

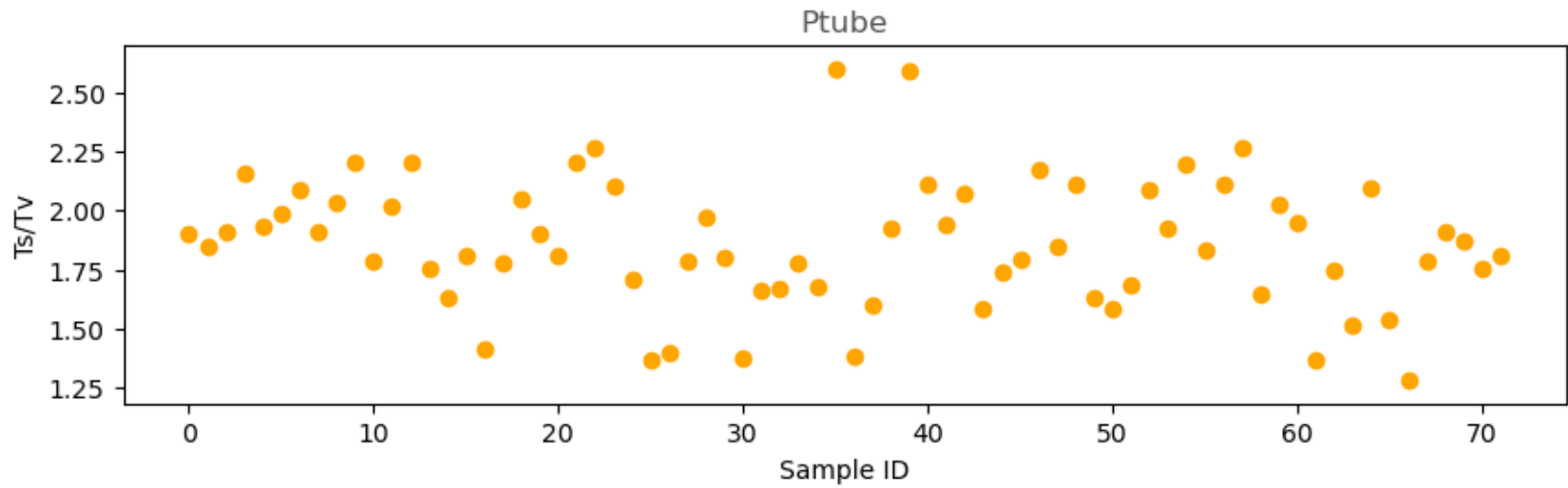
Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
Ptube	2,833	1.73	1.73	0	–	0	0
* frameshift ratio: out/(out+in)							

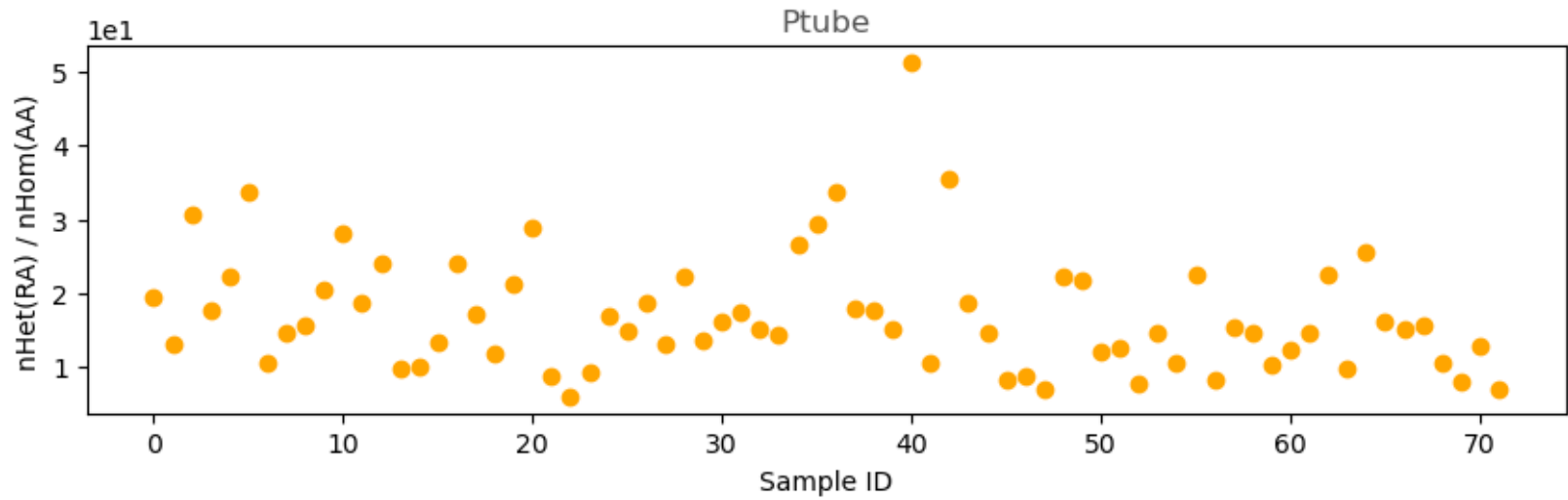
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
Ptube	1.8%	1.17	0.0%	0	0

- Ptube .. Ptuberculosis_filtered_SNPs.vcf

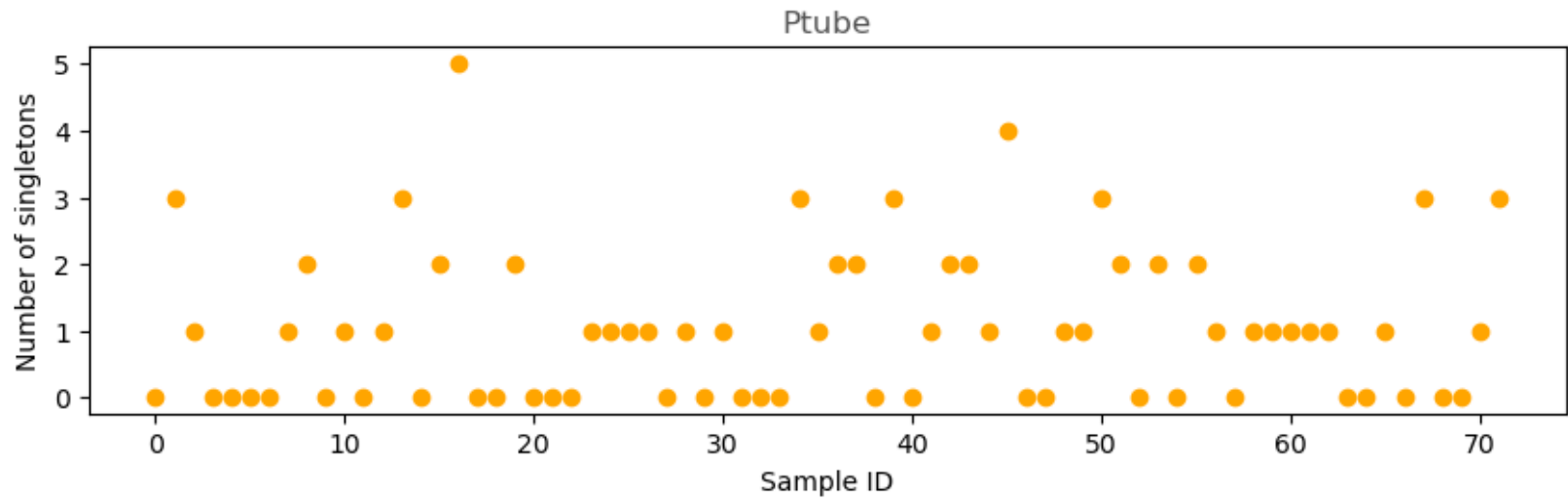
Ts/Tv by sample



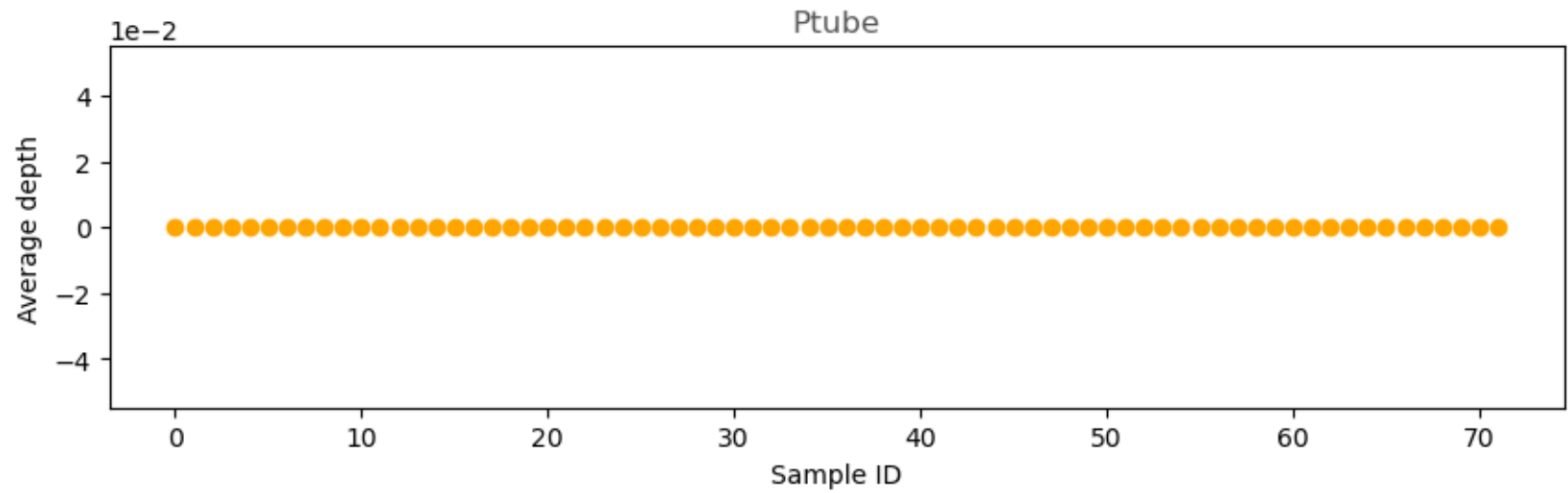
Hets vs non-ref Homs by sample



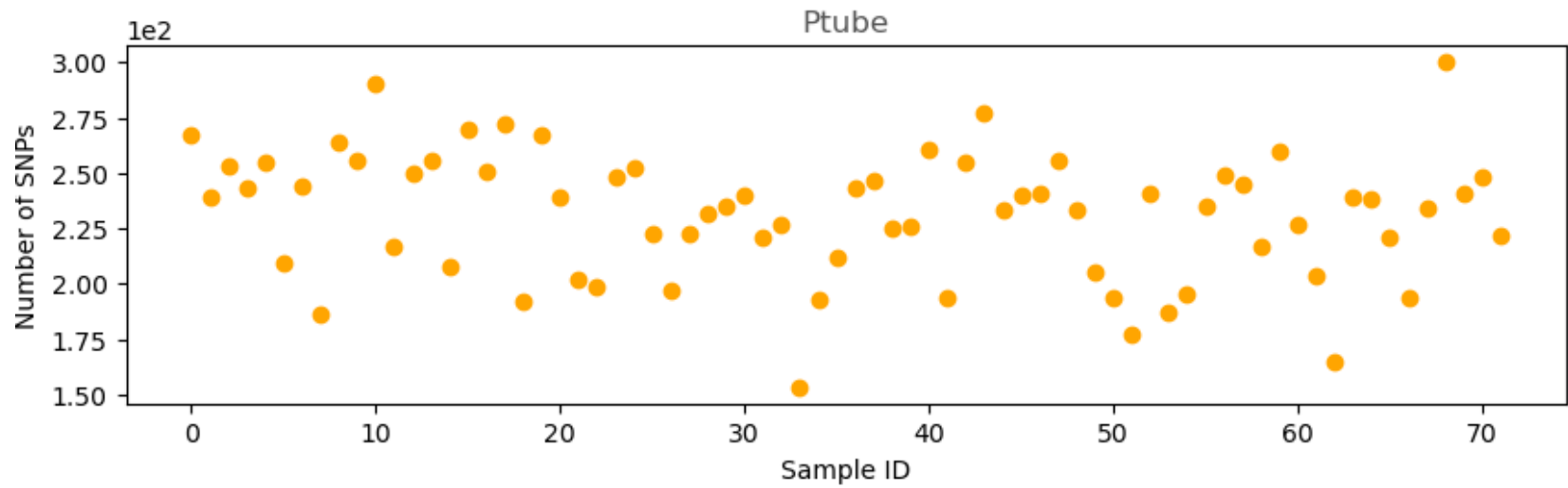
Singletons by sample (hets and homs)



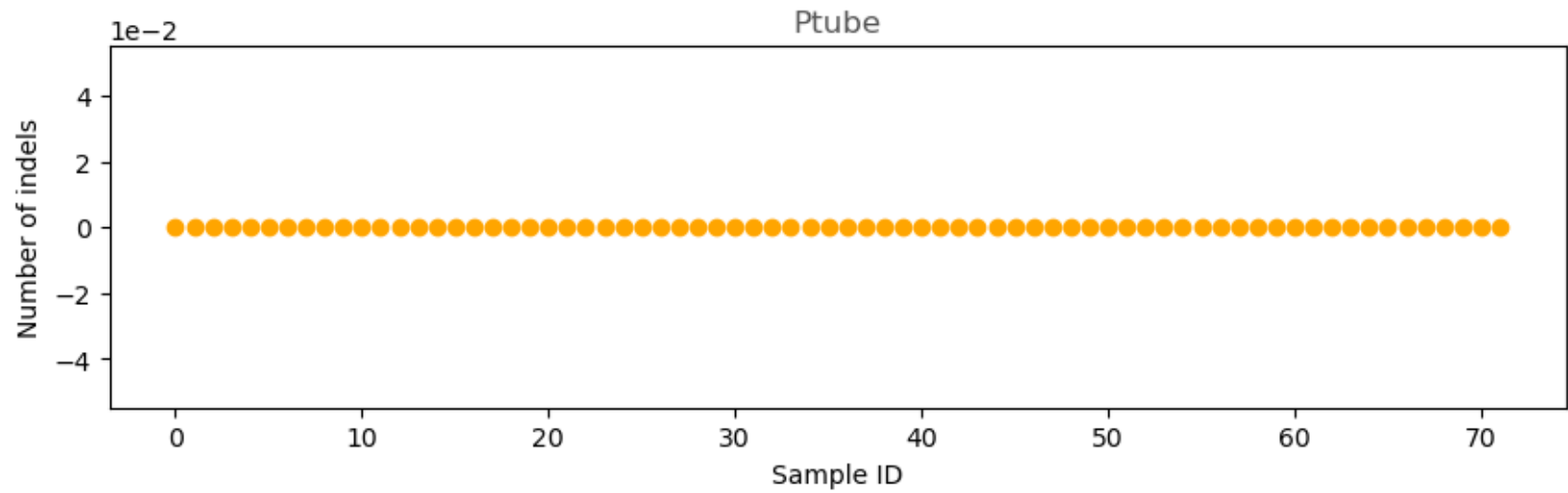
Average depth by sample



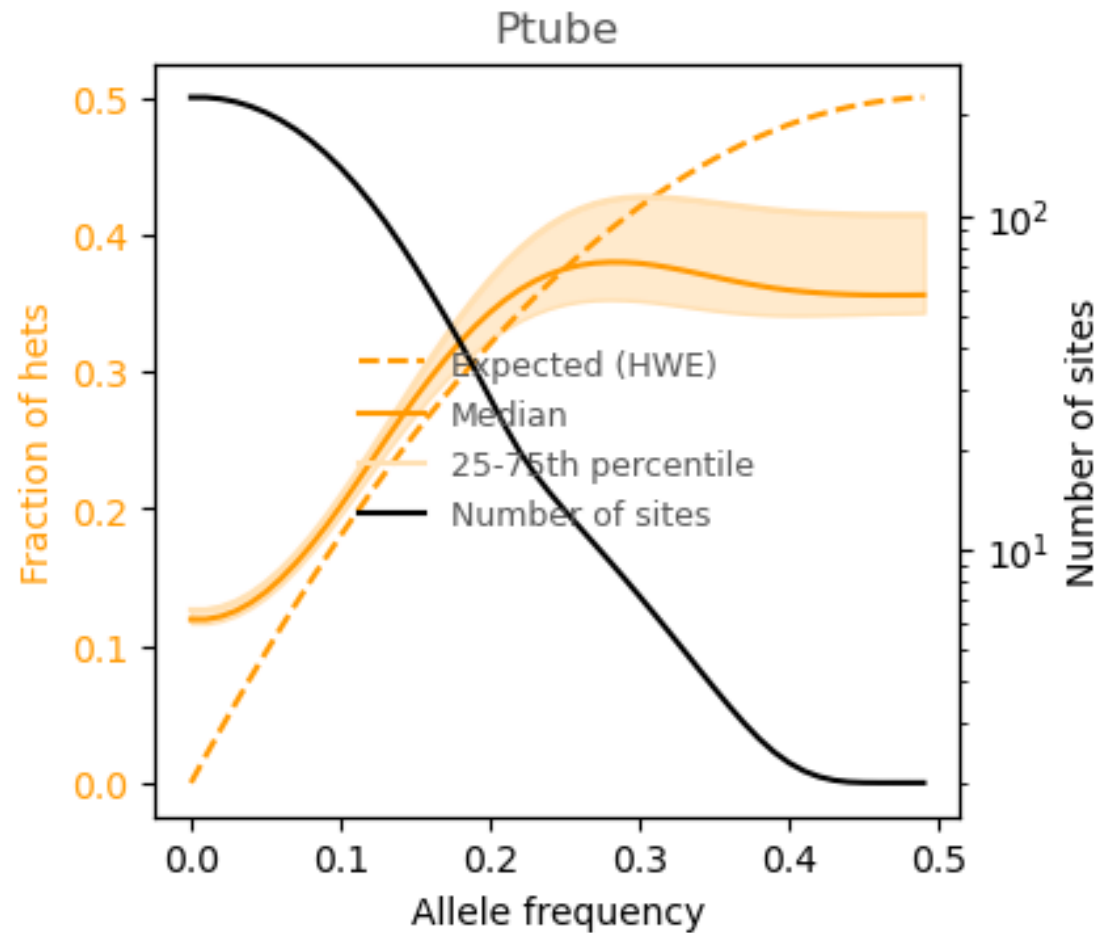
Number of SNPs by sample



Number of indels by sample



Number of HETs by AF



Substitution types

