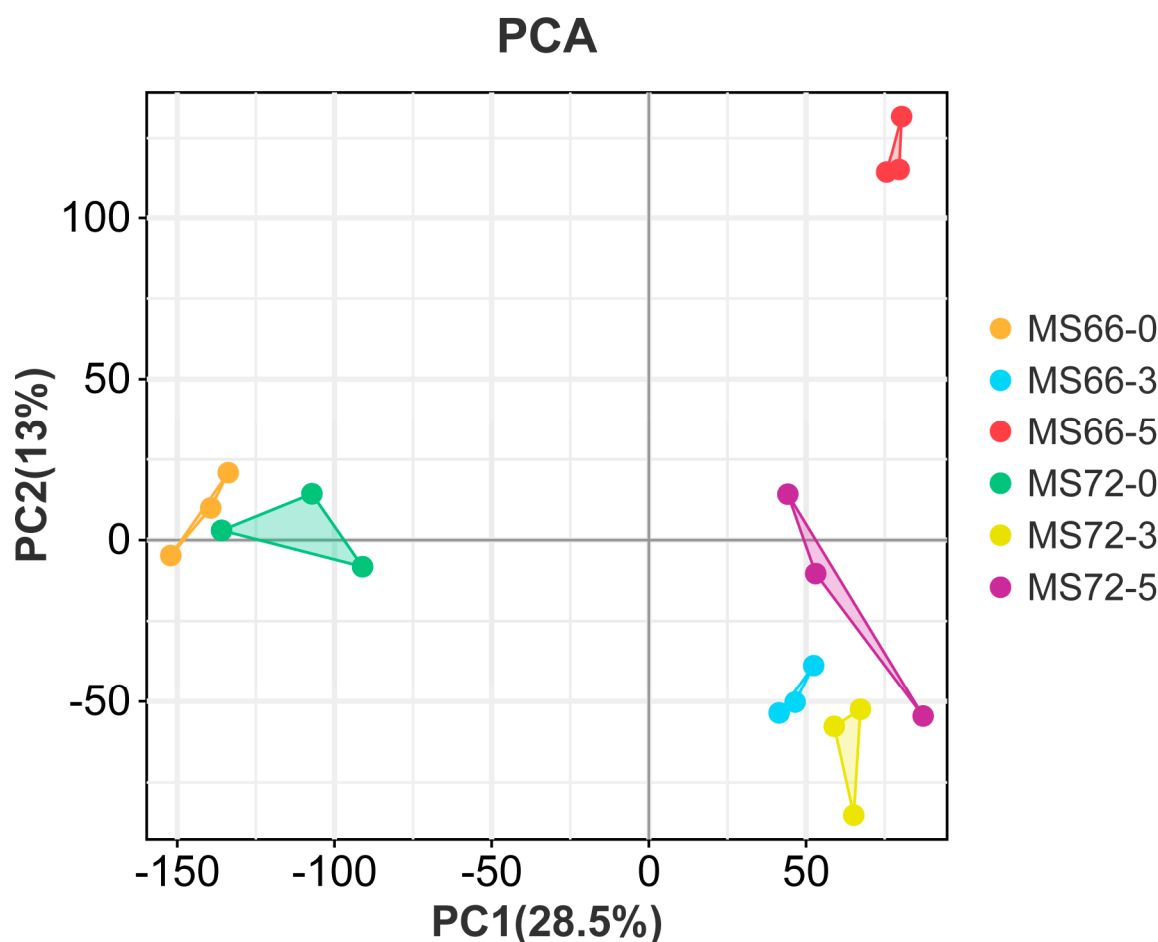




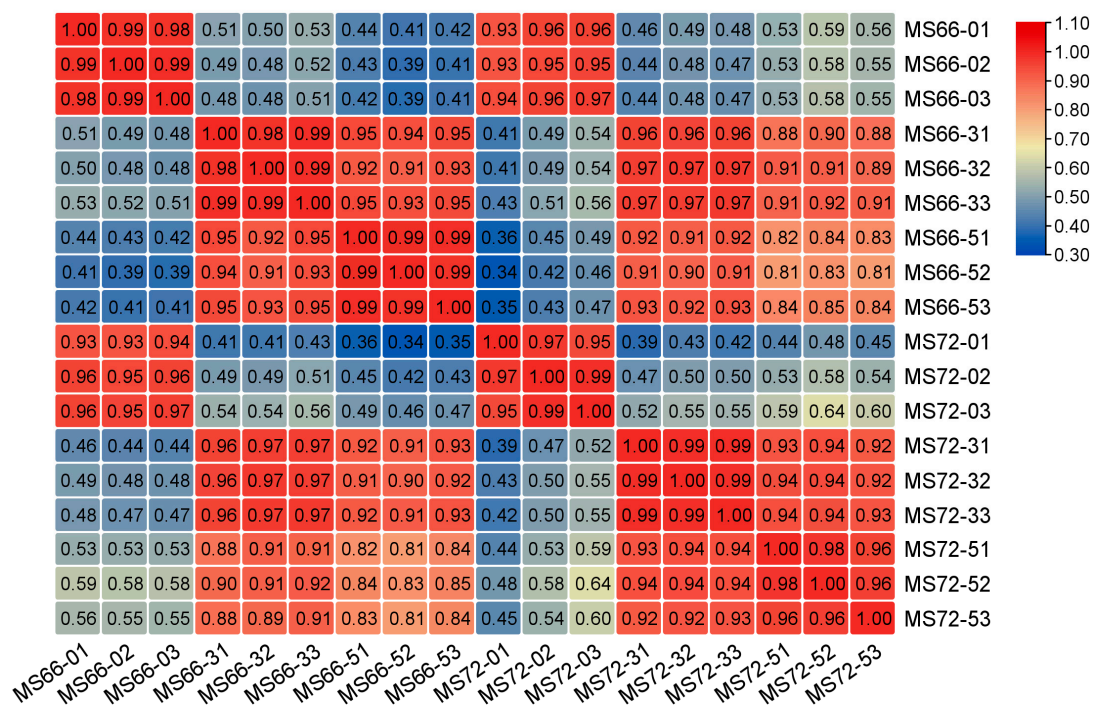
Transcriptome and Physiological Analyses of Resistant and Susceptible Pepper (*Capsicum annuum*) to *Verticillium dahliae* Inoculum

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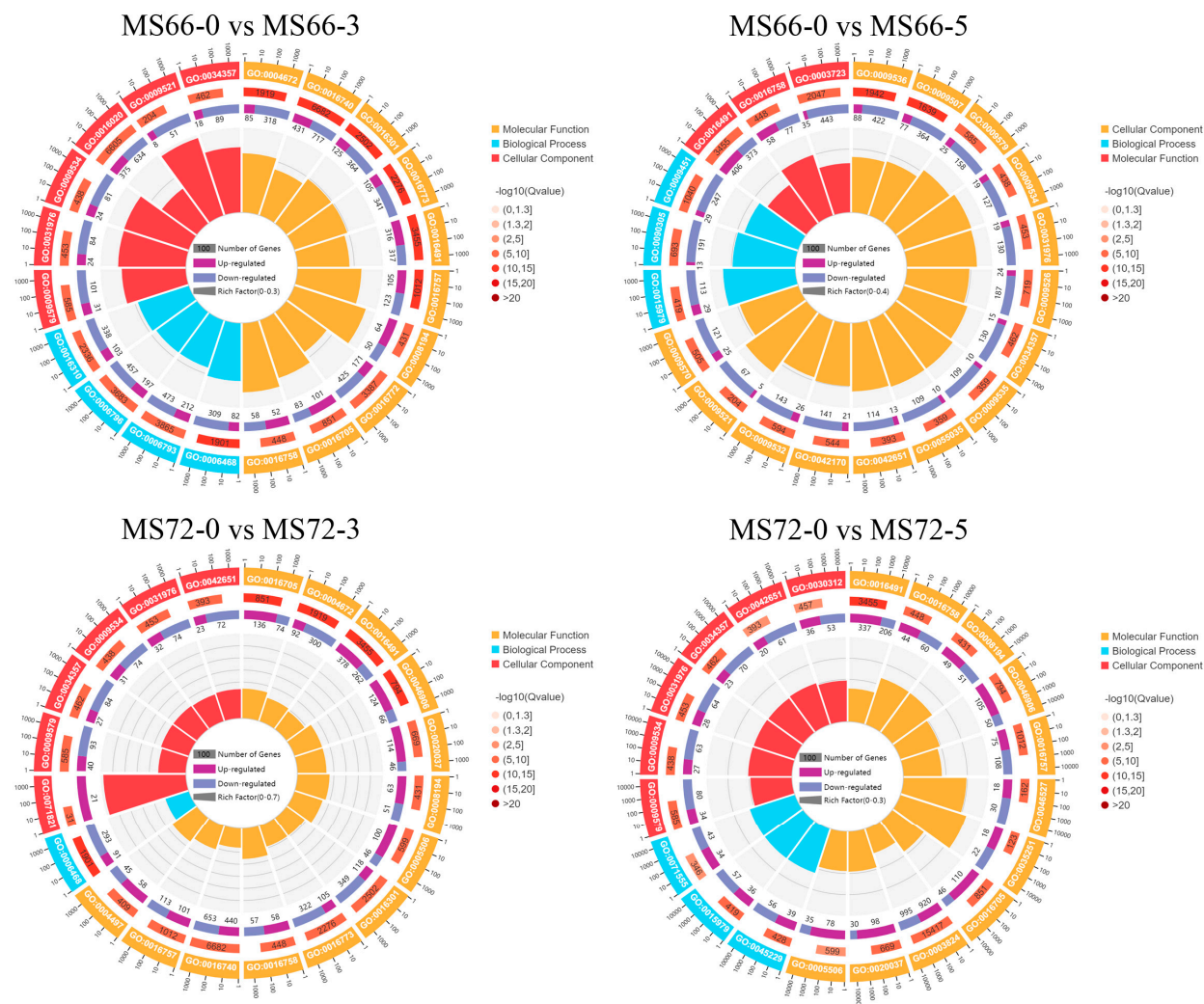
Supplementary Figure



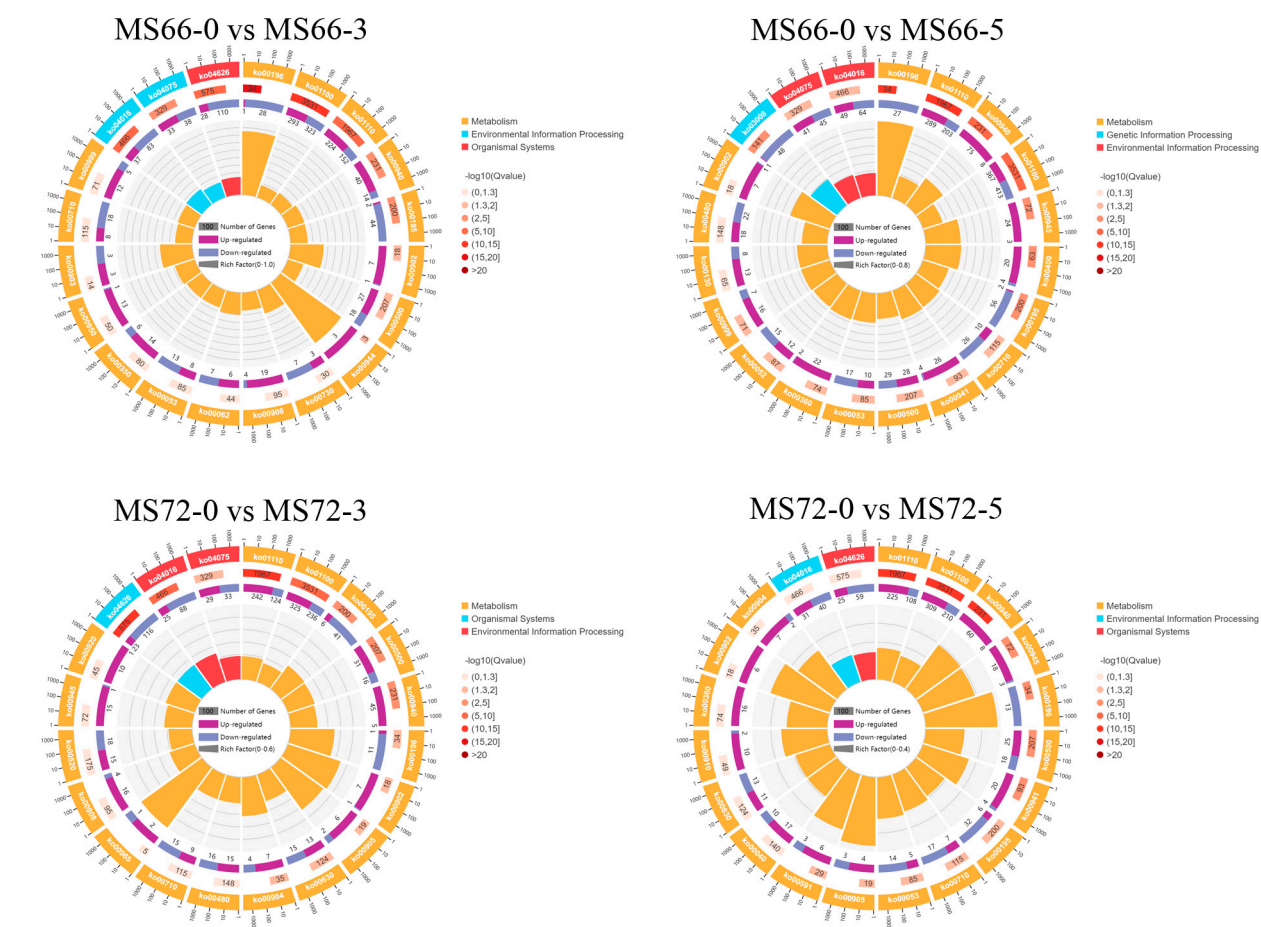
Supplementary Figure S1. Principal component analysis (PCA) of eighteen samples.



Supplementary Figure S2. Pearson correlation coefficient heatmap between eighteen samples. The X and Y axis represent each sample. The color represents the correlation coefficient (The redder the color, the higher the correlation), and the number on the cell represents the corresponding Pearson correlation coefficient.



Supplementary Figure S3. Top twenty significantly enriched GO terms of DEGs in MS66-0 vs. MS66-3, MS66-0 vs. MS66-5, MS72-0 vs. MS72-3 and MS72-0 vs. MS72-5 groups.



Supplementary Figure S4. DEGs enriched in KEGG pathways in MS66-0 vs. MS66-3, MS66-0 vs. MS66-5, MS72-0 vs. MS72-3 and MS72-0 vs. MS72-5 groups.