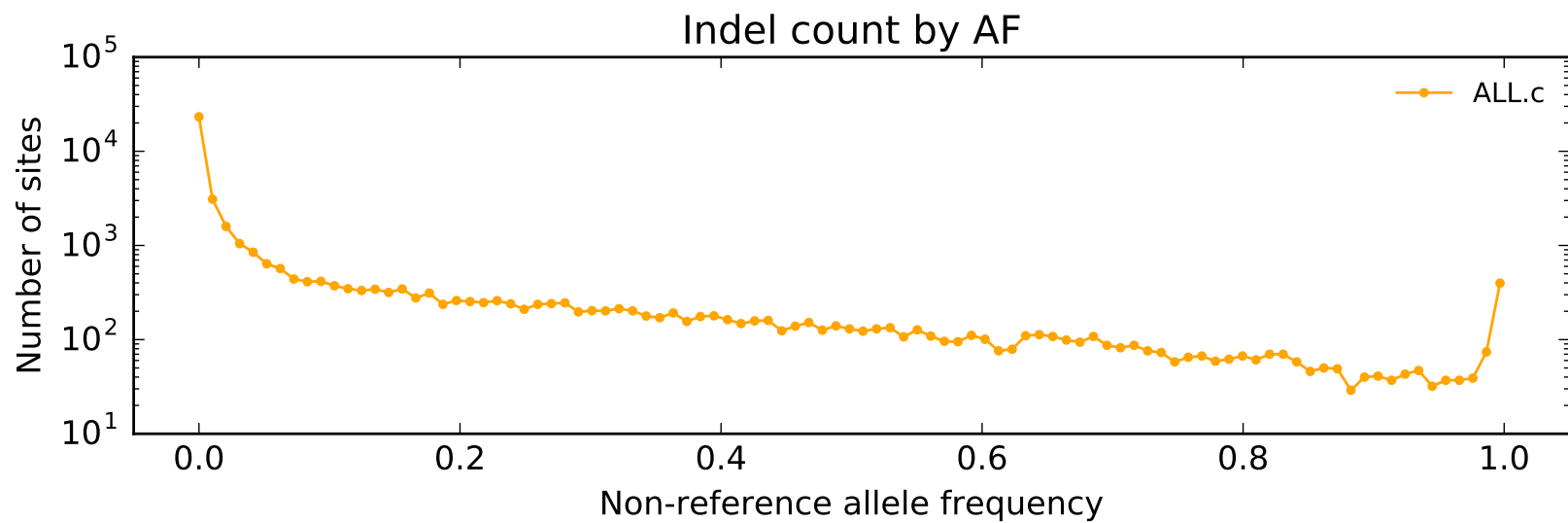
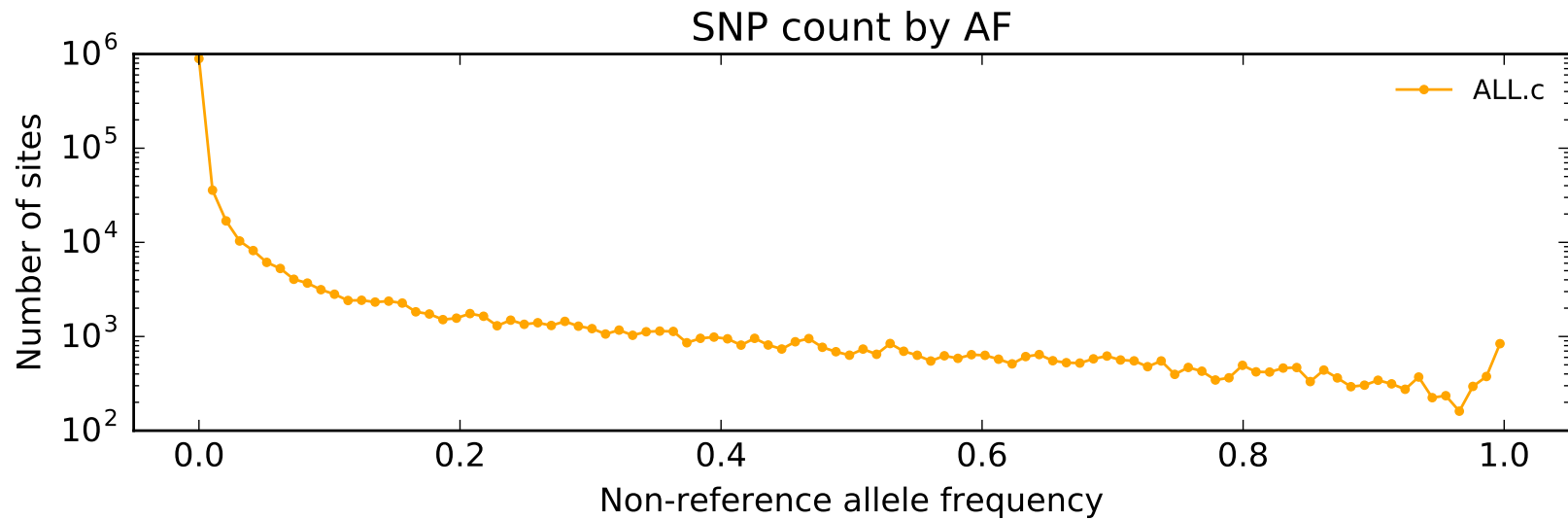


# Summary Numbers

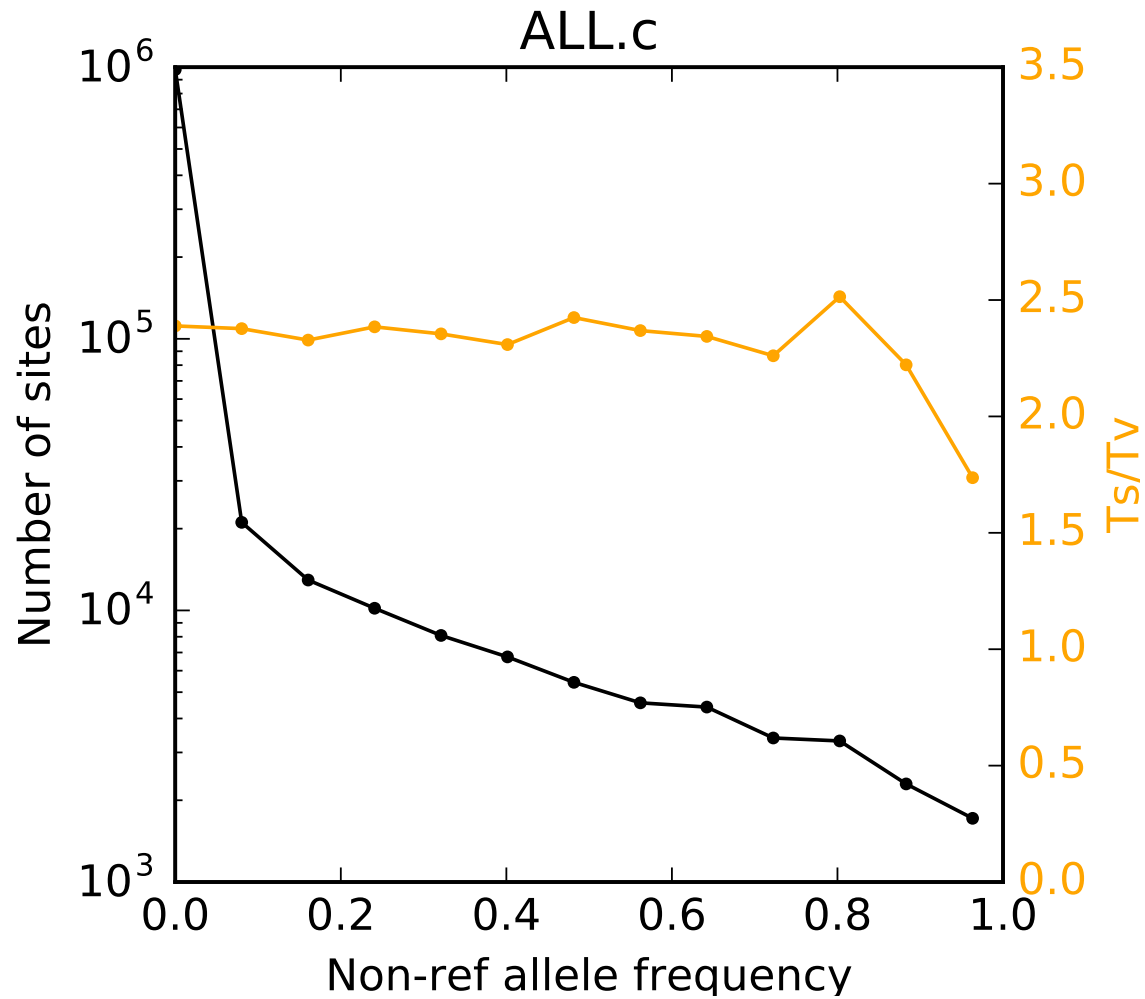
Callset	SNPs			indels			MNPs	others
	n	ts/tv	(1st ALT)	n	frm*	rc**		
ALL.c	1,060,388	2.39	2.40	43,230	–	–	0	801
* frameshift ratio: out/(out+in); ** repeat-consistency								

Callset	singletons (AC=1)				multiallelic	
	SNPs	ts/tv	indels	rc	sites	SNPs
ALL.c	42.6%	2.28	0.2%	0.000	6,348	4,063

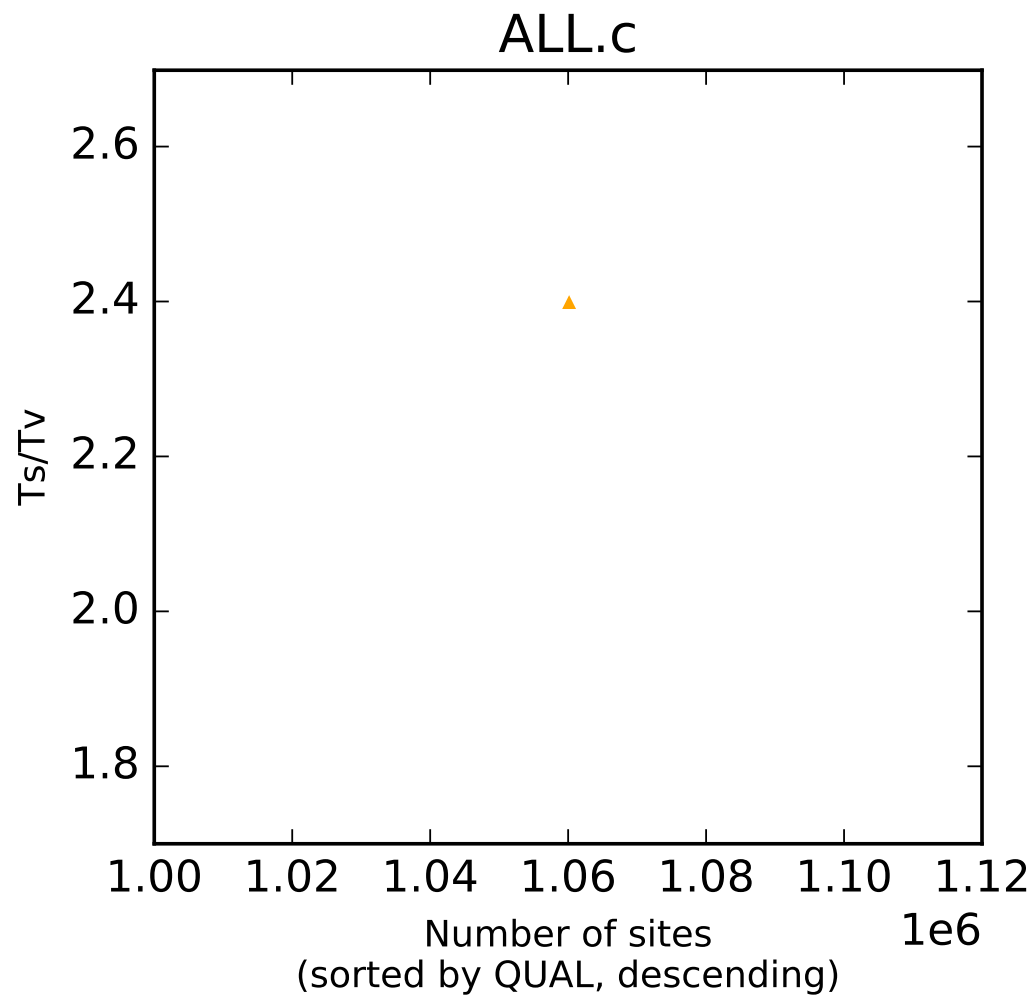
- ALL.c .. data/gz/ALL.chr22.phase3\_shapeit2\_mvncall\_integrated\_v5a.20130502.genotypes.vcf.gz



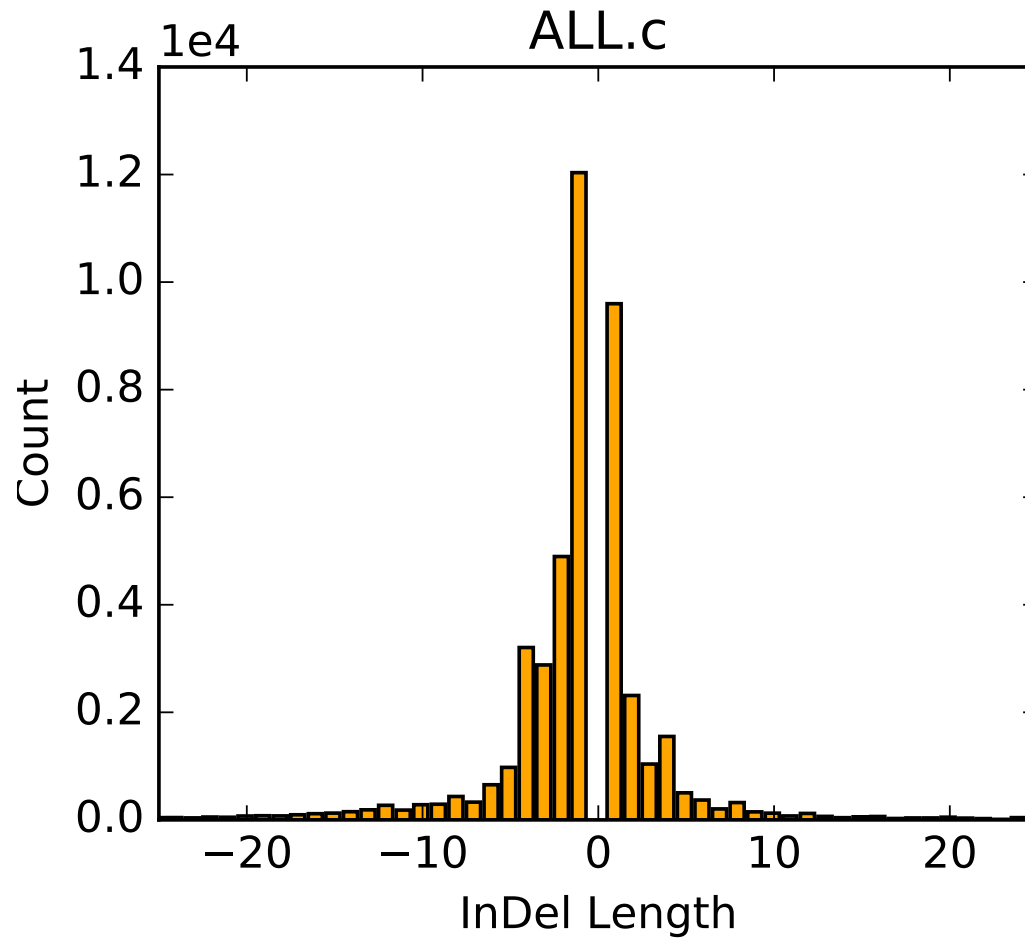
# Ts/Tv by AF



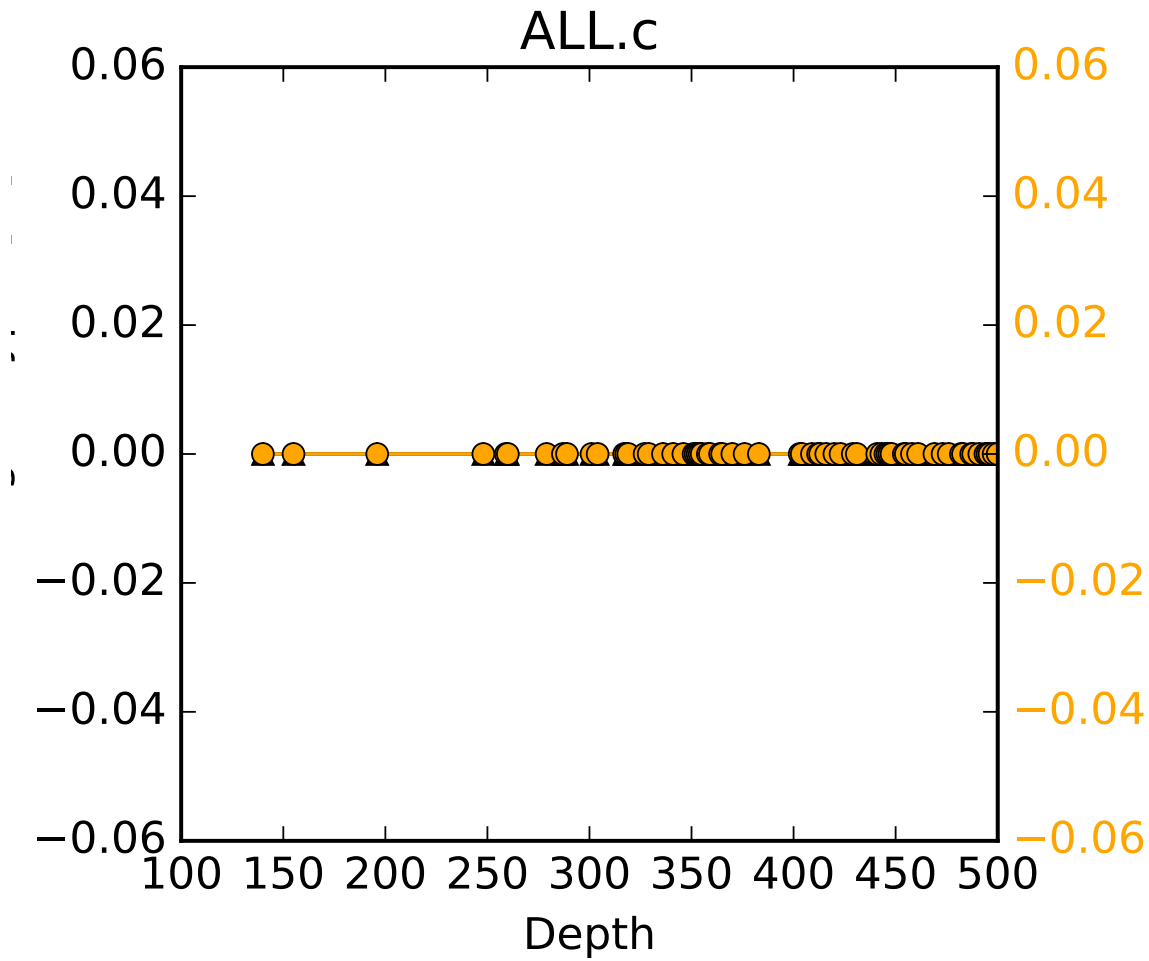
# Ts/Tv stratified by QUAL



# Indel distribution



# Depth distribution



# Substitution types

