MOLECULAR EVIDENCE FOR THE PRESENCE OF *BEMISIA TABACI* BELONGING TO THE MIDDLE EAST-ASIA MINOR 1 AND NEW WORLD SPECIES IN BRAZIL

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Bemisia tabaci is one of the most important insect pests, being vector of the plant virus begomoviruses and causes serious problems in many countries, including Brazil. Based on the mitochondrial cytochrome oxidase I (mtCOI) sequence, the phylogenetic relationships from populations of *B. tabaci* collected from different hosts and locations in São Paulo and Mato Grosso State, Brazil were analyzed. According to the recent classification of *B. tabaci*, the most part of the specimens collected in Brazil belongs to the Middle East- Asia Minor 1 specie, which includes biotypes B and B2. But three specimens collected from Euphorbia heterophylla, Xanthium cavanillesii and Glycine max (soya) respectively, were classified in the New World group/specie and showed higher nucleotide identity with B. tabaci from Colombia (accession number AJ550167 and AJ550168, A biotype). The different species could be found colonizing the same soya plant in commercial area of Mato Grosso, indicating the coexistence of them in Brazil. By RFLP, these species could be easy differentiated using Thru I and Tag I enzymes. Mediterranean specimens could not be found in Brazil. As far as we know this is the first molecular evidence for the presence of the New World specie in Brazil.

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