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Figure 1 is a schematic representation of the experimental design. It shows a flowchart with two main paths. The top path is labeled 'Pretest' and leads to a box labeled 'Main Experiment'. The bottom path is labeled 'Main Experiment' and leads to a box labeled 'Posttest'. The 'Main Experiment' box is further divided into two sub-sections: 'Pretest' and 'Main Experiment'. The 'Pretest' sub-section shows a flow from 'Pretest' to 'Main Experiment'. The 'Main Experiment' sub-section shows a flow from 'Main Experiment' to 'Posttest'.

Age	1998	98	98
1998	1998	98	98
1998	1998	98	98
1998	1998	98	98

Figure 1. *Phylogenetic relationships of the 12 species of the genus *Phyllanthus* based on the chloroplast *trnK* gene. The tree was constructed using the maximum parsimony method. Bootstrap values are shown at the nodes. The scale bar represents 0.1 substitutions per site. The species names are listed at the tips of the branches. The species names are listed at the tips of the branches. The species names are listed at the tips of the branches.*

