

1. *Phylogenetic relationships*—Phylogenetic relationships among the 10 species of *Phrynosoma* were determined using the parsimony method of Farris (1993) with the computer program PAUP (Farris, 1993). The parsimony method was chosen because it is the most commonly used method for determining phylogenetic relationships (Farris, 1993). The parsimony method was used to determine the most parsimonious tree (MPT) for the 10 species of *Phrynosoma*. The MPT was determined by using the computer program PAUP (Farris, 1993). The MPT was determined by using the computer program PAUP (Farris, 1993).

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