Skeletonema

	SNPs			indels		MNPs	others
Callset	n	ts/tv	(1st ALT)	n	frm*		
P8352_102	5,235	1.78	1.78	140	-	0	0
P8352_110	36,640	1.74	1.74	242	_	0	0
P8352_102 + P8352_110	24,053	1.98	1.98	127	_	0	0

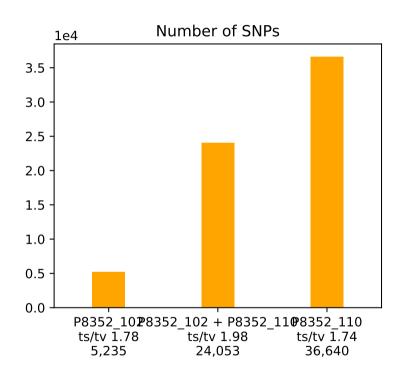
^{*} frameshift ratio: out/(out+in)

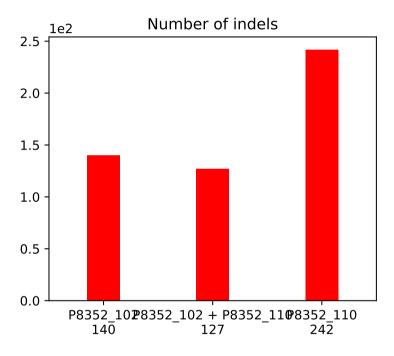
	singletons (AC=1)			multiallelic	
Callset	SNPs	ts/tv	indels	sites	SNPs
P8352_102	0.3%	1.14	4.9%	3	1
P8352_110	0.1%	1.92	1.2%	1	0
P8352_102 + P8352_110	0.2%	0.90	1.6%	2	1

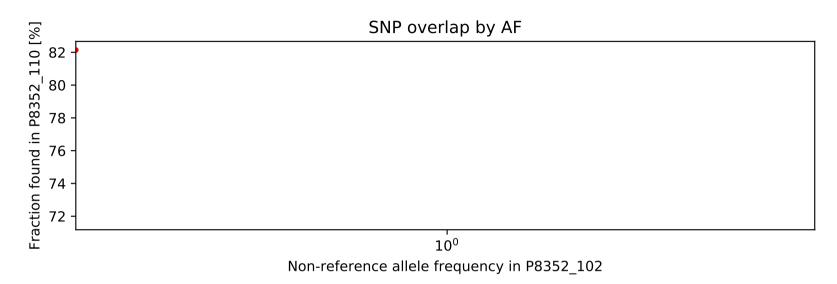
⁻ $P8352_102$.. $P8352_102$ /Vcalling/P8352_102.snpeff_annotated_new.vcf.gz

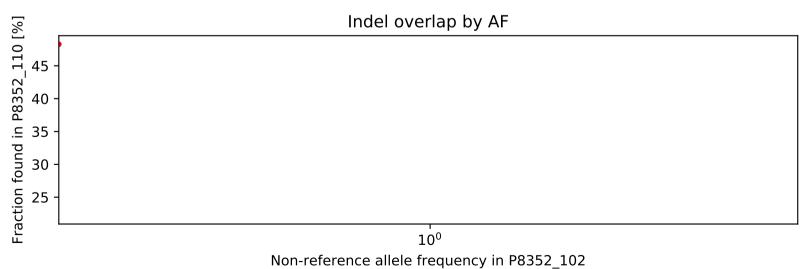
⁻ $P8352_110$.. $P8352_110/V$ calling/ $P8352_110.snpeff_annotated_new.vcf.gz$

Total counts

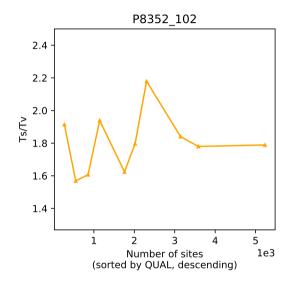


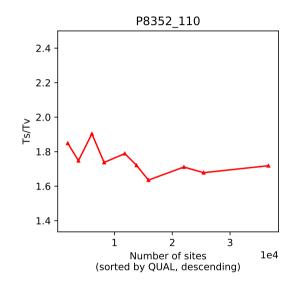


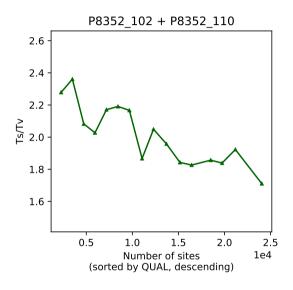




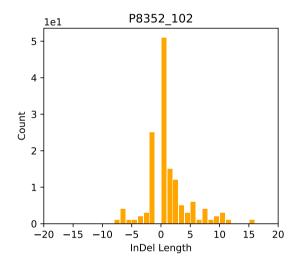
Ts/Tv stratified by QUAL

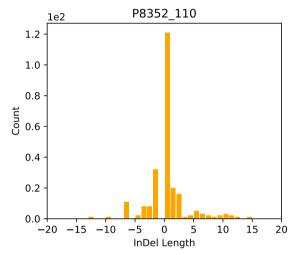


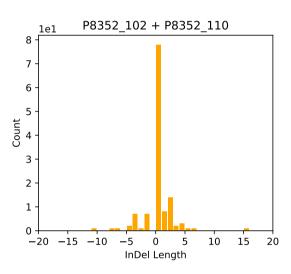




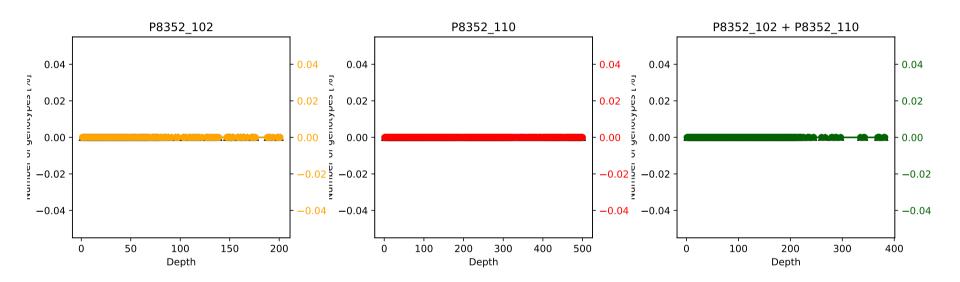
Indel distribution







Depth distribution



Substitution types

