**Supplementary Informations**

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**A simple molecular identification method of the *Thrips tabaci* (Thysanoptera: Thripidae) cryptic species complex**

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**Supplementray Figure 1.** Phylogenetic tree of *T. tabaci* based on 328bp of the mtCOI gene based on the MP analysis using MEGA6 software.

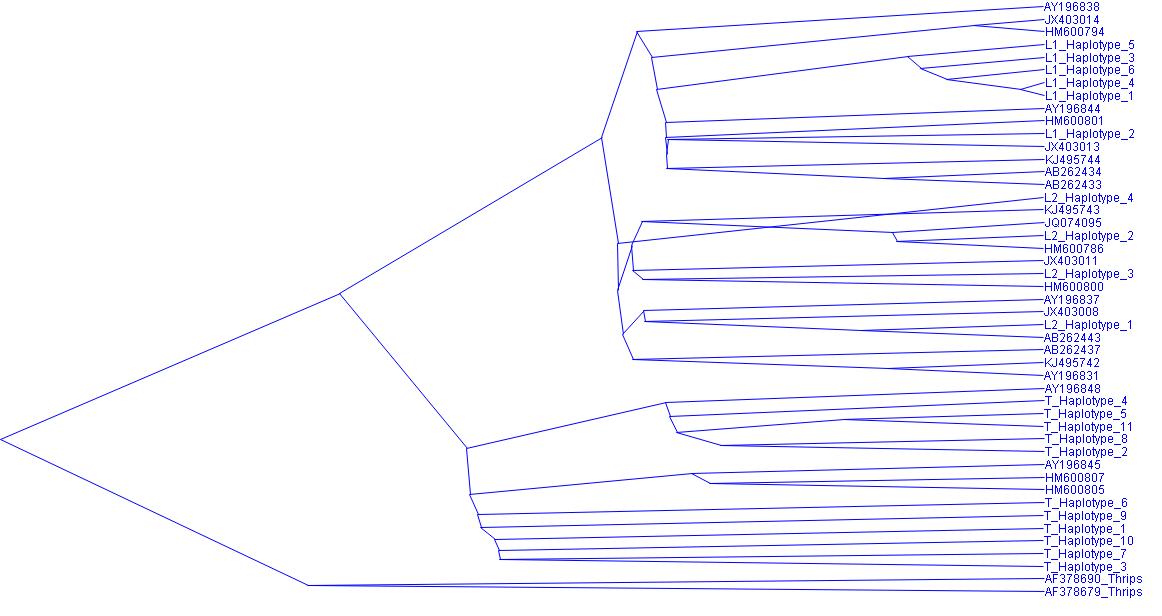
**Supplementray Figure 2.** Phylogenetic tree of *T. tabaci* based on 328bp of the mtCOI gene based on the HKY+G model using JModelTest 2.1.10. and BEAST 2.4.8. Software.

**Supplementray Figure 1.**

Phylogenetic tree of *T. tabaci* based on 328bp of the mtCOI gene*.* The tree is generated by the Maximum Parsimony (MP) method using MEGA6 (Tamura *et al.,* 2013)*.* Bootstrap values (percentage of 1000 replicates) are shown above the branches. Accession numbers shows the sequences data of *T. tabaci* obtained from DNA databases. *Thrips palmi* (AF378690) and *Thrips angusticeps* (AF378679) served as outgroups.

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**Supplementray Figure 2.** Phylogenetic tree of *T. tabaci* based on 328bp of the mtCOI gene*.* The HKY+G model (Hasegawa *et al*., 1985) was selected by JModelTest 2.1.10. (Darriba *et al*., 2012) and phylogenetic tree was generated using BEAST 2.4.8. software ([Bouckaert](http://www.ploscompbiol.org/article/info%3Adoi%2F10.1371%2Fjournal.pcbi.1003537) *et al.,* 2014). *Thrips palmi* (AF378690) and *Thrips angusticeps* (AF378679) served as outgroups.



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