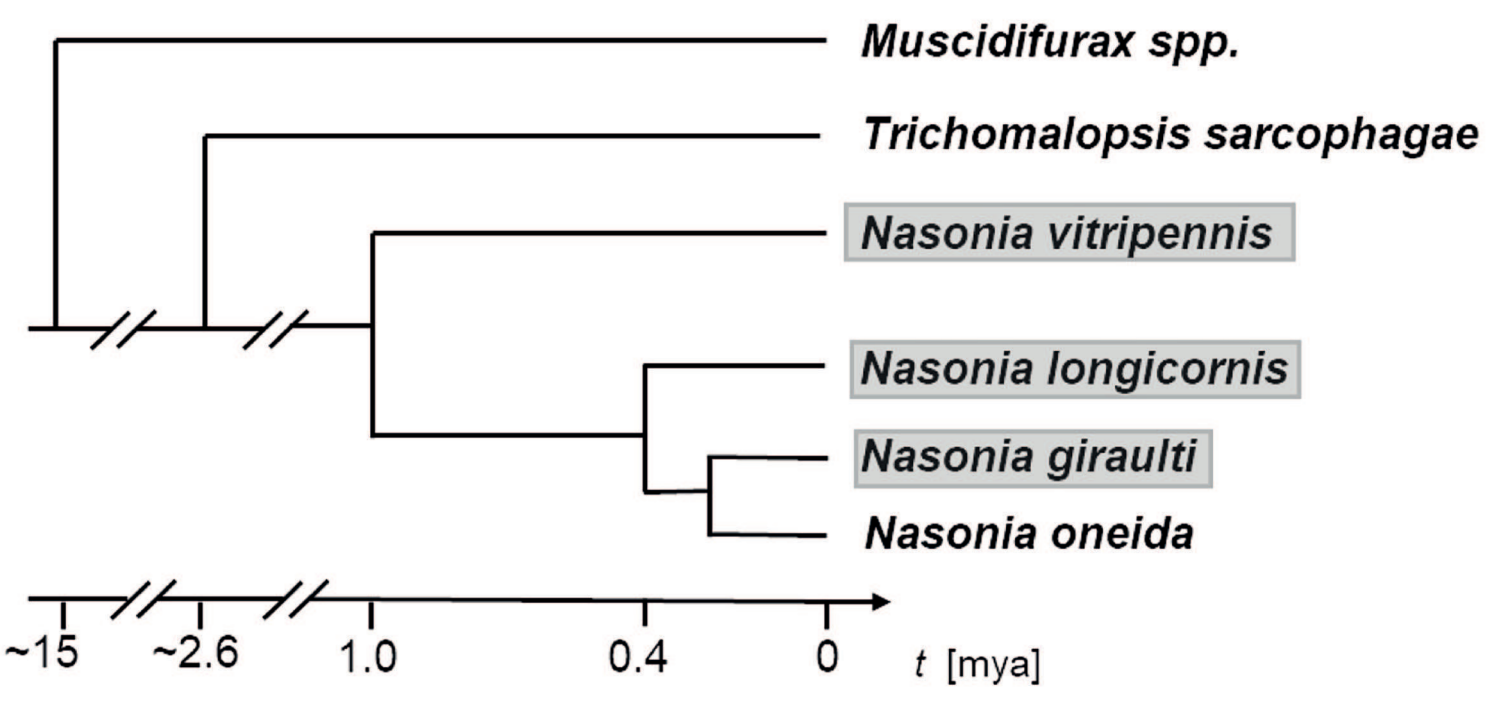

Figure S1: Phylogenetic relationship of the *Nasonia* species complex rooted by representatives of its two most closely related genera, *Trichomalopsis* and *Muscidifurax*, based on mitochondrial DNA sequences [20] and on the D2 expansion region of their 28S rDNA sequences [43]. The three *Nasonia* species used in the current study are highlighted in grey.