



Figure S2. Comparative genomics reveals evolutionary adaptations in *Lonicera oblata*. (a) Analysis of LTR insertion times in the *L. oblata* genome. The abscissa represents the timing of LTR insertions. (b) Whole-genome duplication (WGD) events. The solid line shows the K_s distribution within the species, while the dashed line shows the K_s distribution among different species. γ -WGD events shared by all core eudicots are highlighted. (c) Cross-species orthogroup clustering across 14 angiosperm species. (d) Patterns of core gene conservation among limestone-endemic species. Red indicates the number of orthogroups that are commonly conserved among the four limestone-endemic plant species.