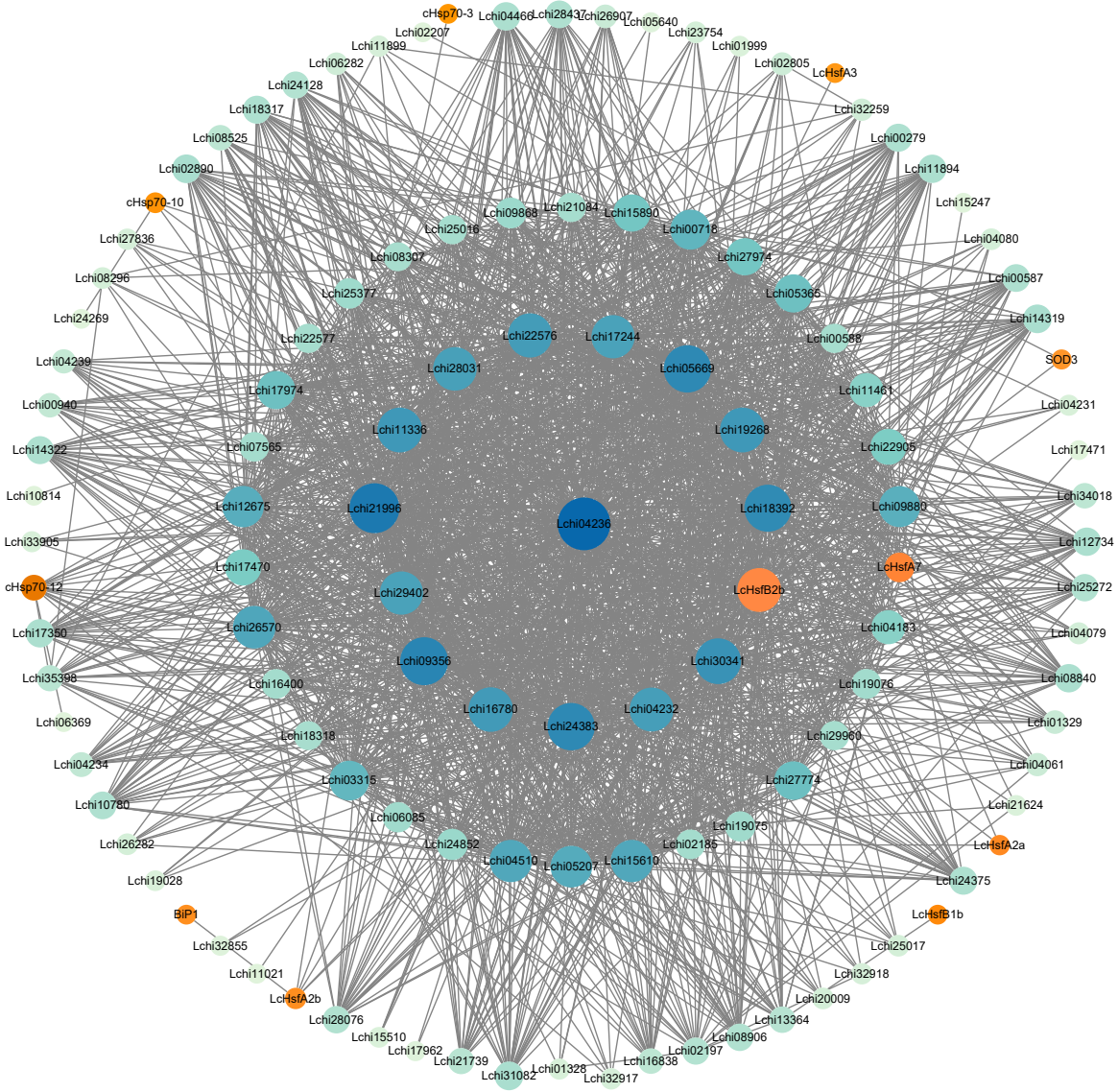
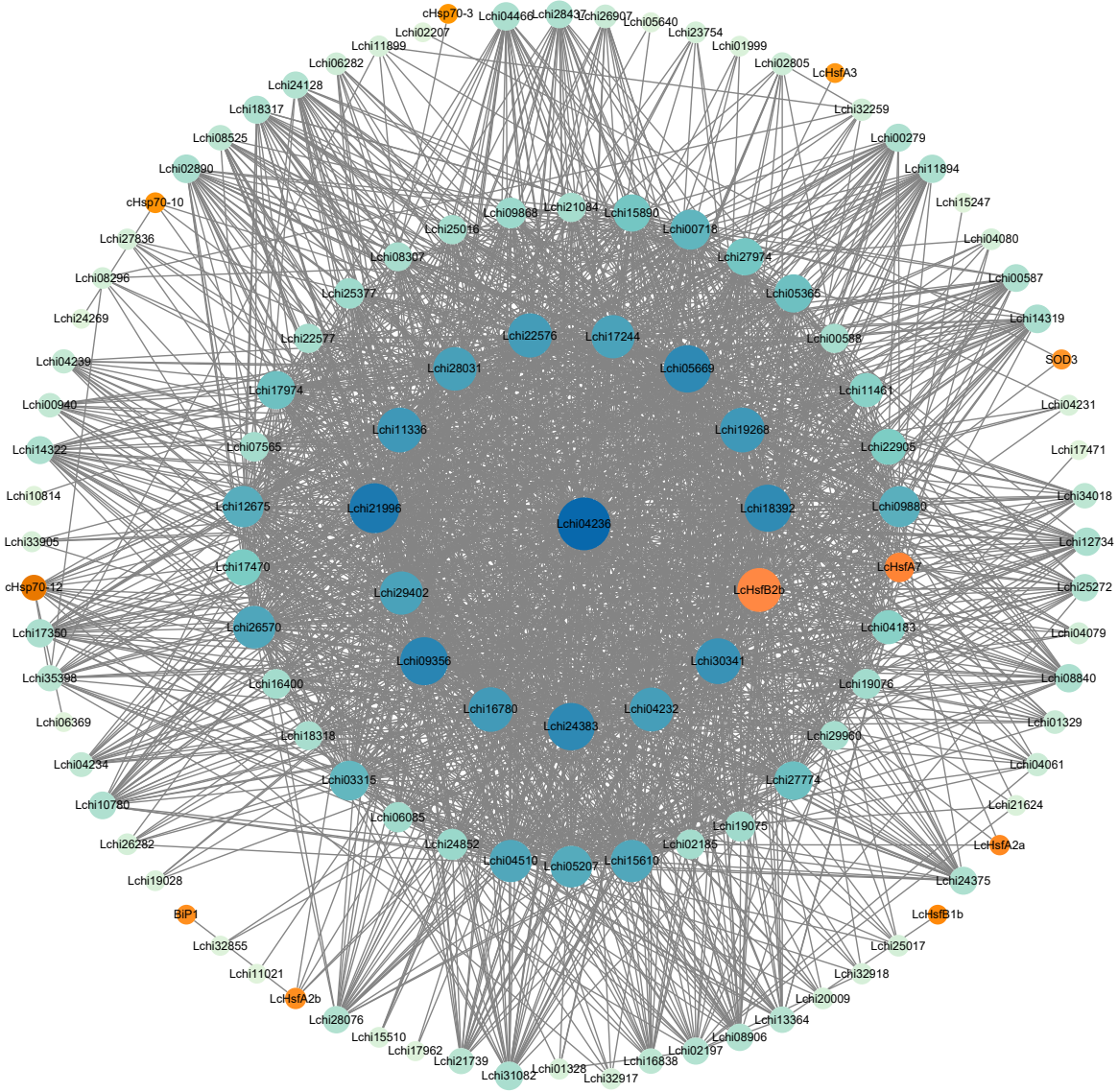


Figure S3 The correlation network of blue4 module.

Note: The network was visualized in a concentric circle diagram using Cytoscape software (v3.8.2) and was distributed using the degree algorithm. Genes with the edge weight higher than 0.60 are visualized. The LcHsf, LcHsp70 and redox genes in this figure are marked with yellow circles.



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