

HYPOTHESES DOMINGUEZ

Phylogeny - Swallow

The following hypotheses were tested in the present study. The first hypothesis was that the phylogeny of the Swallow family is well supported by the morphological data. The second hypothesis was that the phylogeny of the Swallow family is well supported by the molecular data. The third hypothesis was that the phylogeny of the Swallow family is well supported by the combined morphological and molecular data.

The phylogenetic relationships among the members of the Swallow family were investigated using morphological and molecular data. The morphological data were analyzed using parsimony analysis, and the molecular data were analyzed using maximum likelihood analysis. The results of the analyses are presented in the following sections.

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The combined morphological and molecular data were analyzed using parsimony analysis. The results of the analysis are presented in the following sections. The phylogenetic relationships among the members of the Swallow family are well supported by the combined morphological and molecular data.

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