Figure S1. Expression profiles of LcWUS, LcWOX1, LcWOX2A, LcWOX3, LcWOX4, LcWOX5, LcWOX9, and LcWOX14 in different tissues of L. chinense determined by qRT-PCR. The PCR signals were normalized to ubiquitin and 18S transcript levels. Data were calculated using three replicates. Roots, stems, leaves, and leaf buds were harvested from 2-month-old regenerated seedlings. Gynoecium, stamen, petal, and sepal samples were collected from a mature tree (Lushan 1`) growing in a breeding orchard in Xiaoshu.