**Table S8. Functional enrichments in the network of Lj0g3v0072079.1 proteins**

|  |  |  |
| --- | --- | --- |
| Gene Ontology | cluster | Description |
| Biological Process | GO:0009788 | Negative regulation of abscisic acid-activated signaling pathway |
| GO:0009738 | Abscisic acid-activated signaling pathway |
| GO:0009414 | Response to water deprivation |
| GO:0048585 | Negative regulation of response to stimulus |
| GO:0009737 | Response to abscisic acid |
| Molecular Function | GO:0004724 | Magnesium-dependent protein serine/threonine phosphatase activity |
| GO:0043565 | Sequence-specific dna binding |
| GO:0003700 | DNA-binding transcription factor activity |
| GO:0003677 | DNA binding |
| GO:0003676 | Nucleic acid binding |
| GO:0005488 | Binding |
| Cellular Component | GO:0005634 | Nucleus |

Functional enrichments in the network originate from Gene Ontology.