

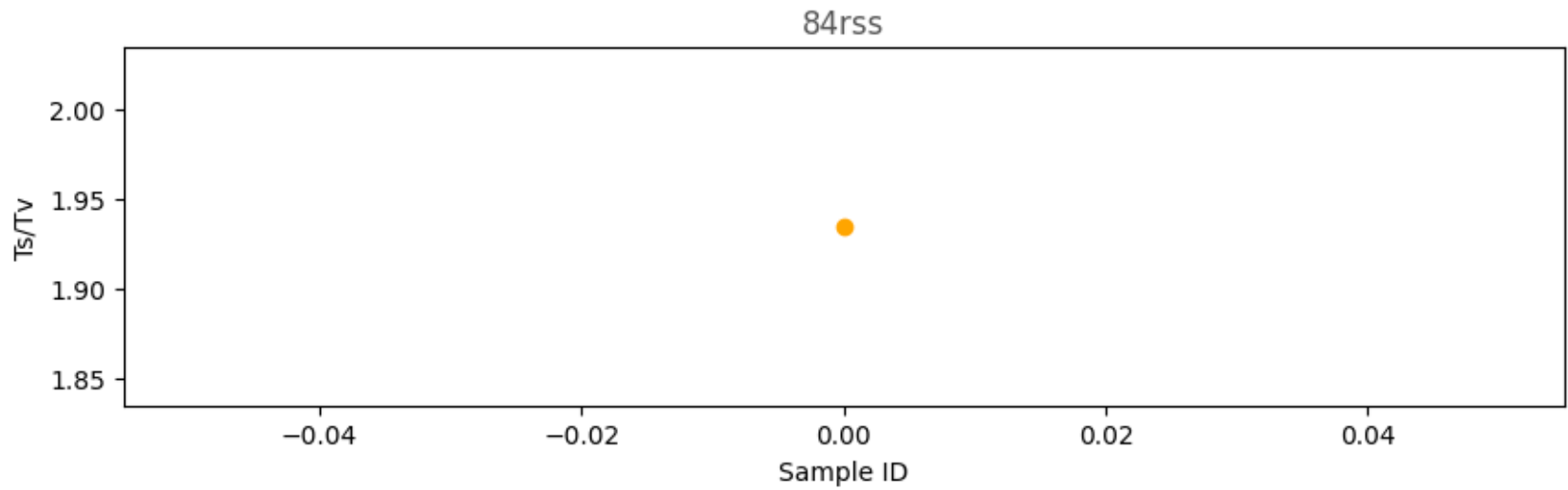
# Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
84rss	4,119,976	1.93	1.94	951,199	–	0	0
* frameshift ratio: out/(out+in)							

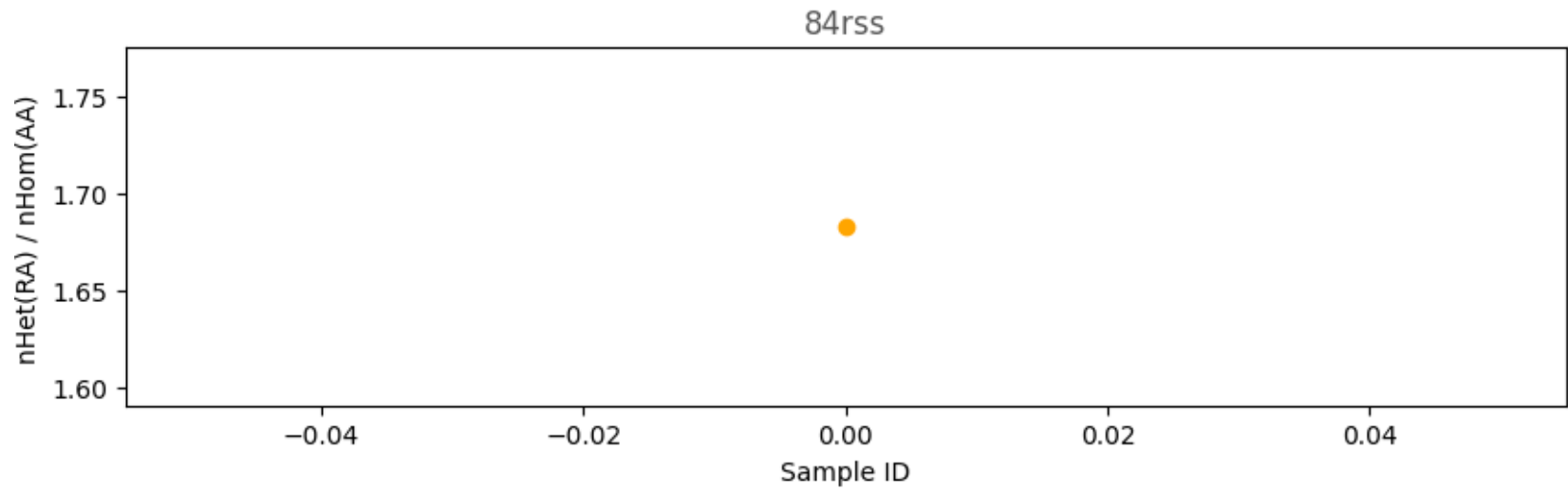
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
84rss	62.7%	1.91	67.3%	97,888	1,928

- 84rss .. /ngc/projects2/gm/data/archive/2022/variants/snv/84rssaldm-103893007134-Normal\_Blood\_noinfo-WGS\_v1\_IlluminaDNAPCRFree\_X-220720\_A00559\_AHMHLDX3-RHGM\_LABKA\_WGSA\_KUT-WGSAKUT04654\_22RKG016676x01\_snv\_germline\_raw.haplotype\_caller.vcf.gz

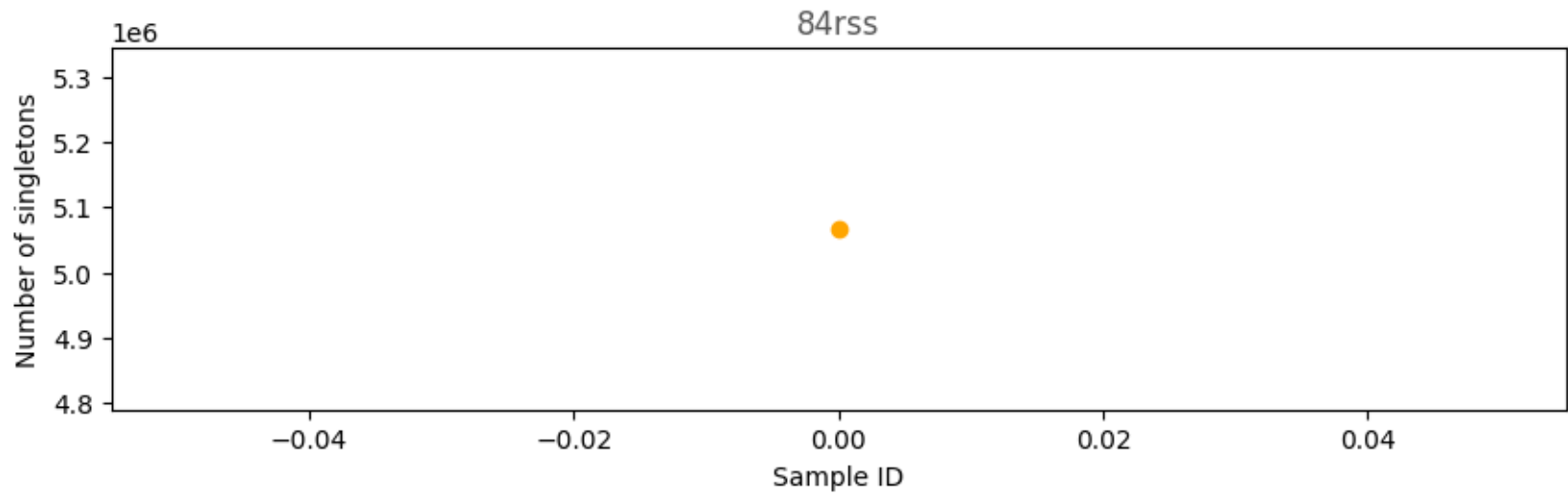
# Ts/Tv by sample



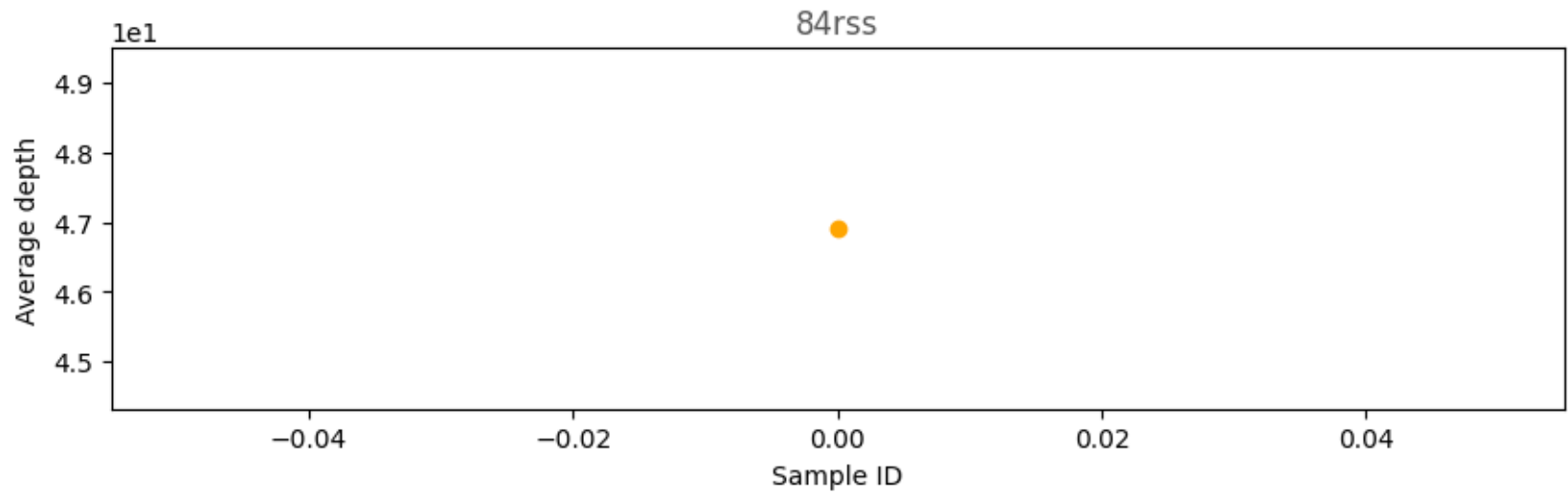
# Hets vs non-ref Homs by sample



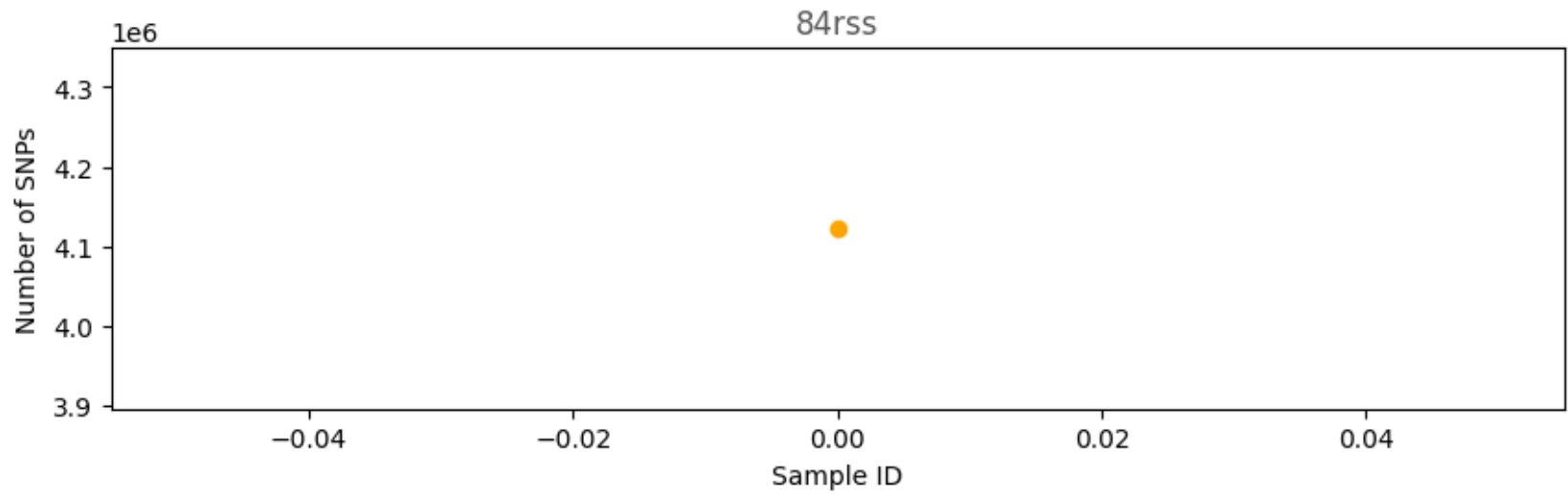
# Singletons by sample (hets and homs)



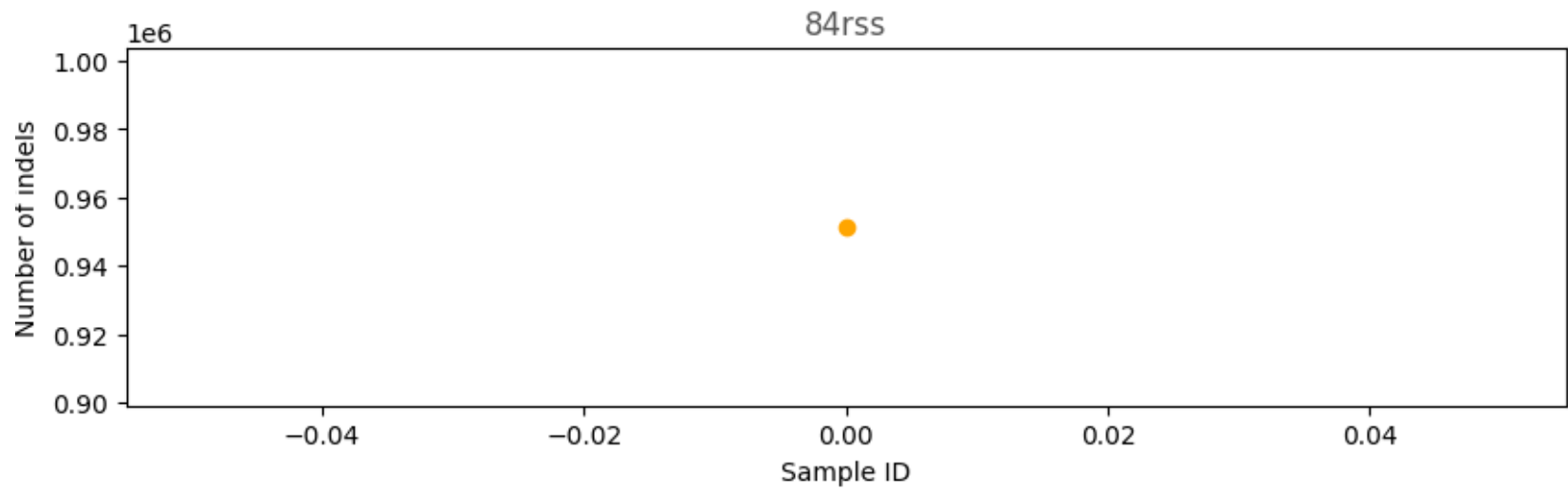
# Average depth by sample



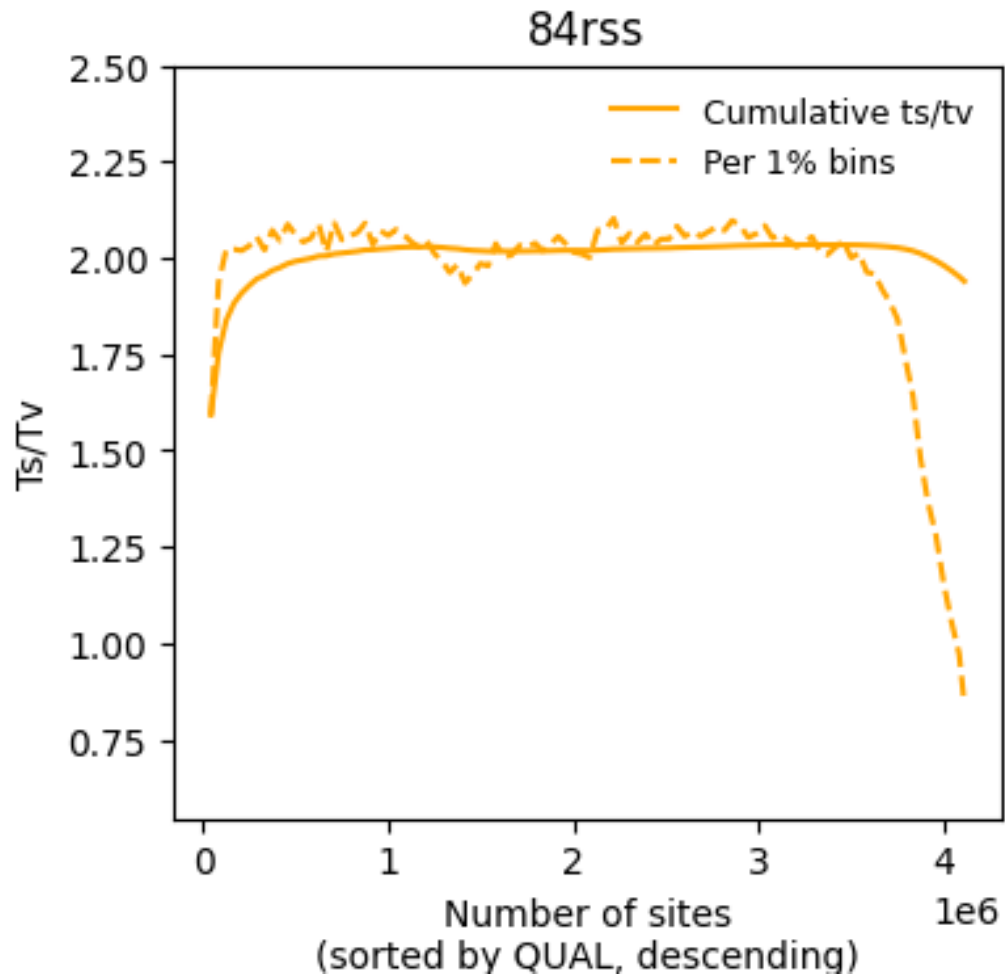
# Number of SNPs by sample



# Number of indels by sample

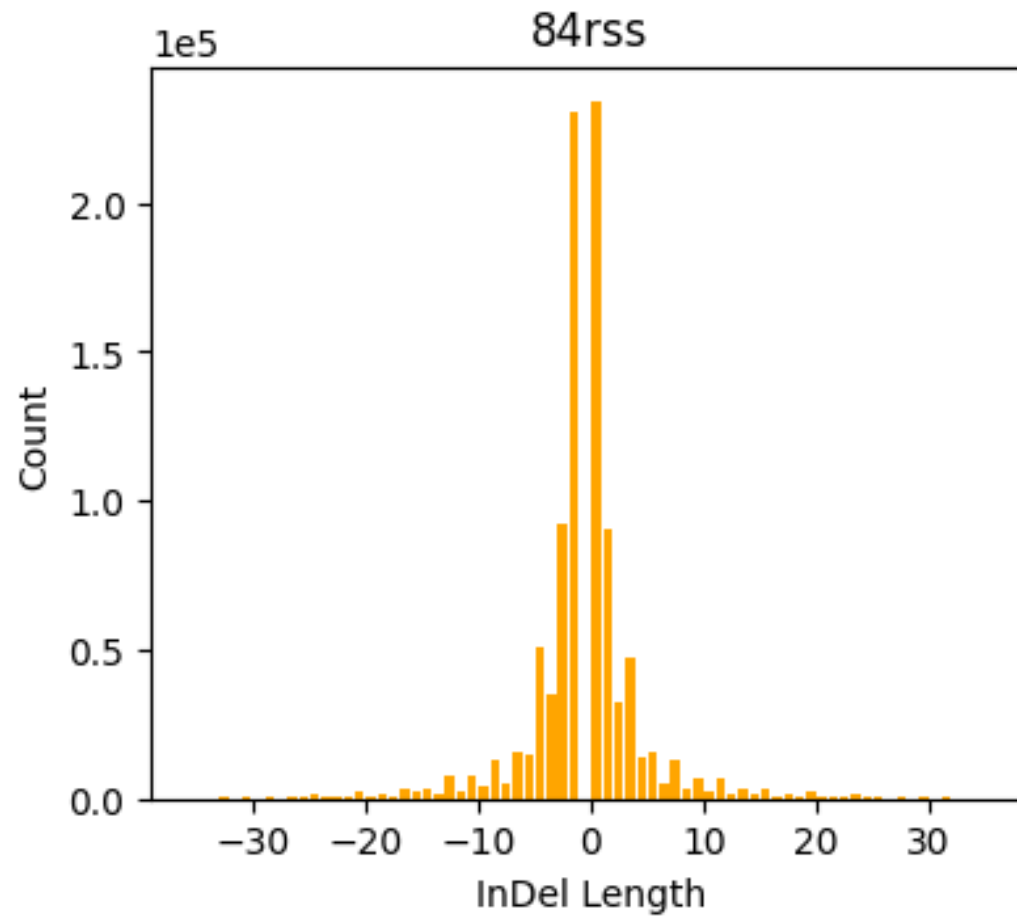


# Ts/Tv stratified by QUAL

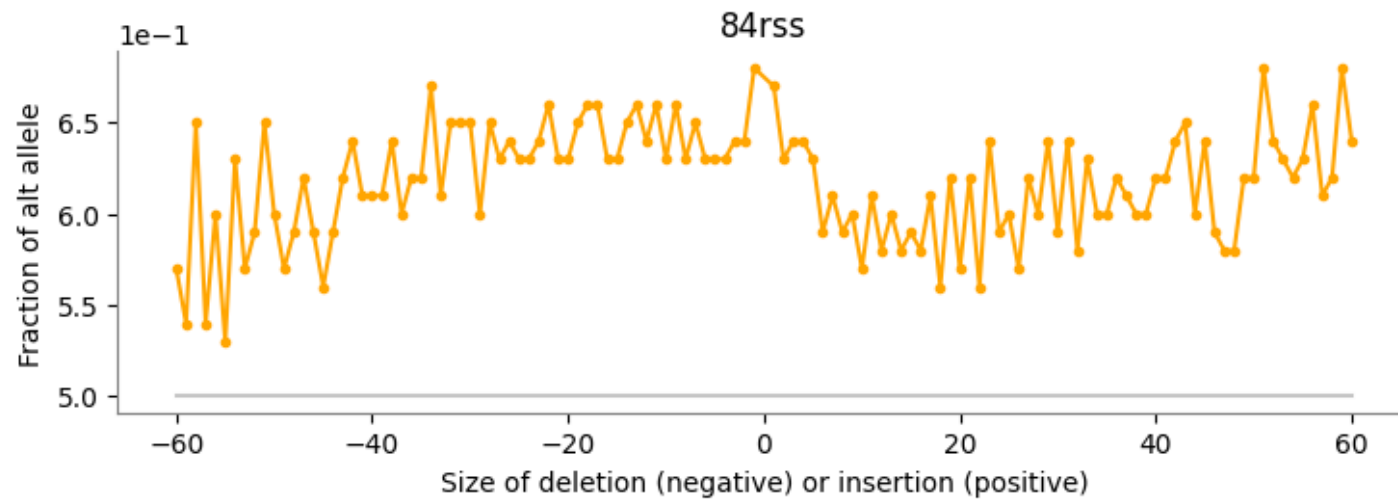




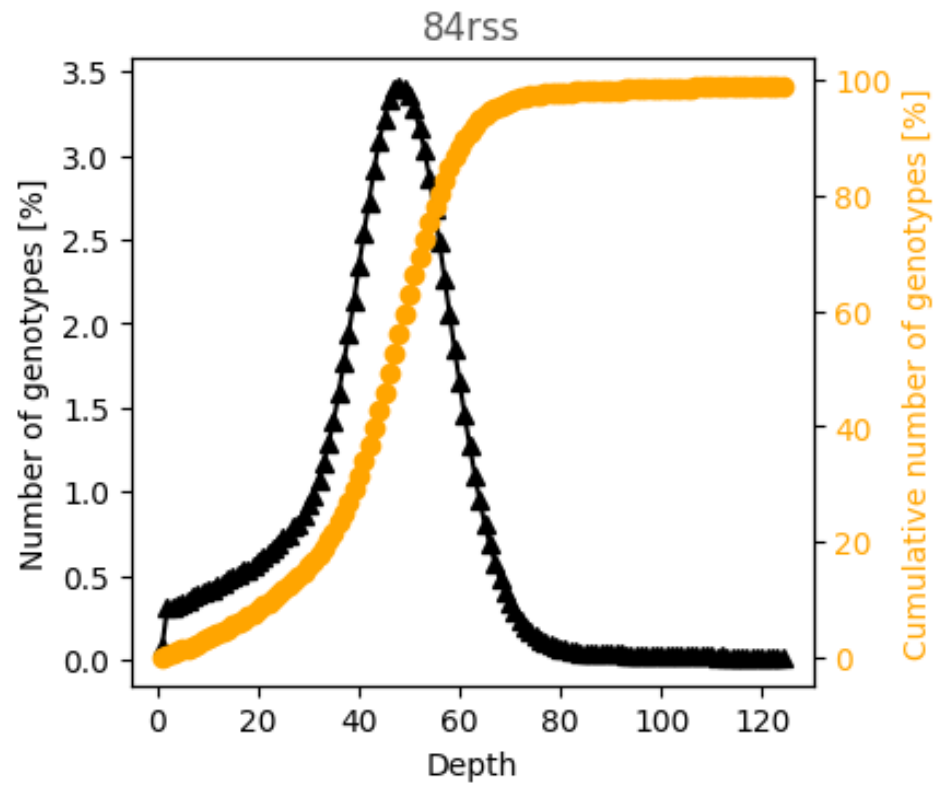
# Indel distribution



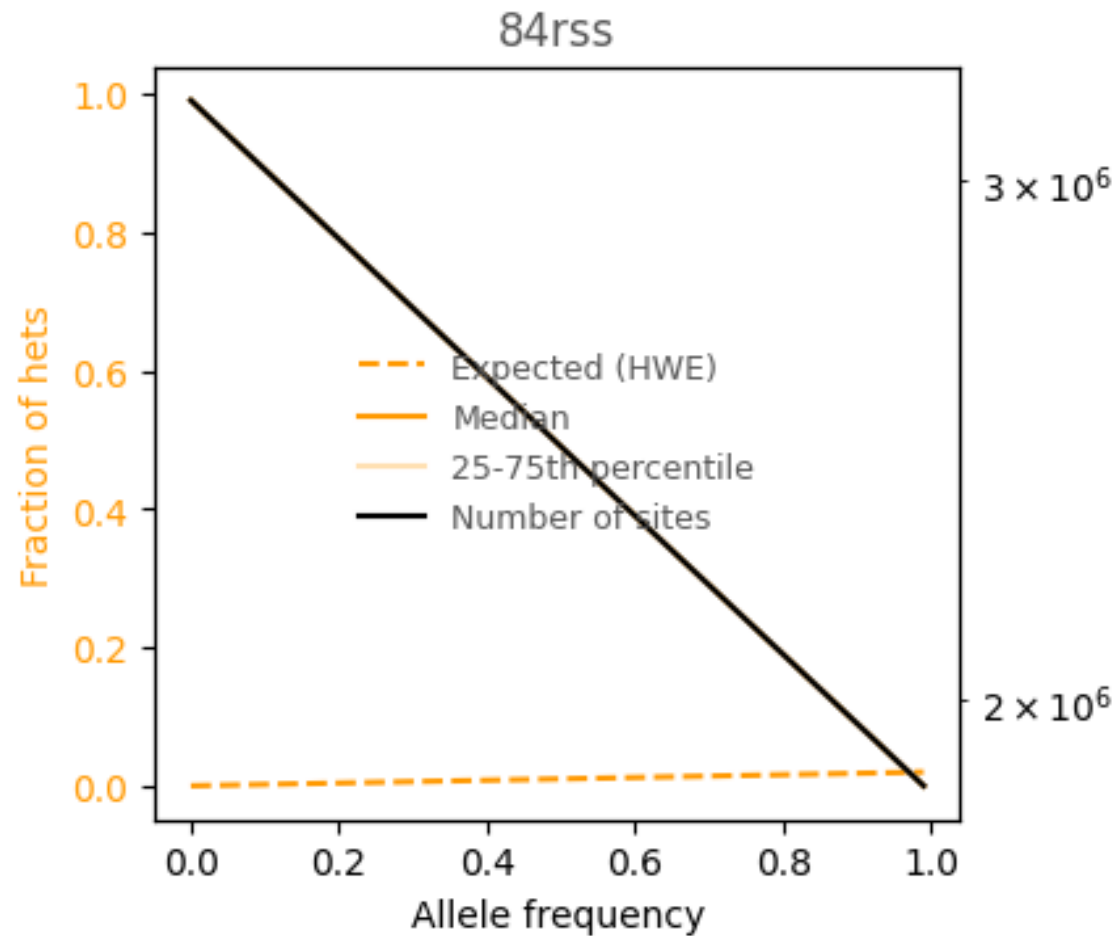
# Fraction of alternate indel allele



# Depth distribution



# Number of HETs by AF



# Substitution types

