

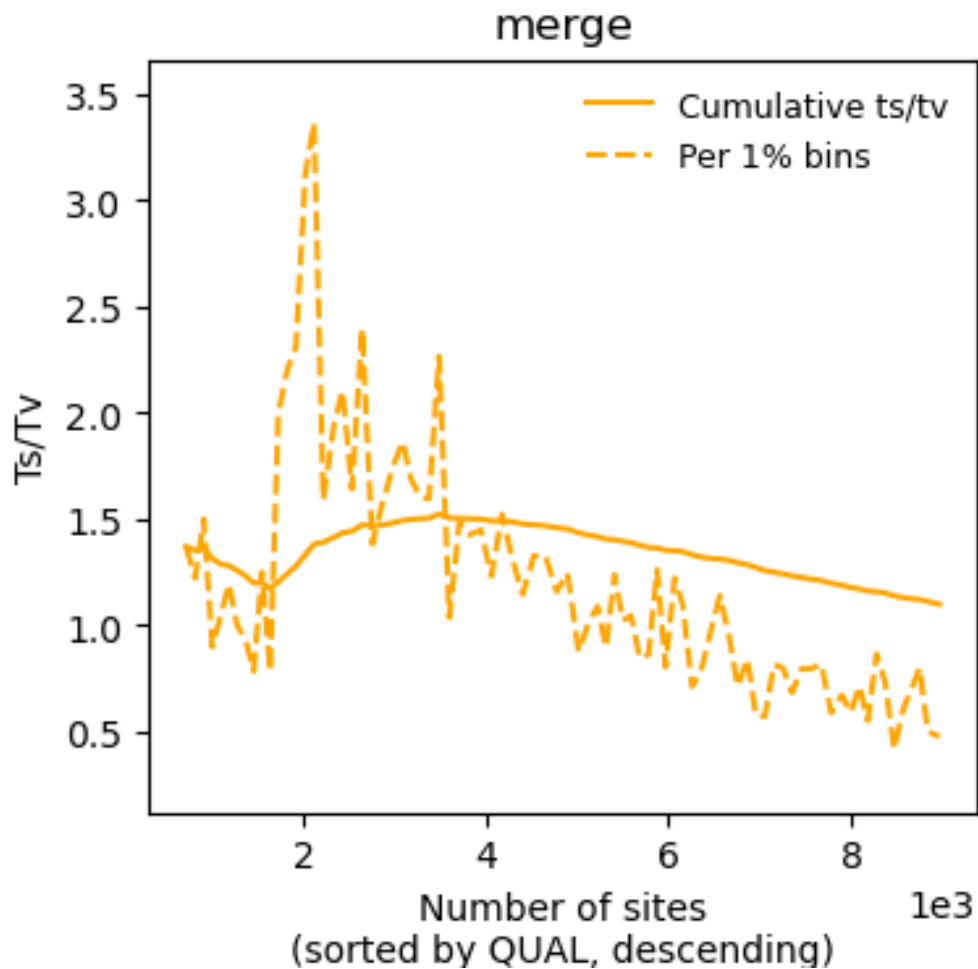
# Summary Numbers

Callset	SNPs			indels		MNPs	others
	n	ts/tv	(1st ALT)	n	frm*		
merge	8,991	1.10	1.10	1,114	–	0	0
* frameshift ratio: out/(out+in)							

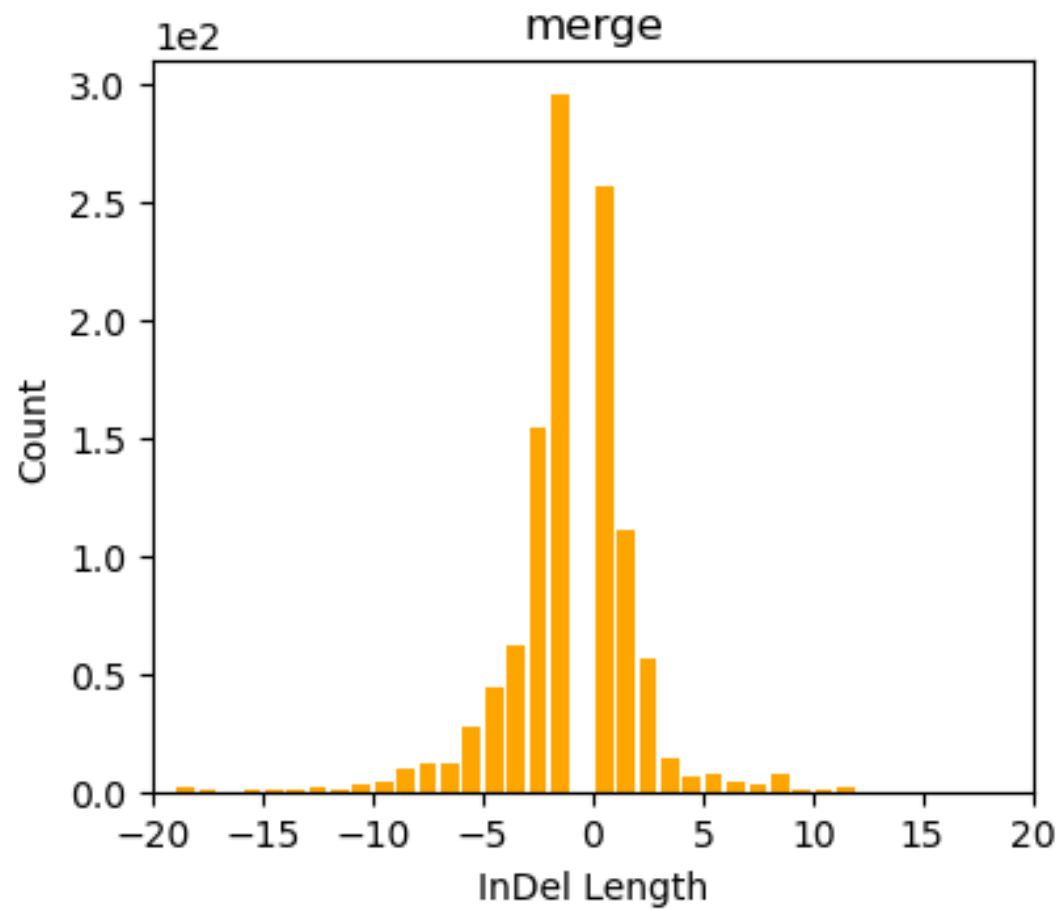
Callset	singletons (AC=1)			multiallelic	
	SNPs	ts/tv	indels	sites	SNPs
merge	100.0%	1.10	100.0%	44	14

- merge .. merge\_output.vcf

# Ts/Tv stratified by QUAL



# Indel distribution



# Substitution types

