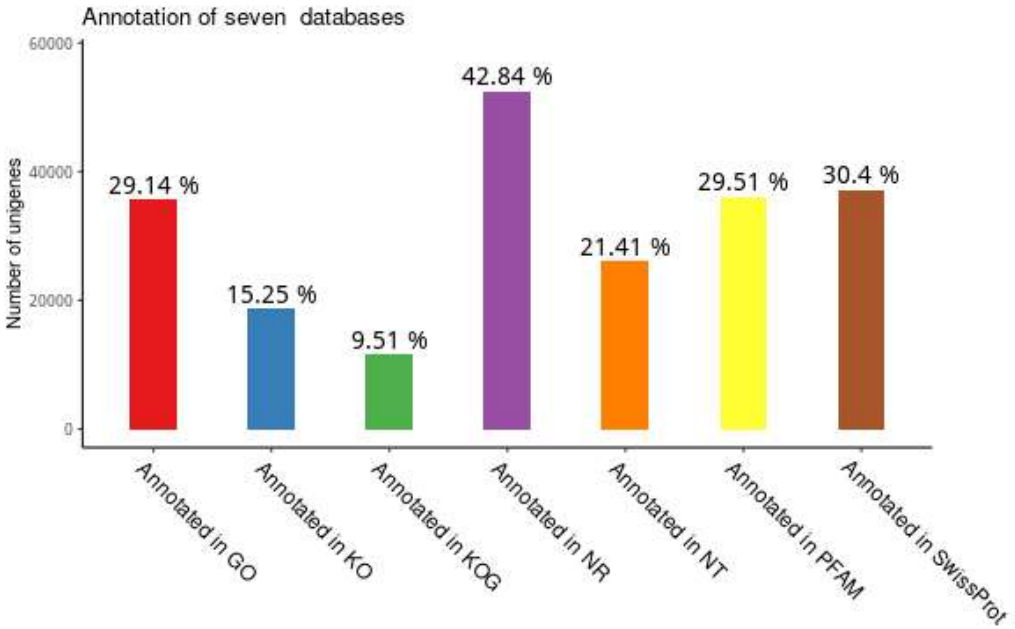


Figure S1. Annotation of the lily unigenes in databases.

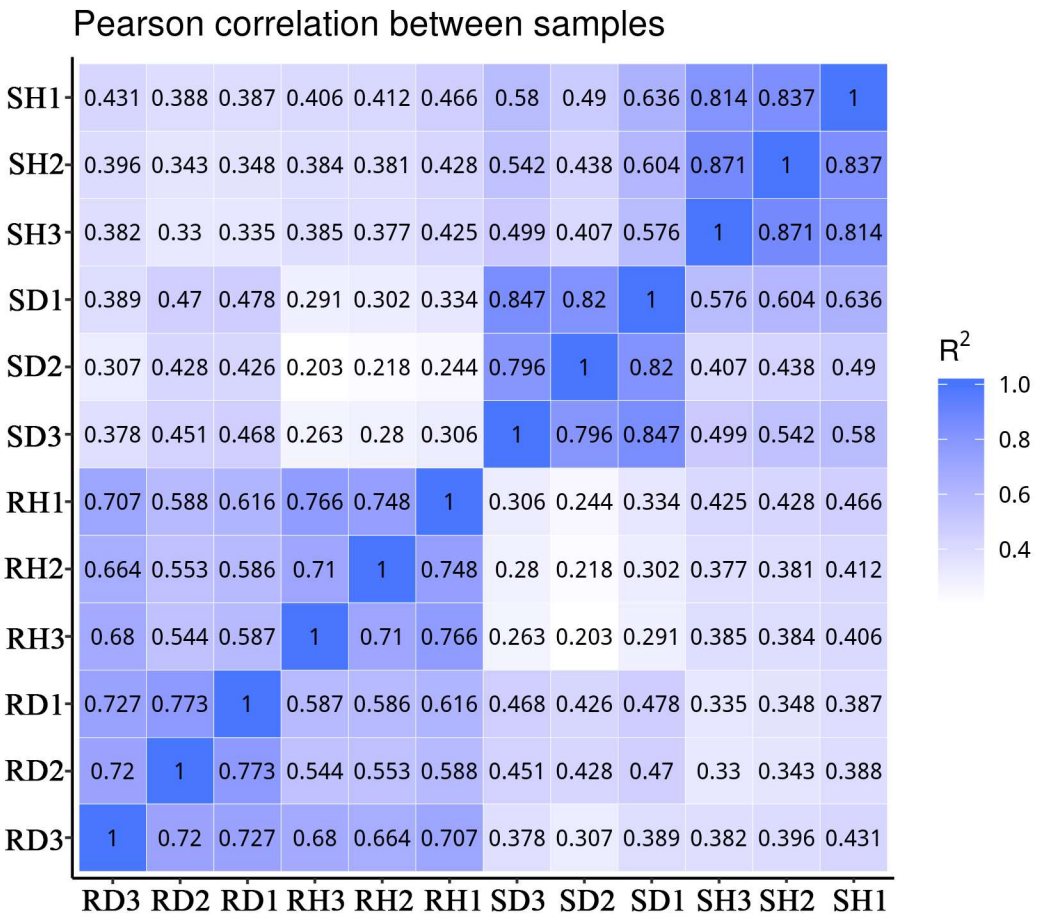


Figure S2. Repeatability evaluation of transcriptome data from lily after using Pearson's correlation analysis.

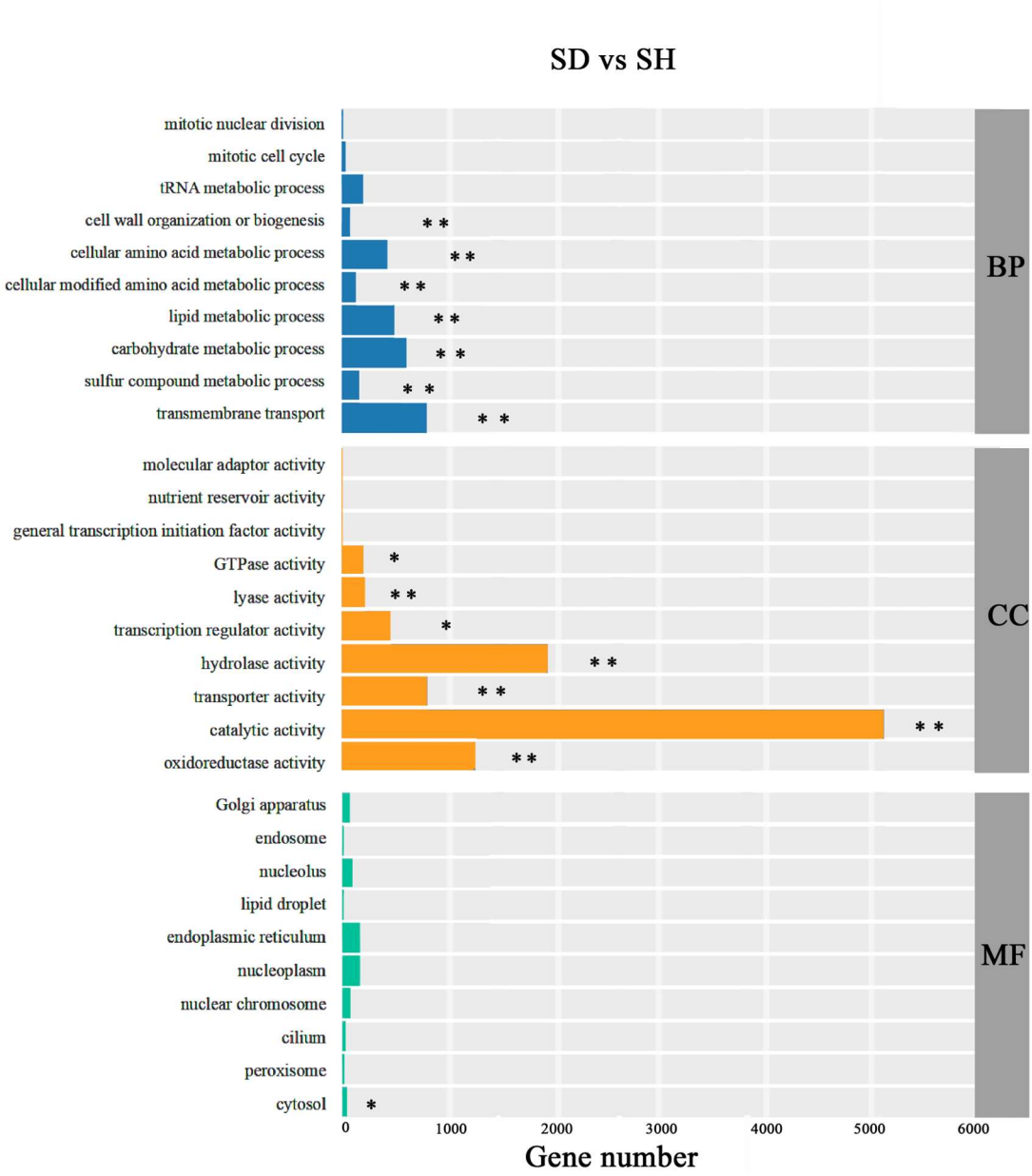


Figure S3. GO enrichment analysis of the DEGs detected in *B. elliptica*-infected 'Tresor' compared with its control. * $p < 0.05$, ** $p < 0.01$.

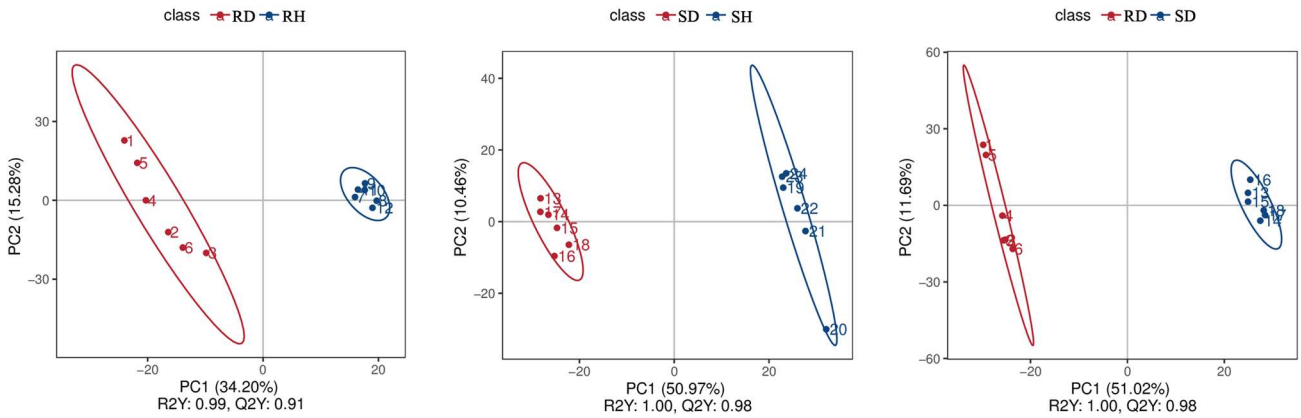


Figure S4. Partial least squares discriminant analysis (PLS-DA) for the samples based on the identified metabolites.

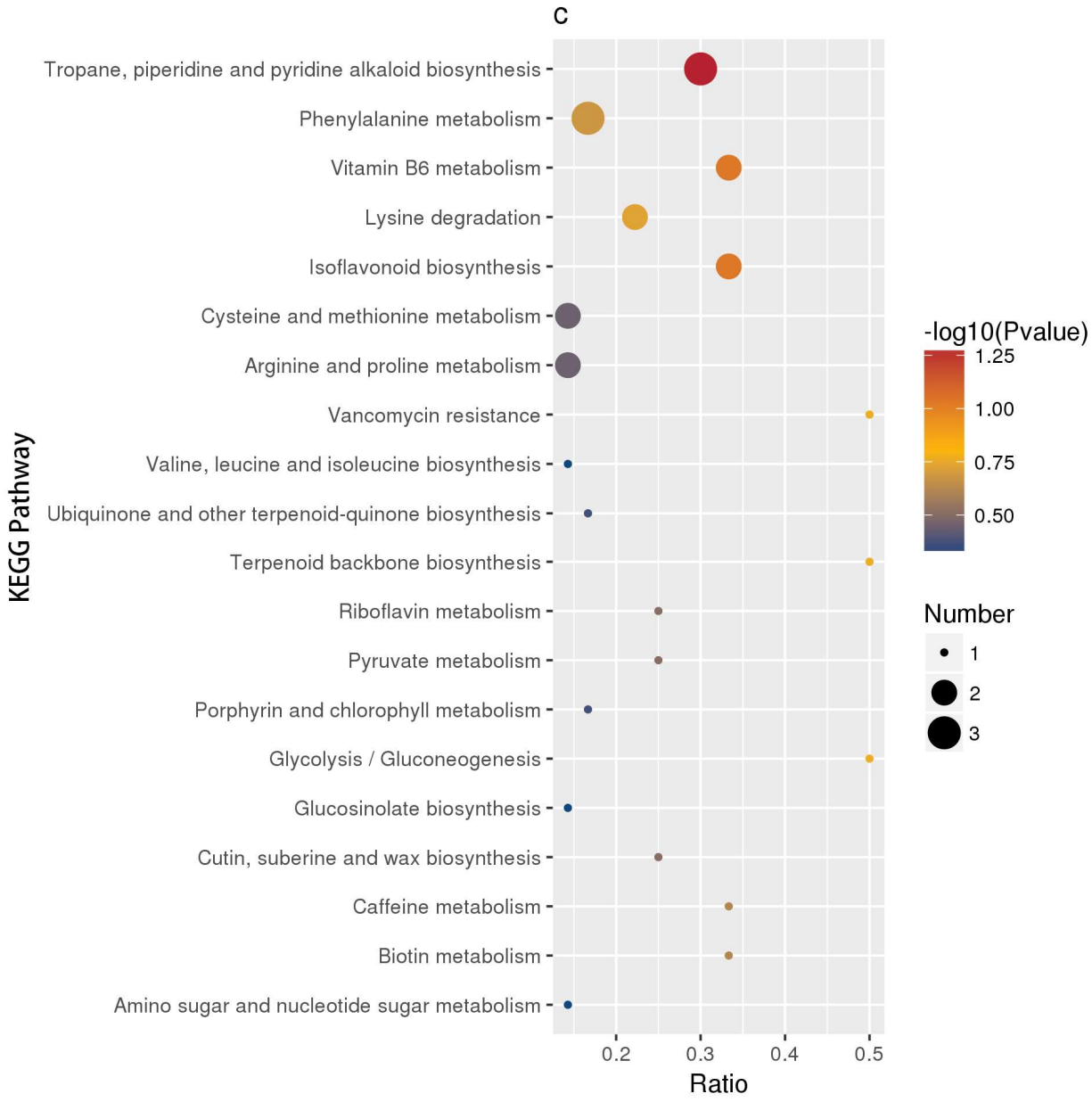


Figure S5. KEGG enrichment analysis for the 179 common DAMs.

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