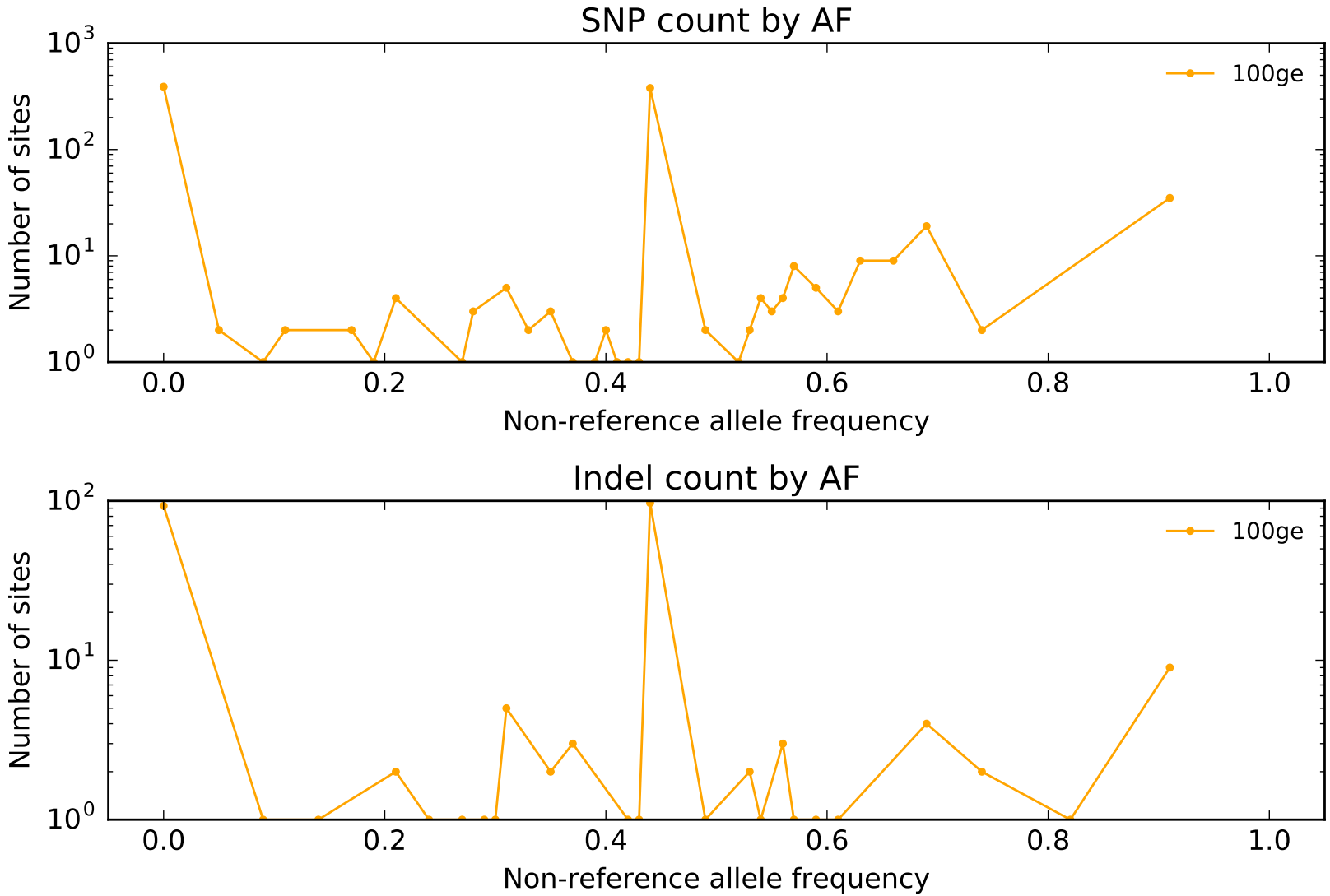
Summary Numbers

	SNPs			indels			MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	rc**		
100ge	874	1.67	1.70	213	–	–	0	0

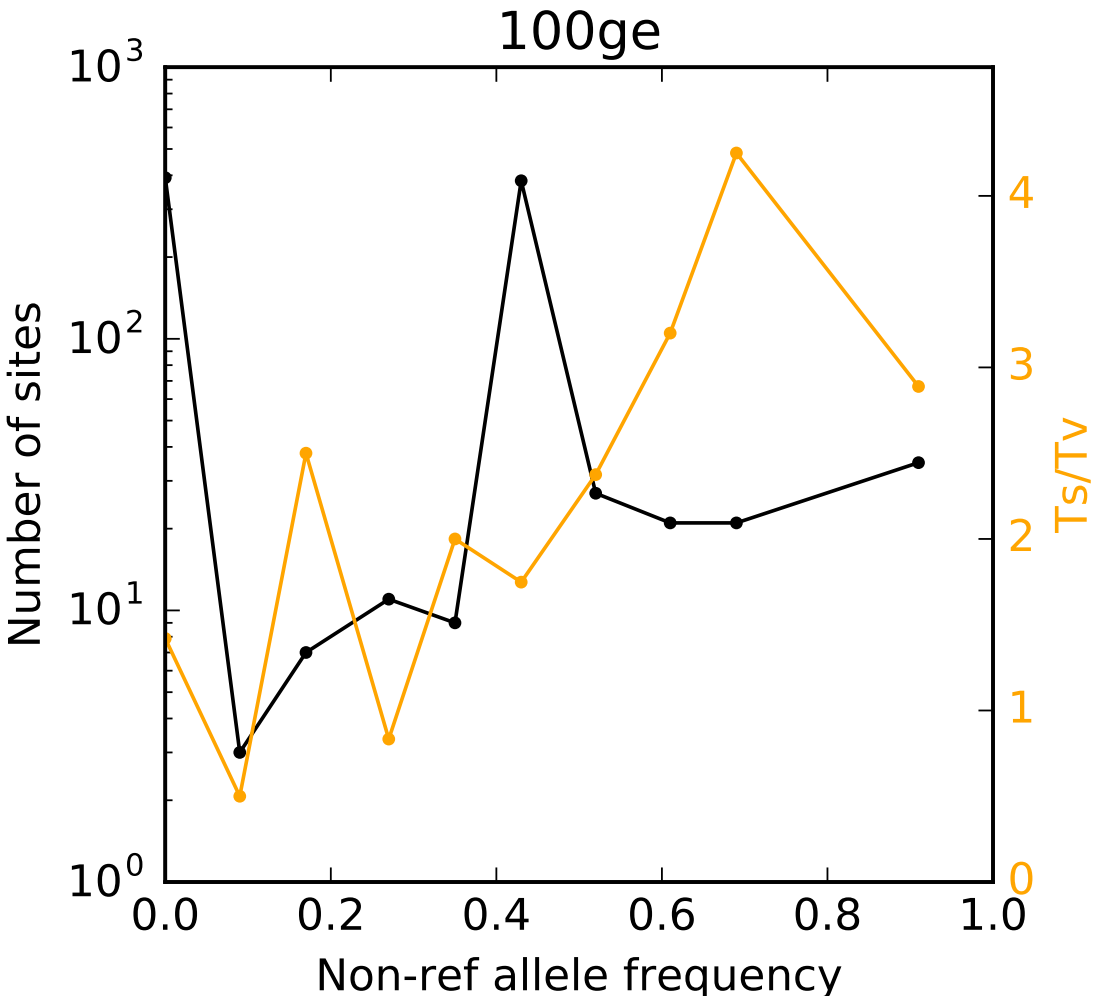
* frameshift ratio: out/(out+in); ** repeat-consistency

	singletons (AC=1)				multiallelic	
Callset	SNPs	ts/tv	indels	rc	sites	SNPs
100ge	43.0%	1.44	39.0%	0.000	54	34

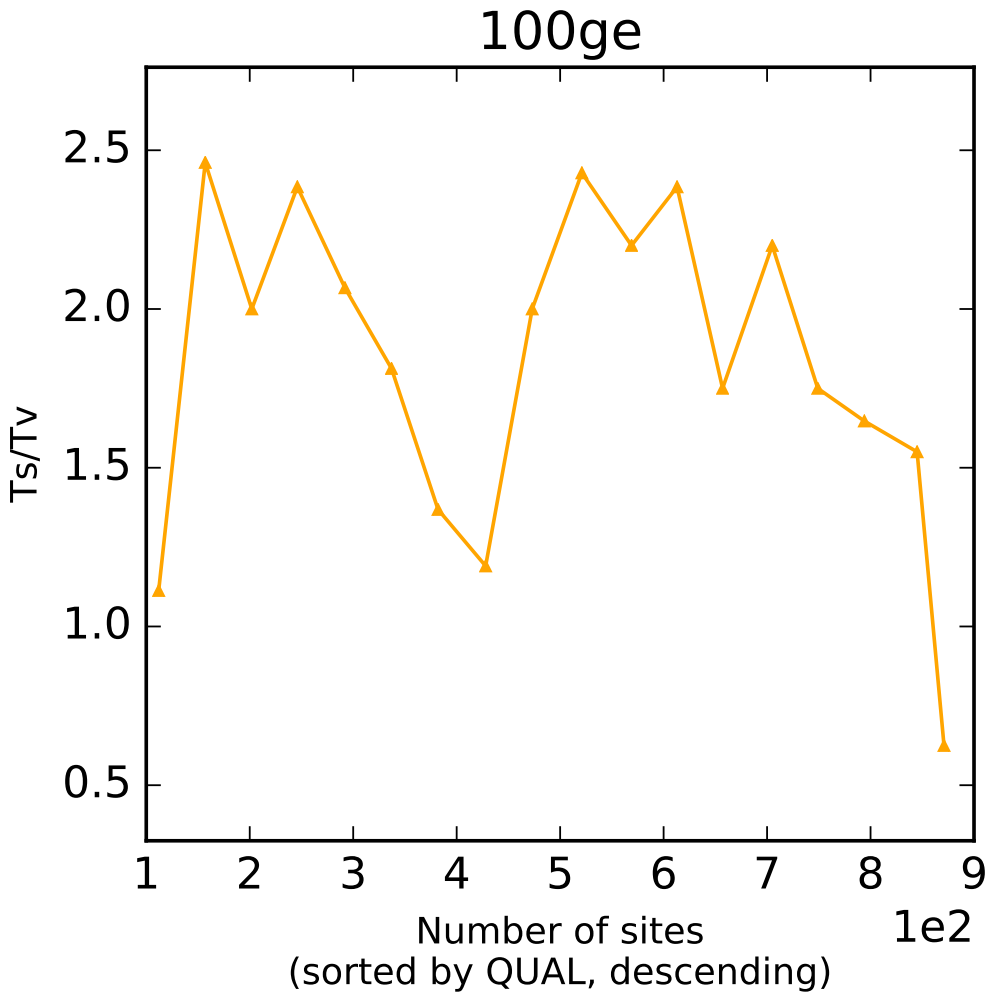
- 100ge .. `100genomes_merged.vcf`


SNP count by AF
Number of sites
10^3
10^2
10^1
10^0
0.0
0.2
0.4
0.6
0.8
1.0
Non-reference allele frequency
100ge
Indel count by AF
Number of sites
10^2
10^1
10^0
0.0
0.2
0.4
0.6
0.8
1.0
Non-reference allele frequency
100ge

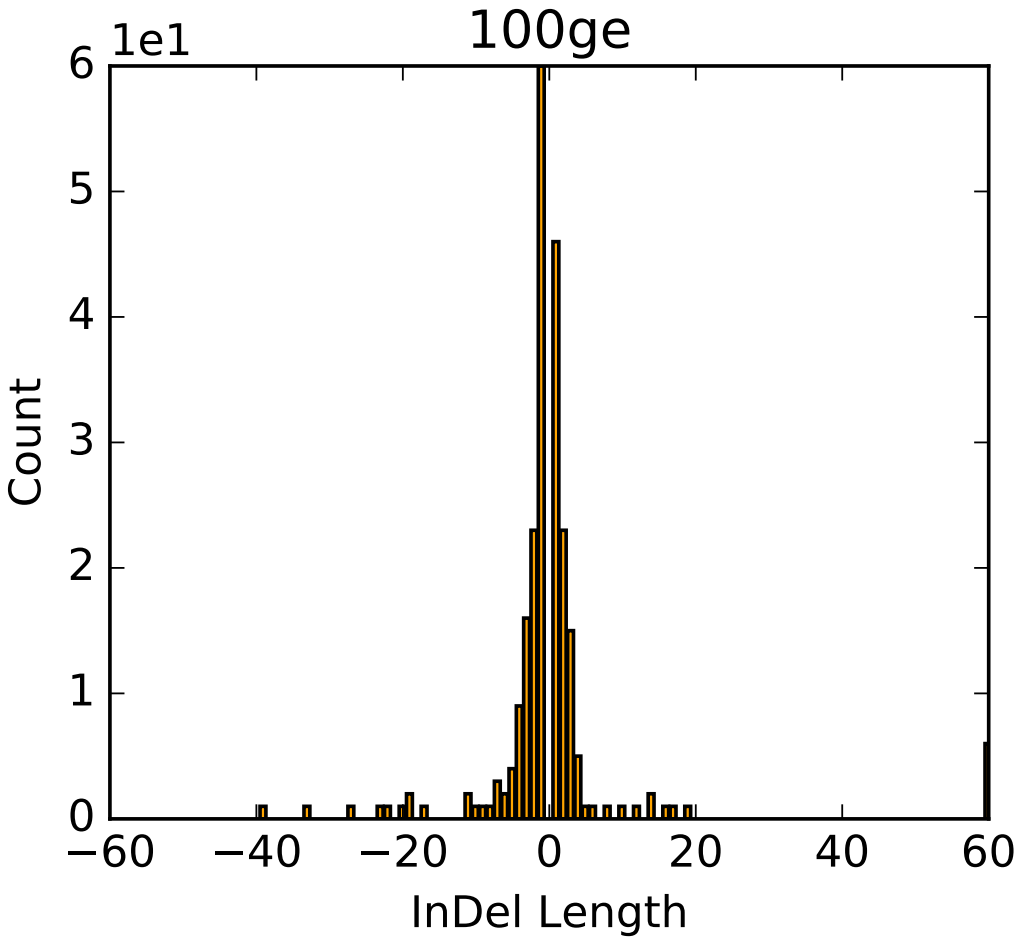
Ts/Tv by AF


100ge
Number of sites
Ts/Tv
Non-ref allele frequency
10^3
10^2
10^1
10^0
0.0
0.2
0.4
0.6
0.8
1.0
0
1
2
3
4

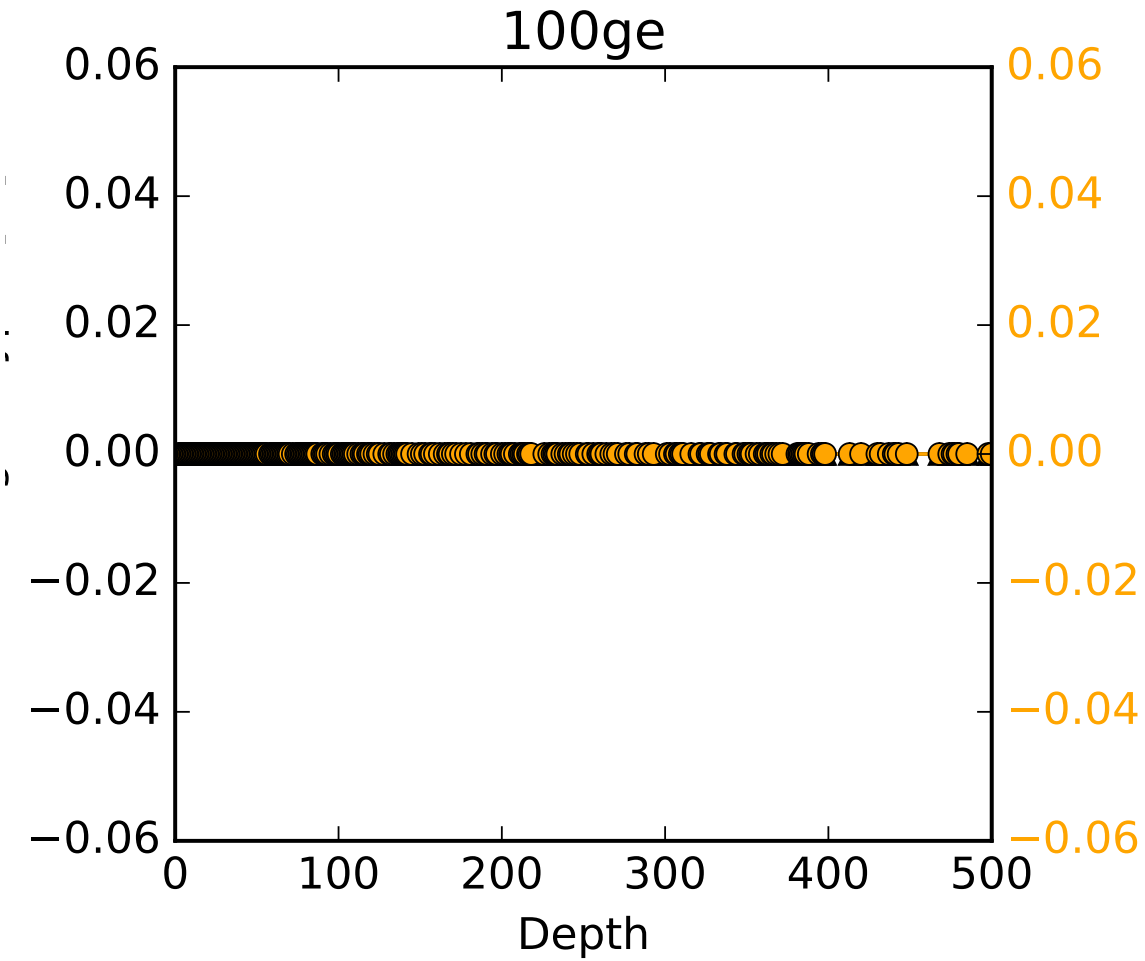
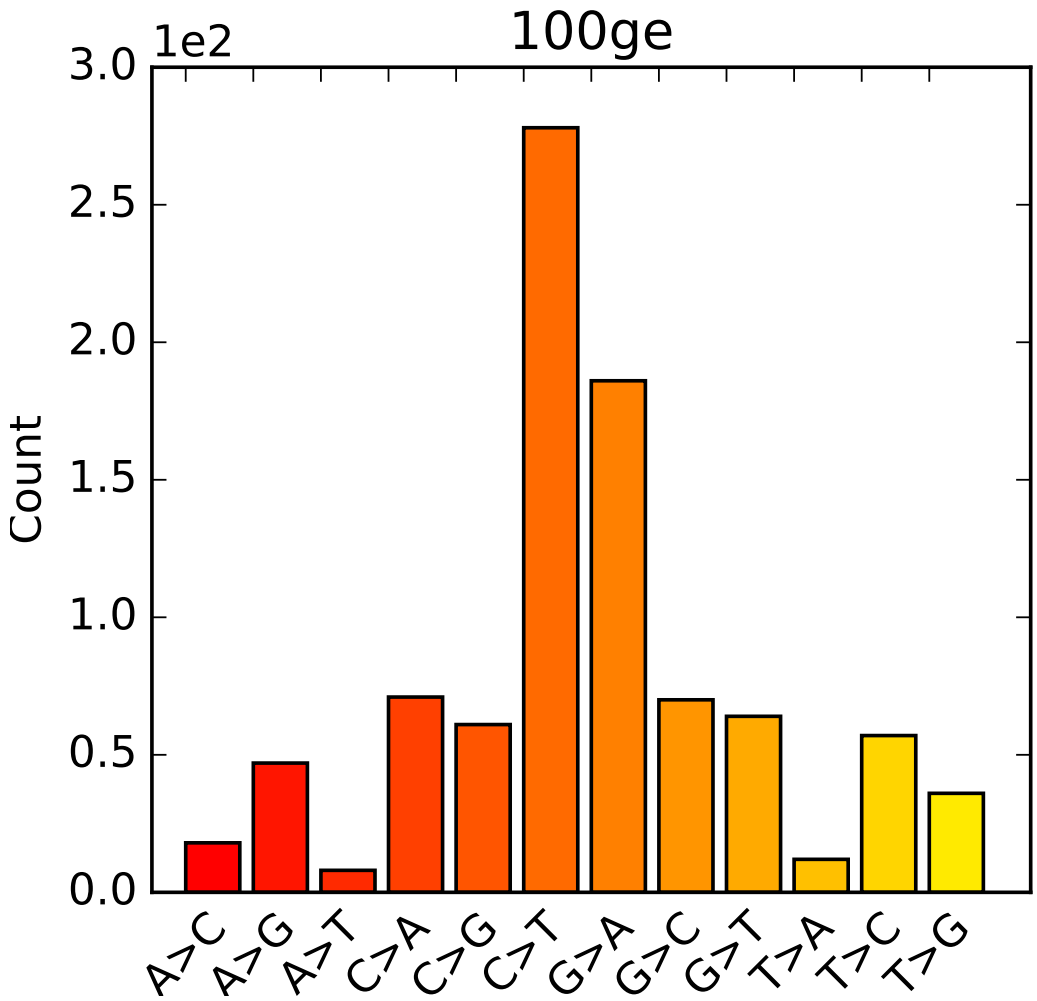
Ts/Tv stratified by QUAL



Indel distribution



Depth distribution



Substitution types