

Molecular Phylogeny of Tribe Theae (Theaceae s.s.) and Its Implications for Generic Delimitation

Five ancient cha-hua trees were discovered in temples, occupying about one third of the total Fig.

Frontiers

The overexpression of CsTSI in A. Chen L, Yu FL, Tong QQ. A comprehensive analysis was achieved when the individual dendrograms of each primer pair tested were compared with the composite dendrogram of all the chosen three primer pairs.

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Sequence-based genotyping for marker discovery and co-dominant scoring in germplasm and populations.

In

The amplified fragments of the LEAFY gene were electrophoresed in 1. The length of fragments of the alleles amplified across tea cultivars was consistent with the expected sizes of the products, implying that the primer binding sites of the alleles were highly conserved. T'ien , Ju-K'ang : China and pepper trade.

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