

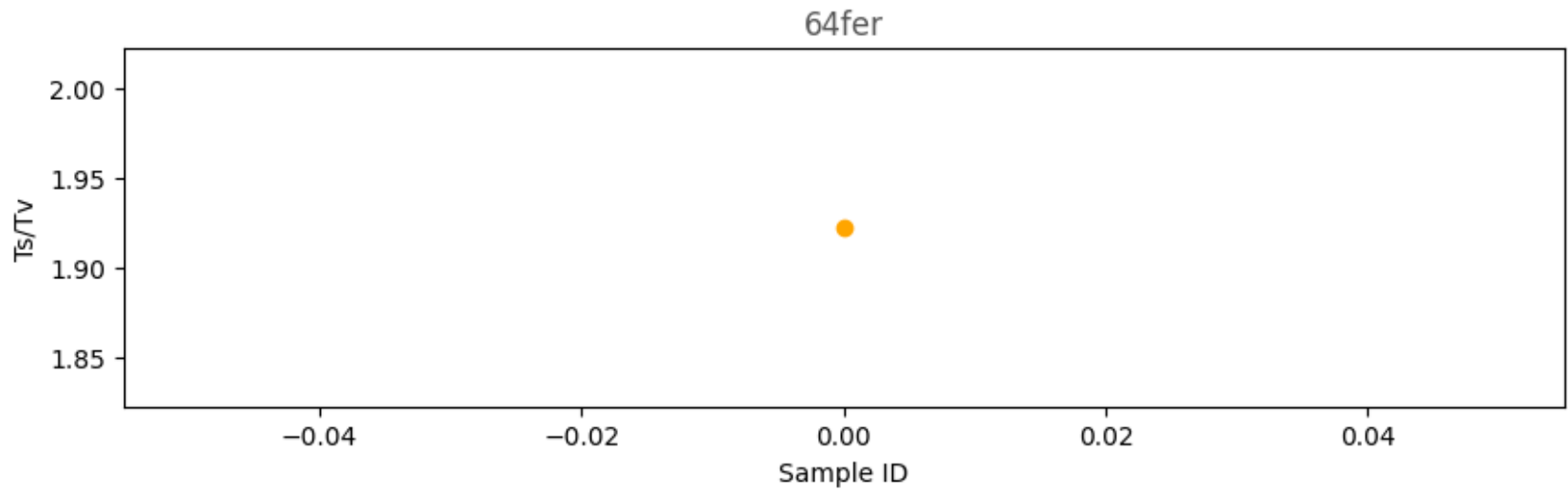
# Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
64fer	4,160,956	1.92	1.93	978,625	–	0	0
* frameshift ratio: out/(out+in)							

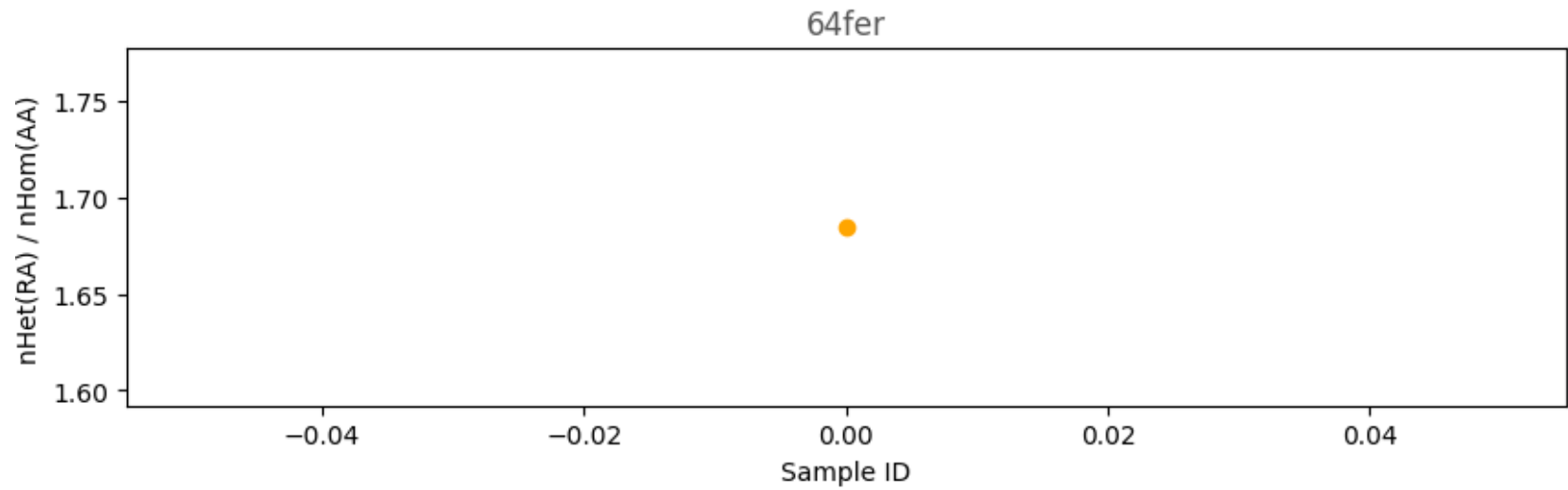
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
64fer	62.8%	1.89	68.3%	110,143	2,023

- 64fer .. /ngc/projects2/gm/data/archive/2022/variants/snv/64ferndrm-103736023985-Normal\_Blood\_noinfo-WGS\_v1\_IlluminaDNAPCRFree\_RHGM00133-210510\_A01176\_AH5727DSX2-EXT\_LAB  
KA\_NGCWGS-NGCWGS00048\_64xk8wmk3-snv\_germline\_raw.haplotype\_caller.vcf.gz

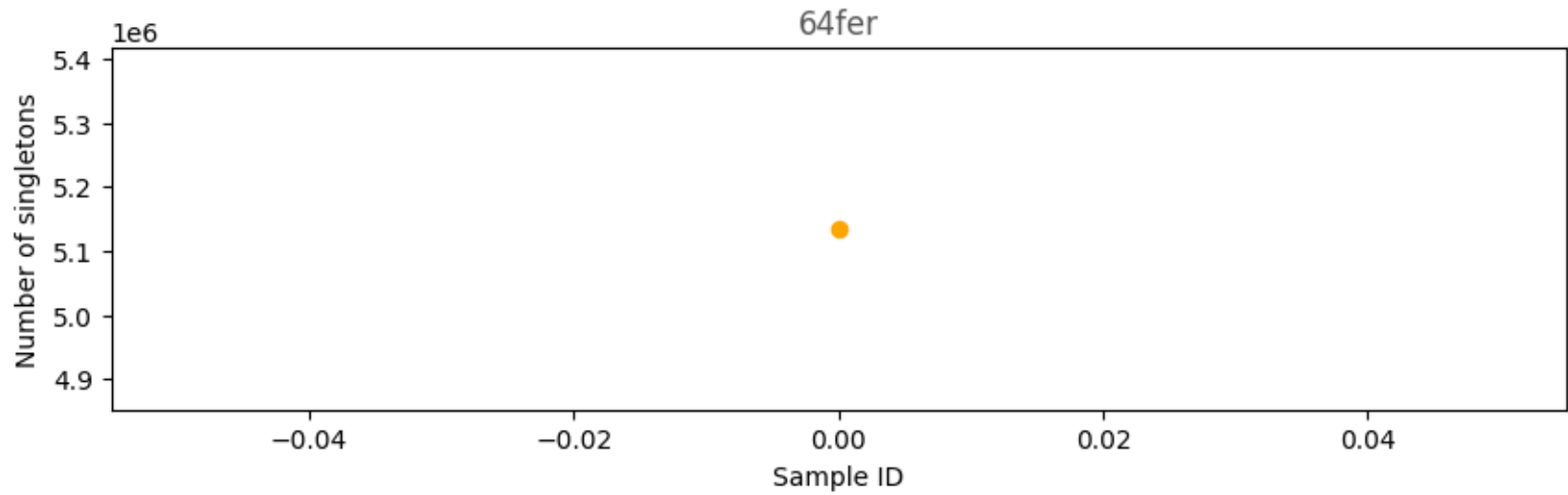
# Ts/Tv by sample



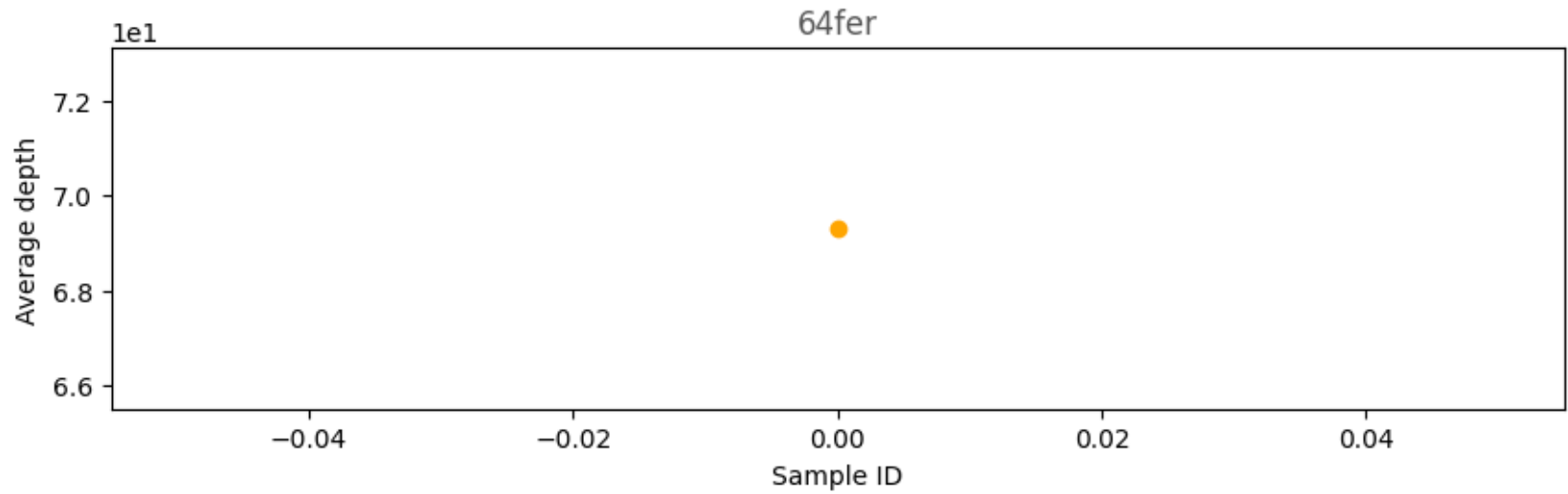
# Hets vs non-ref Homs by sample



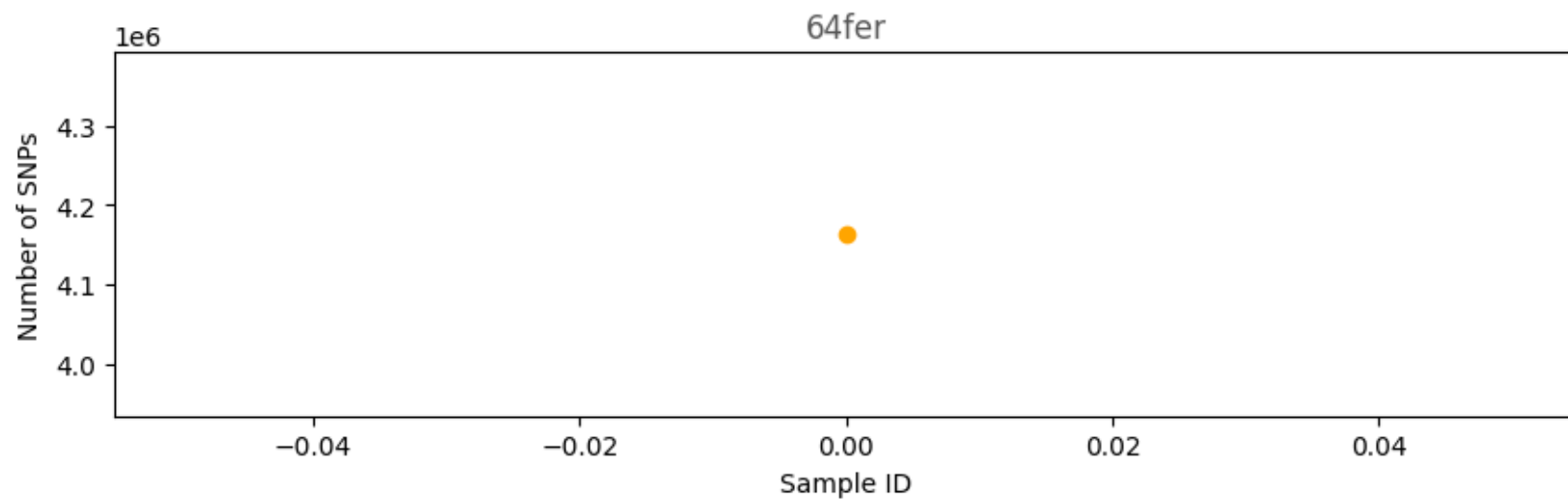
# Singletons by sample (hets and homs)



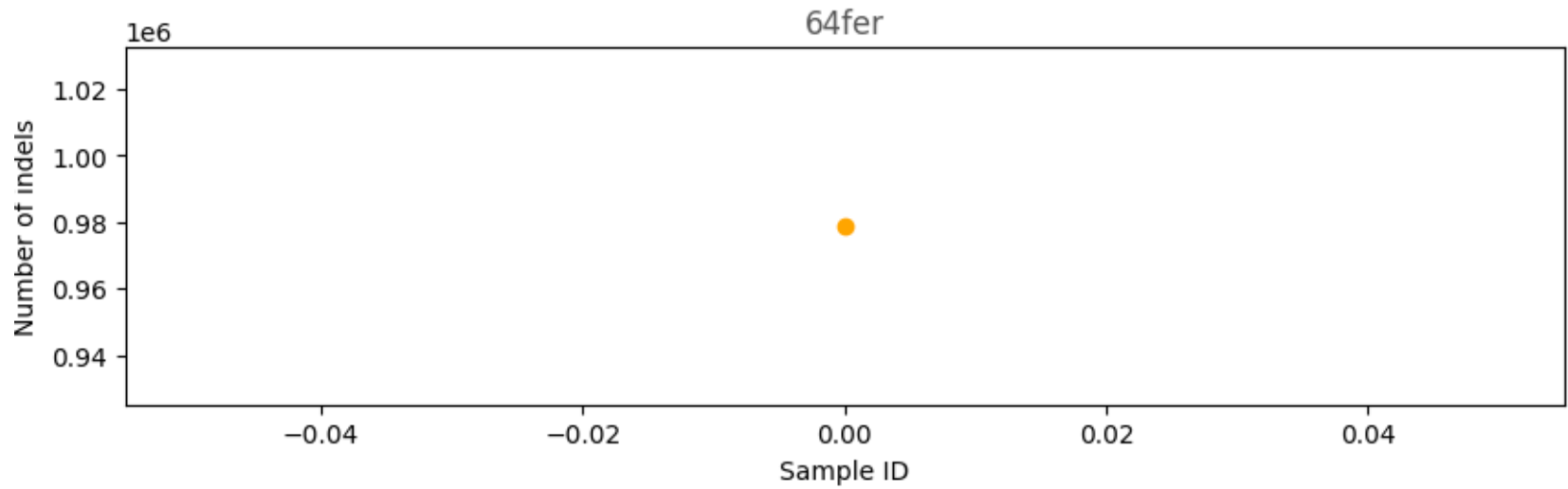
# Average depth by sample



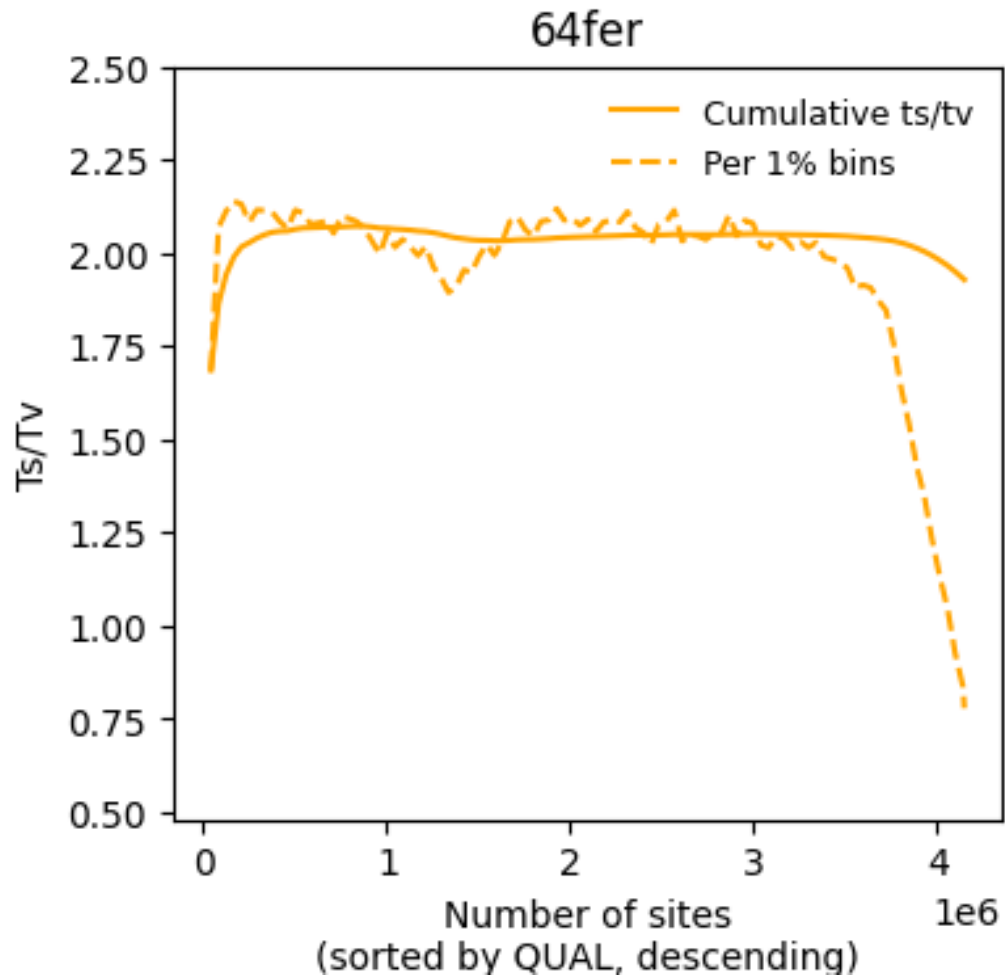
# Number of SNPs by sample



# Number of indels by sample

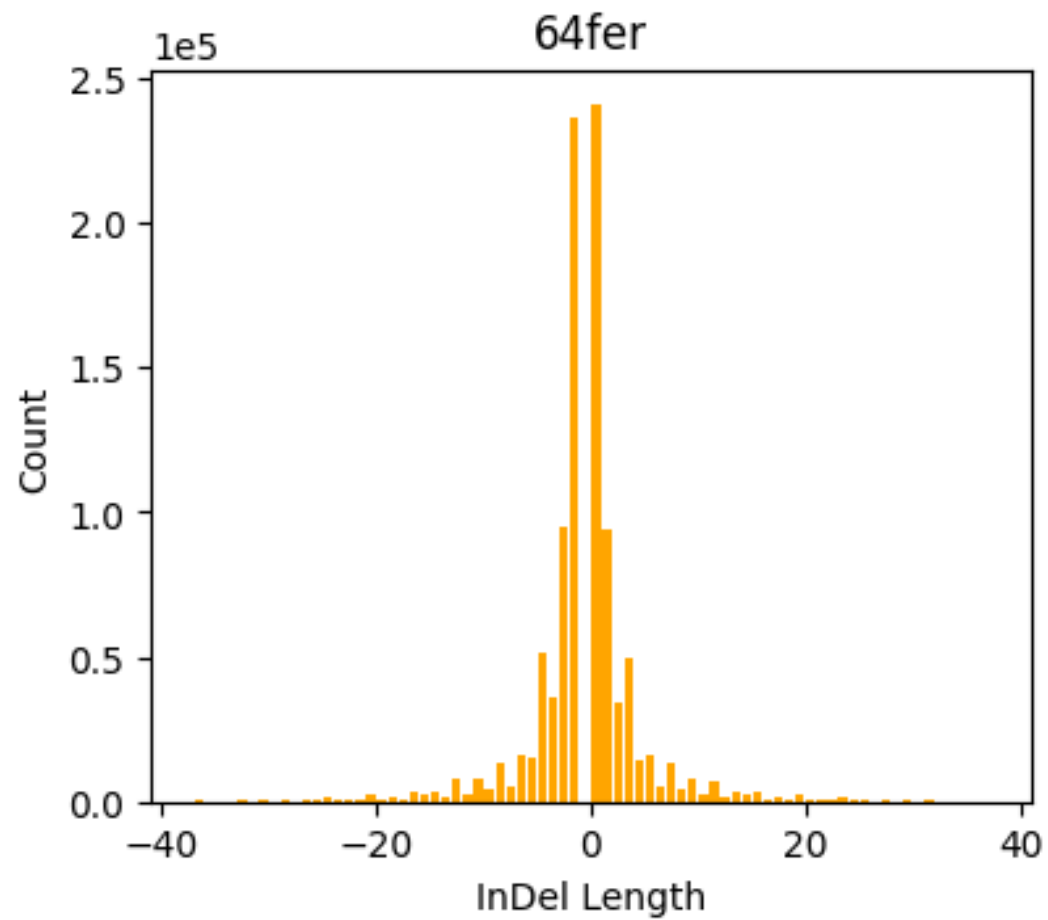


## Ts/Tv stratified by QUAL

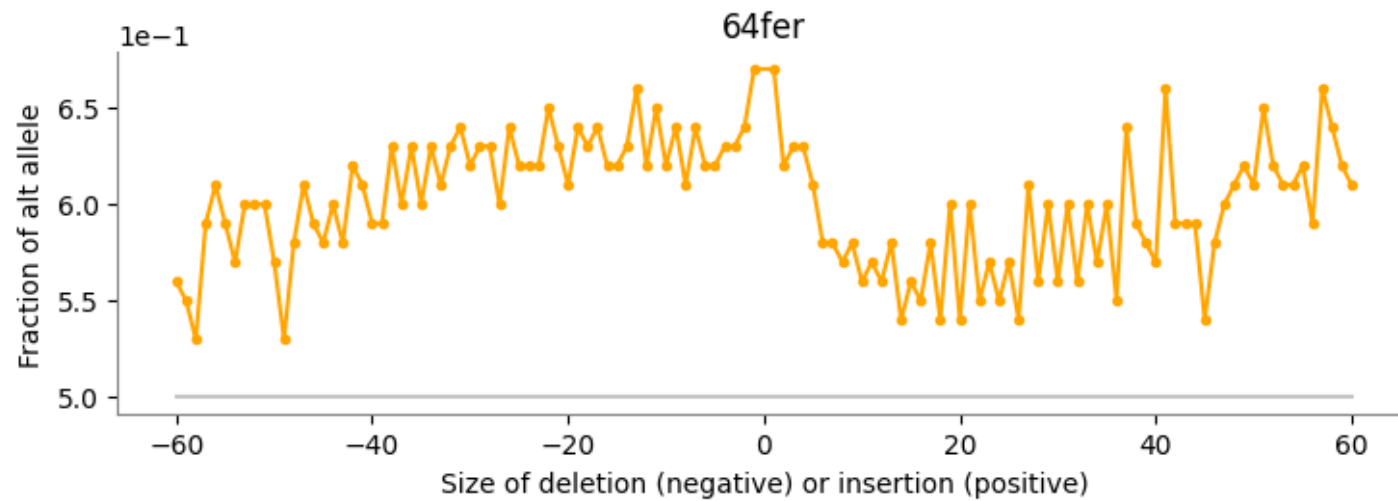




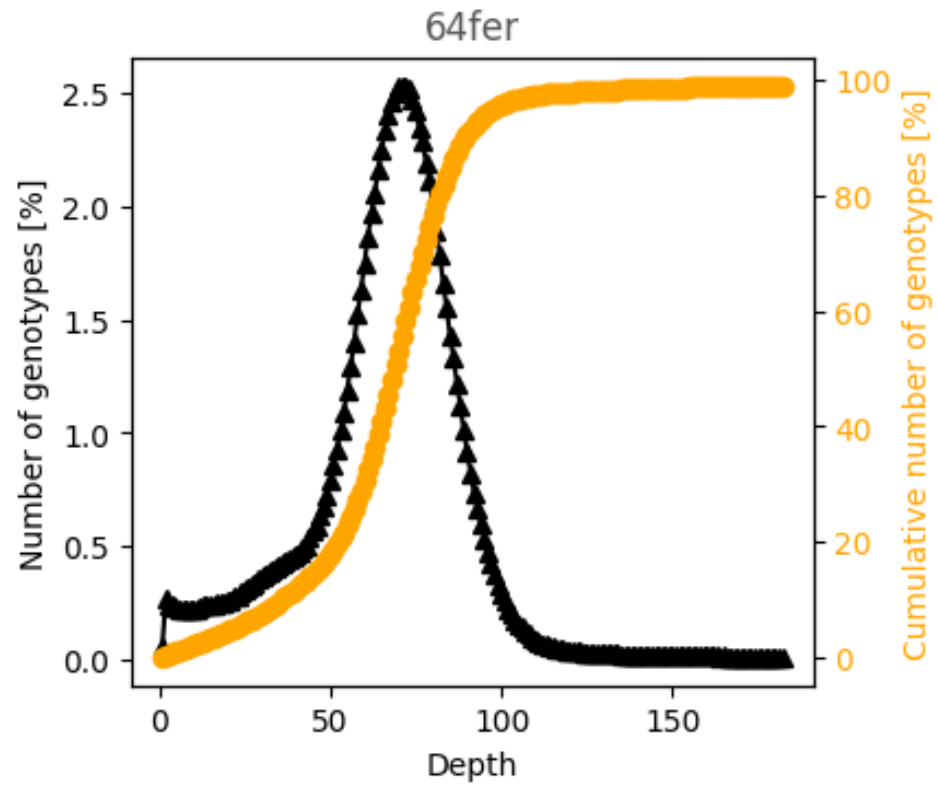
# Indel distribution



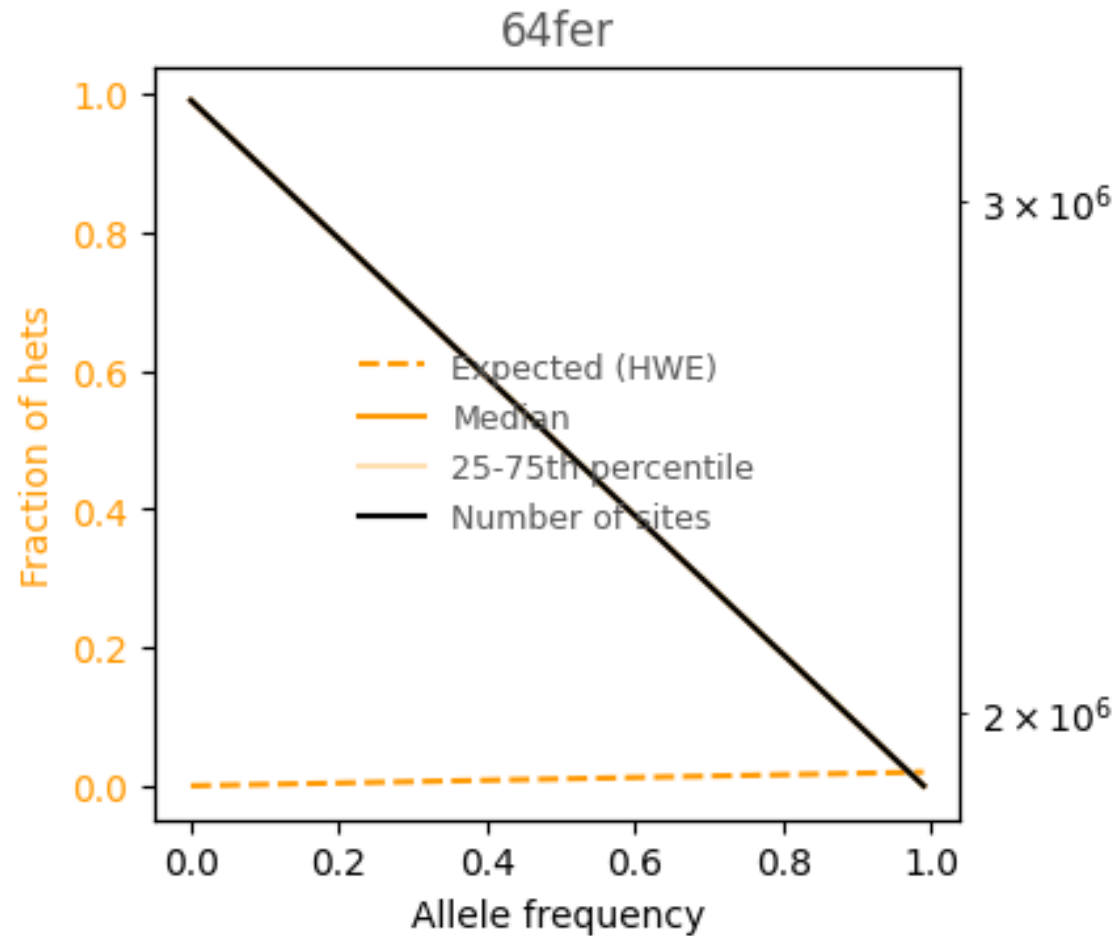
# Fraction of alternate indel allele



# Depth distribution



# Number of HETs by AF



# Substitution types

