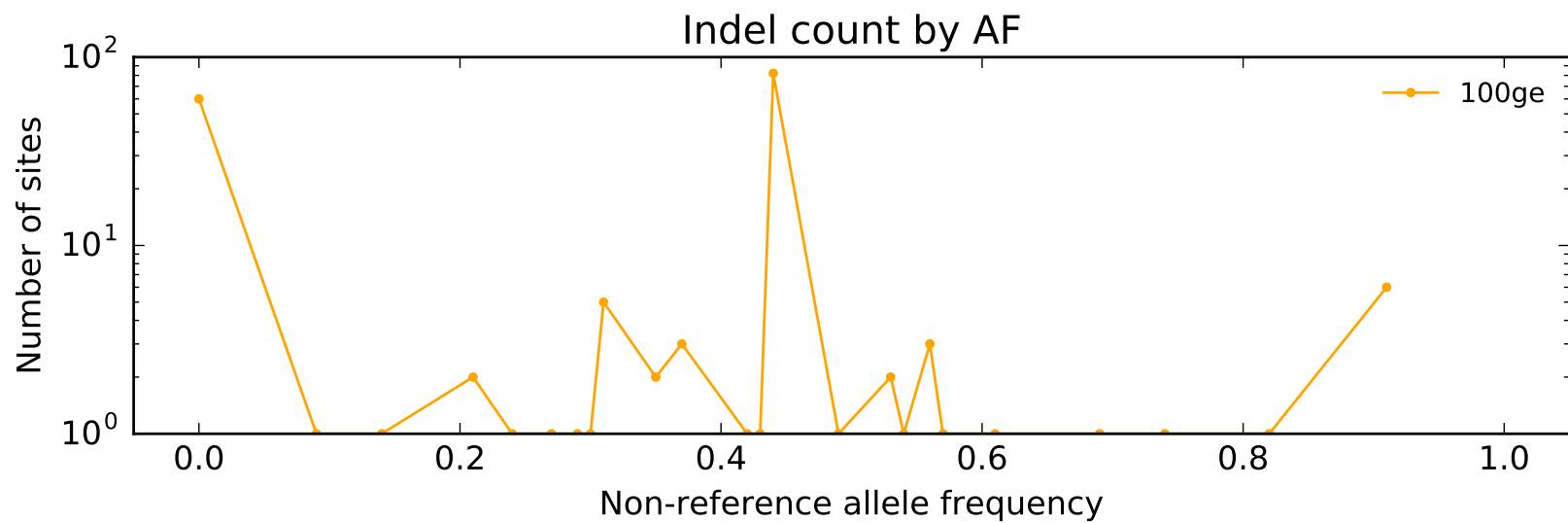
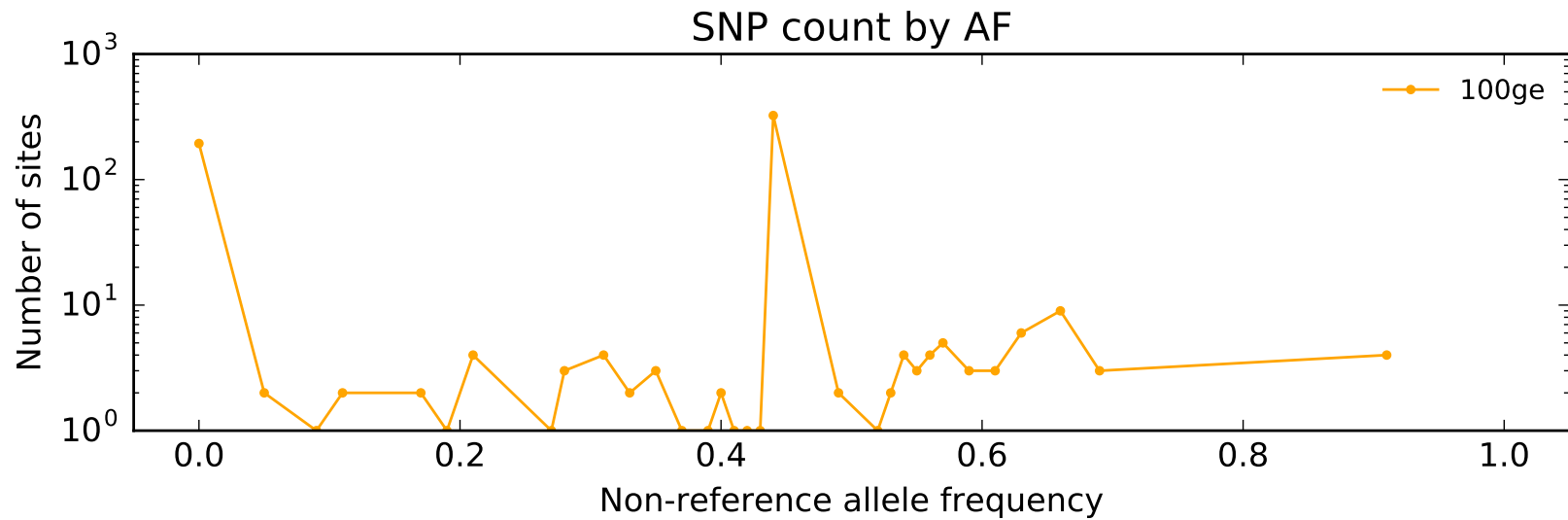


Summary Numbers

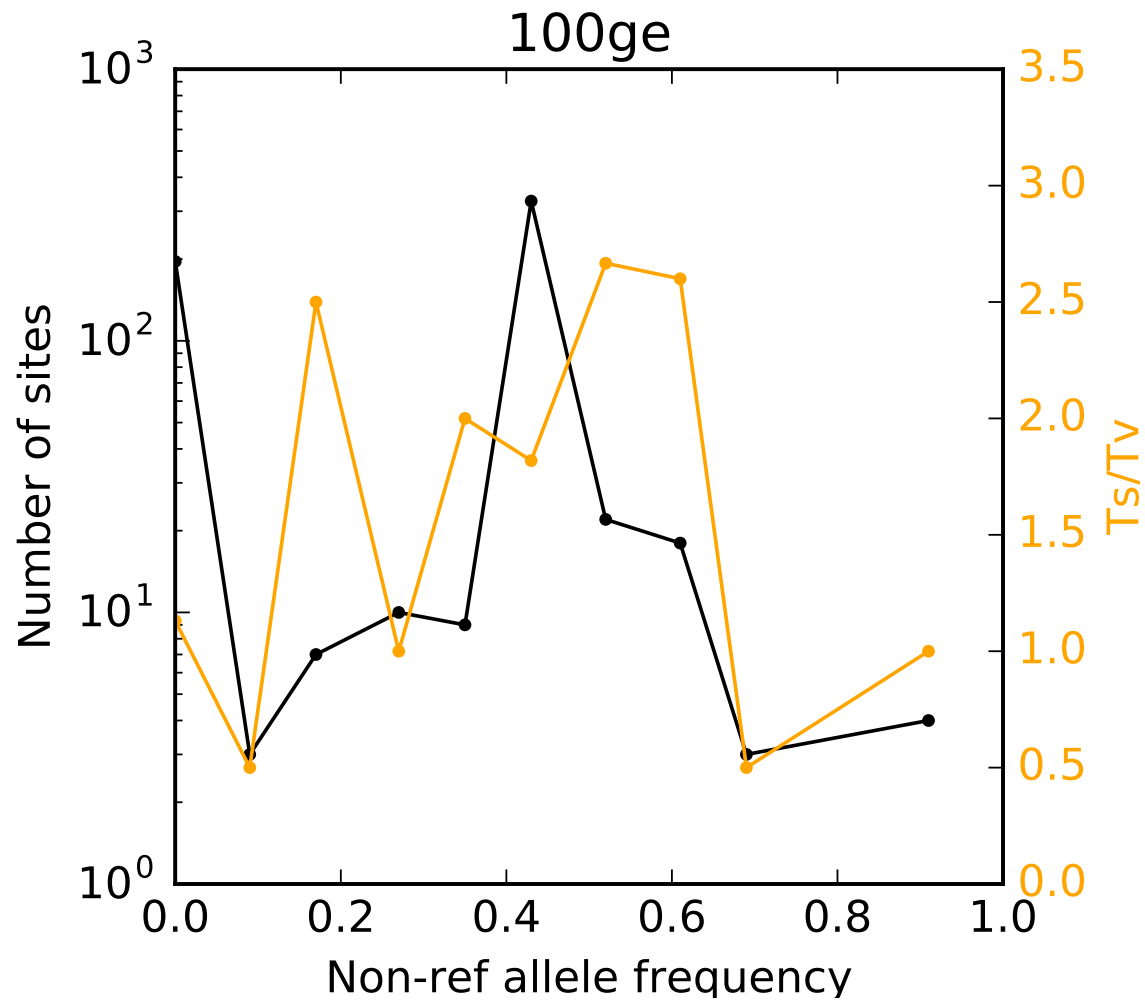
	SNPs			indels			MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	rc**		
100ge	569	1.55	1.61	157	–	–	0	0
* frameshift ratio: out/(out+in); ** repeat-consistency								

	singletons (AC=1)				multiallelic	
Callset	SNPs	ts/tv	indels	rc	sites	SNPs
100ge	32.4%	1.16	32.8%	0.000	50	30

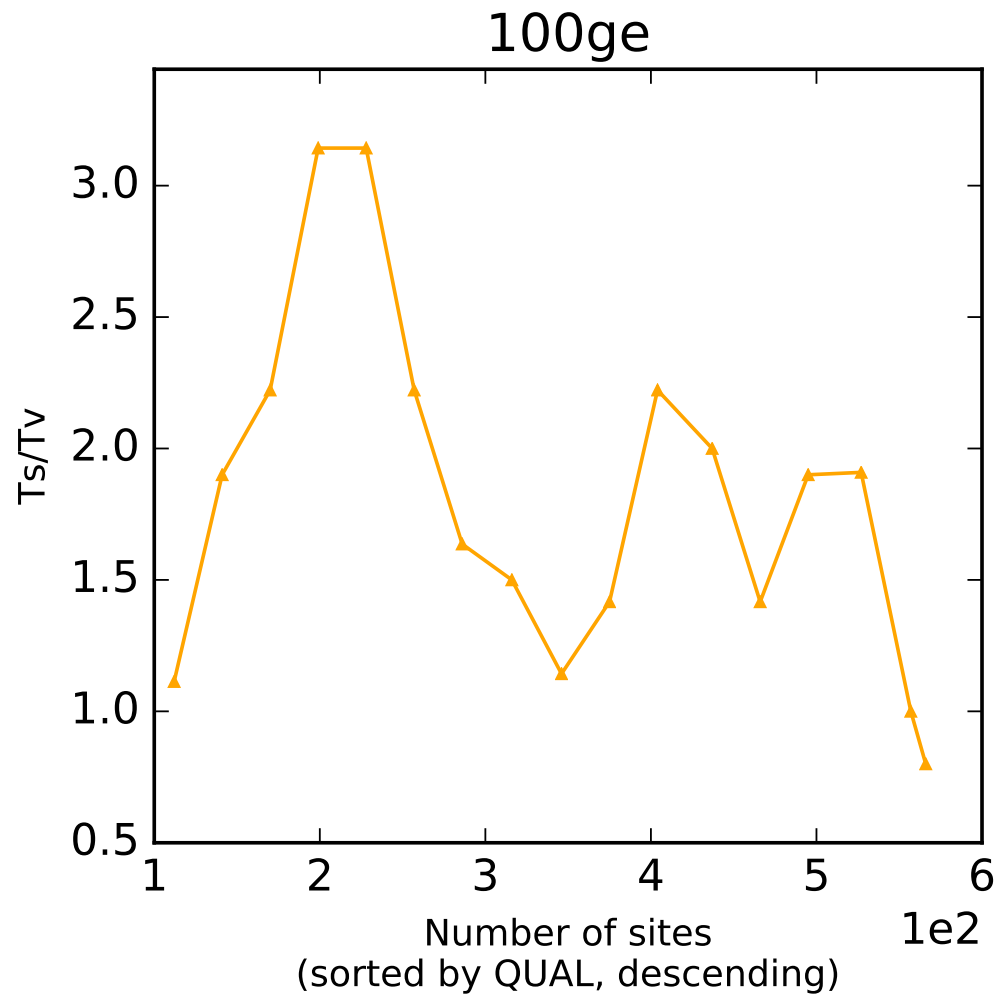
- 100ge .. 100genomes_DP30.vcf



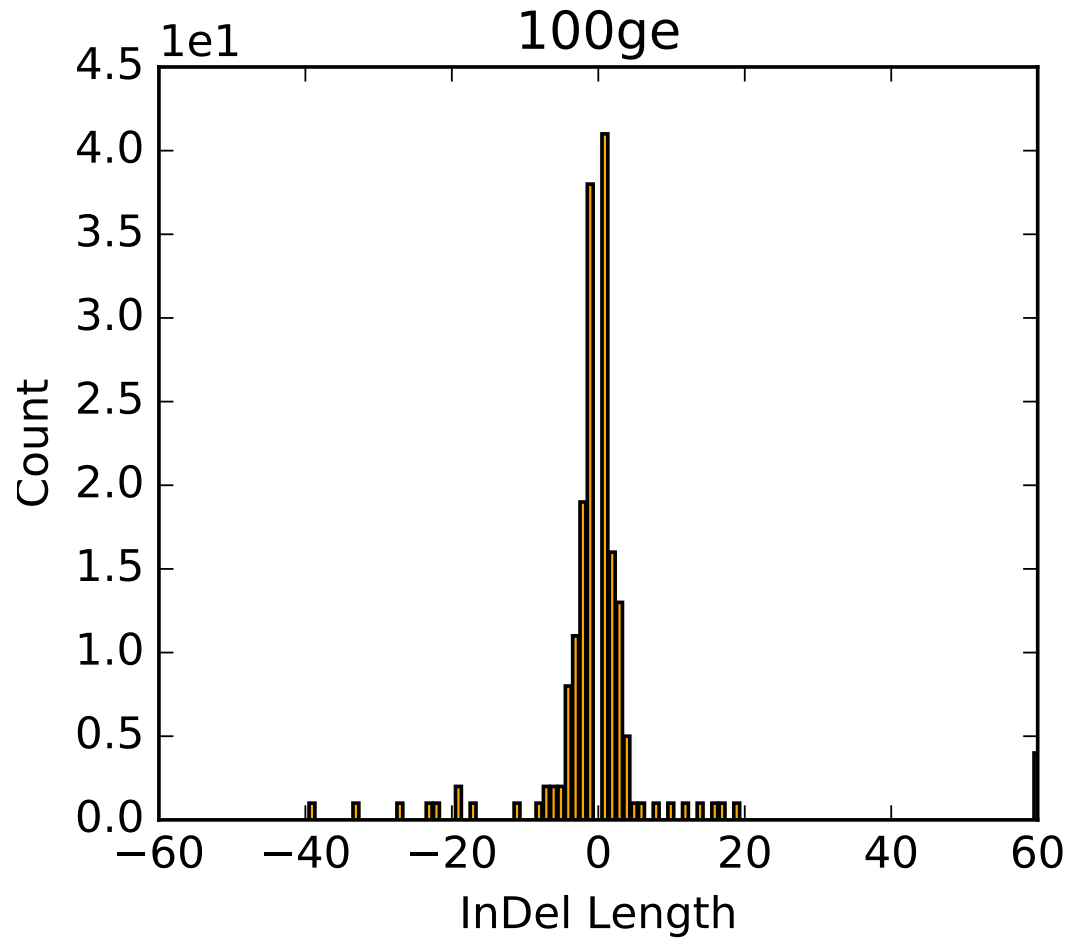
Ts/Tv by AF



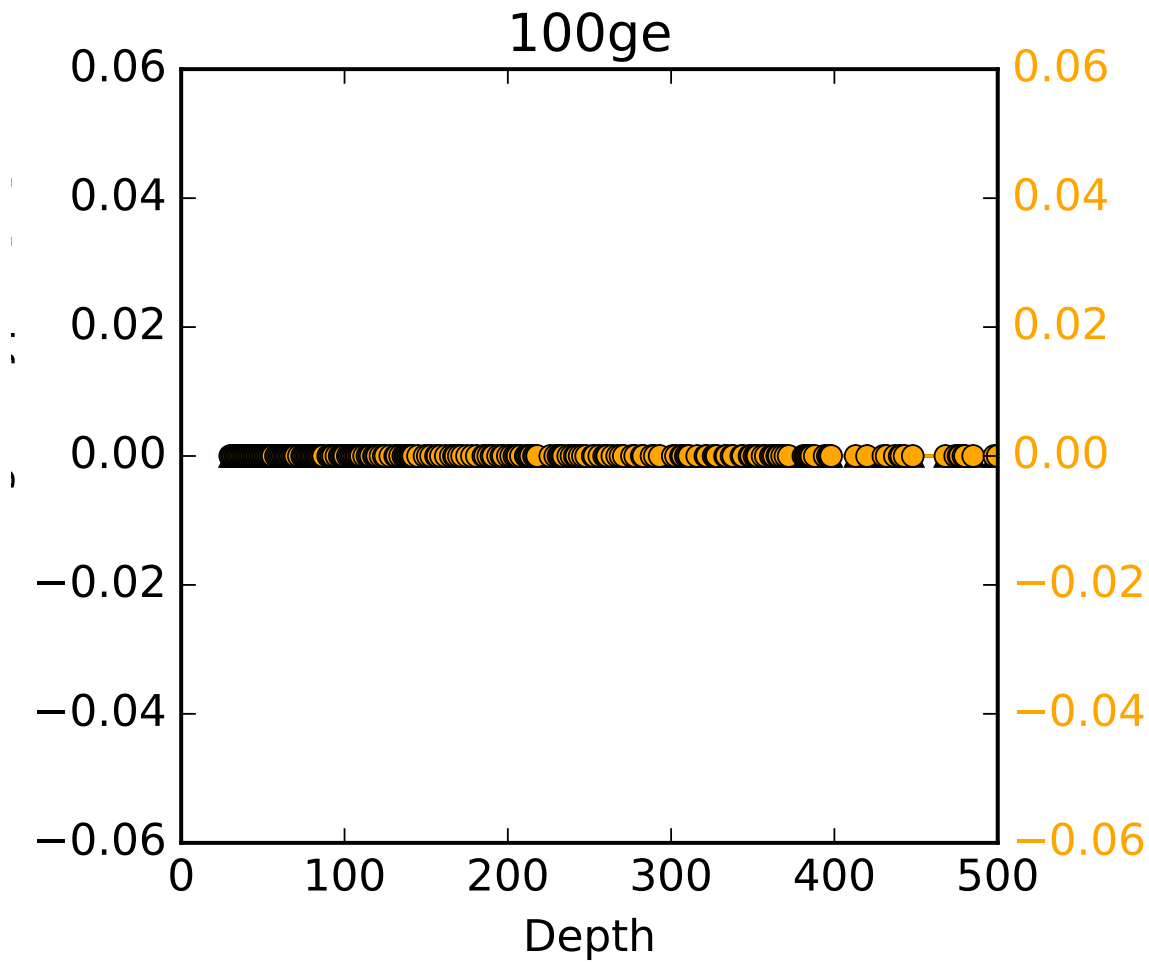
Ts/Tv stratified by QUAL



Indel distribution



Depth distribution



Substitution types

