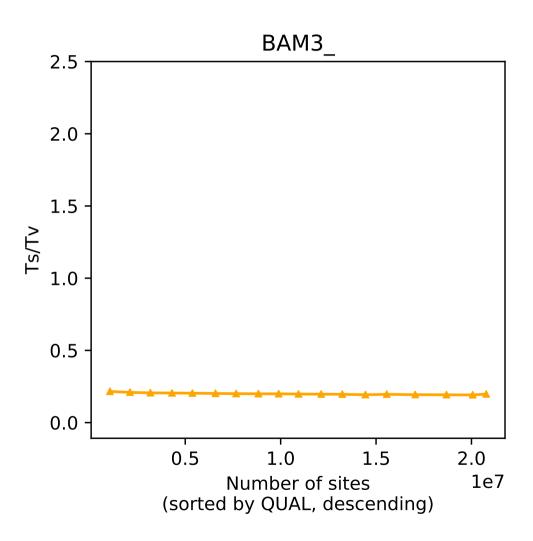
## **BAM3**<sub>5</sub>

	SNPs			indels		MNPs	others		
Callset	n	ts/tv	(1st ALT)	n	frm*				
BAM3_	20,865,599	0.20	0.20	1	-	0	2,164,326		
* frameshift ratio: out/(out+in)									

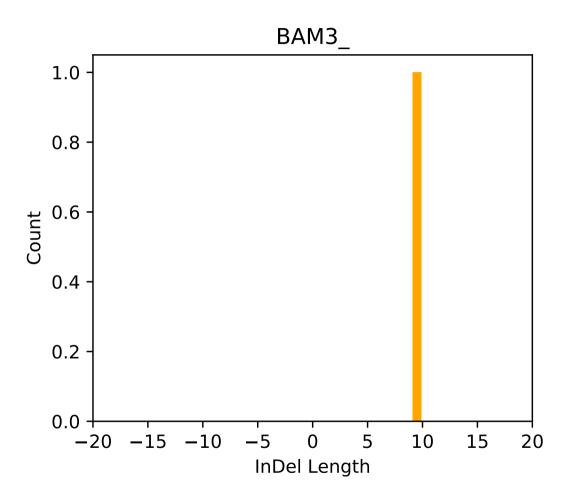
	singletons (AC=1)			multiallelic		
Callset	SNPs	ts/tv	indels	sites	SNPs	
BAM3_	100.0%	0.20	100.0%	515,777	214,203	

<sup>-</sup>  $BAM3_{-}...$  BAM3\_5.fixed.vcf.gz

## Ts/Tv stratified by QUAL



## Indel distribution



## Substitution types

