

Figure S1. KEGG enrichment analysis. (A) DS vs control, (B) DS+SL vs control, and (C) DS+SL vs DS.

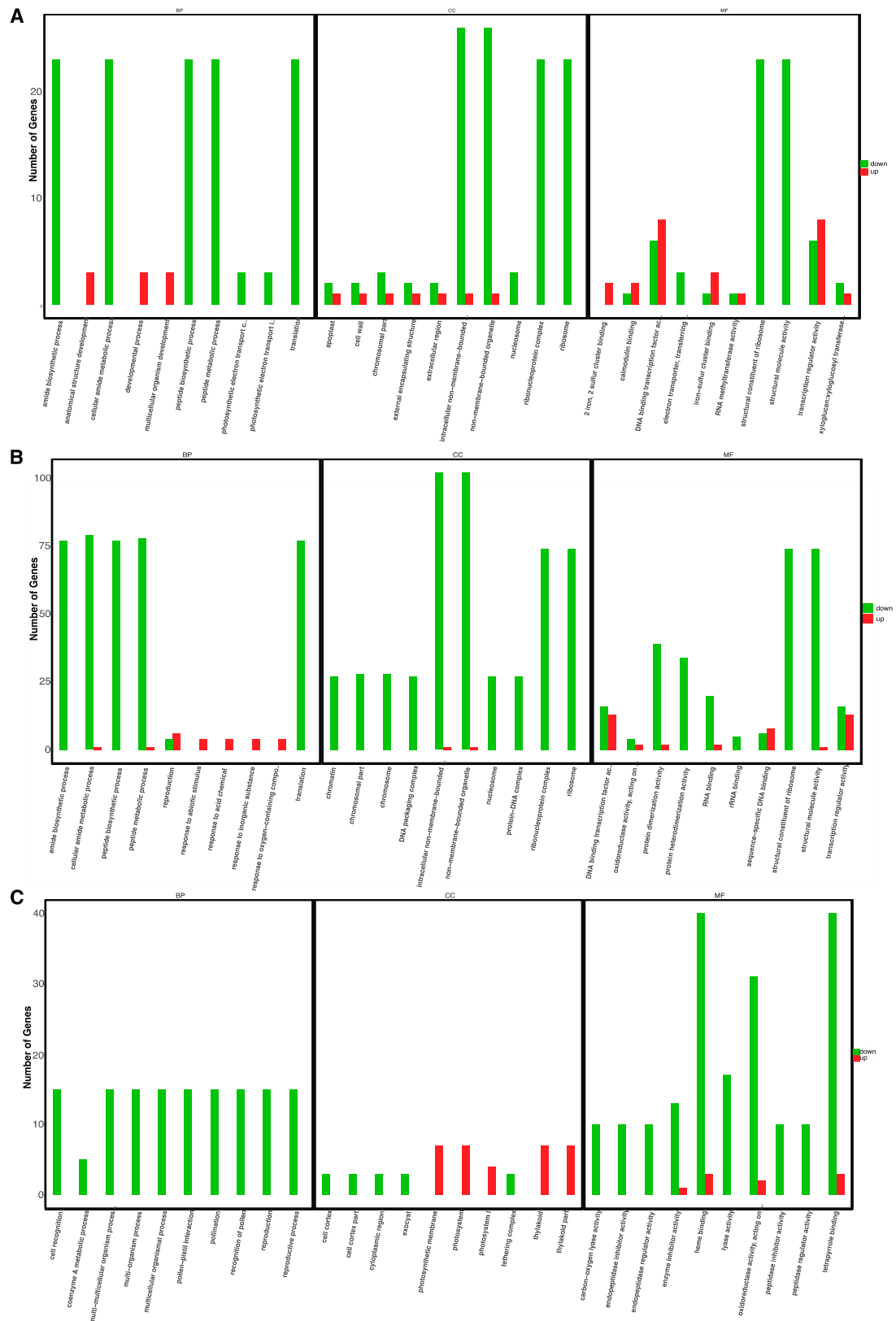


Figure S2. GO enrichment analysis. (A) DS vs control, (B) DS+SL vs control, and (C) DS+SL vs DS.

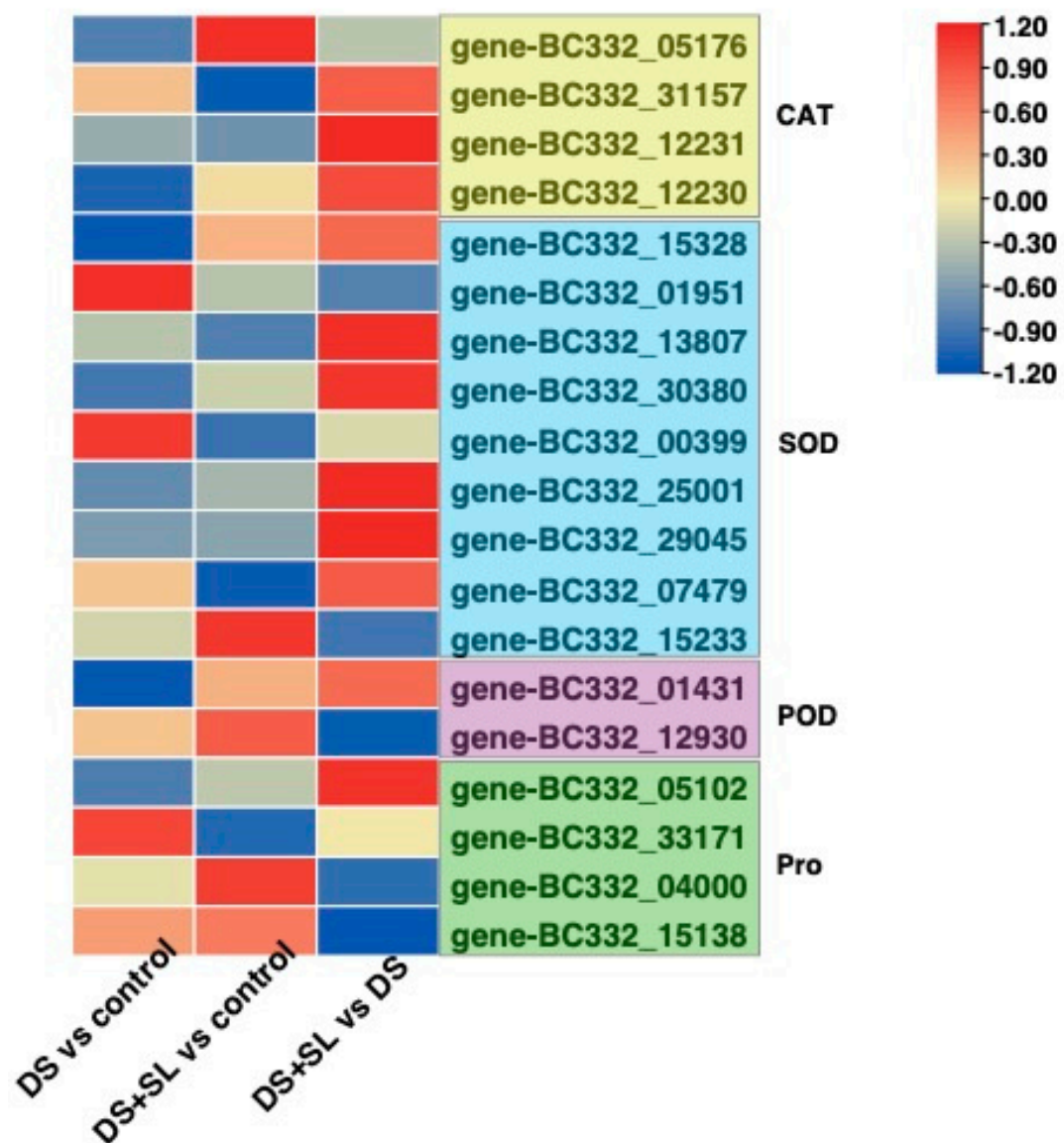


Figure S3. Heat-map representing the transcription level of differentially expressed proline and antioxidant enzymes related genes in pepper seedlings under drought stress. Heatmap colors show the log2FoldChange values.