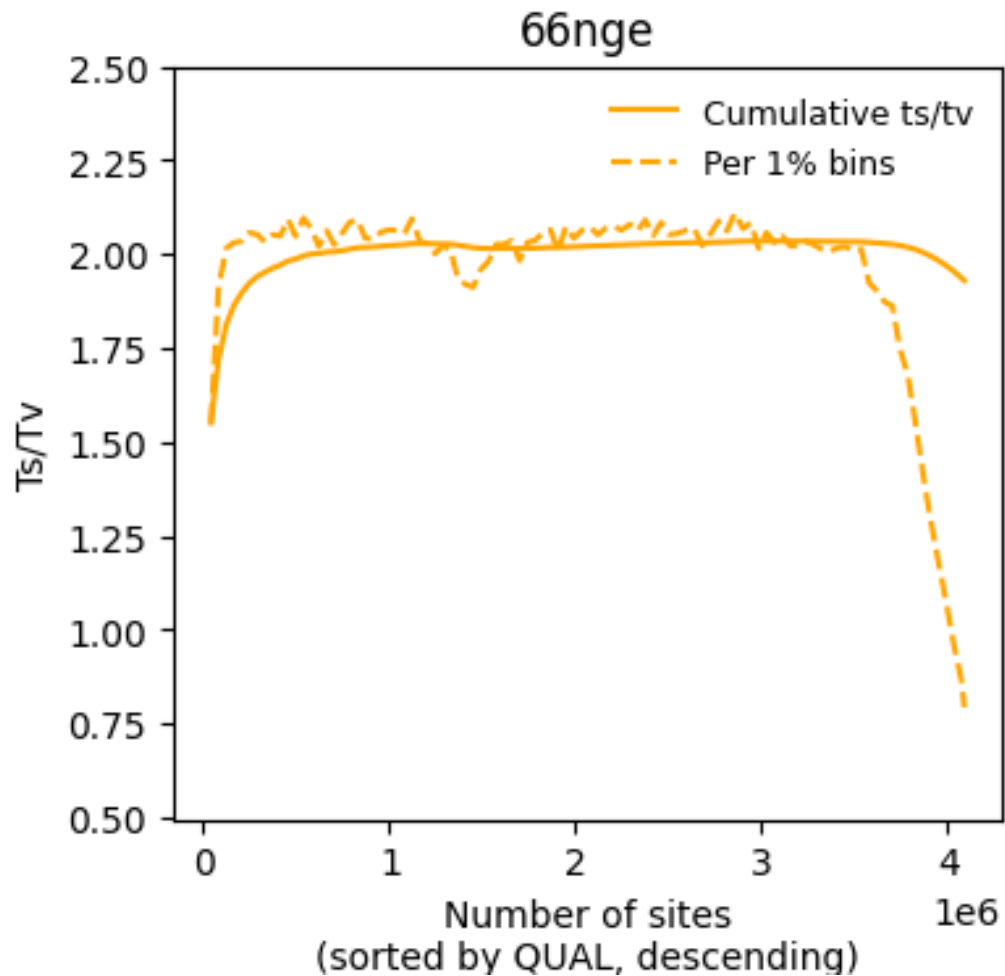
Number of SNPs by sample



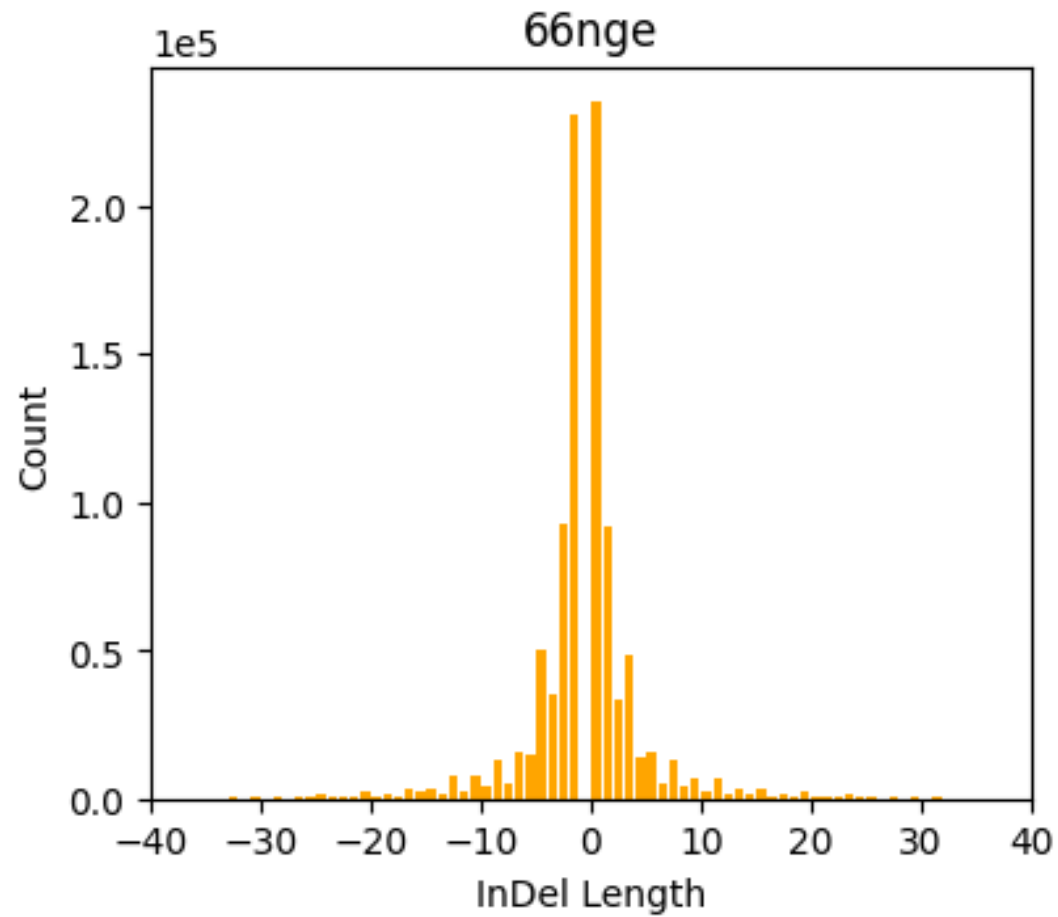
Number of indels by sample



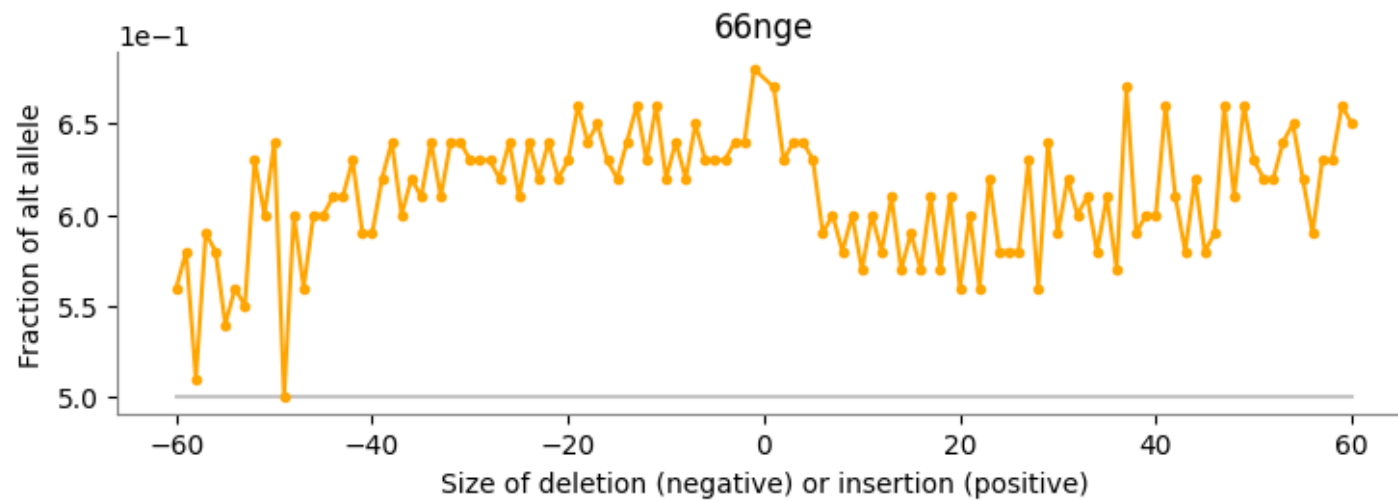
Ts/Tv stratified by QUAL



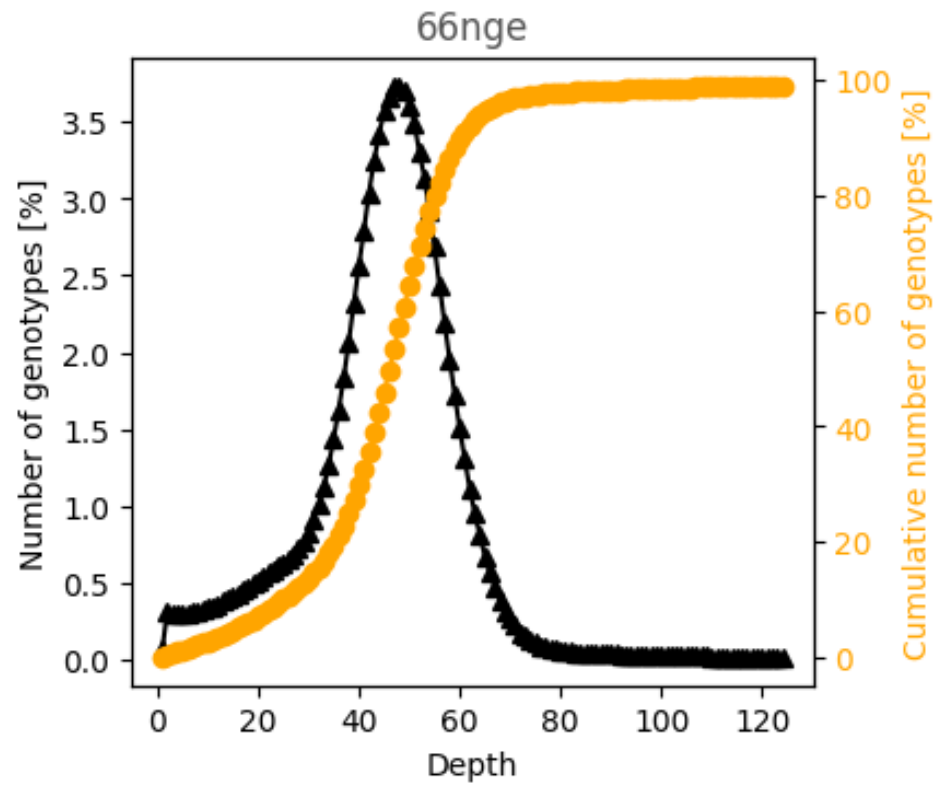
Indel distribution



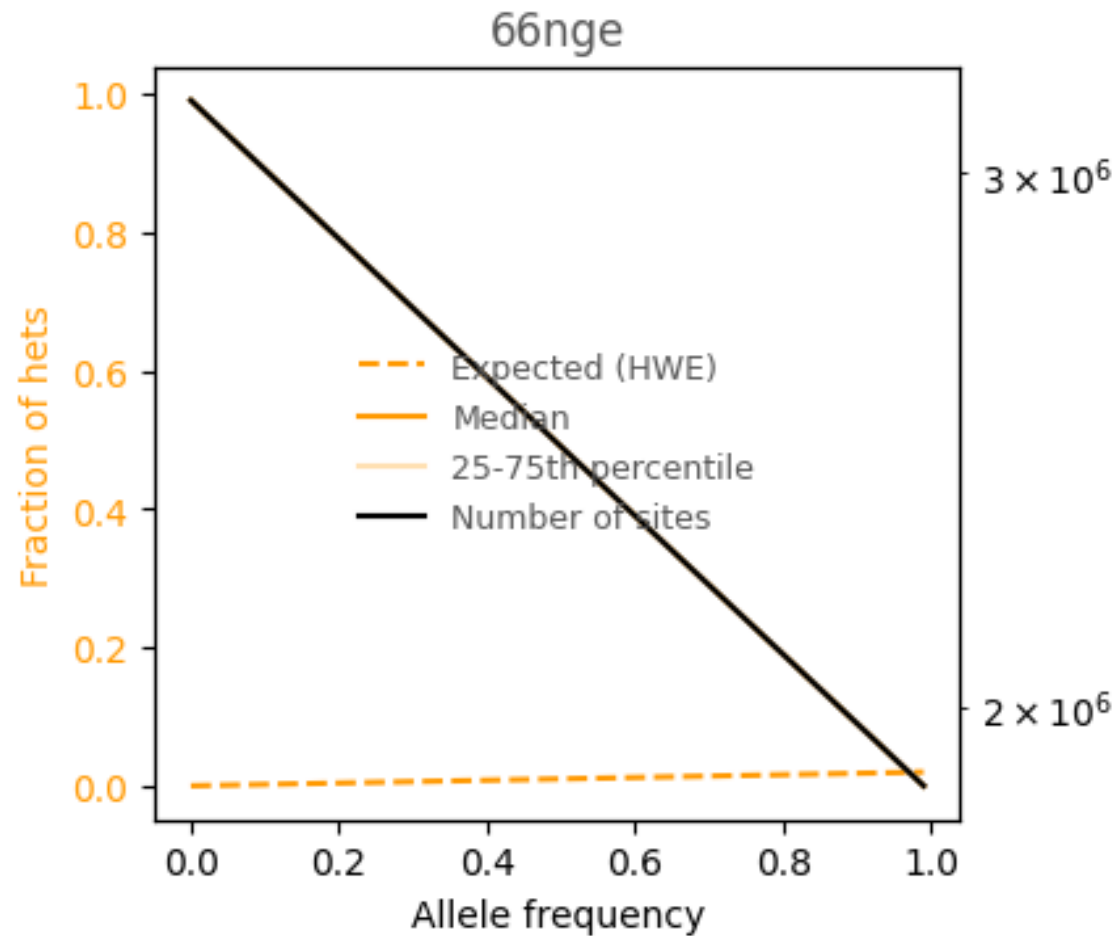
Fraction of alternate indel allele



Depth distribution



Number of HETs by AF



Substitution types

