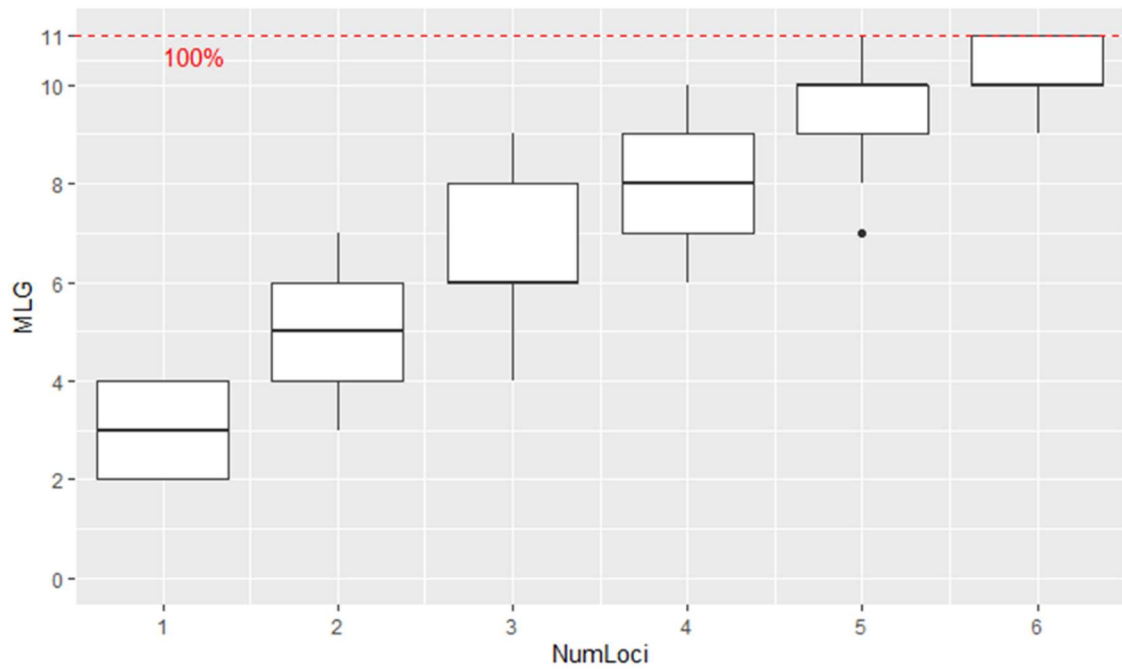


Supplementary Material

Supplementary Table 1 – Parameters used in phylogenetic analysis

Parameters		ITS data set	BT data set	Combined data set
No. of taxa		44	44	44
No. of characters		458	792	1250
ML	Substitution model	TIM3ef	TPM3uf+G	GTR+I+G
	Gamma shape	N/A	0.132 for BT1	0.132 for BT1
			0.308 for BT2	0.308 for BT2
BI	Substitution model	TIM3ef	TPM3uf+G	TIM3ef for ITS
				TPM3uf+G for BT1 and BT2

N/A: not applicable



Supplementary Figure 1 – Genotype accumulation curve with seven loci. Box plots were drawn using 1,000 times random sampling of isolates in each locus.