

# Summary Numbers

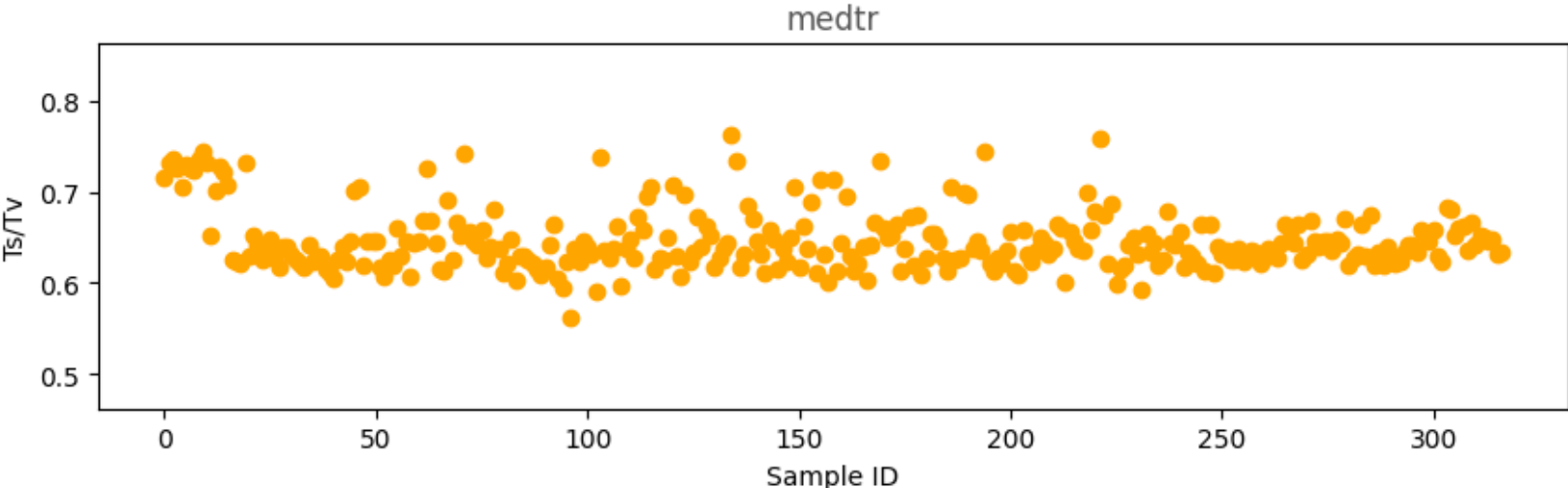
Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
medtr	48,997,264	1.30	1.37	4,845,927	–	8,064,865	2,191,312

\* frameshift ratio: out/(out+in)

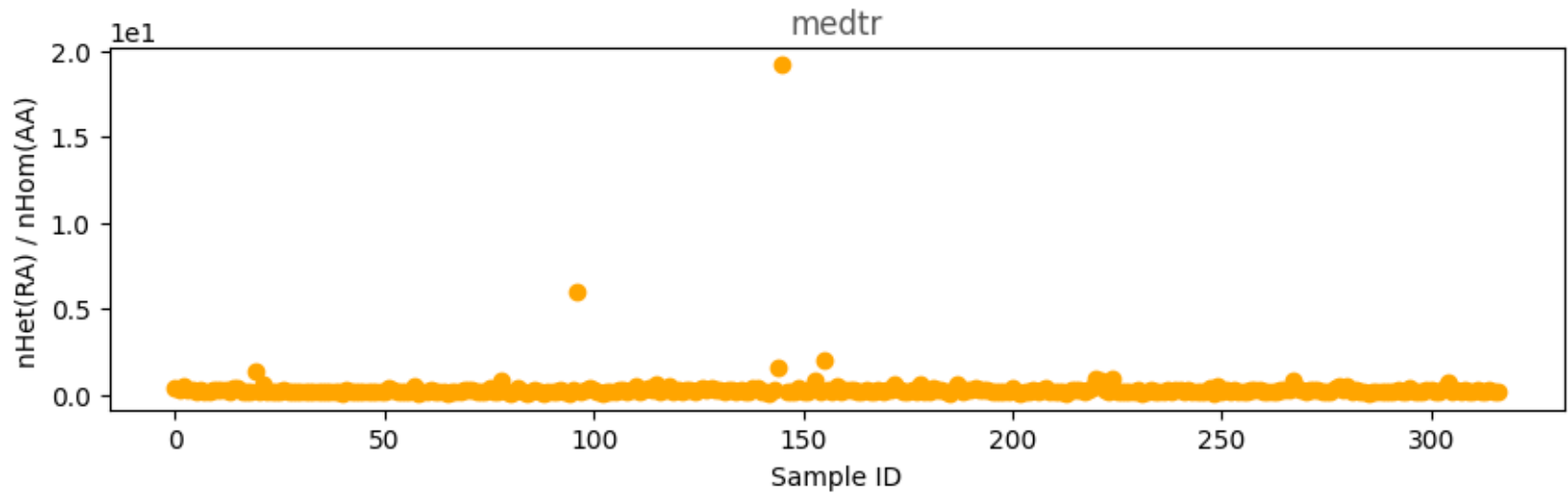
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
medtr	12.8%	0.79	6.1%	12,789,268	2,636,418

- medtr .. medtr.A17.gnm5.div.Epstein\_Burghardt\_2022.Mt5\_raw\_variant\_calls.2021-09-29.vcf.g  
z

# Ts/Tv by sample

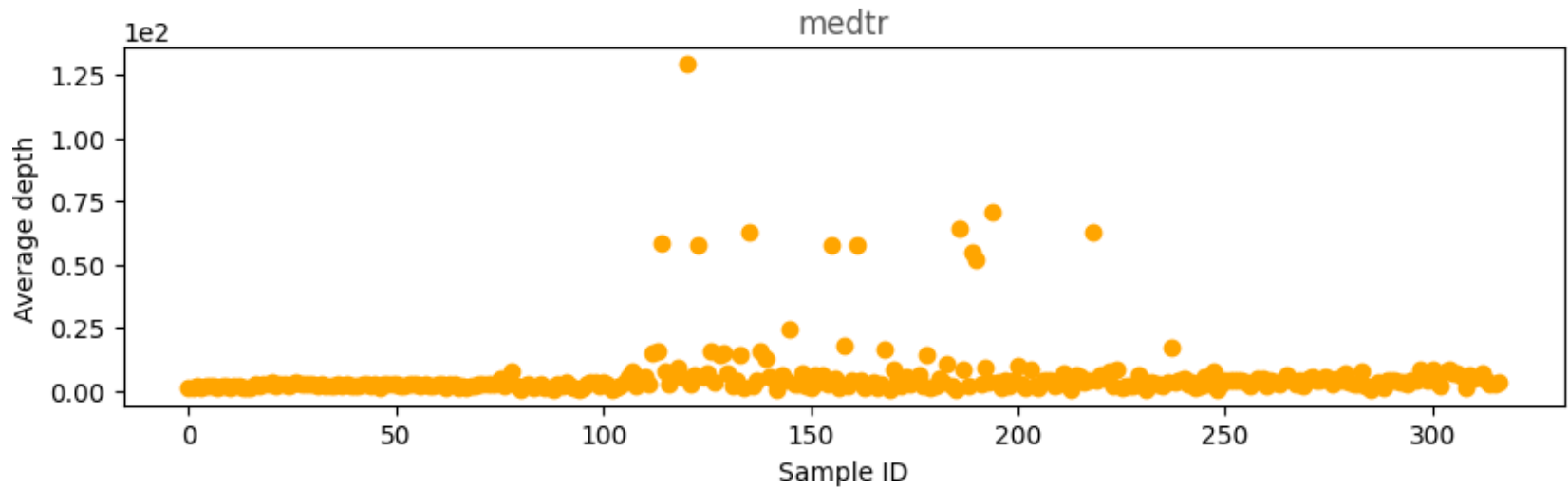


# Hets vs non-ref Homs by sample

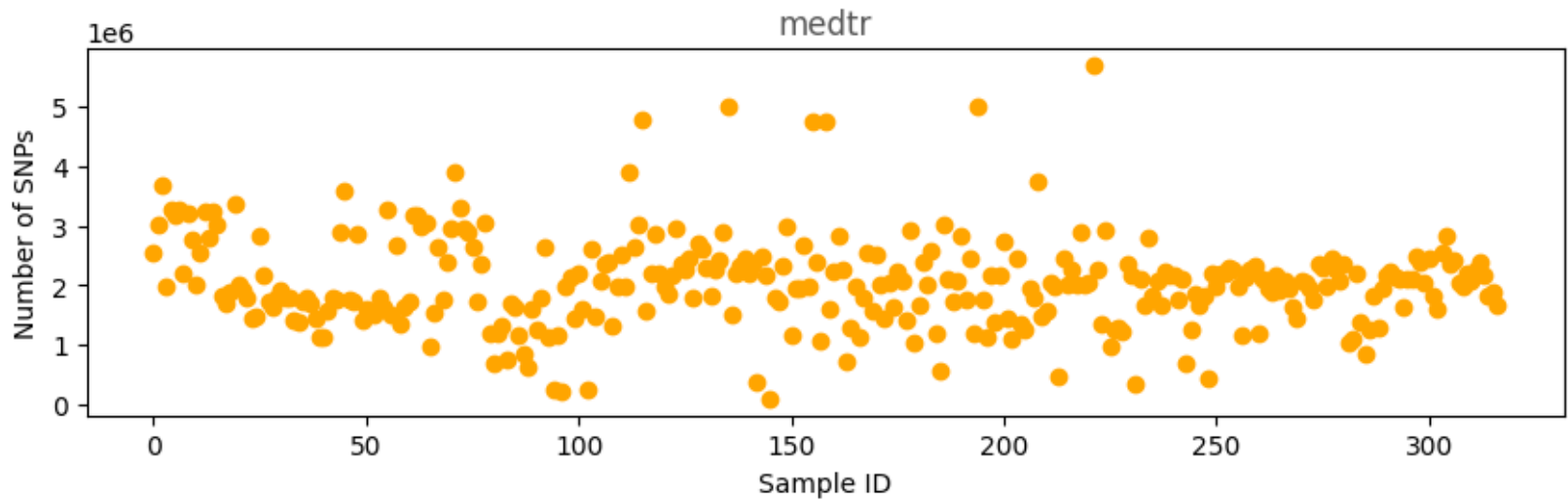




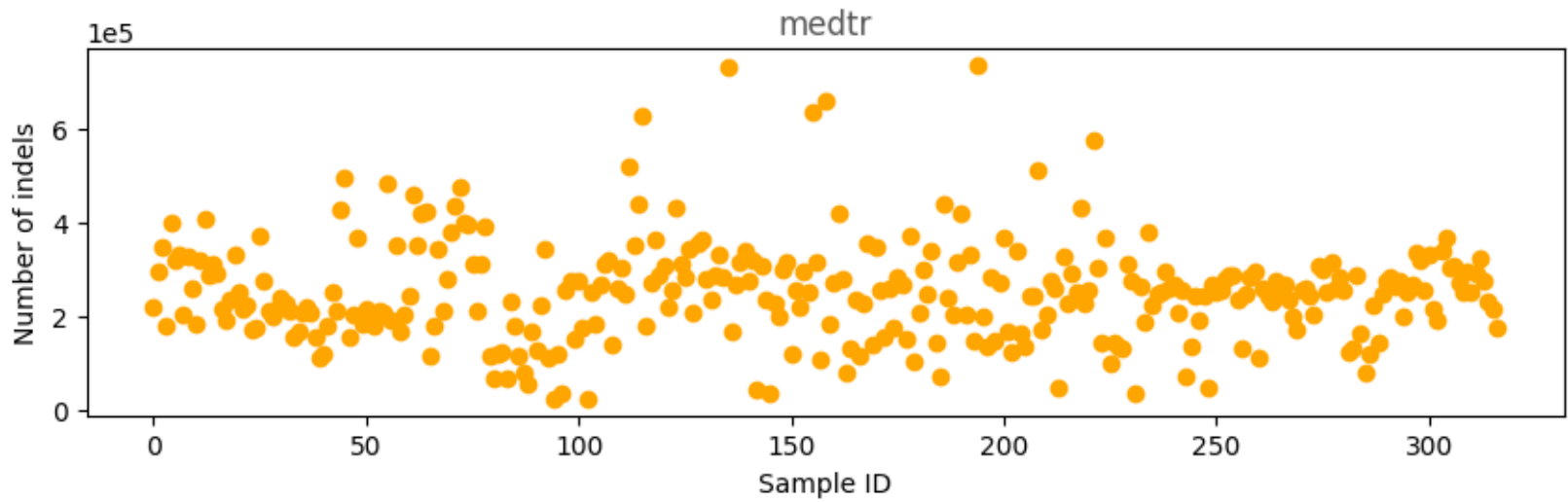
# Average depth by sample



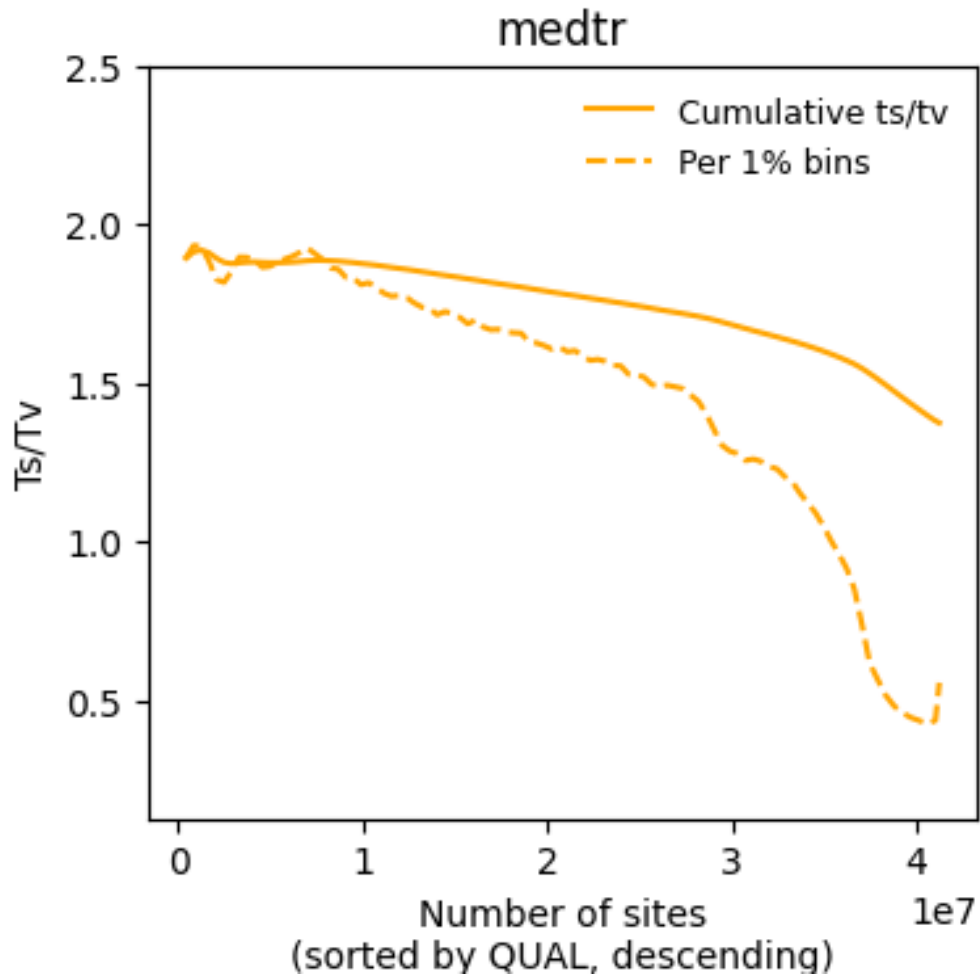
# Number of SNPs by sample



# Number of indels by sample

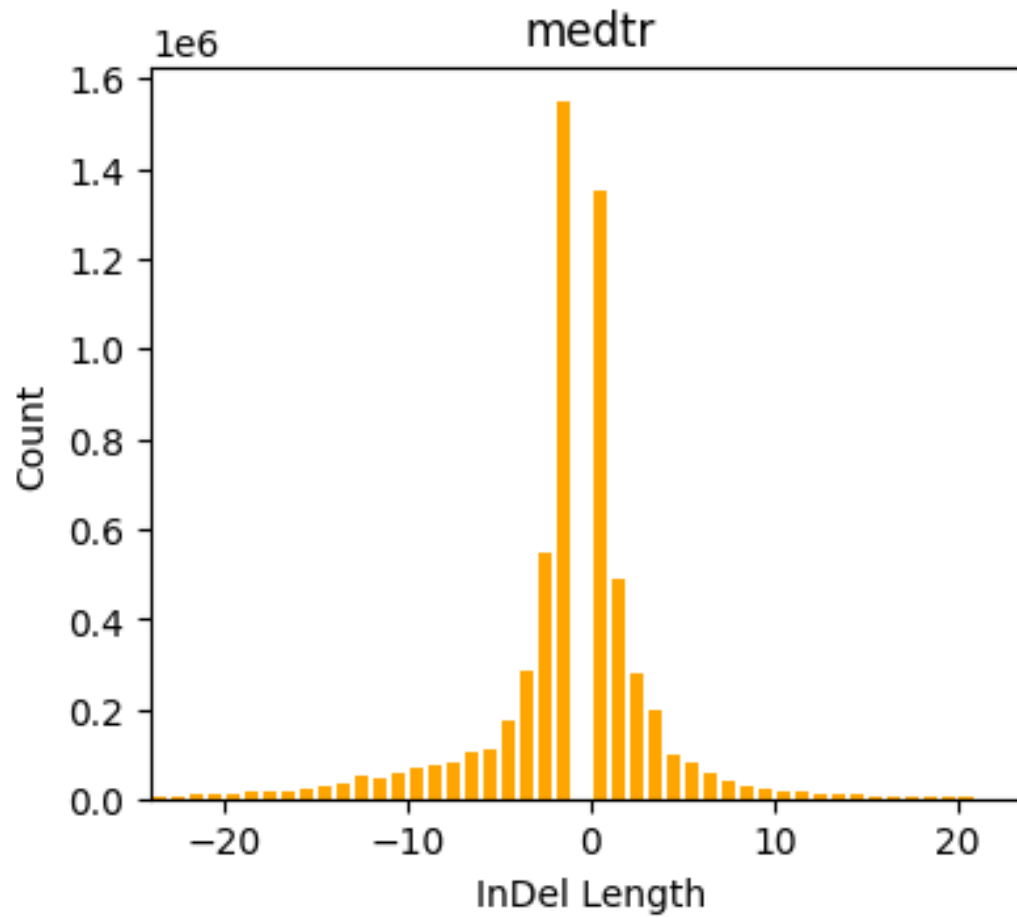


# Ts/Tv stratified by QUAL





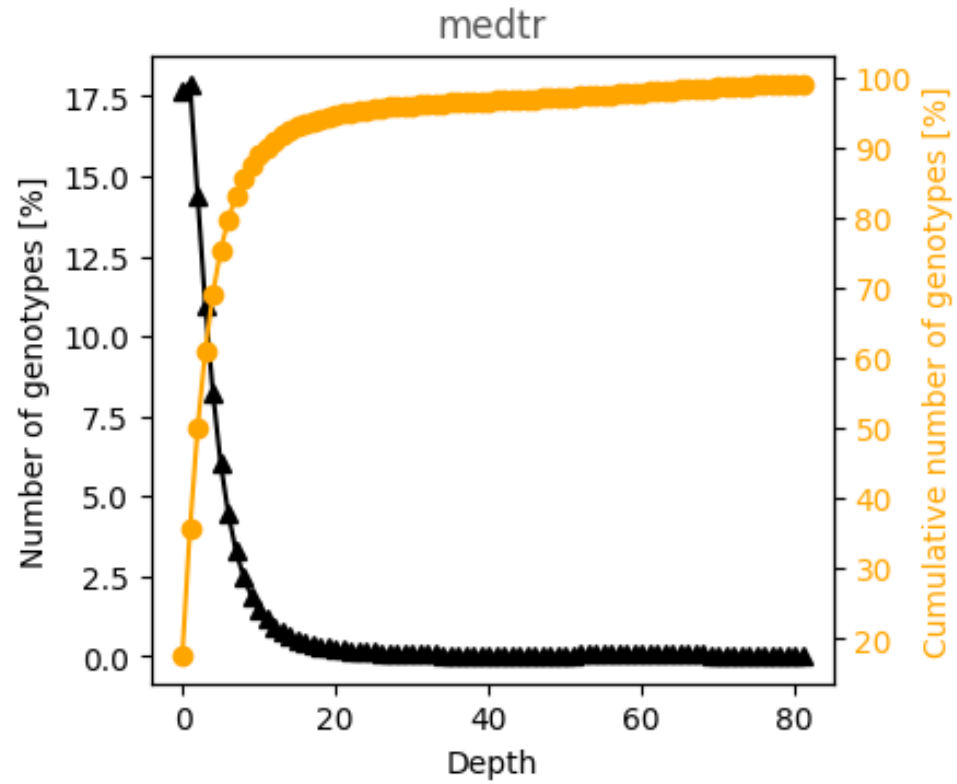
# Indel distribution



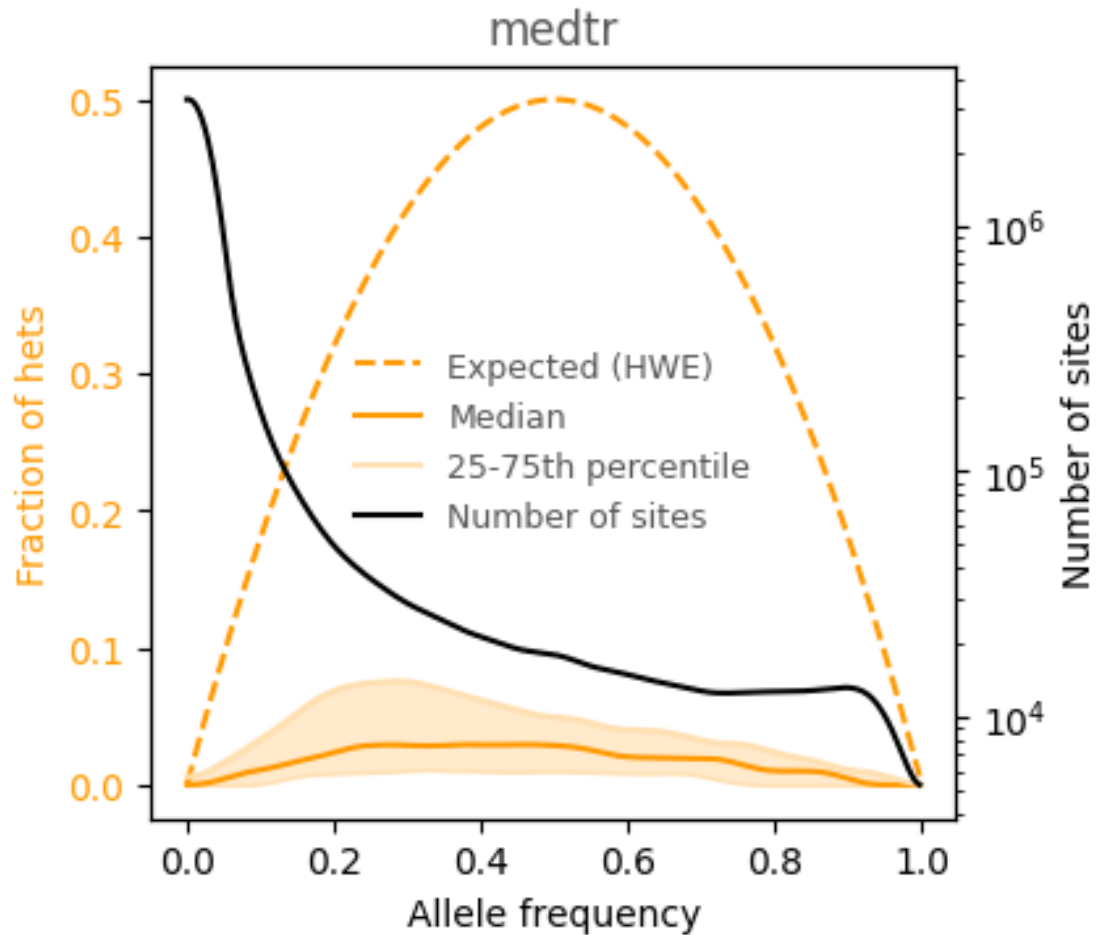
# Fraction of alternate indel allele



# Depth distribution



# Number of HETs by AF



# Substitution types

