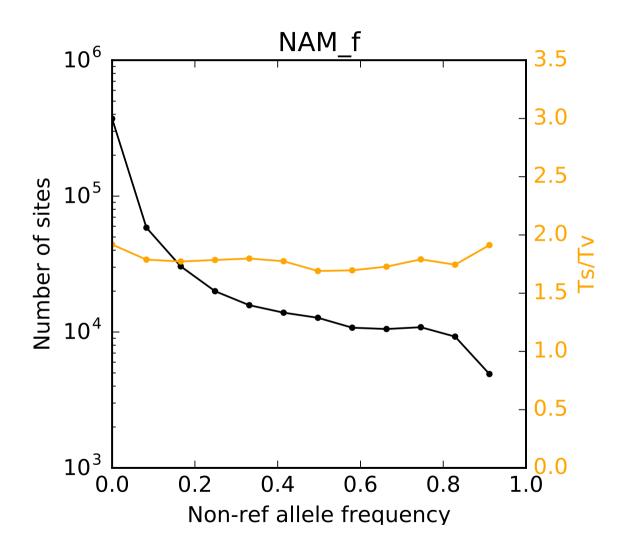
Summary Numbers

	SNPs			indels			MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*	rc**		
NAM_f	569,146	1.87	1.87	0	_	_	0	0
* frameshift ratio: out/(out+in); ** repeat-consistency								

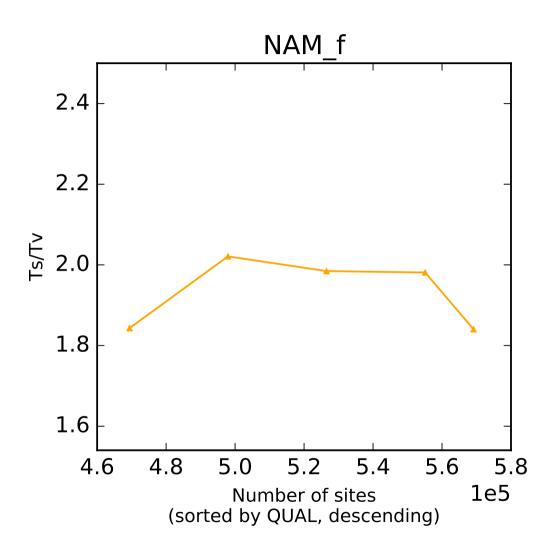
		singleto	multiallelic			
Callset	SNPs	ts/tv	indels	rc	sites	SNPs
NAM_f	1.3%	1.81	0.0%	0.000	0	0

⁻ NAM_f .. /panfs/roc/groups/9/morrellp/shared/Projects/Barley_NAM_Parents/sequence_handling/Variant_Filtering/NAM_final.vcf

Ts/Tv by AF



Ts/Tv stratified by QUAL



Substitution types

