

# Summary Numbers

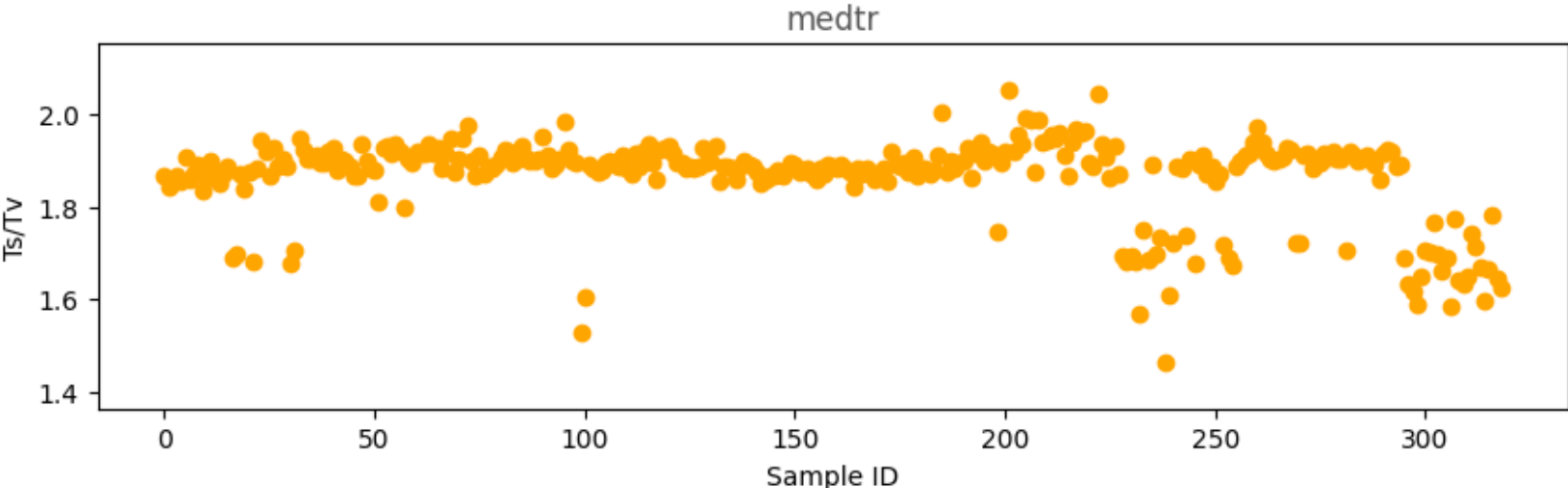
Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
medtr	44,505,624	1.65	1.73	5,195,606	-	0	0

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

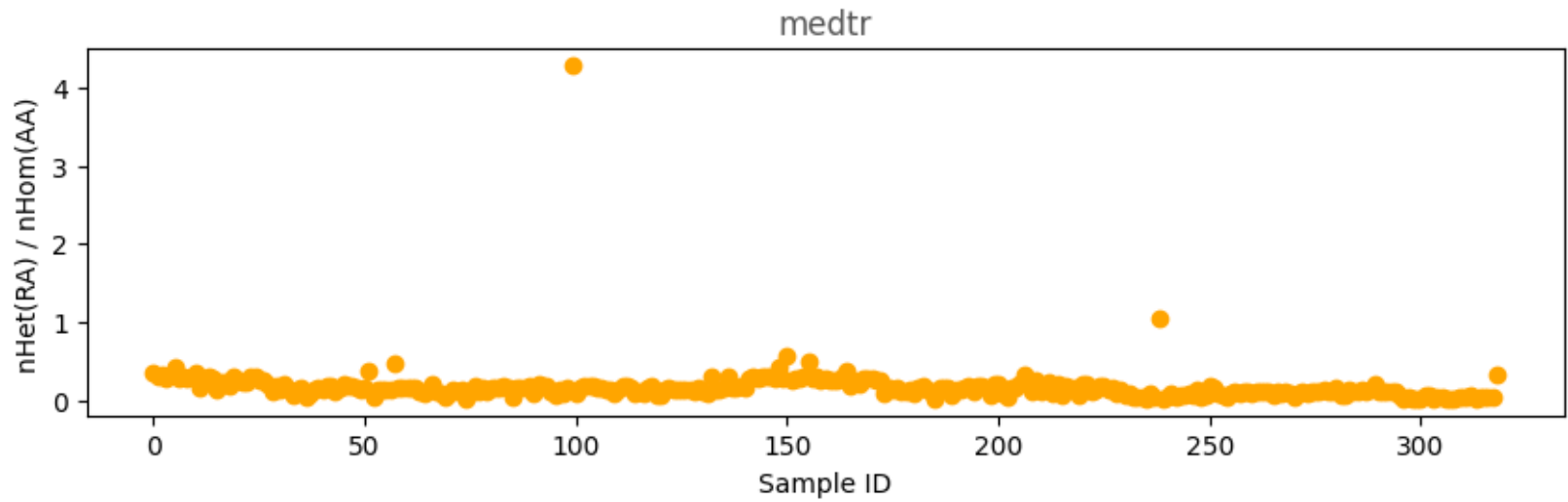
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
medtr	2.4%	1.13	0.9%	2,930,348	2,397,624

- medtr .. medtr.A17\_HM341.gnm4.div.Stanton-Geddes\_Paape\_2013.everything-2014Apr20.vcf.gz

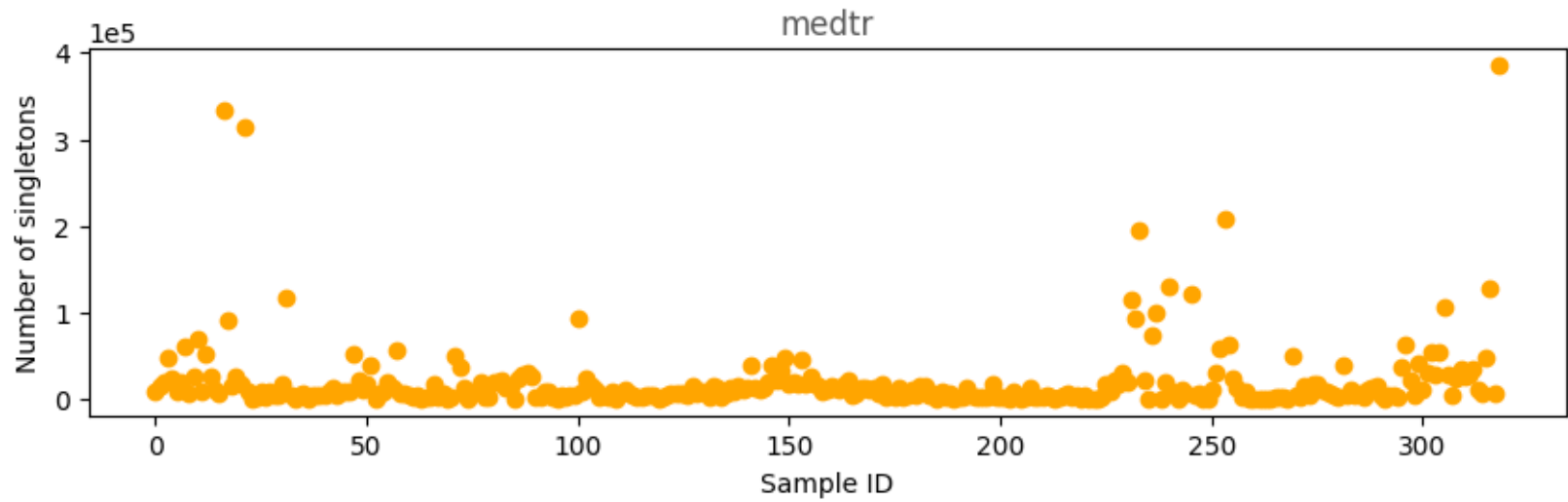
# Ts/Tv by sample



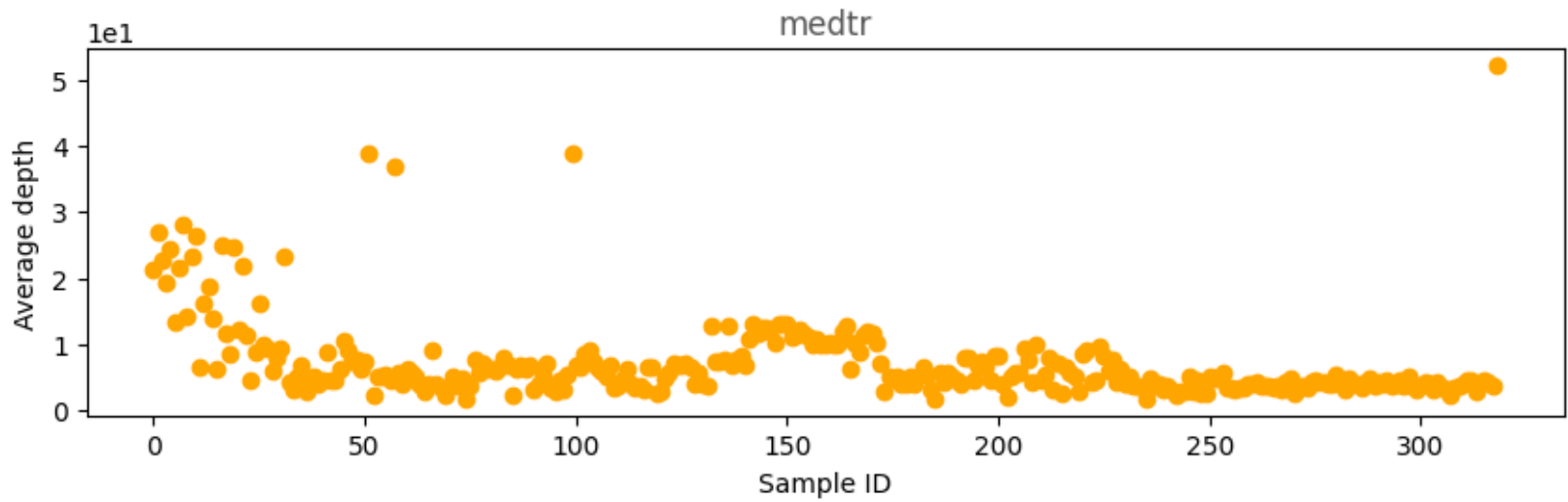
# Hets vs non-ref Homs by sample



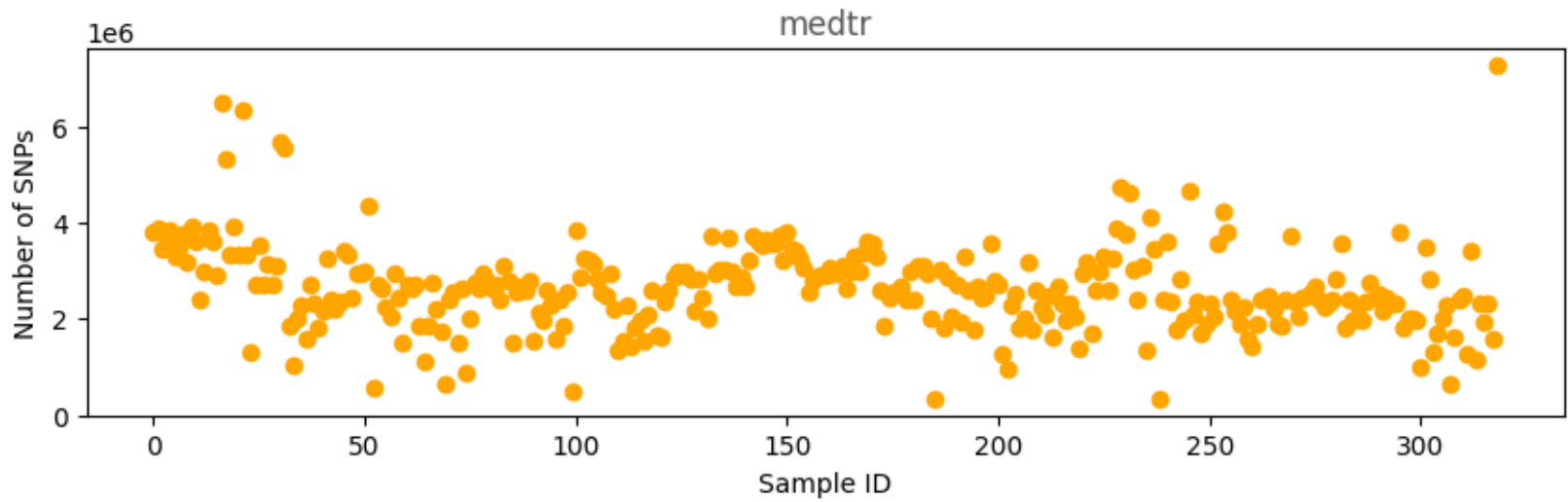
# Singletons by sample (hets and homs)



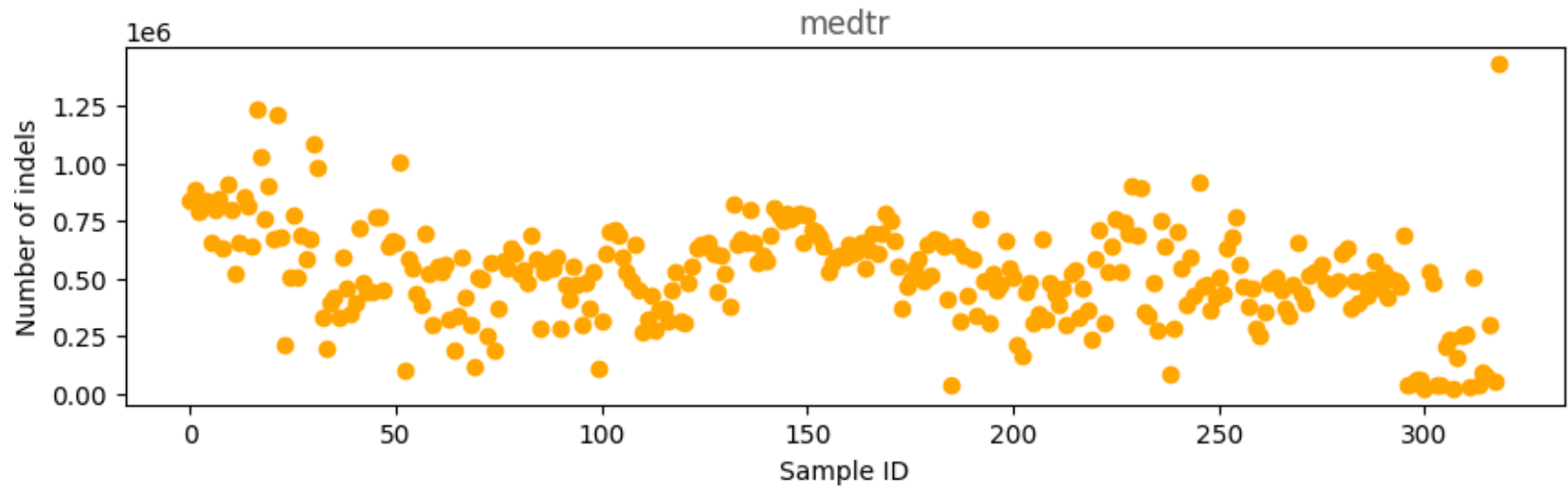
# Average depth by sample



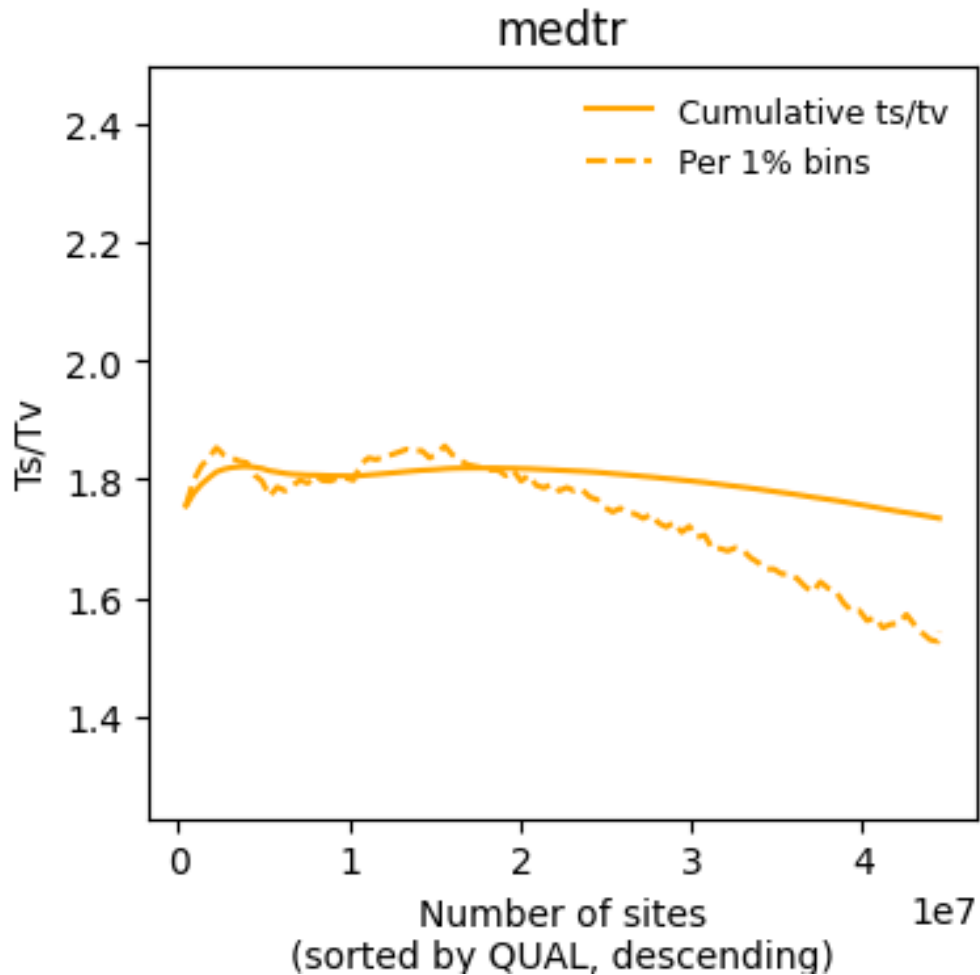
# Number of SNPs by sample



# Number of indels by sample

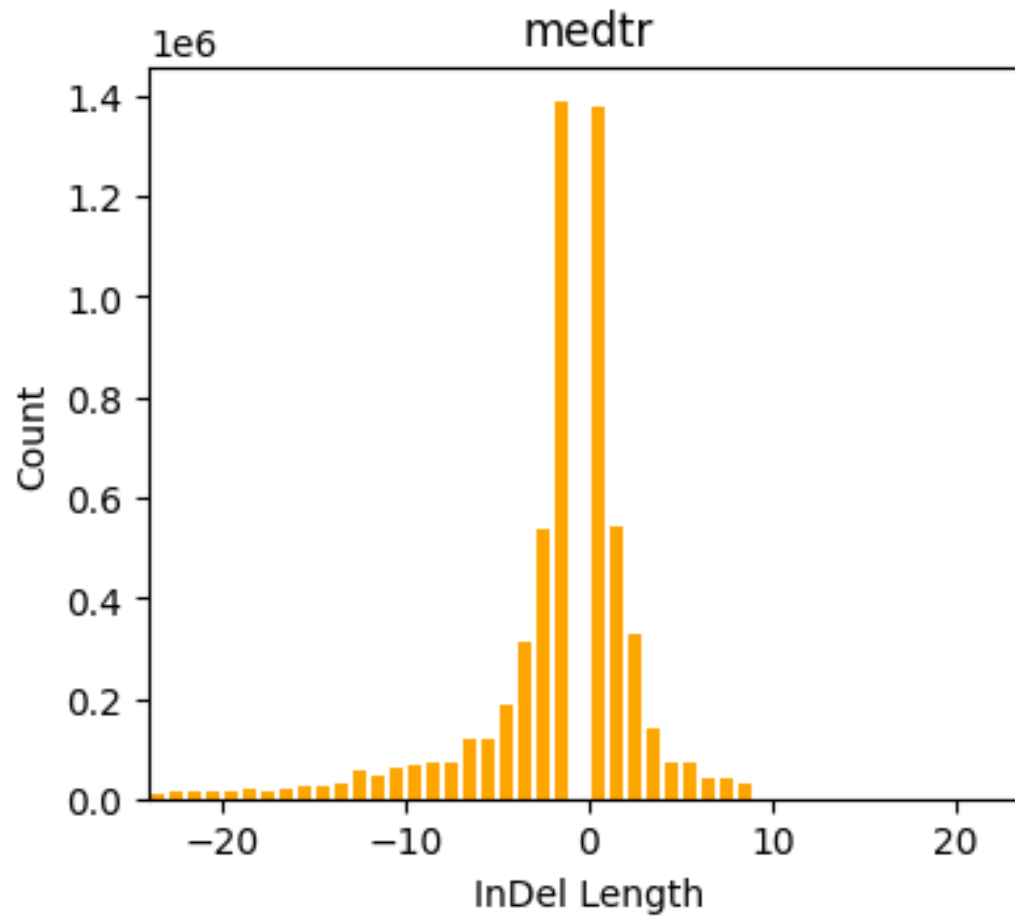


# Ts/Tv stratified by QUAL

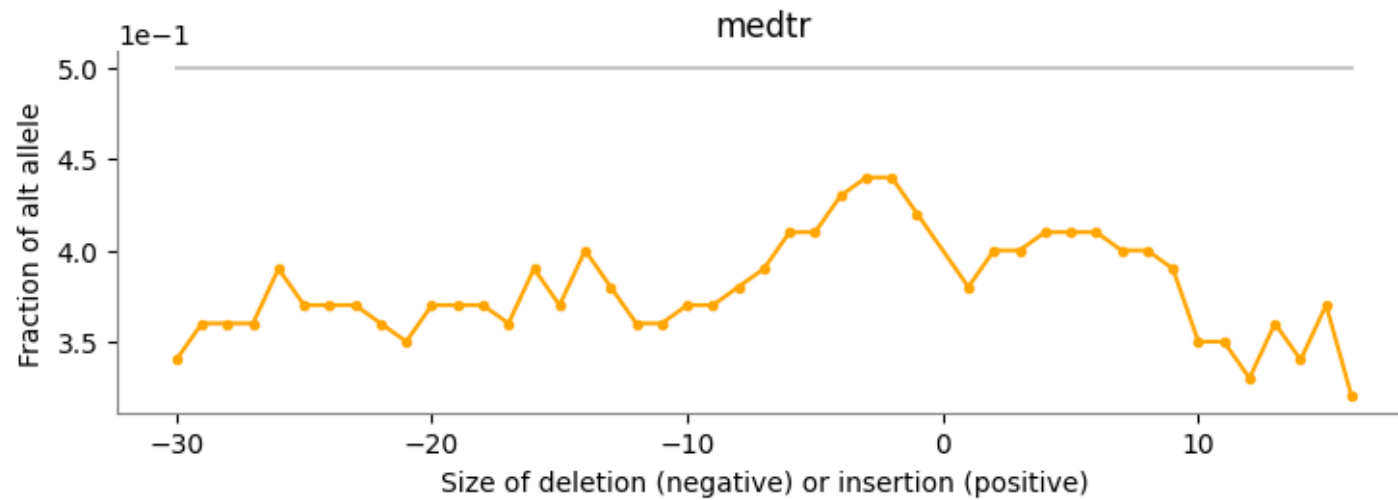




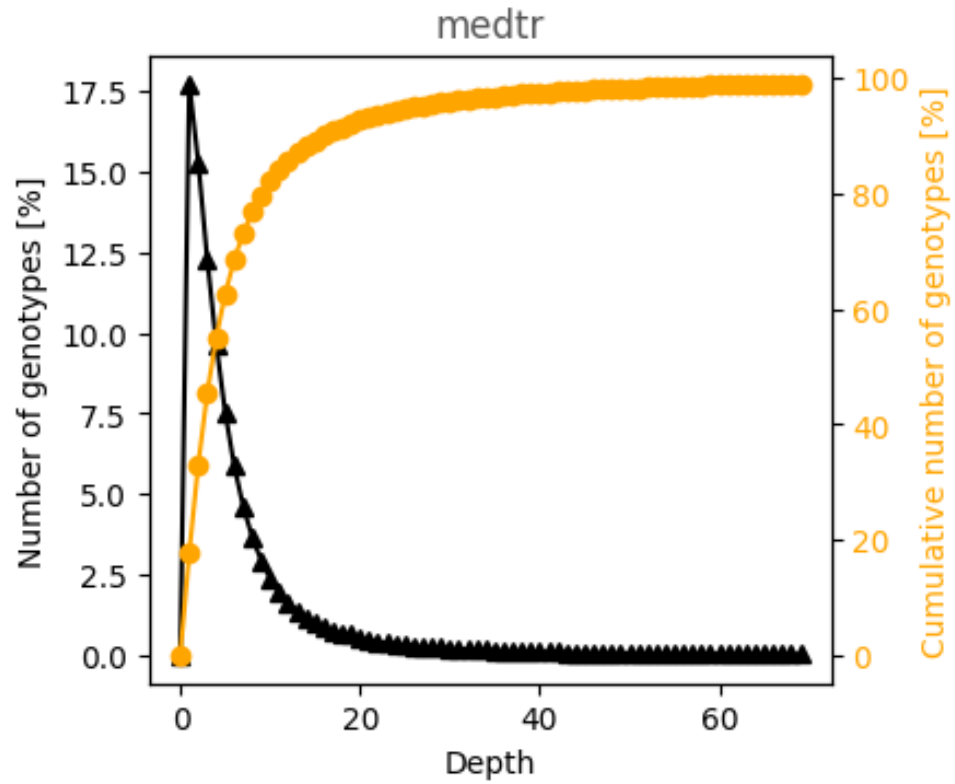
# Indel distribution



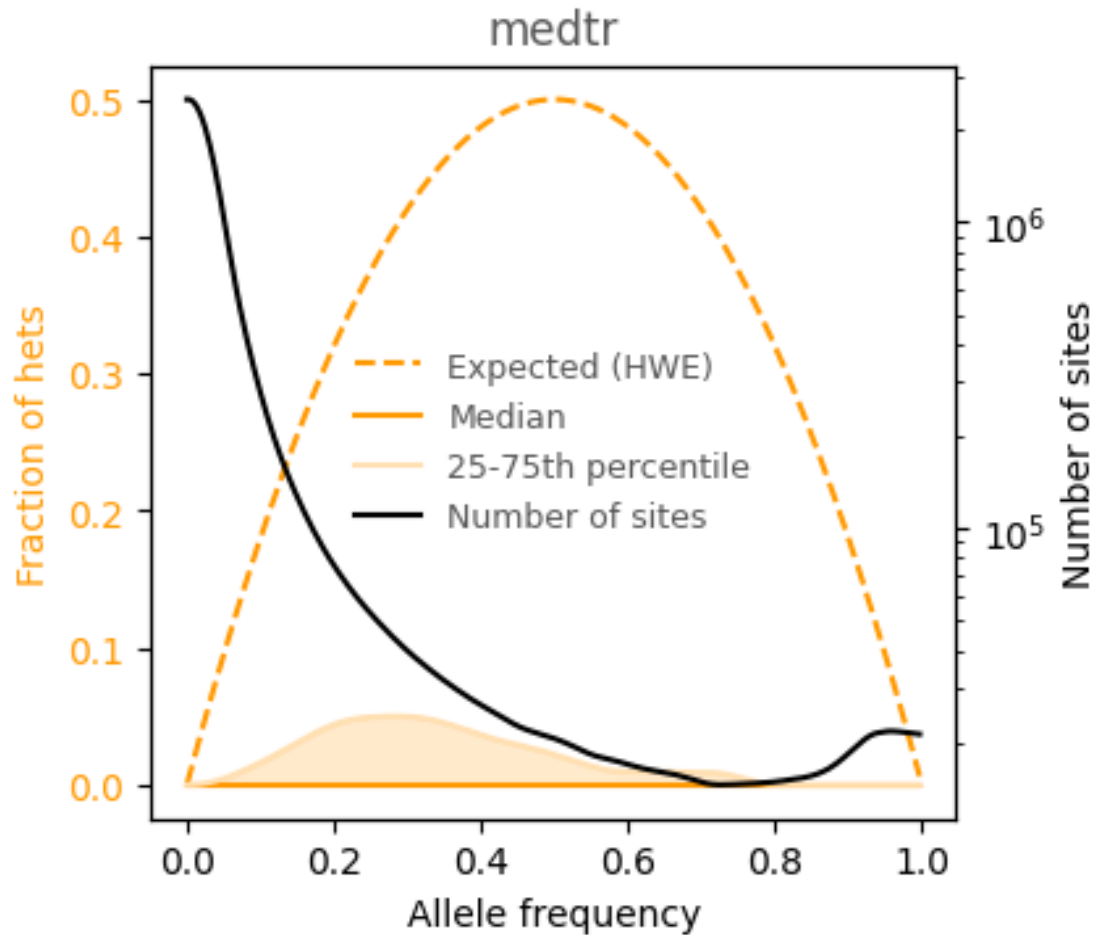
# Fraction of alternate indel allele



# Depth distribution



# Number of HETs by AF



# Substitution types

