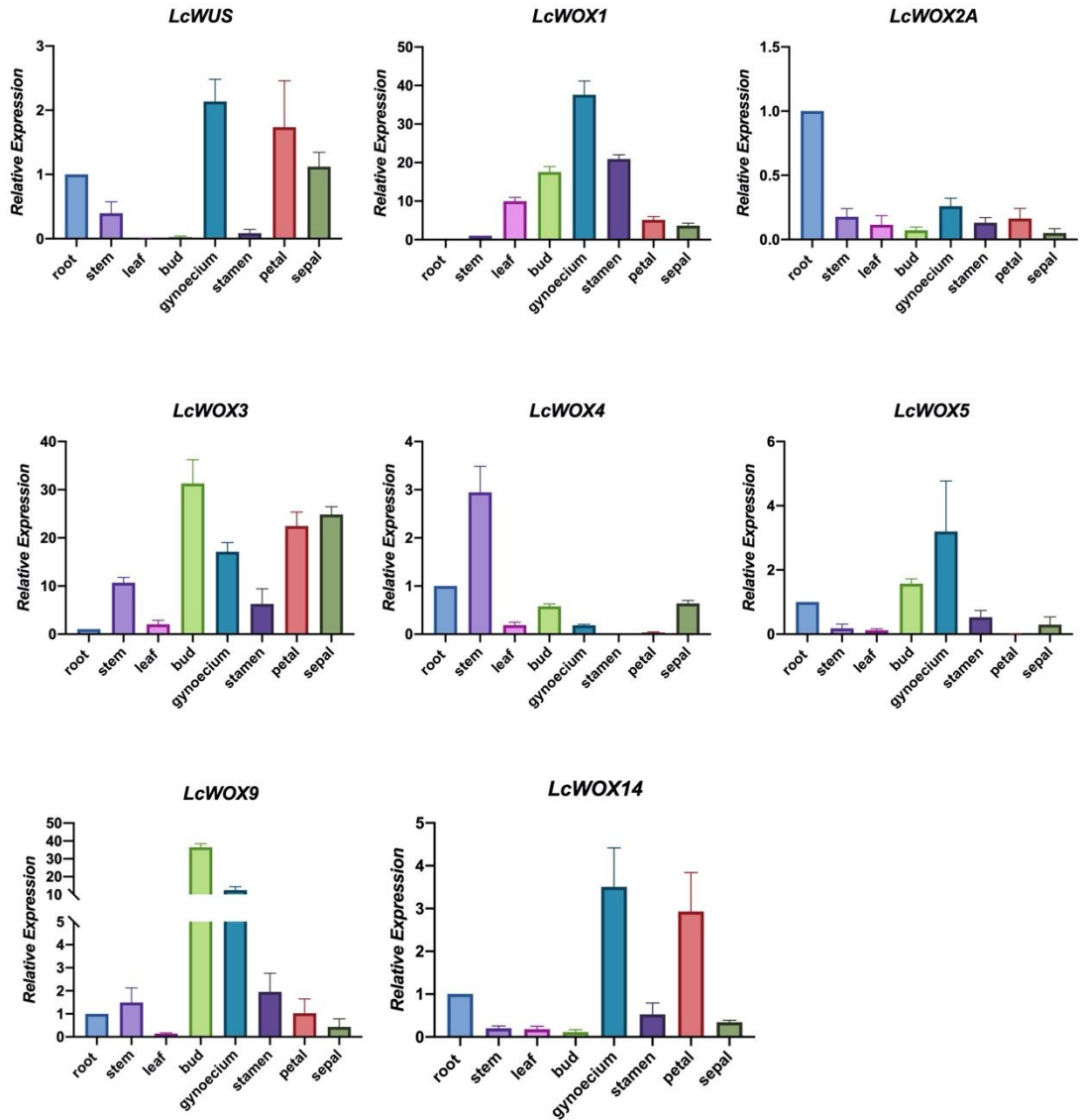


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636 Figure S1. Expression profiles of *LcWUS*, *LcWOX1*, *LcWOX2A*, *LcWOX3*, *LcWOX4*, *LcWOX5*, *LcWOX9*,

637

and *LcWOX14* in different tissues of *L. chinense* determined by qRT-PCR.

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The PCR signals were normalized to ubiquitin and 18S transcript levels. Data were calculated using three

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replicates. Roots, stems, leaves, and leaf buds were harvested from 2-month-old regenerated seedlings.

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Gynoecium, stamen, petal, and sepal samples were collected from a mature tree (Lushan 1[#]) growing in a

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breeding orchard in Xiashu.

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