

Top 20 of KEGG Enrichment

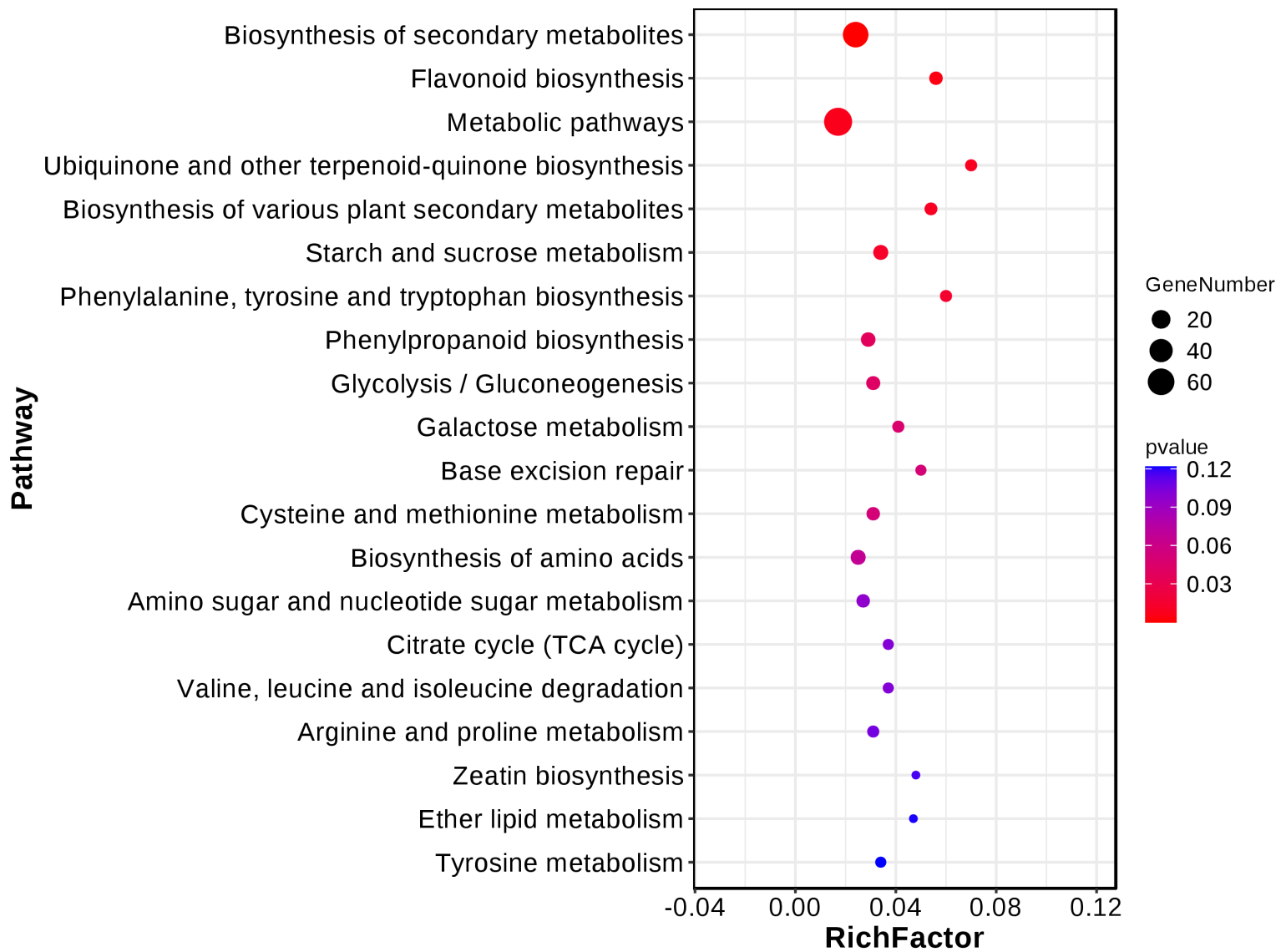


Fig.S8 The top 20 pathways of KEGG enrichment were identified for the 428 differentially expressed genes in stipule spine (AGT-Ss, LC110-Ss) compared to leaf (BL01-Le). Ss: stipule spine, Le: leaf.