

Summary Numbers

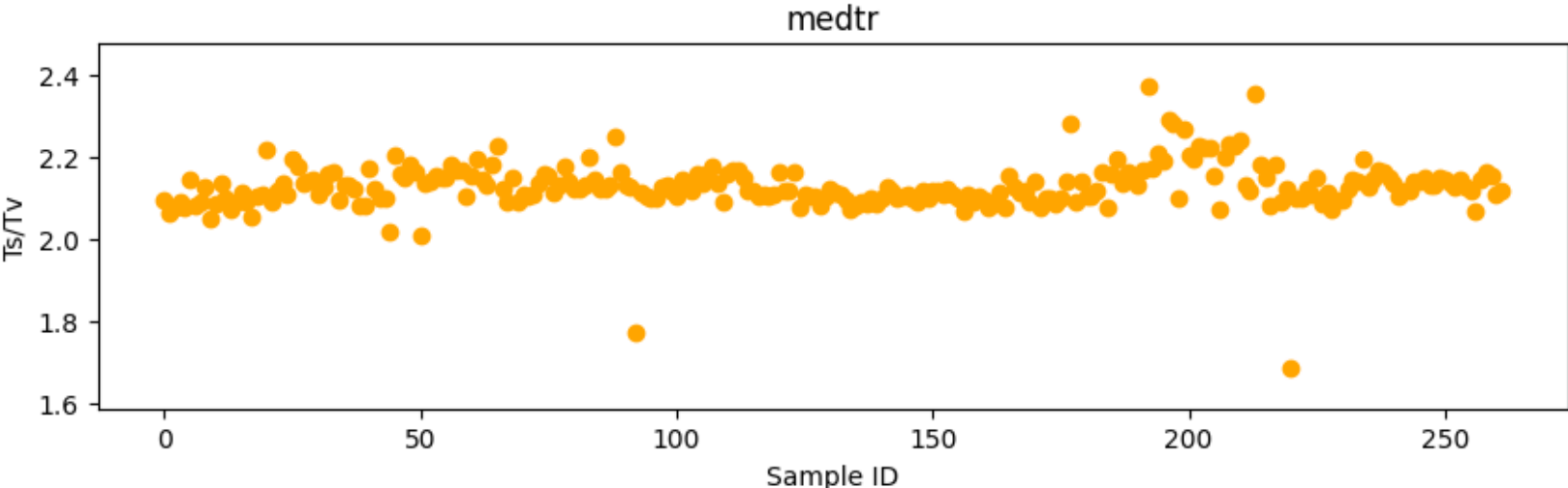
Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
medtr	40,065,843	1.81	1.81	0	-	0	0

* frameshift ratio: out/(out+in)

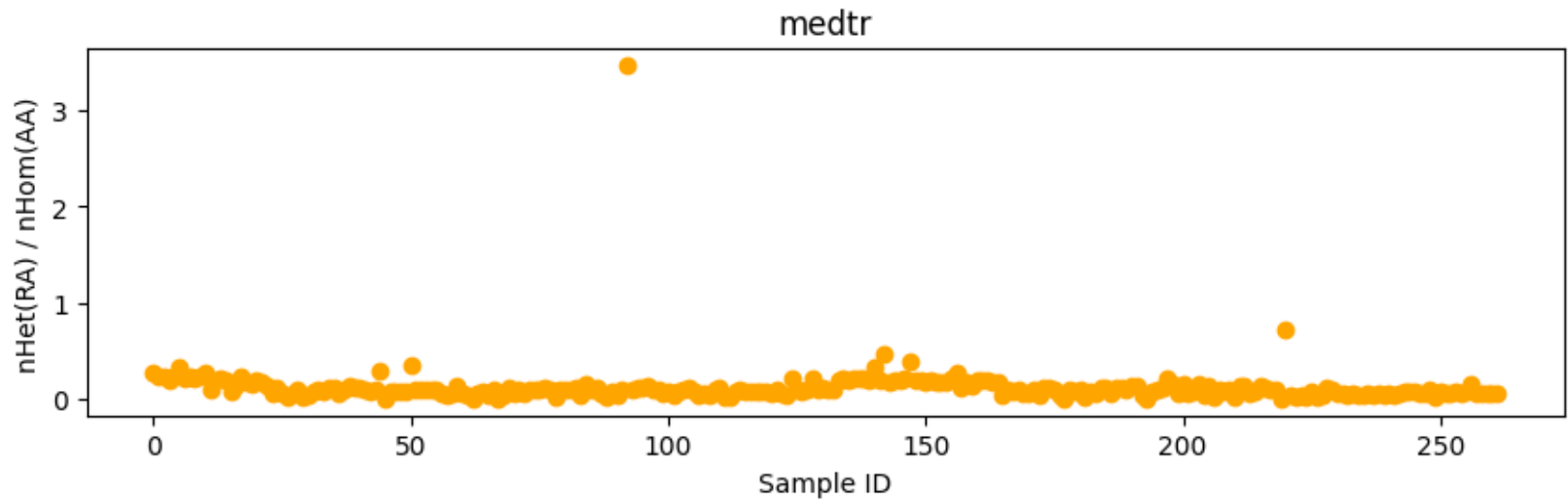
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
medtr	2.8%	1.74	0.0%	0	0

- medtr .. medtr.A17_HM341.gnm4.div.Stanton-Geddes_Paape_2013.filtered-set-2014Apr12.vcf.gz

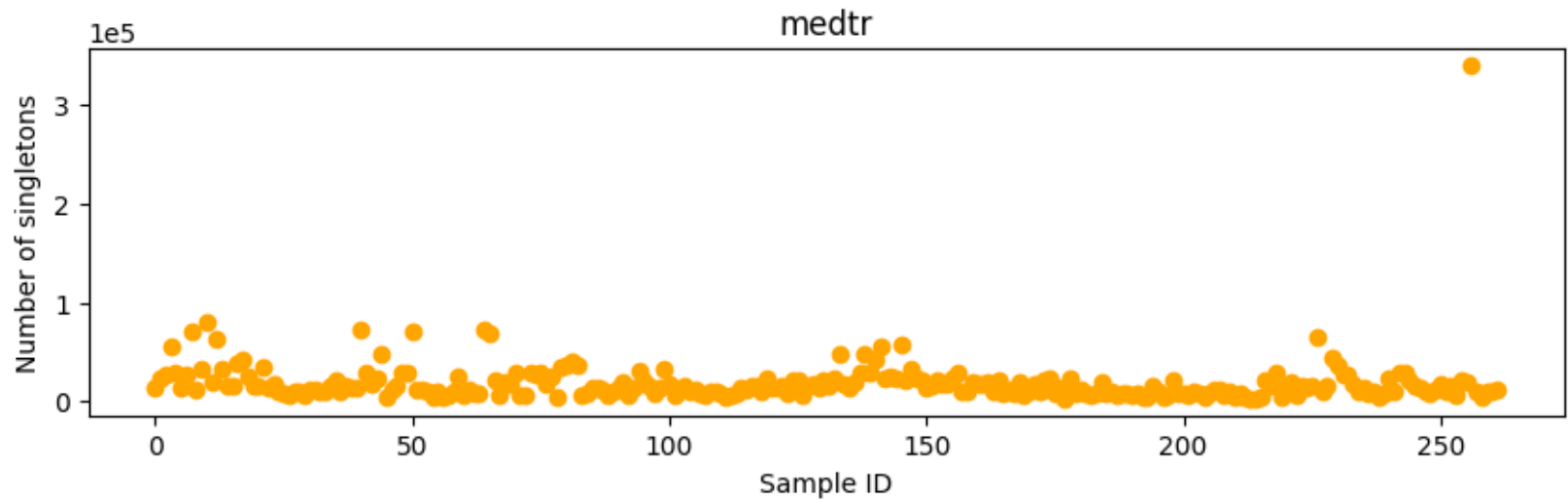
Ts/Tv by sample



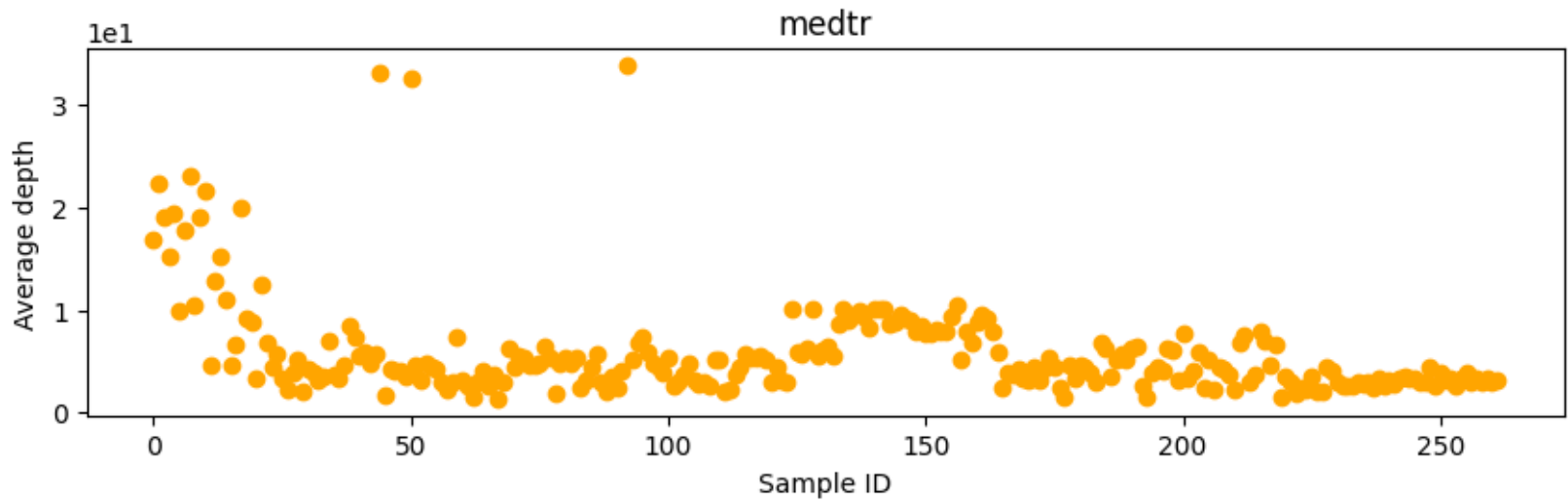
Hets vs non-ref Homs by sample



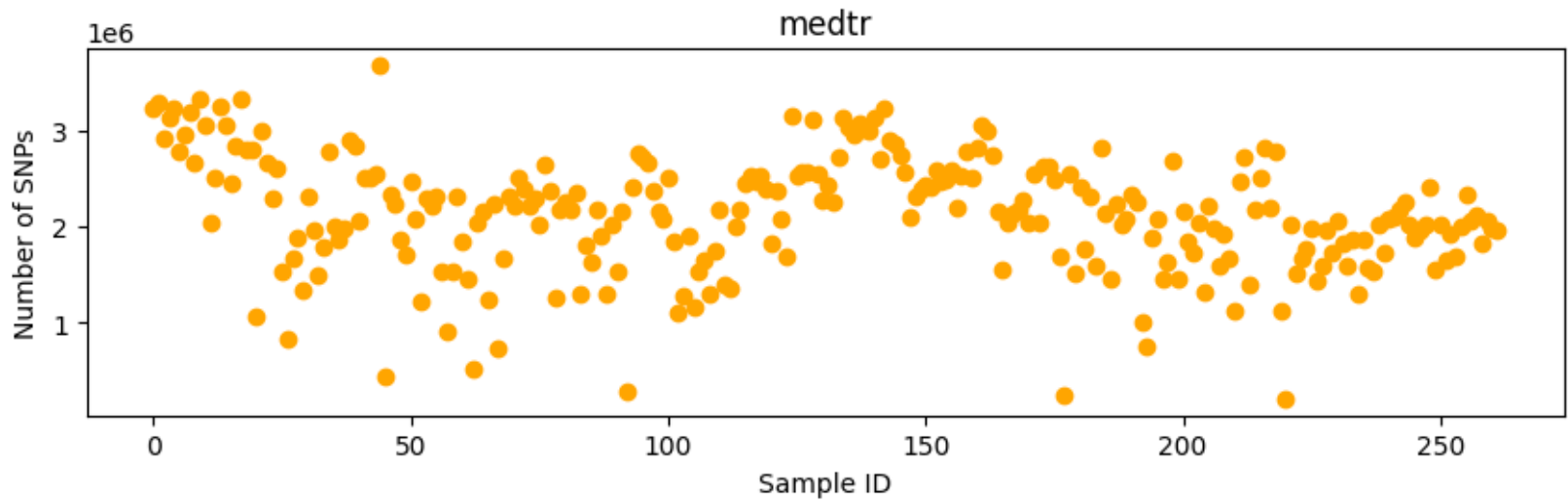
Singletons by sample (hets and homs)



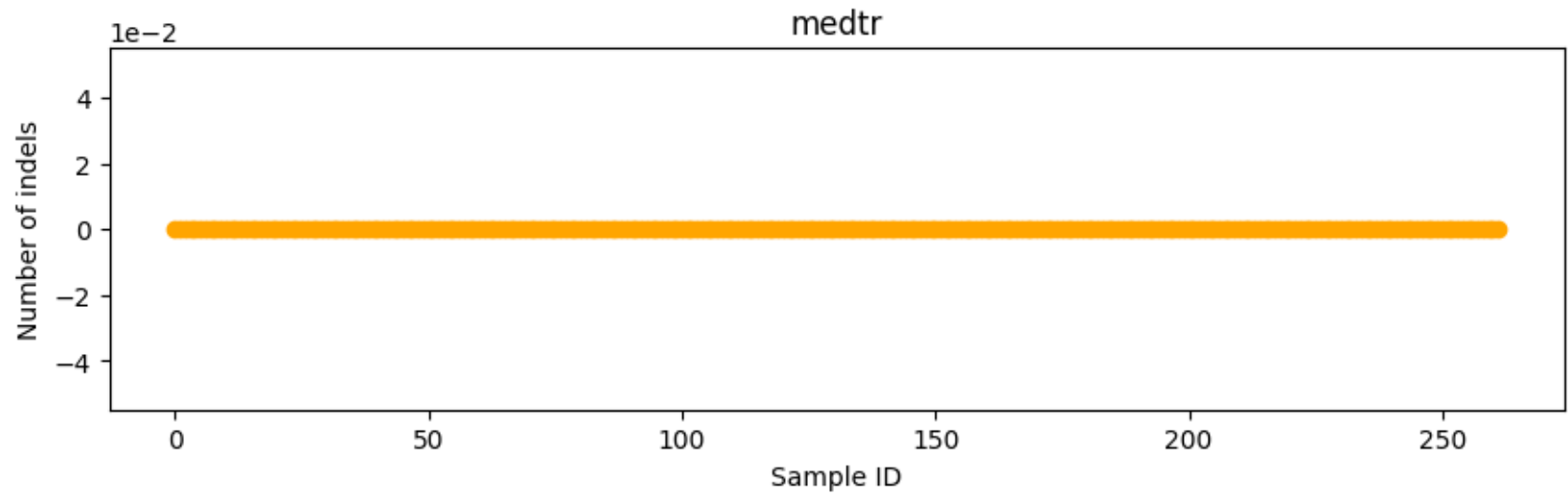
Average depth by sample



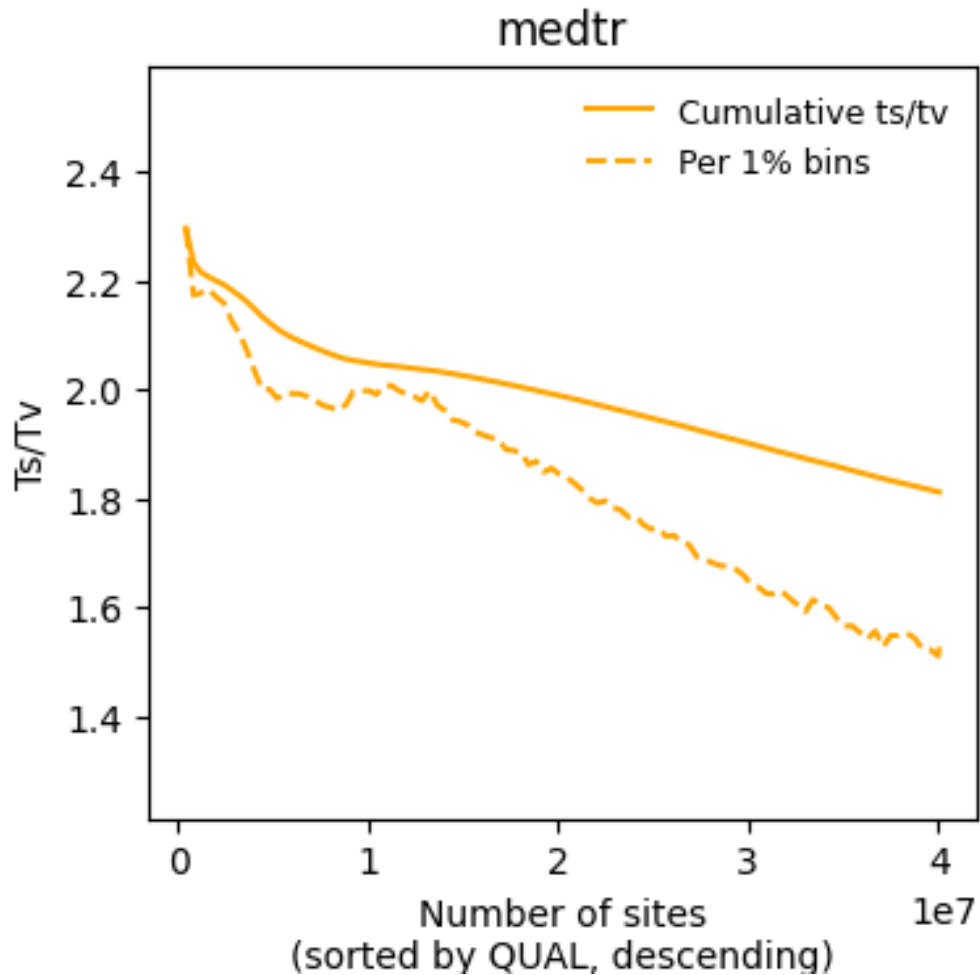
Number of SNPs by sample



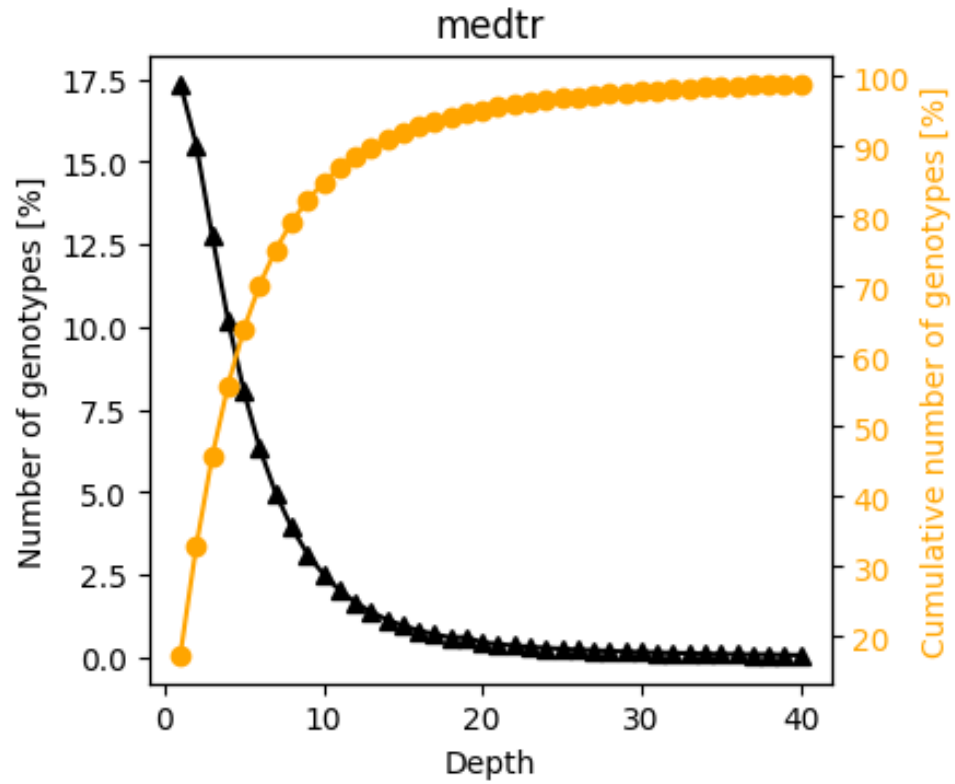
Number of indels by sample



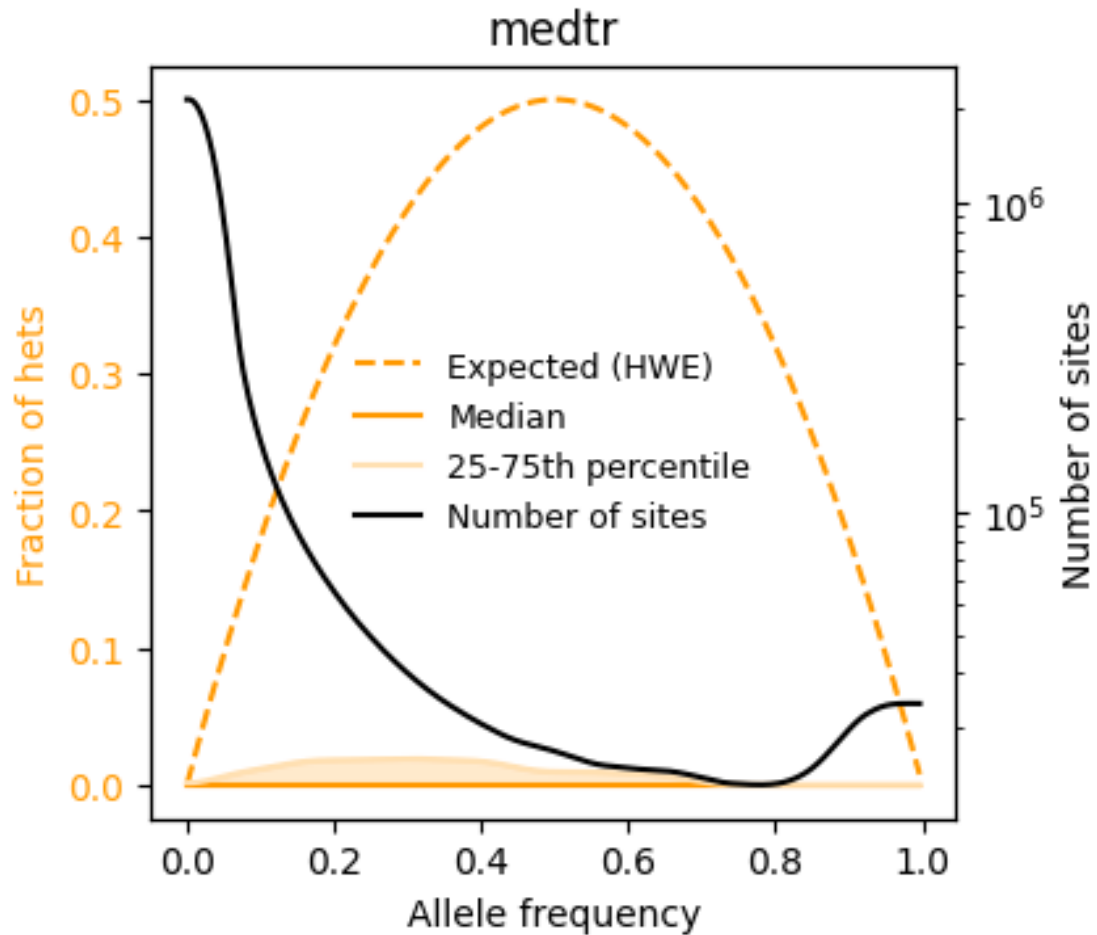
Ts/Tv stratified by QUAL



Depth distribution



Number of HETs by AF



Substitution types

