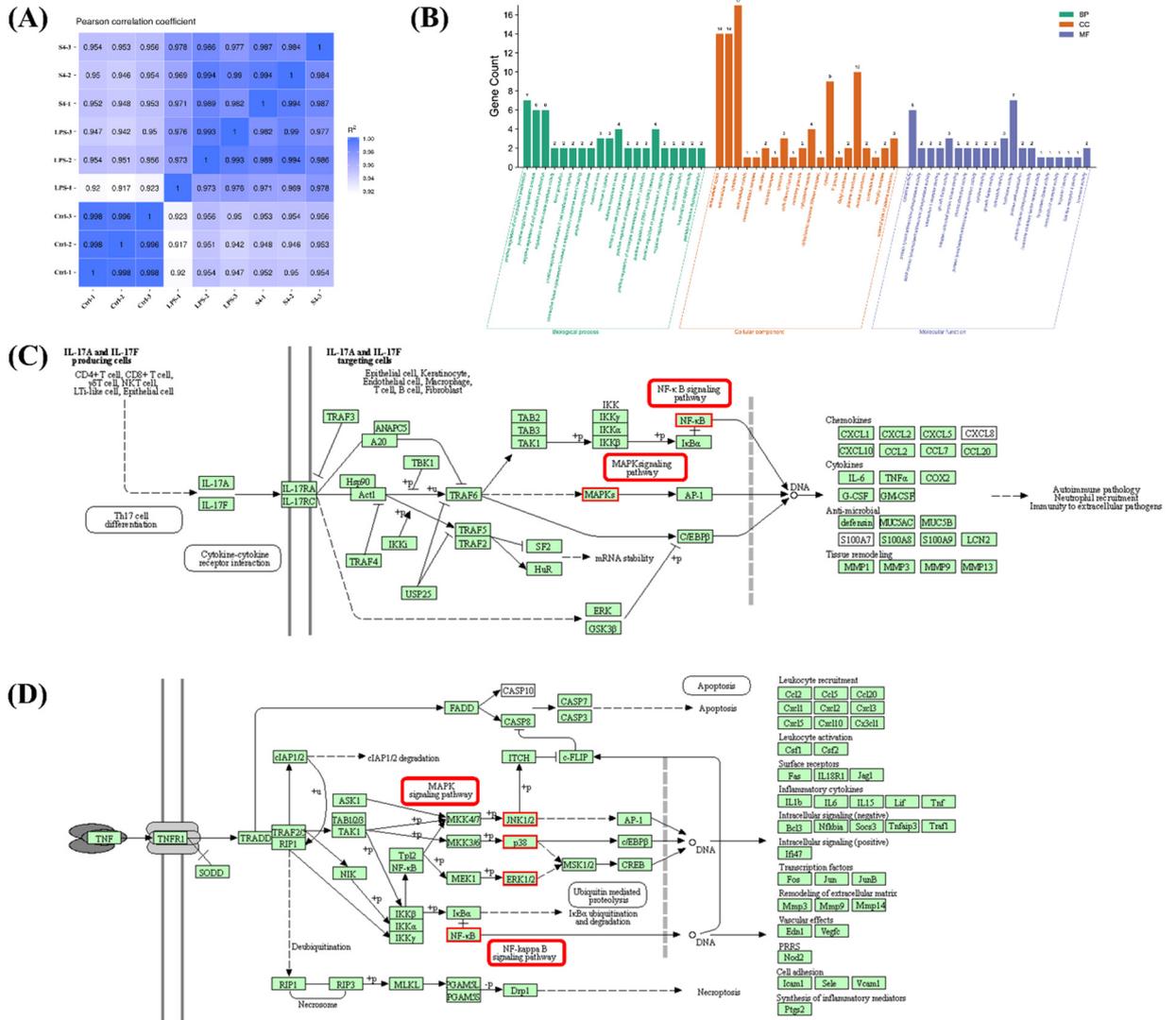


## Supplementary Materials

### Tables and Figures

**Table S1.** Geographic information of *Z. striolatum* collected from seven different habitats

| <b>Sample<br/>NO</b> | <b>Province</b> | <b>City</b> | <b>Locality</b>                      | <b>Collection<br/>Time</b> | <b>Storage<br/>Location</b> |
|----------------------|-----------------|-------------|--------------------------------------|----------------------------|-----------------------------|
| S1                   | Sichuang        | Yaan        | Mingshan                             | 2018.09                    | School of                   |
| S2                   | Hunan           | Huaihua     | Yuanling                             | 2018.09                    | Biomedical and              |
| S3                   | Jiangsu         | Nantong     | Tongzhou                             | 2018.10                    | Pharmatheutical             |
| S4                   | Hunan           | Zhangjiajie | Sangzhi                              | 2018.10                    | Sciences,                   |
| S5                   | Hubei           | Enshi       | Tujia and Miao Autonomous Prefecture | 2018.10                    | Guangdong                   |
| S6                   | Hubei           | Shiyan      | Yunxi                                | 2018.09                    | University of               |
| S7                   | Anhui           | Liuan       | Dabieshan                            | 2018.09                    | Technology                  |



**Figure S1.** Analysis of RNA-seq data in Ctrl, LPS and S4 groups. **(A)** Heatmap showed Pearson correlation coefficient between each sample. **(B)** GO enrichment analysis of 45 DEGs upregulated by LPS and downregulated by S4. **(C,D)** Part of IL-17 signaling pathway and TNF signaling pathway obtained from Kyoto Encyclopedia of Genes and Genomes (KEGG, <https://www.kegg.jp/>, accessed on 21 September 2021).