

Summary Numbers

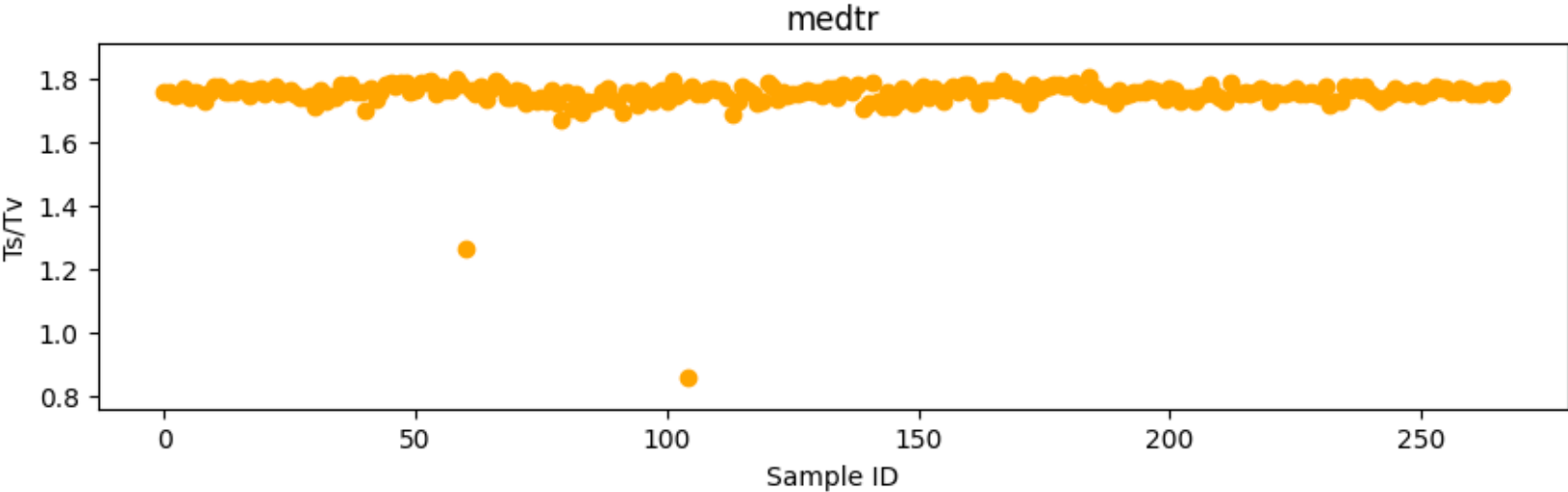
Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
medtr	37,046,405	1.43	1.56	4,955,793	-	0	9,647

* frameshift ratio: out/(out+in)

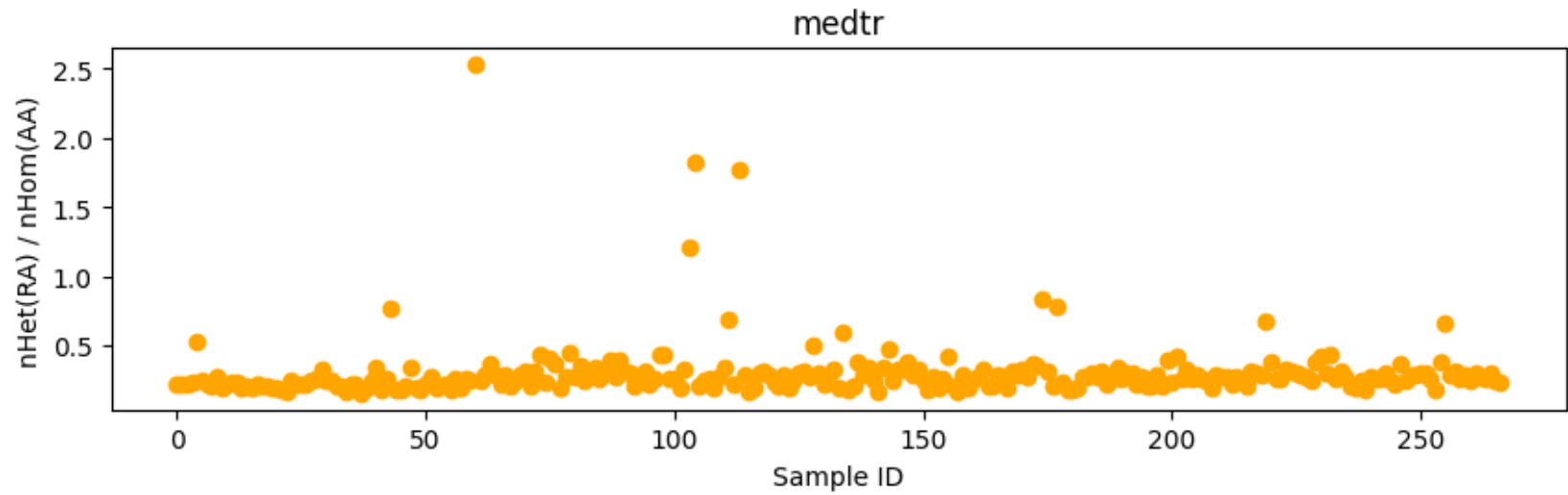
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
medtr	17.1%	1.13	9.2%	5,745,776	4,124,966

- medtr .. medtr.A17.gnm5.div.Epstein.Burghardt.2022.Mt5_qual30_primitives_imputed.2021-09-29.vcf.gz

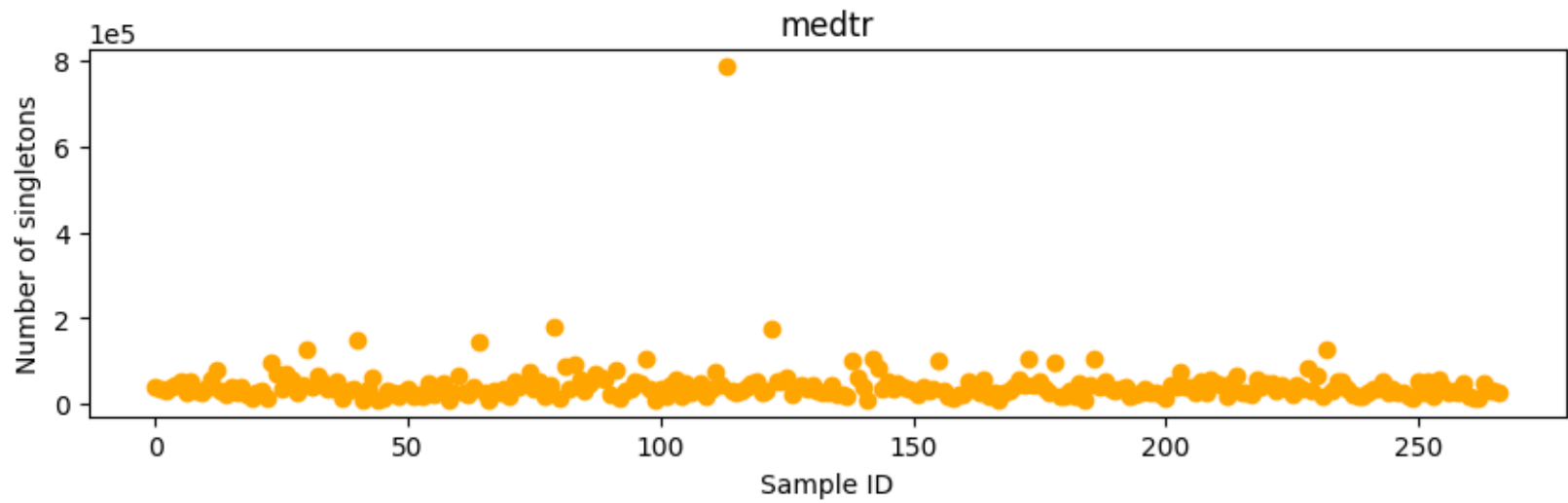
Ts/Tv by sample



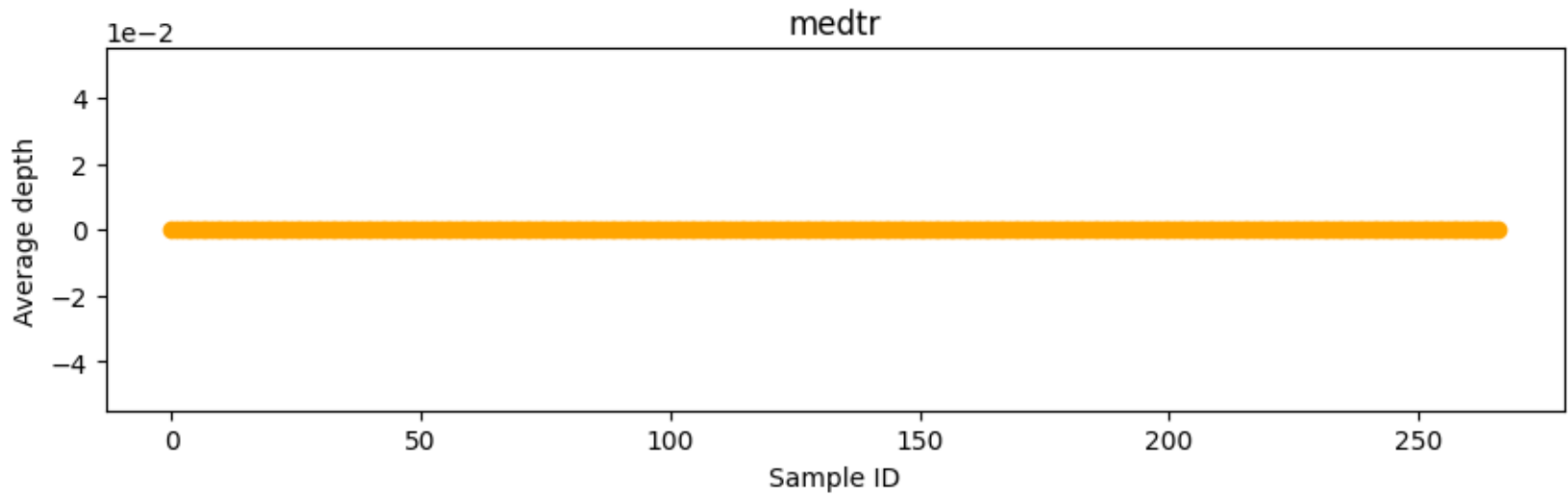
Hets vs non-ref Homs by sample



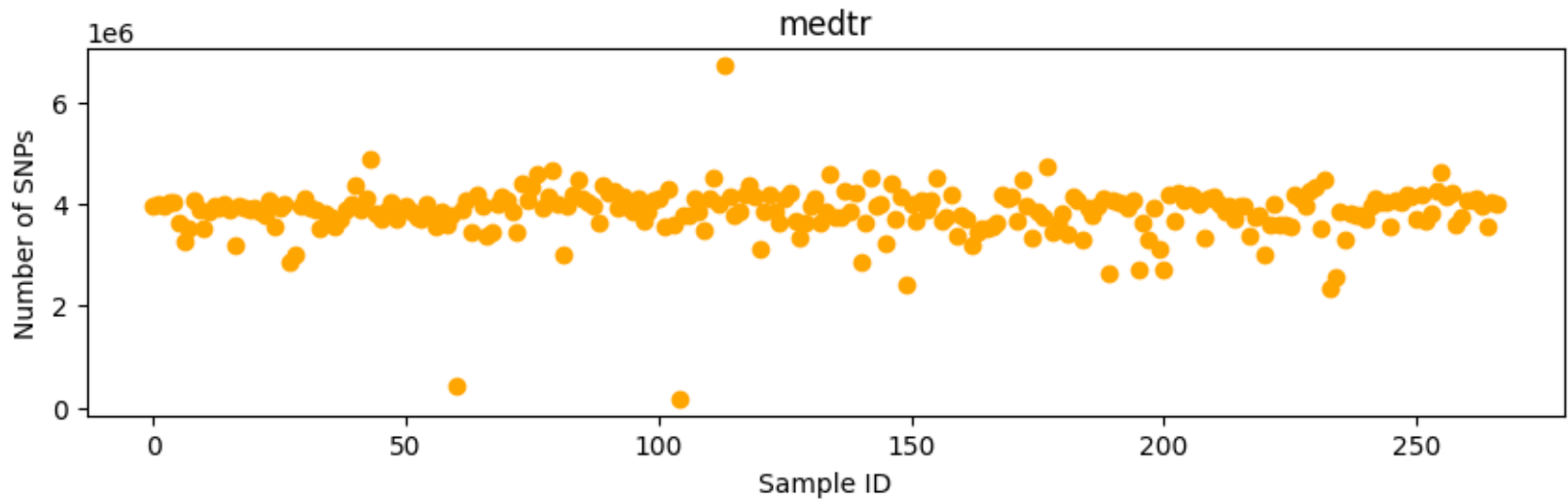
Singletons by sample (hets and homs)



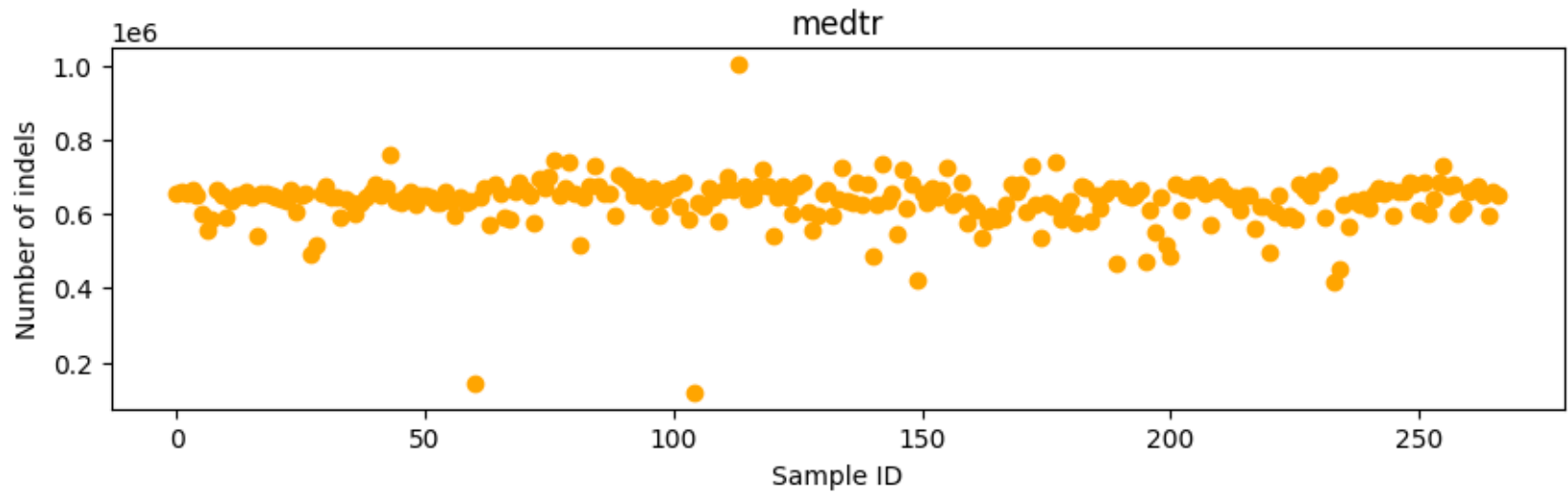
Average depth by sample



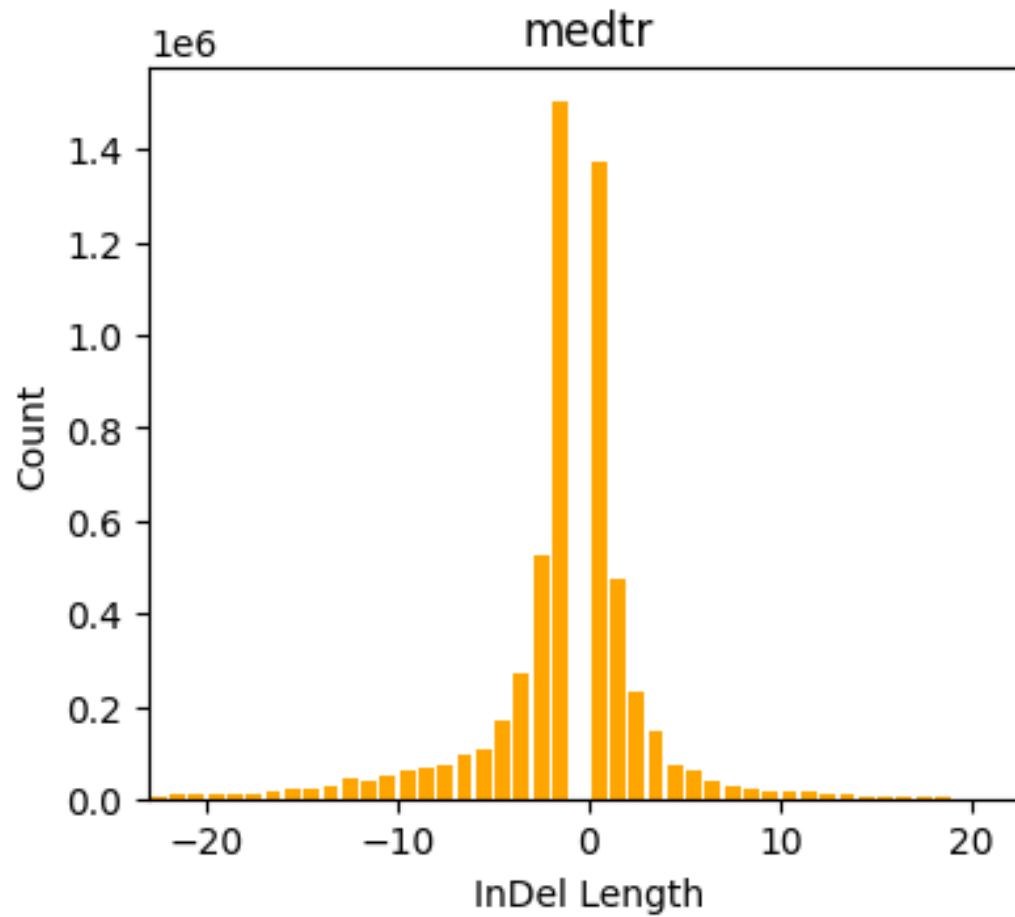
Number of SNPs by sample



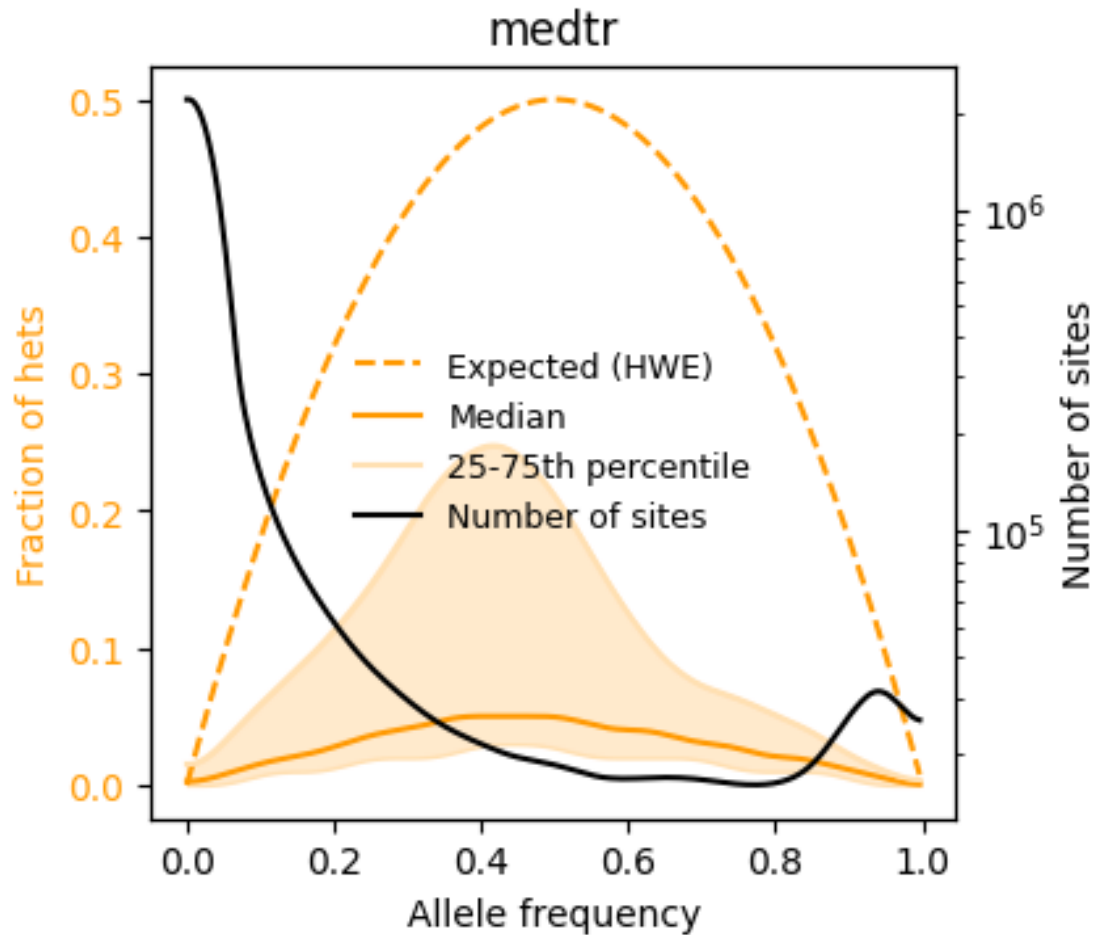
Number of indels by sample



Indel distribution



Number of HETs by AF



Substitution types

