



Figure S7. Transcriptome analysis of *Lonicera oblata* under calcium stress treatment. (a) PCA of transcriptome samples across six time points under calcium stress. Each time point is represented by a distinct color. (b) Numbers of upregulated and downregulated genes in five pairwise comparison groups. The total number of differentially expressed genes (DEGs) is shown. (c) Venn diagram showing the overlap between DEGs, expanded genes in *L. oblata*, and genes shared among four lithophytic species.