



Fig. 6-3. Strict consensus tree of the analyzed species of Heliotropiaceae, computed from the 2,604 most parsimonious trees of the combined *trnL* and ITS1 datasets (1=950 steps, CI=0.622, RI=0.832, HI=0.378). Important clades are indicated. The numbers above the branches are bootstrap percentages. Percentages <50% are not shown.