

Summary Numbers

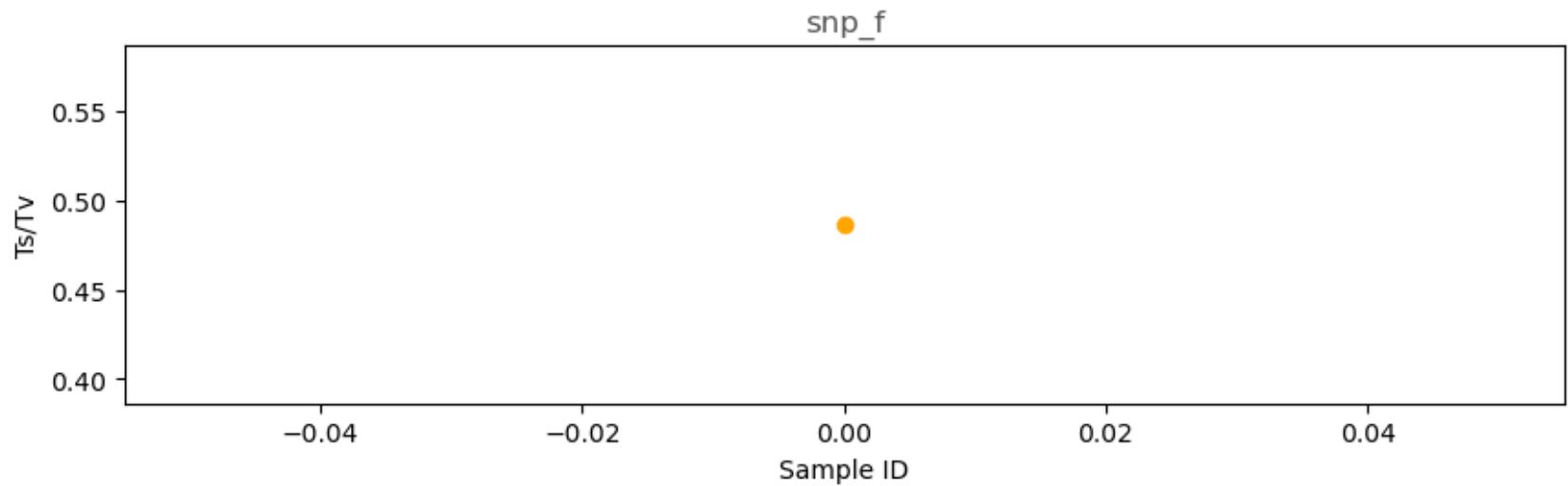
Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
snp_f	633	0.49	0.49	77	—	9	1

* frameshift ratio: out/(out+in)

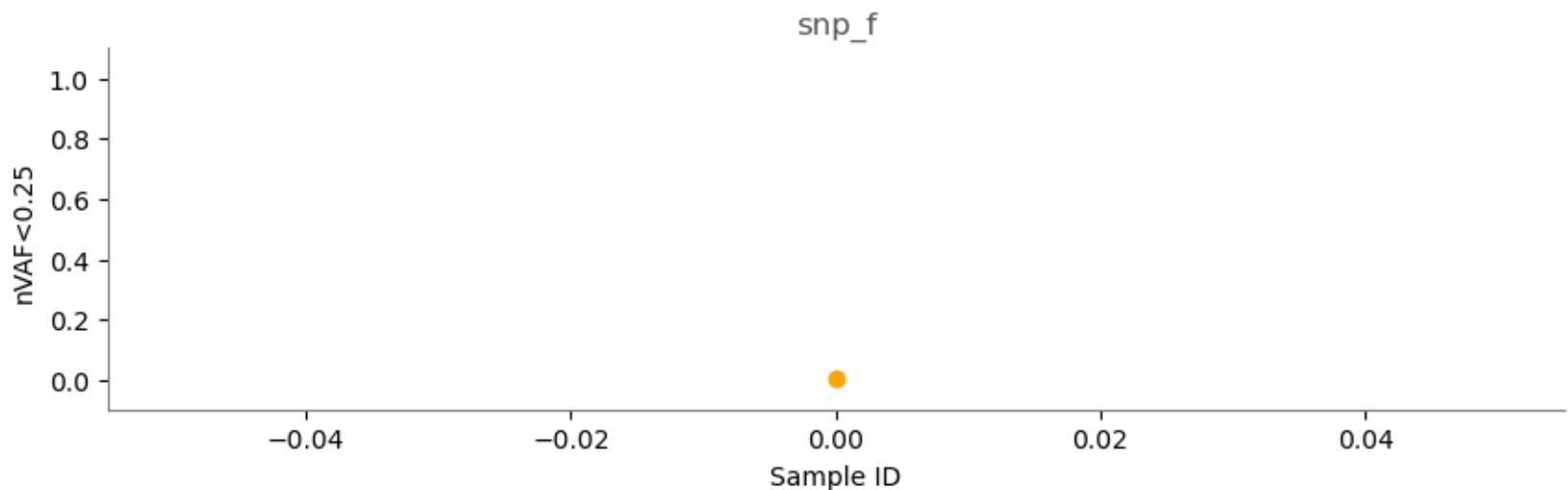
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
snp_f	28.4%	0.57	35.9%	6	0

-.snp_f ...snp_freebayes.vcf.gz

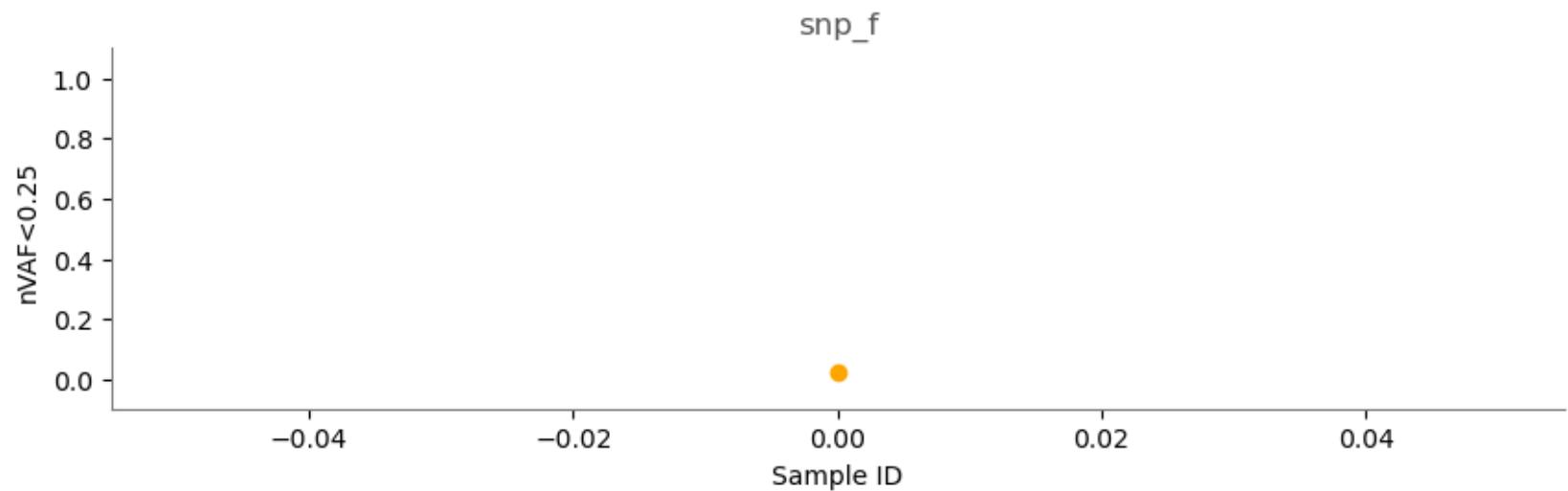
T_s/T_v by sample



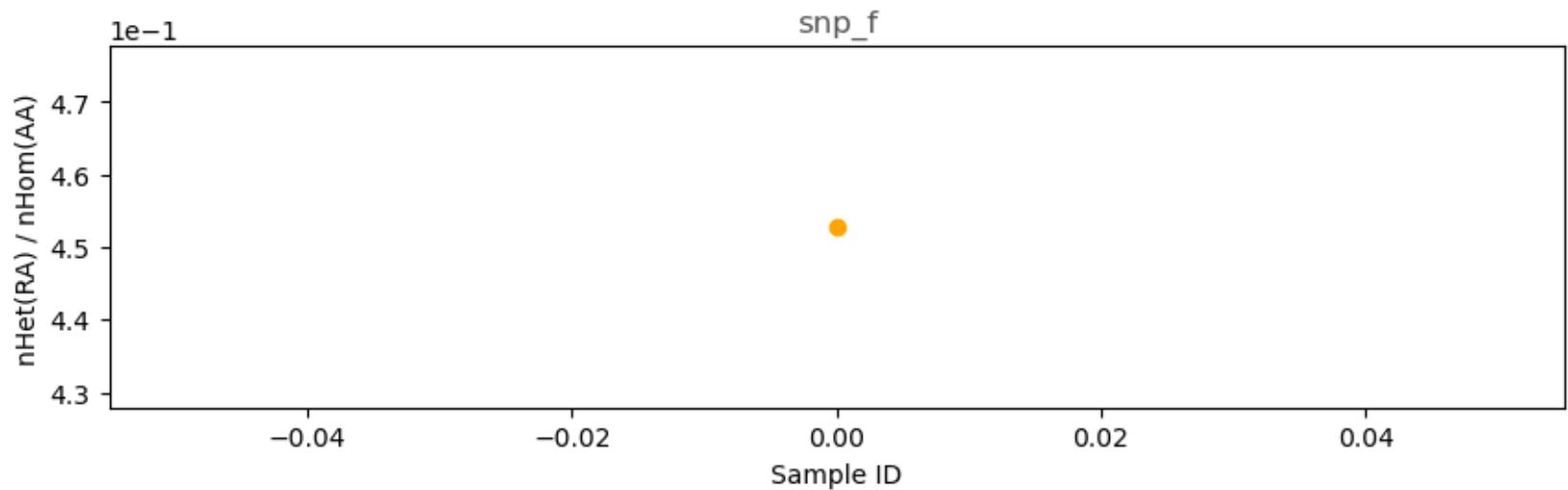
Fraction of SNVs with VAF<25% by sample



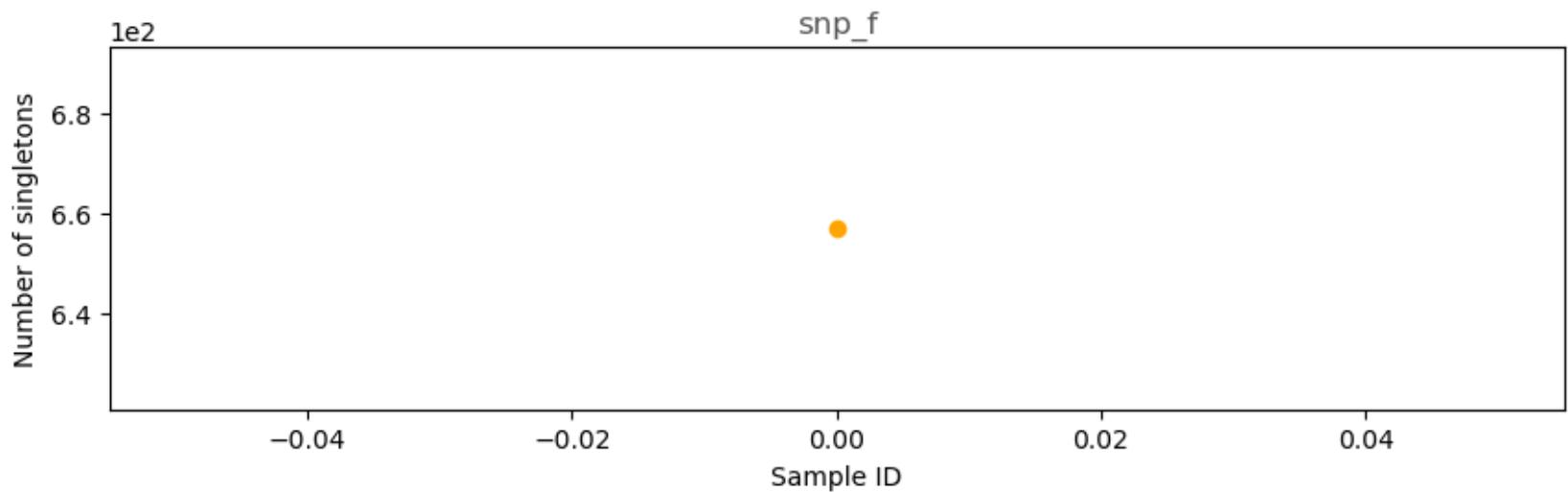
Fraction of indels with VAF<25% by sample



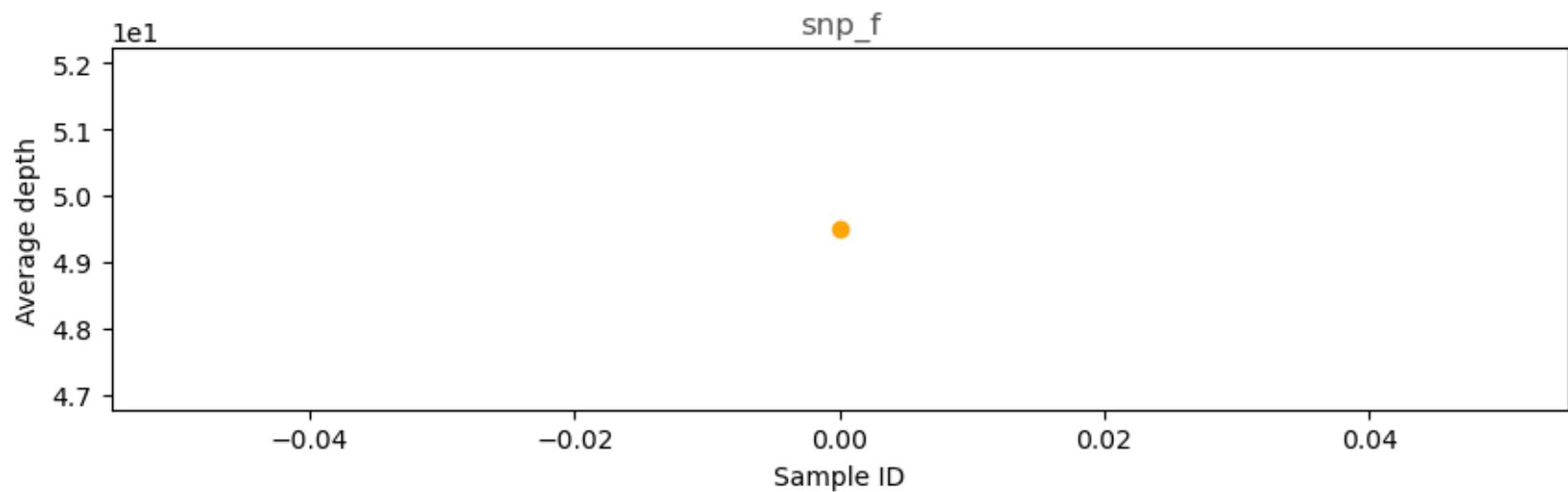
Hets vs non-ref Homs by sample



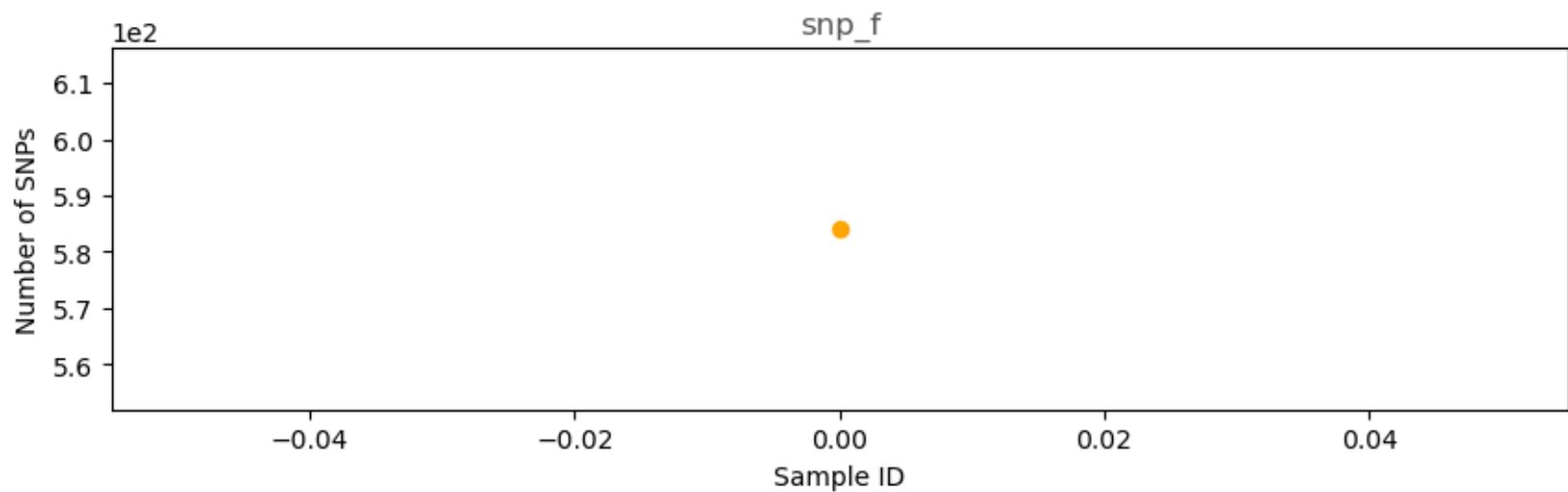
Singletons by sample (hets and homs)



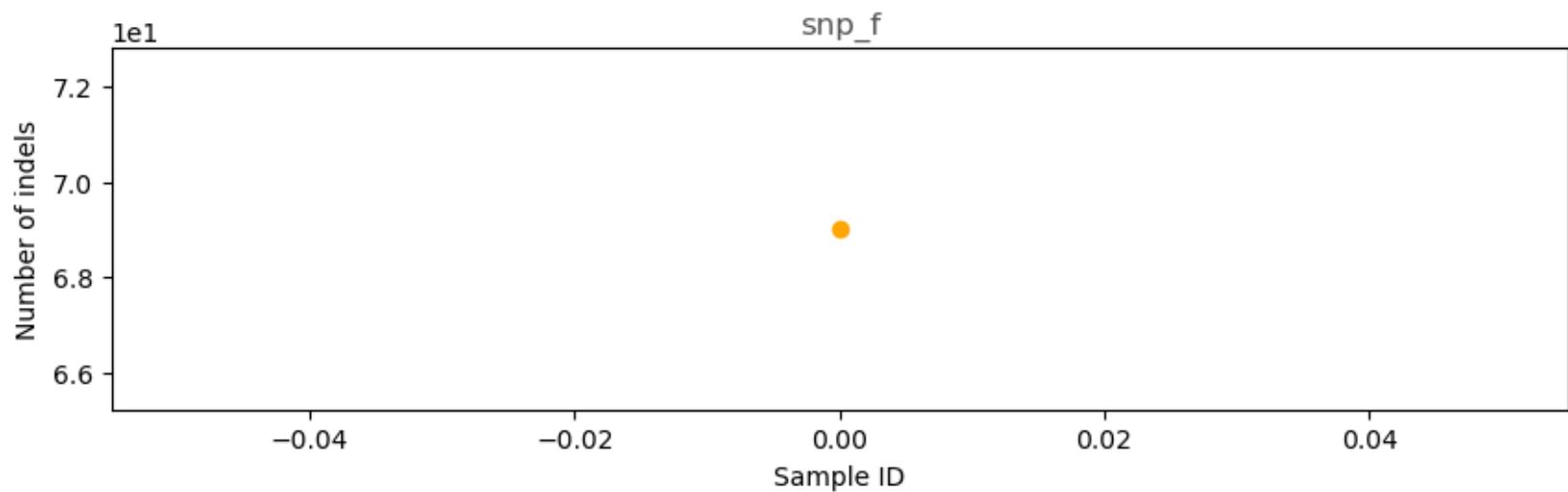
Average depth by sample



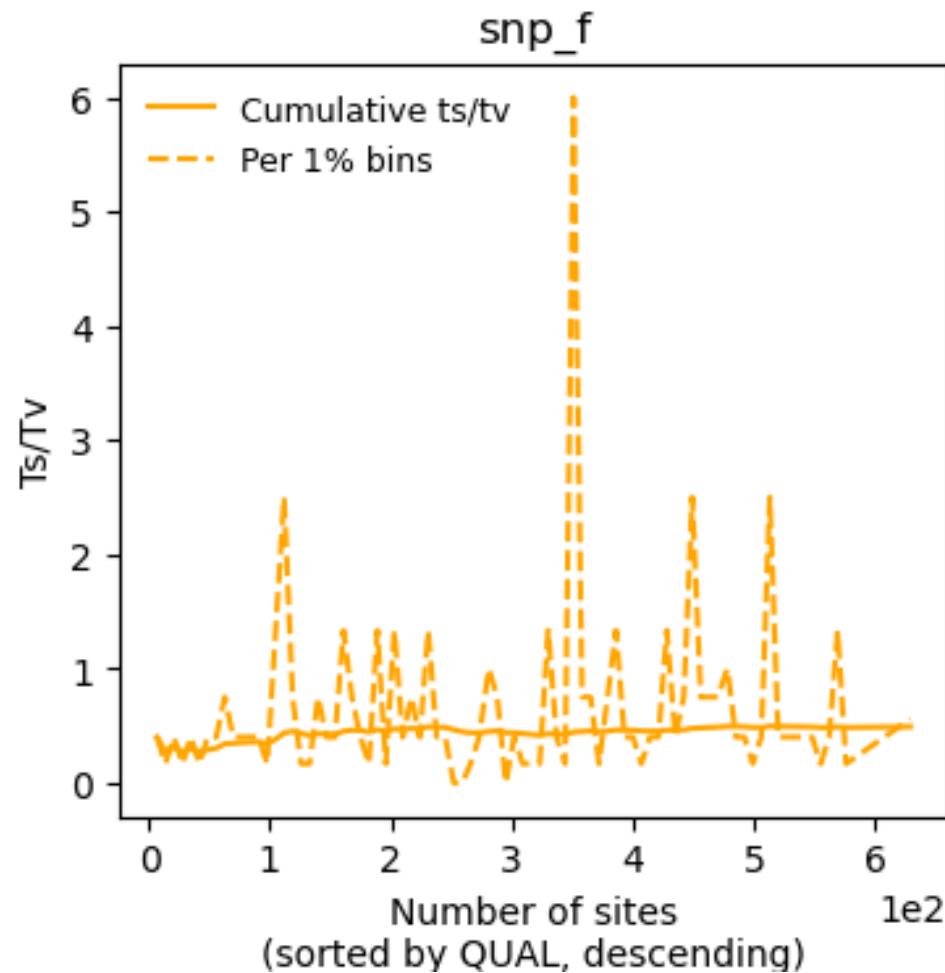
Number of SNPs by sample



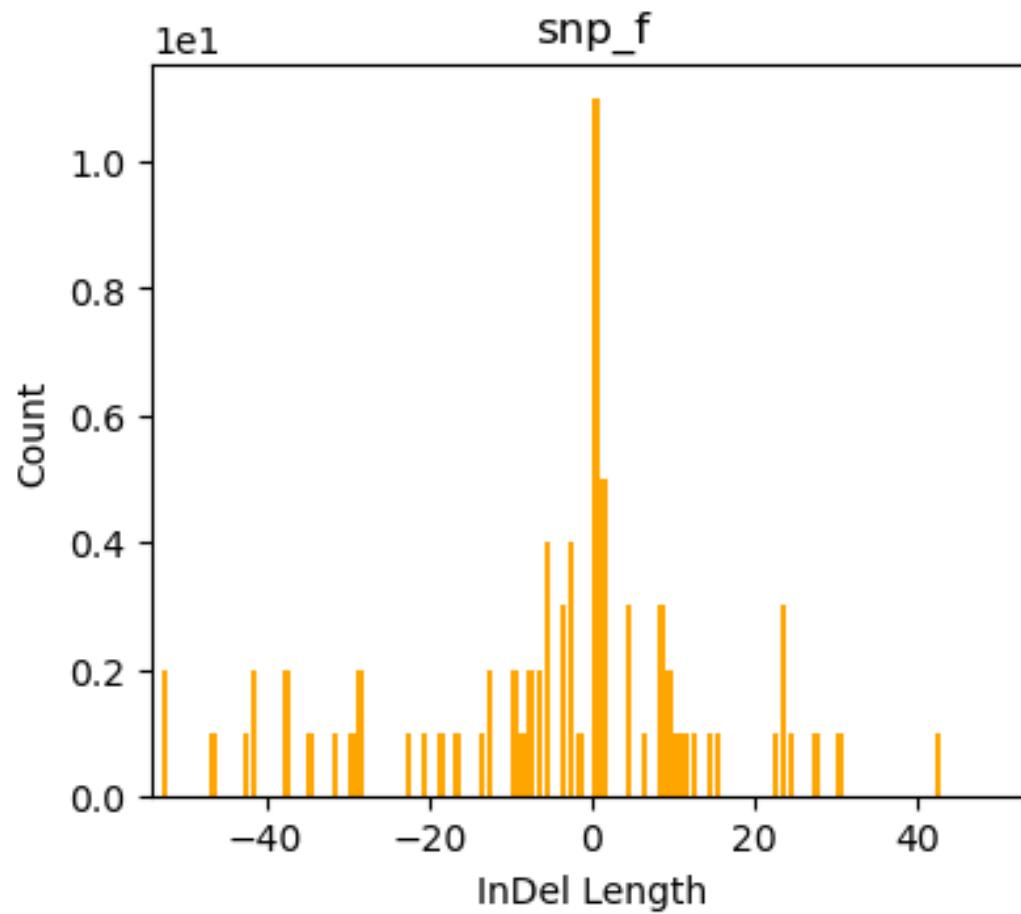
Number of indels by sample



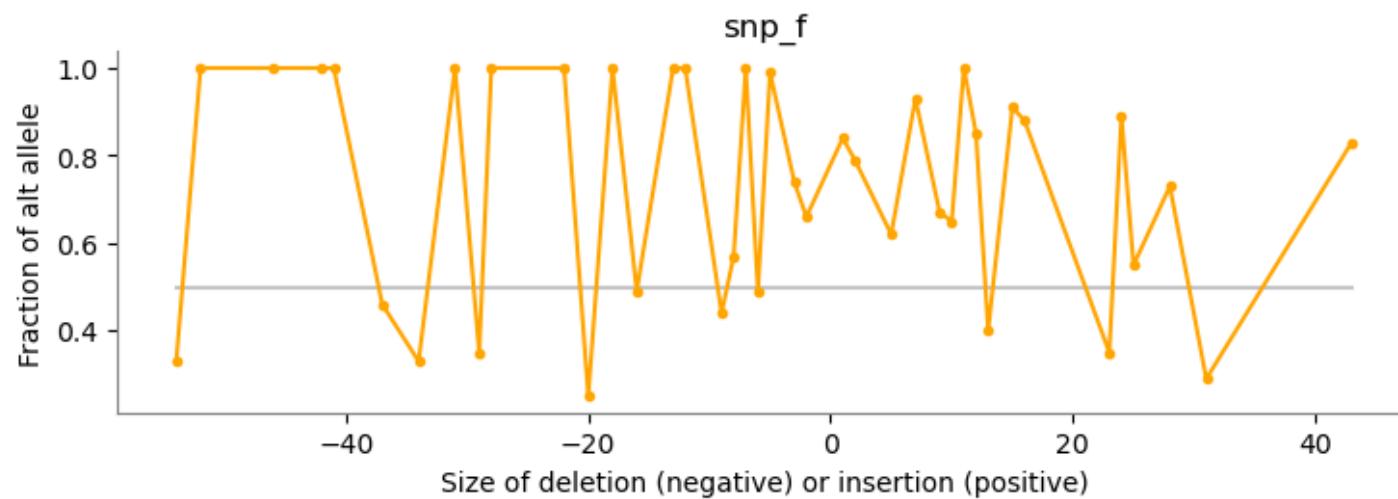
Ts/Tv stratified by QUAL



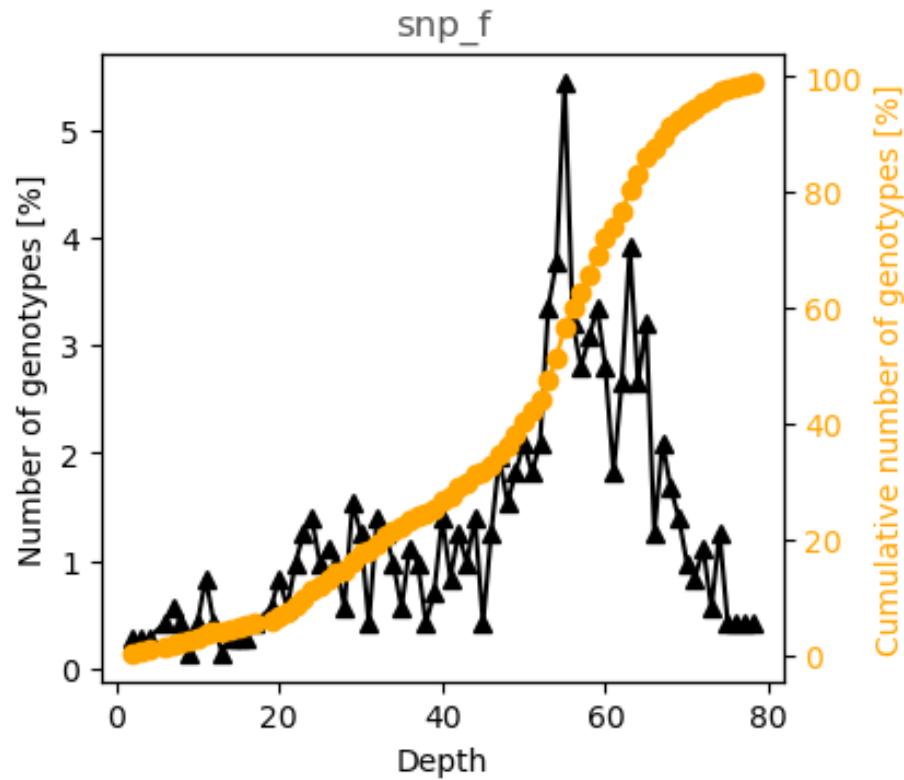
Indel distribution



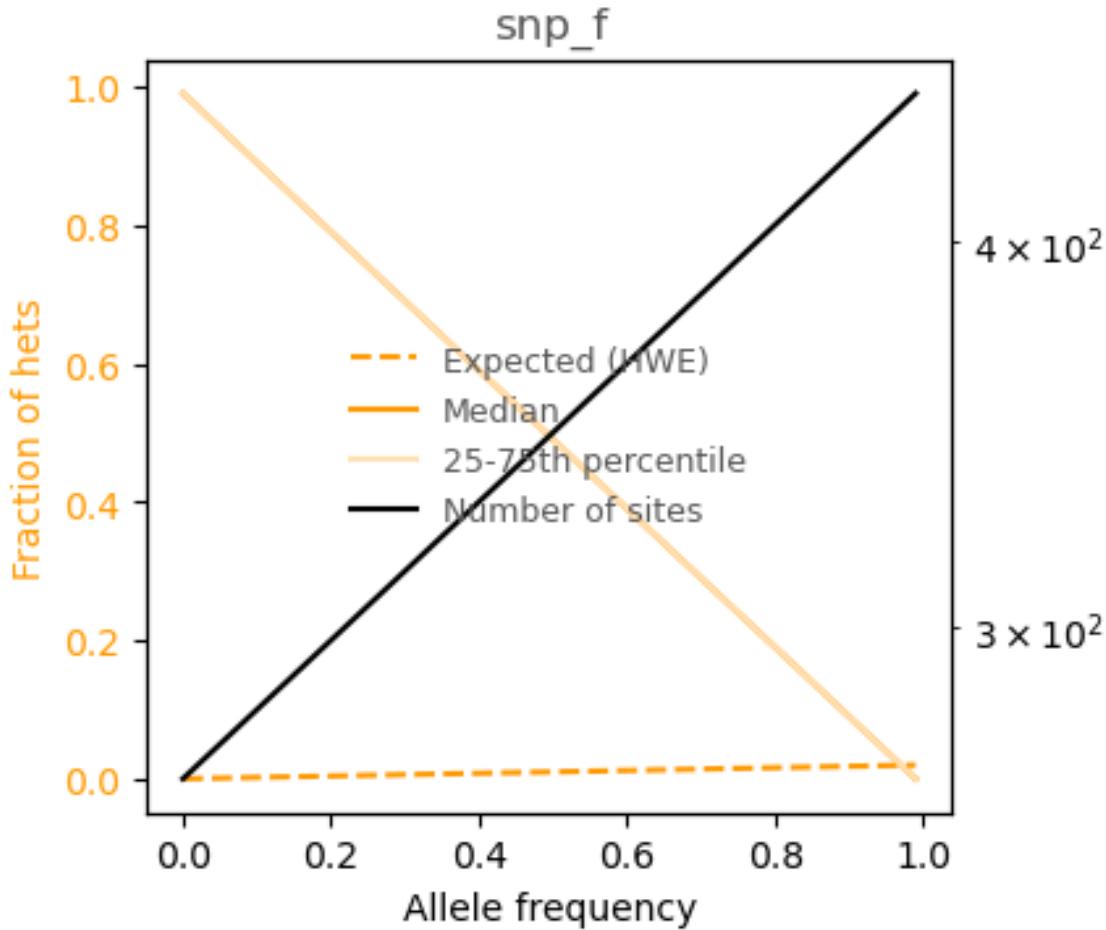
Fraction of alternate indel allele



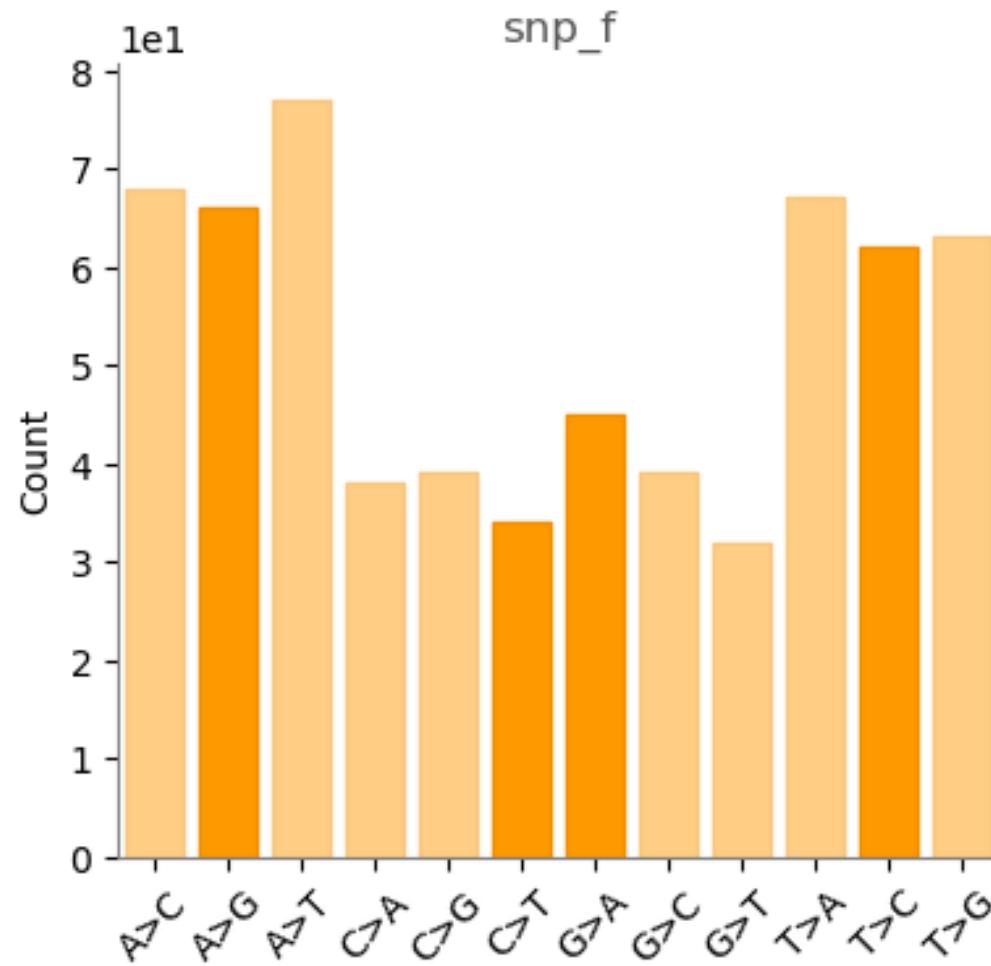
Depth distribution



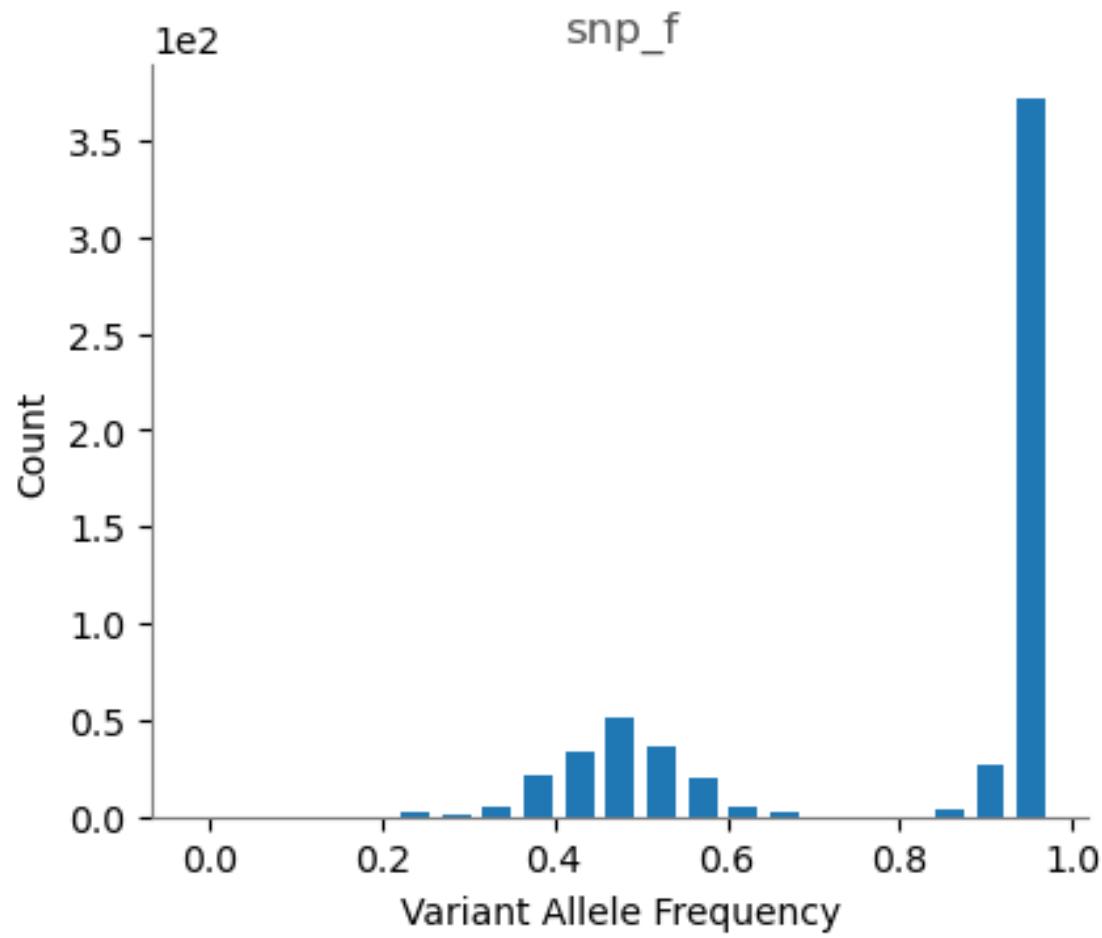
Number of HETs by AF



Substitution types



SNV Variant Allele Frequency



Indel Variant Allele Frequency

