

Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
medtr	62,667,460	1.40	1.50	8,206,269	-	0	15,566

* frameshift ratio: out/(out+in)

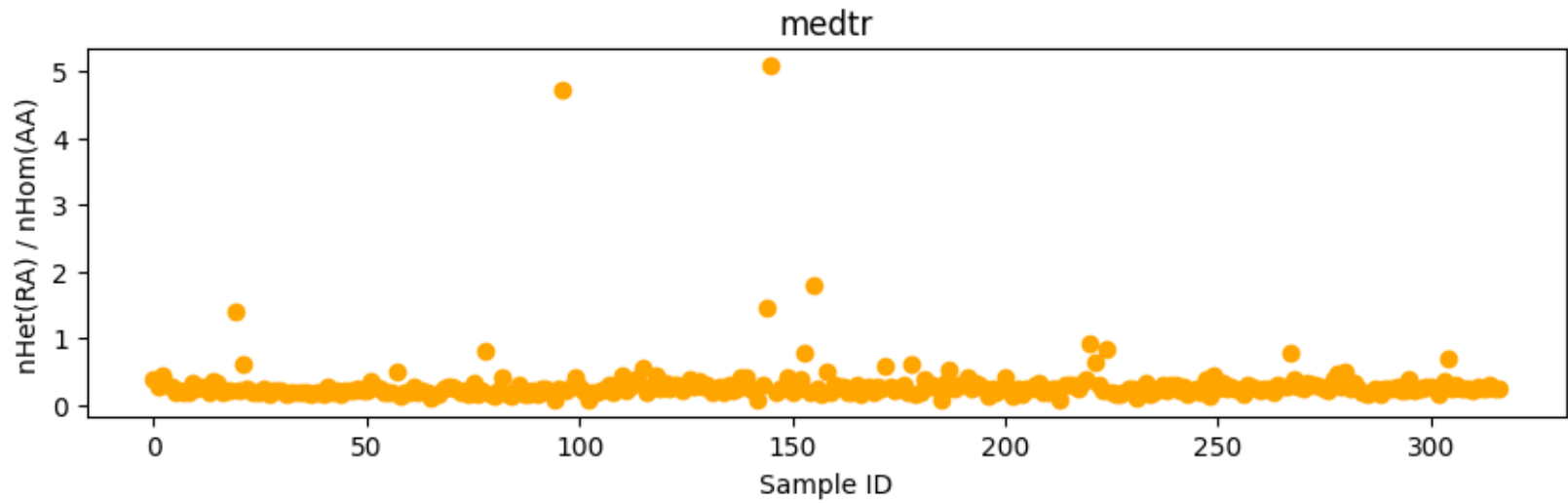
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
medtr	7.6%	0.91	5.4%	7,365,100	5,327,417

- medtr .. medtr.A17.gnm5.div.Epstein_Burghardt_2022.Mt5_qual30_primitives.2021-09-29.vcf.g
z

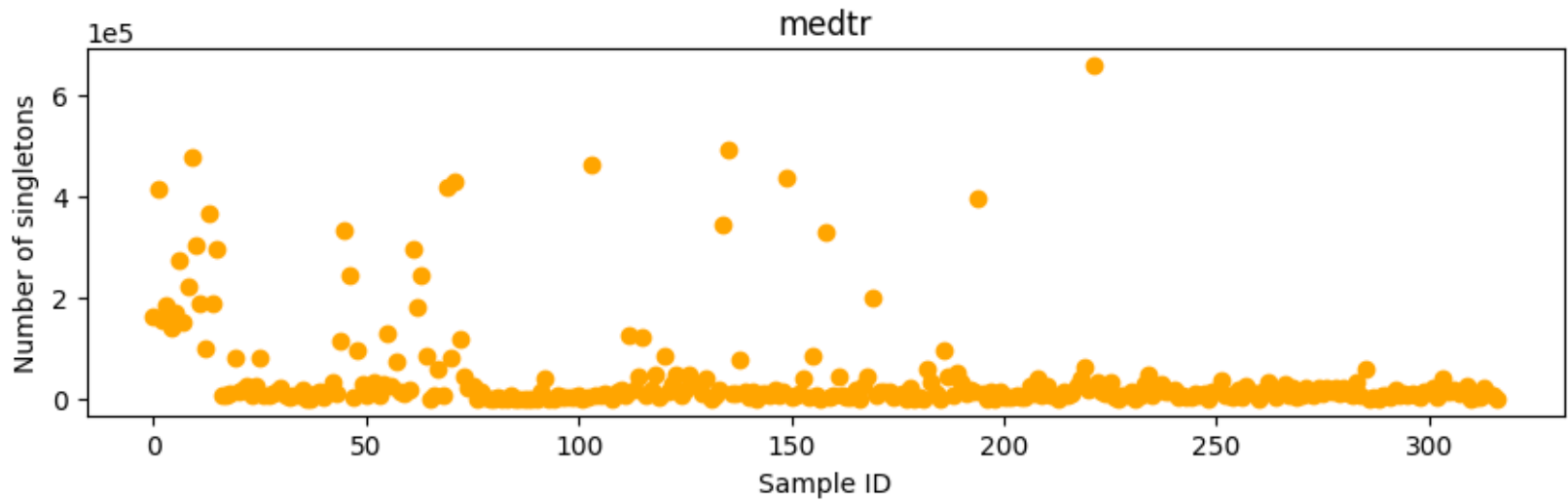
Ts/Tv by sample



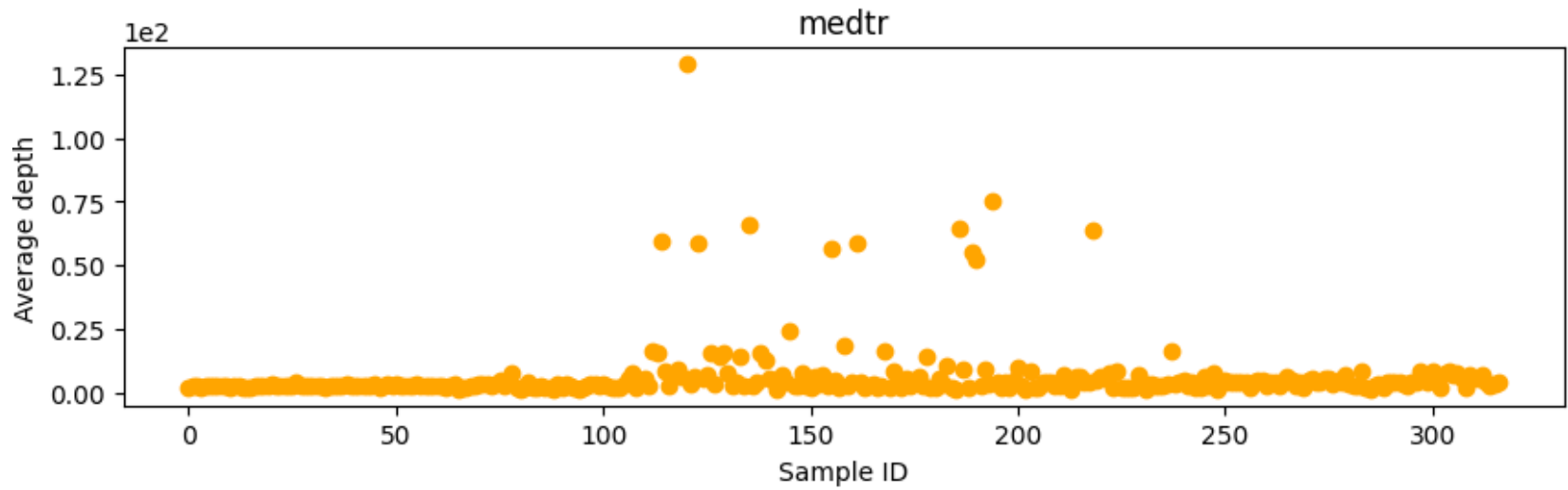
Hets vs non-ref Homs by sample



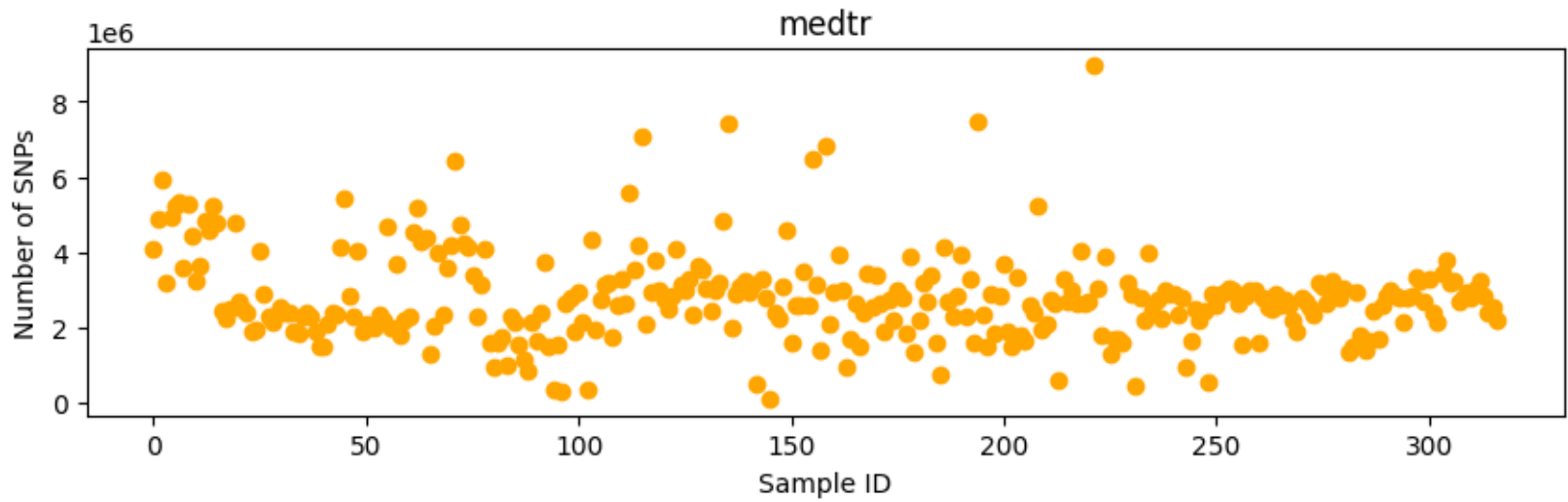
Singletons by sample (hets and homs)



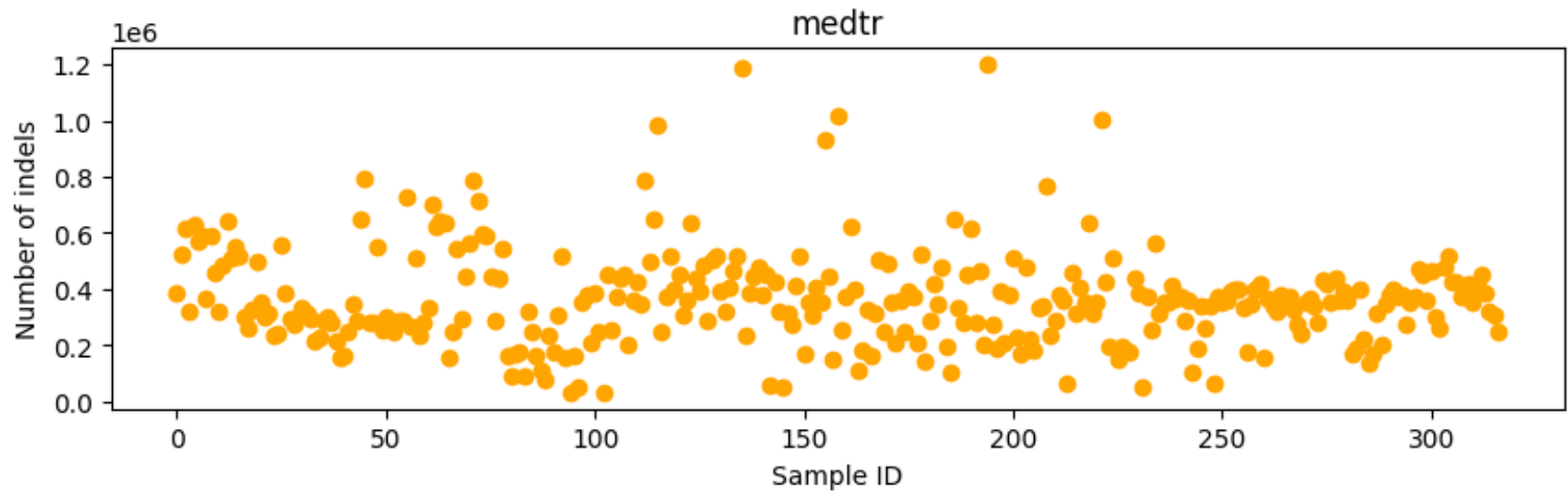
Average depth by sample



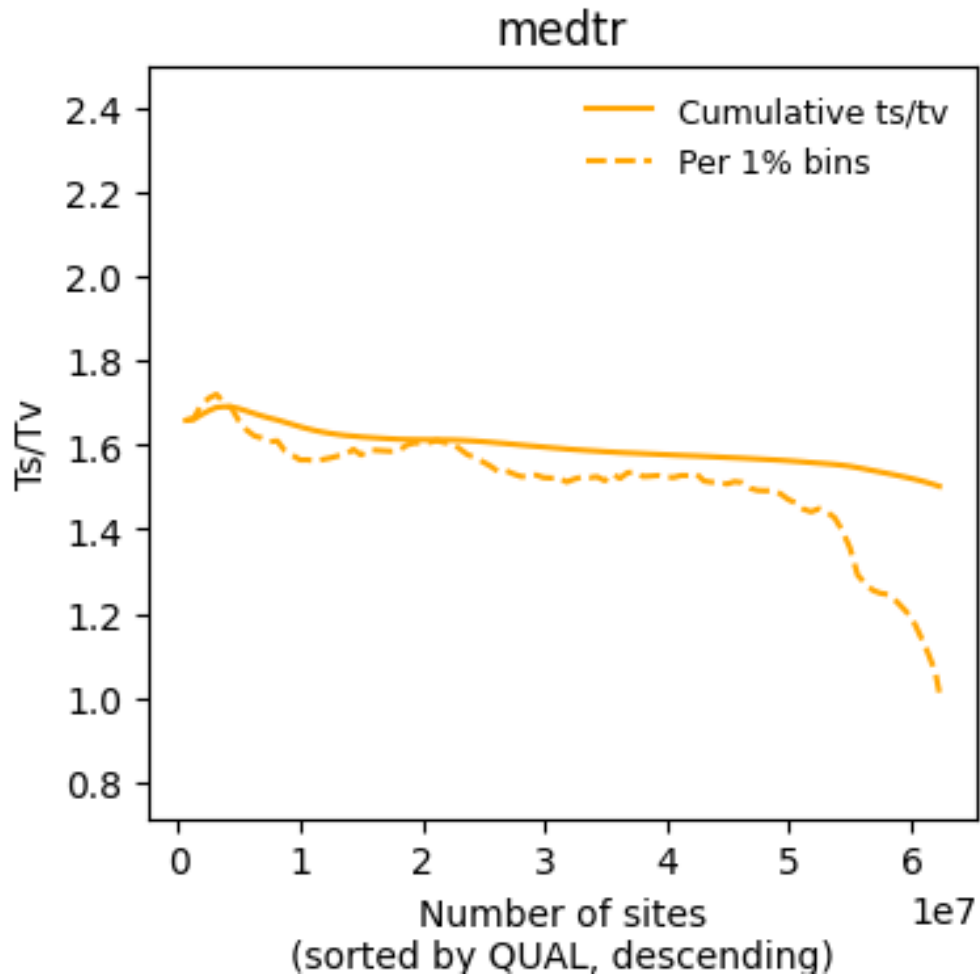
Number of SNPs by sample



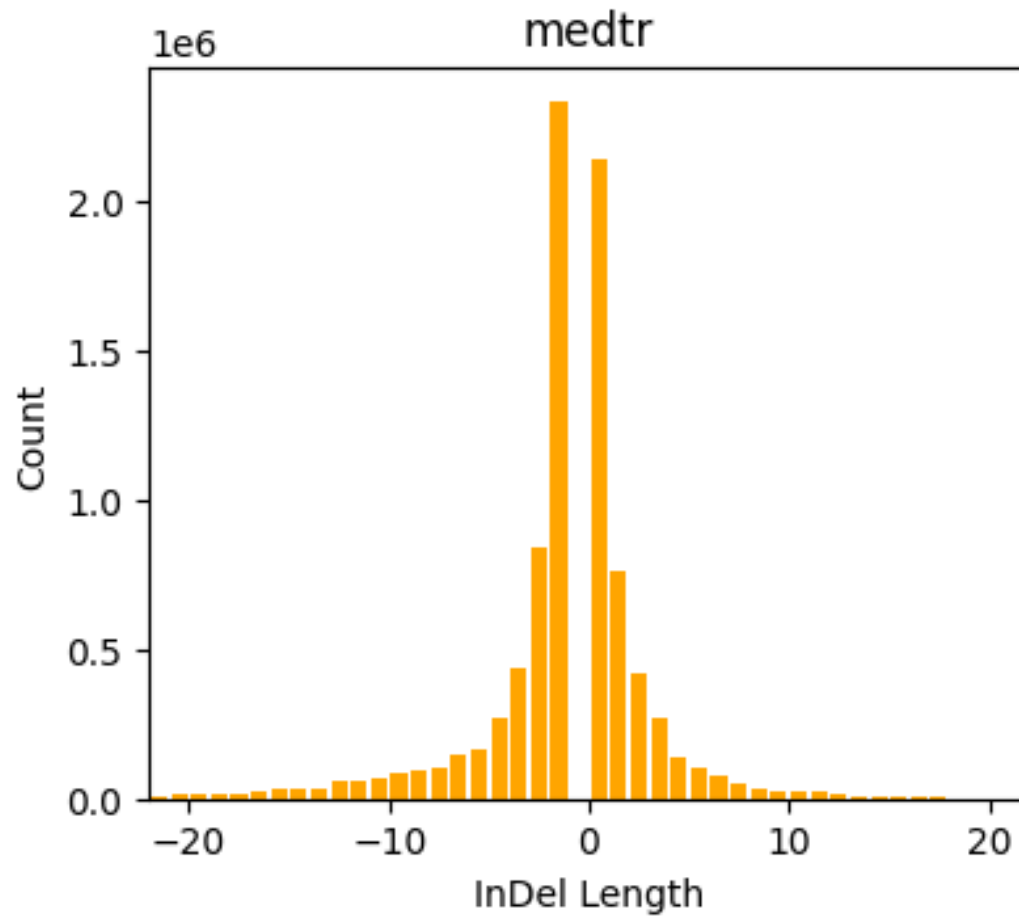
Number of indels by sample



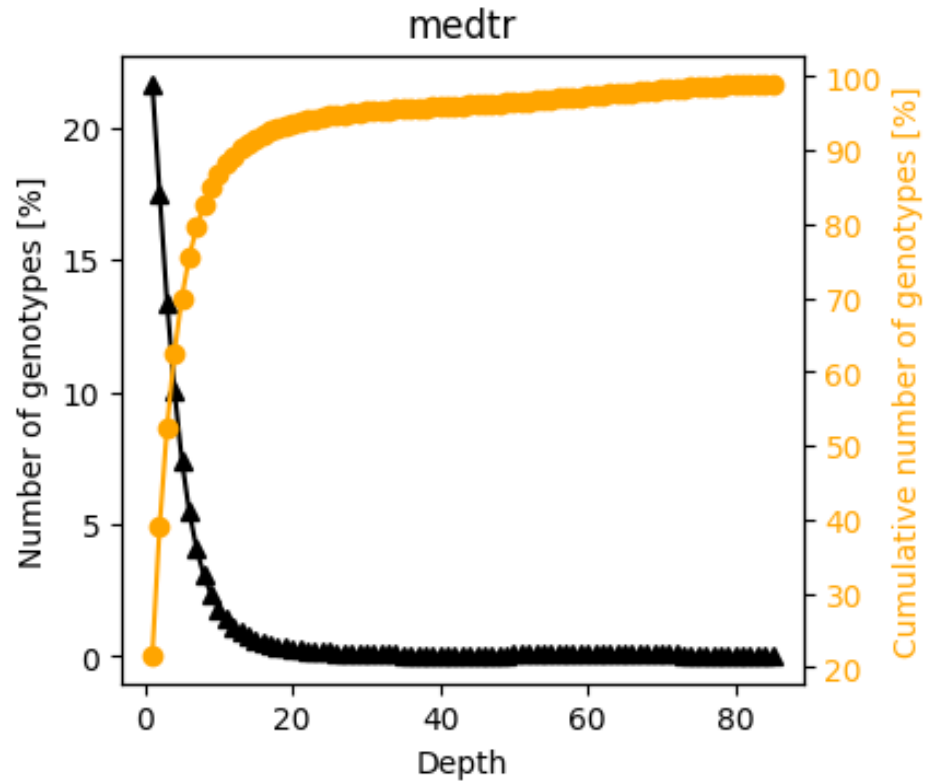
Ts/Tv stratified by QUAL



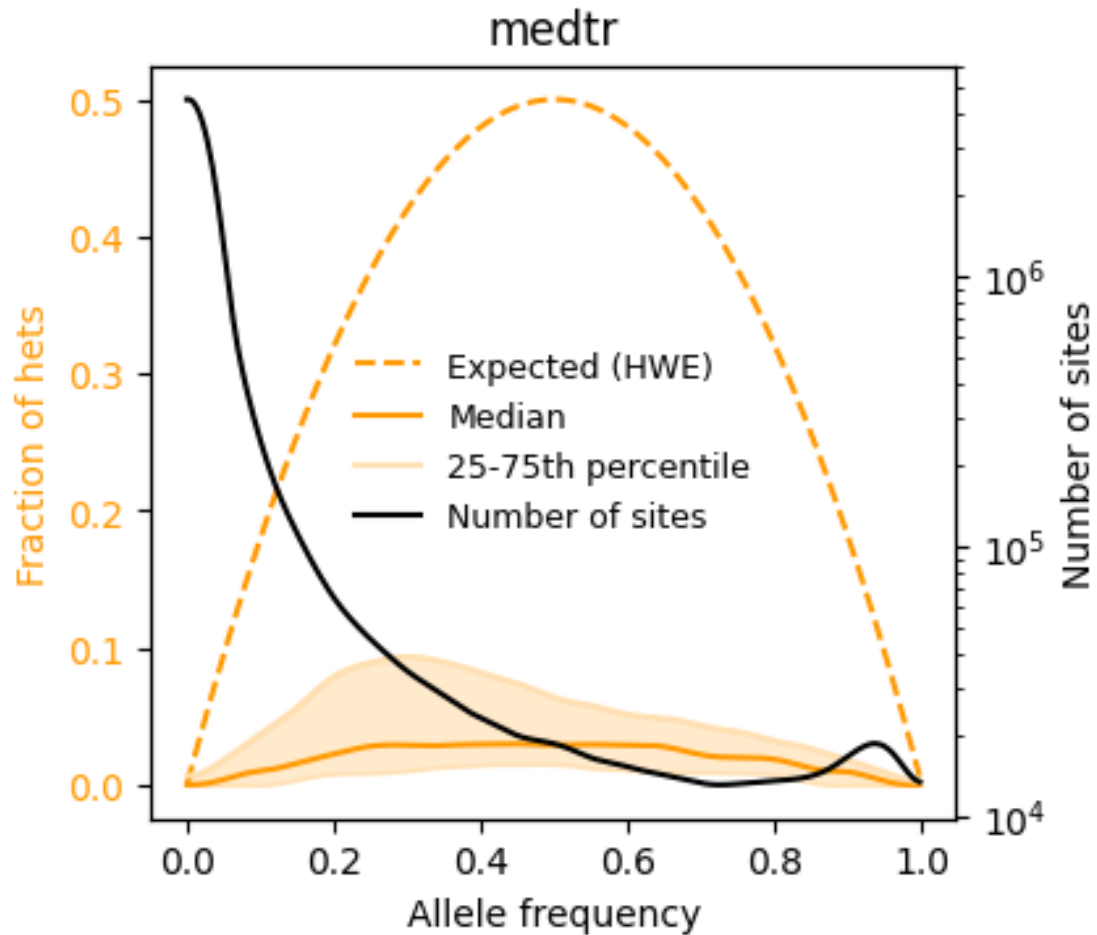
Indel distribution



Depth distribution



Number of HETs by AF



Substitution types

