

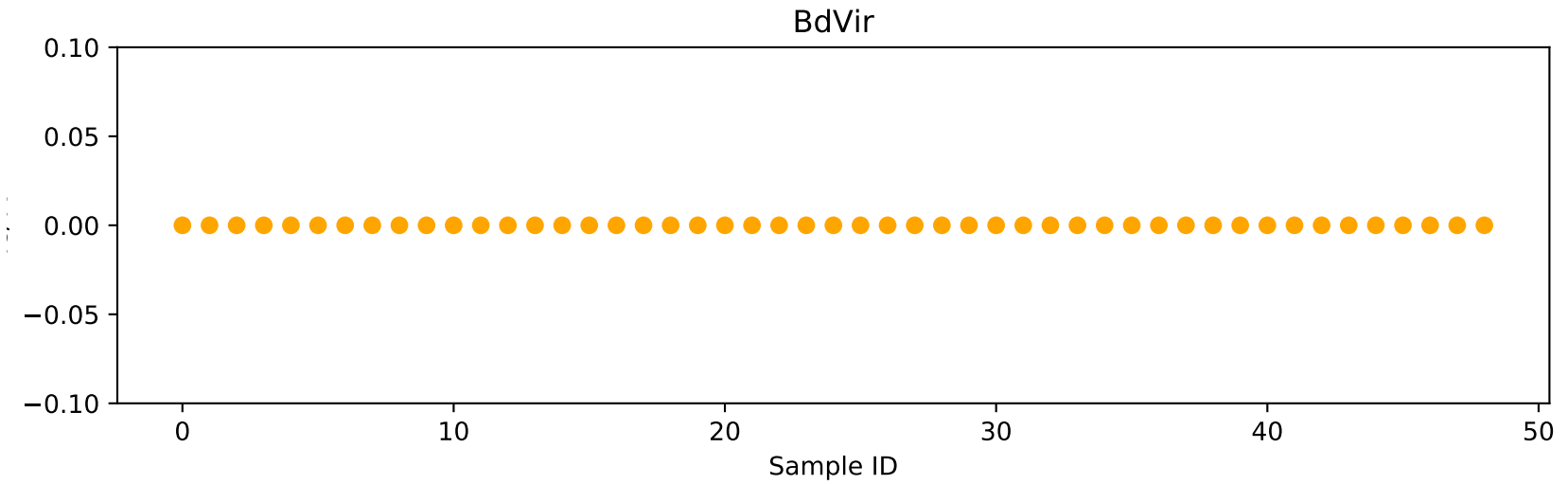
# Summary Numbers

	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
BdVir	194	1.98	2.03	2	–	0	0
* frameshift ratio: out/(out+in)							

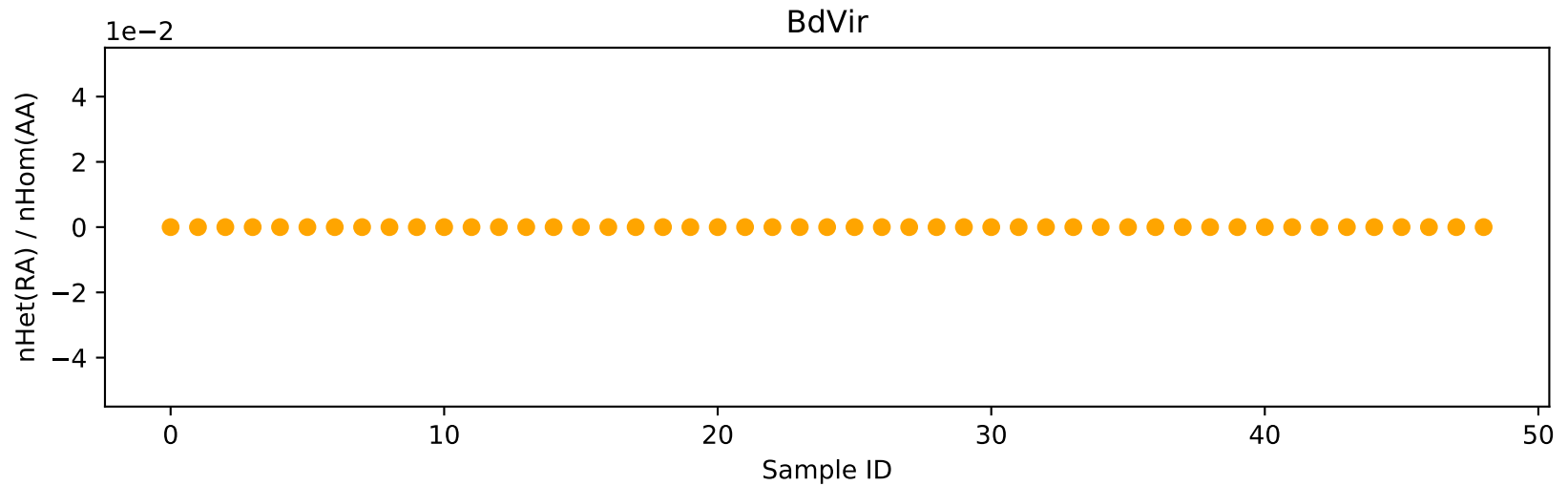
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
BdVir	18.3%	1.77	0.0%	3	3

- BdVir .. vcf/BdVirus.vcf.gz

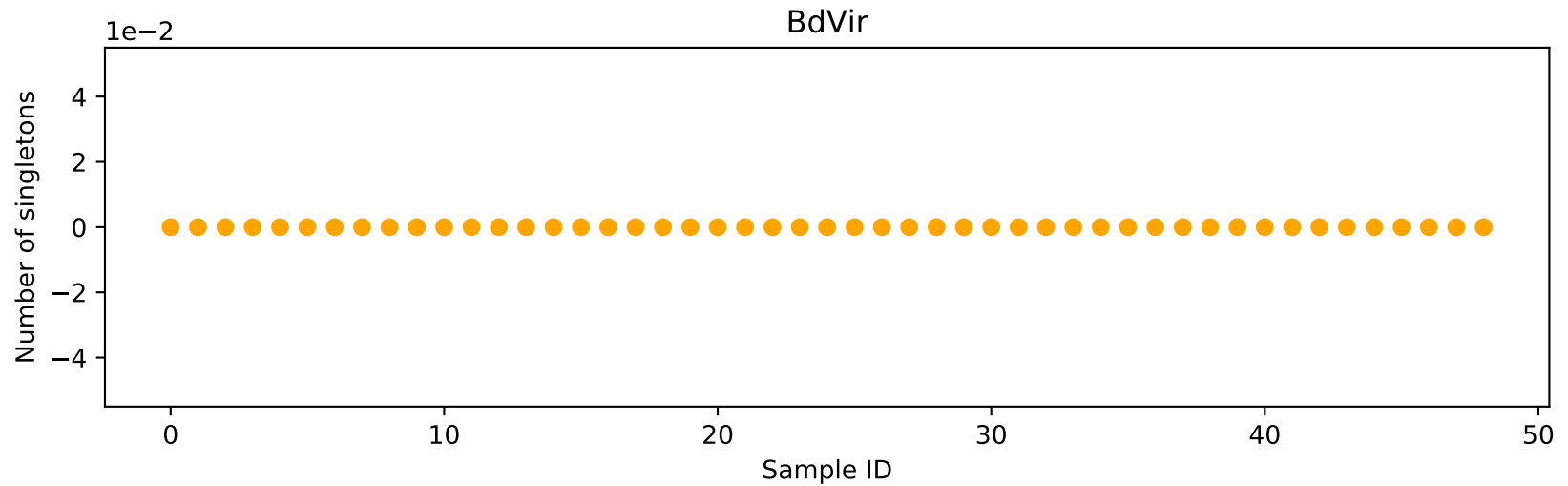
# Ts/Tv by sample



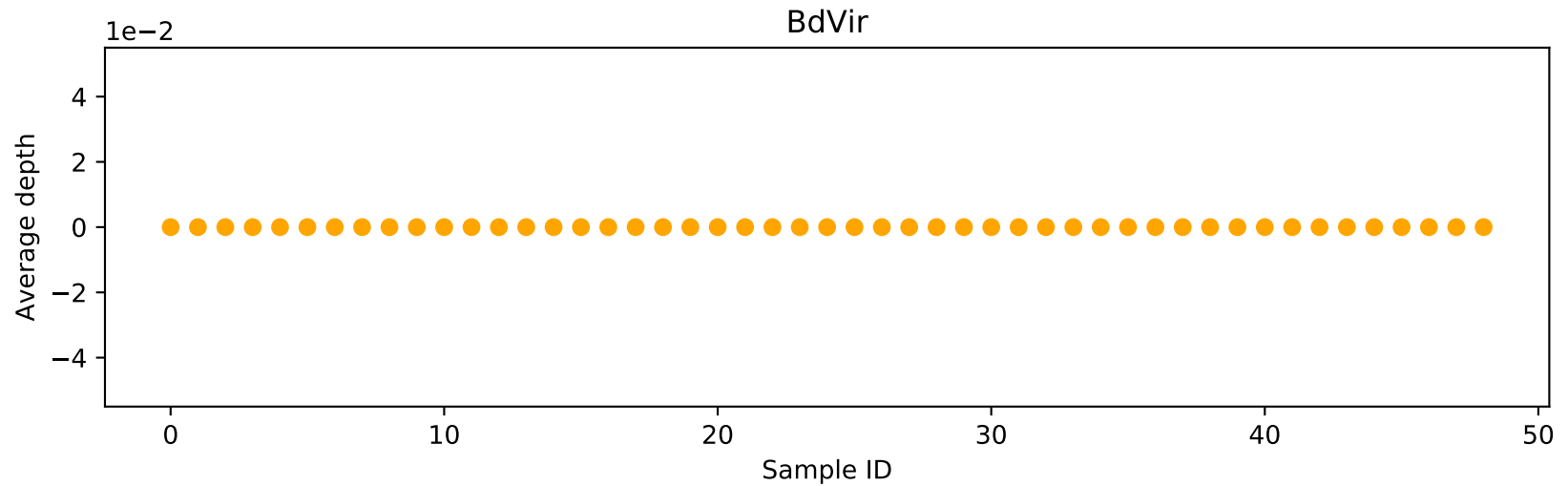
# Hets vs non-ref Homs by sample



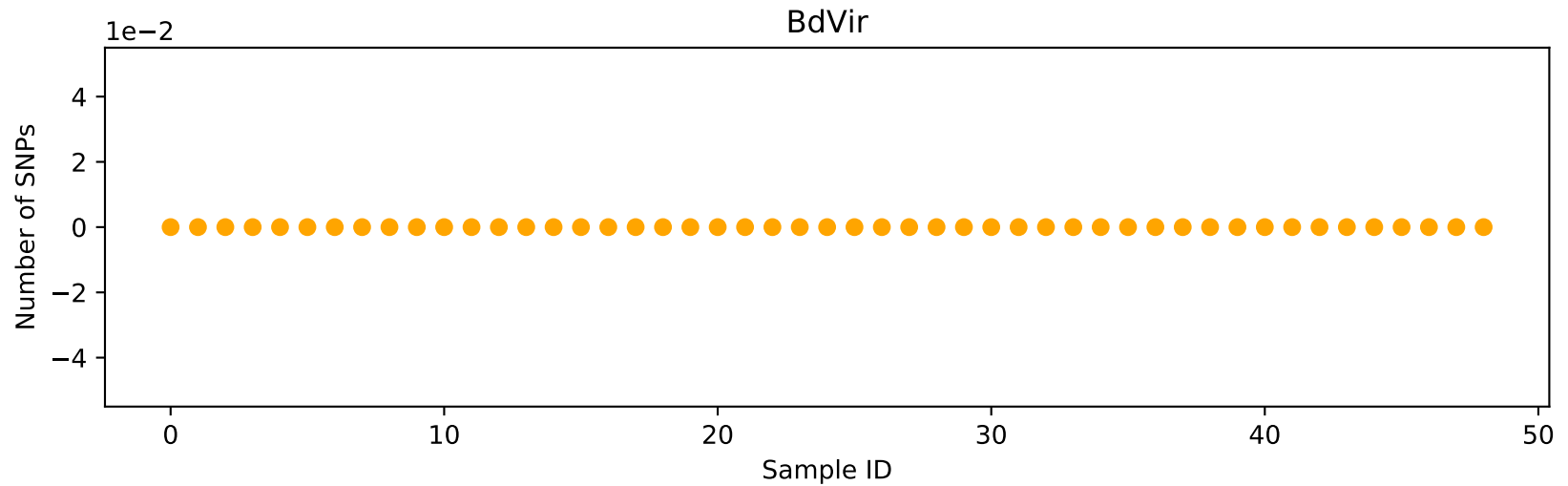
# Singletons by sample (hets and homs)



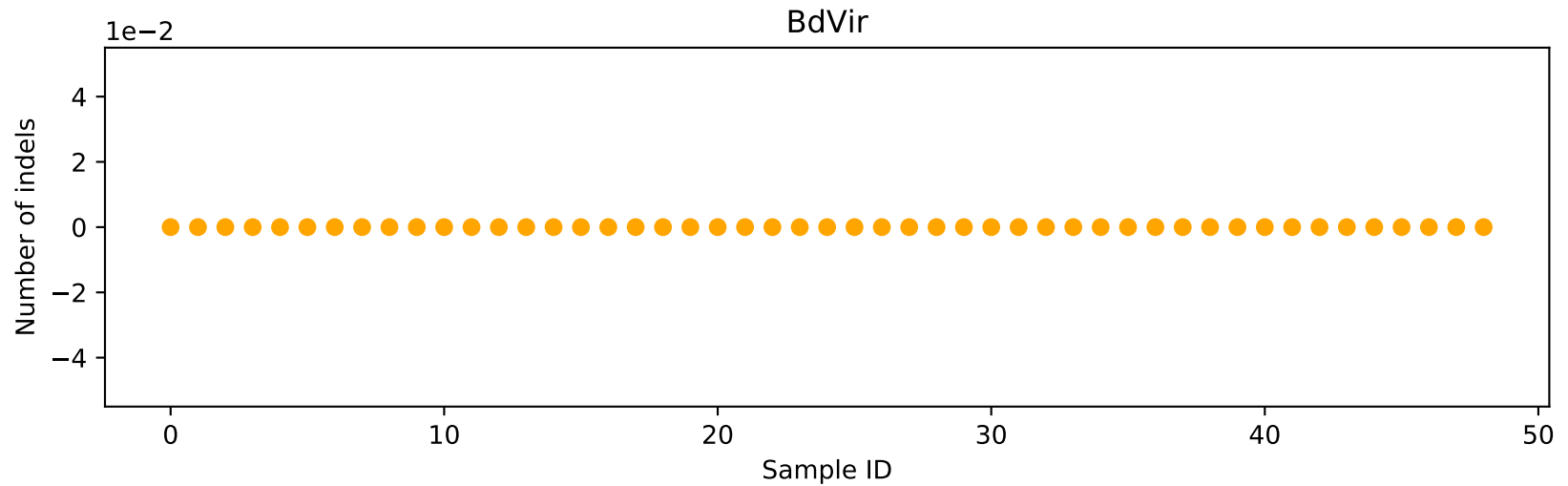
# Average depth by sample



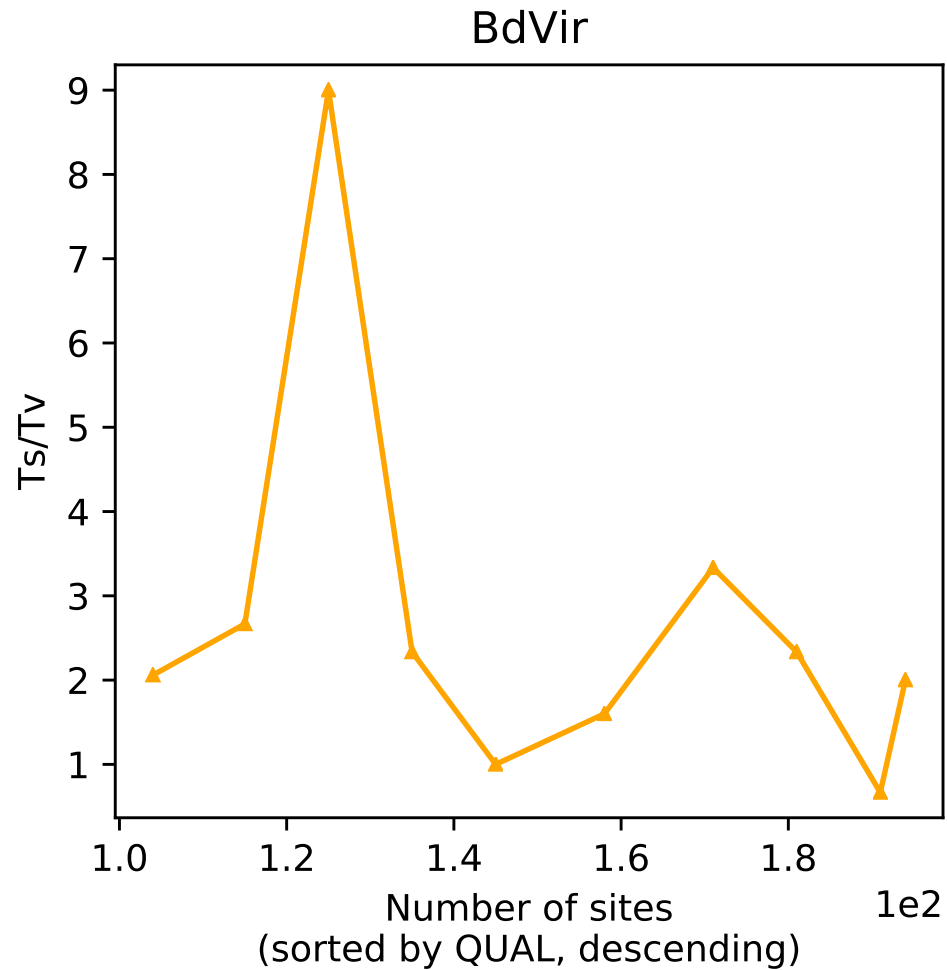
# Number of SNPs by sample



# Number of indels by sample

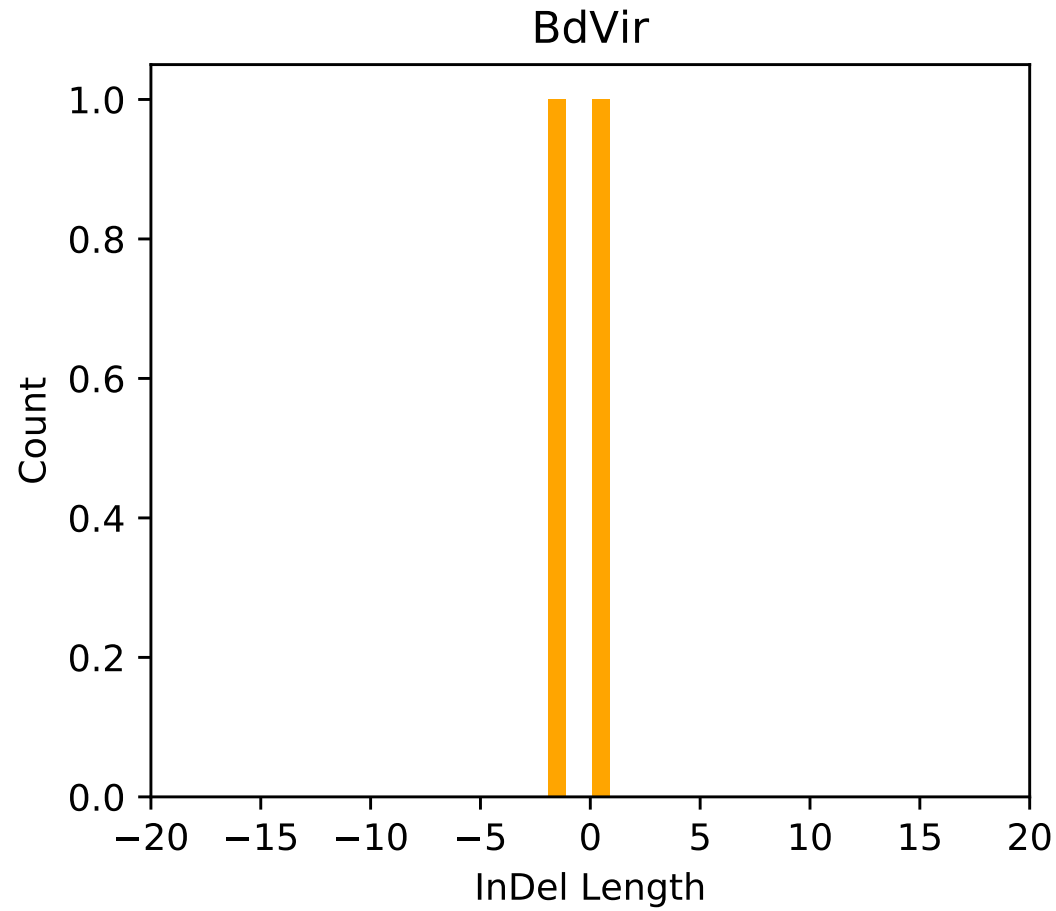


# Ts/Tv stratified by QUAL

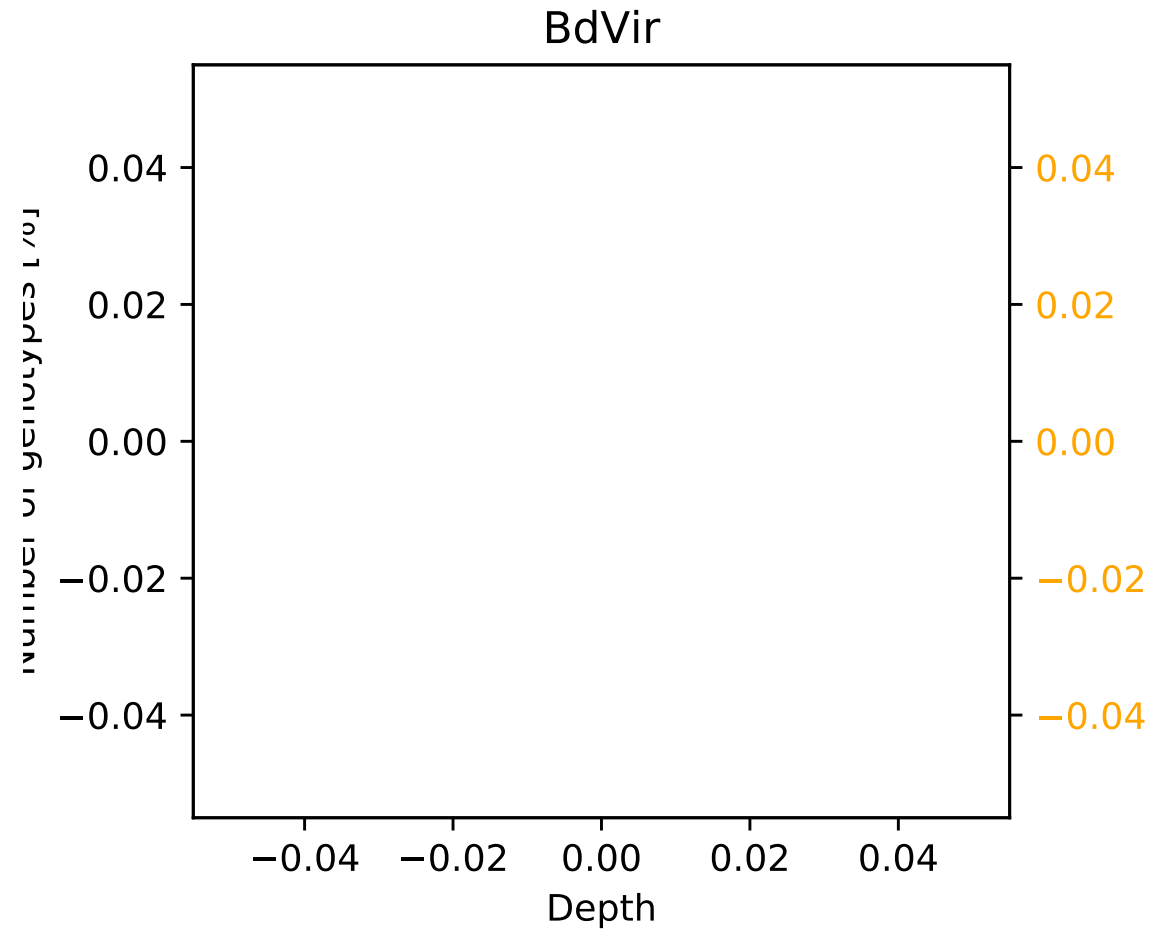




# Indel distribution



# Depth distribution



# Substitution types

