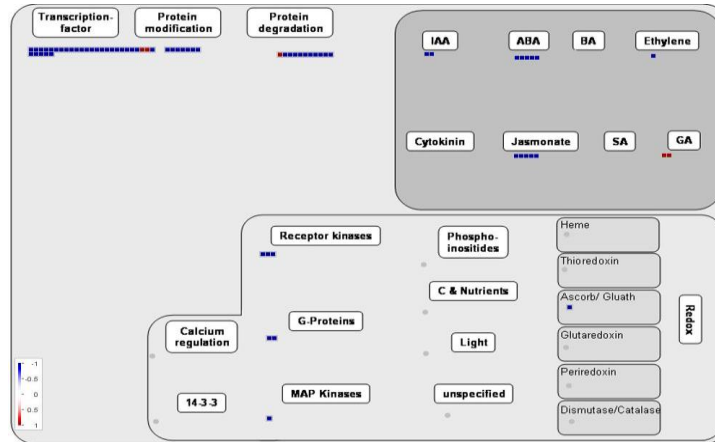


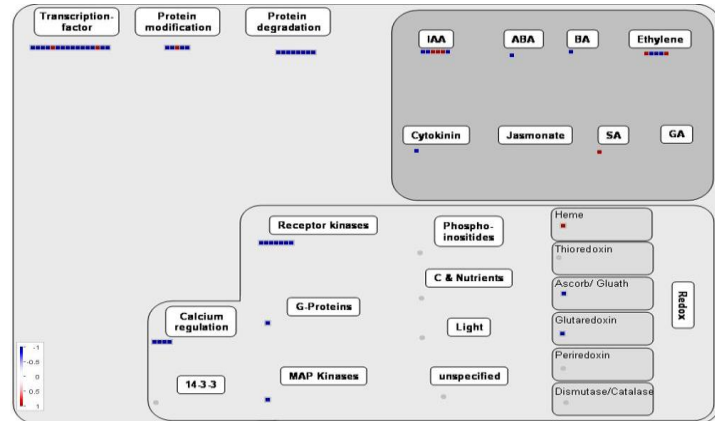
Supplemental Figure 2. MAPMAN pathways showing categories of genes that were differentially-expressed in *C. sativa* grown in soil treated with *P. migulae* 8R6 or transgenic lines expressing *acdS* under the control of the *35S* or *rolD* promoters relative to the wild type line under salt stress. Output from MAPMAN software version 3.5.1R2 (<http://mapman.gabipd.org/>).

Regulation

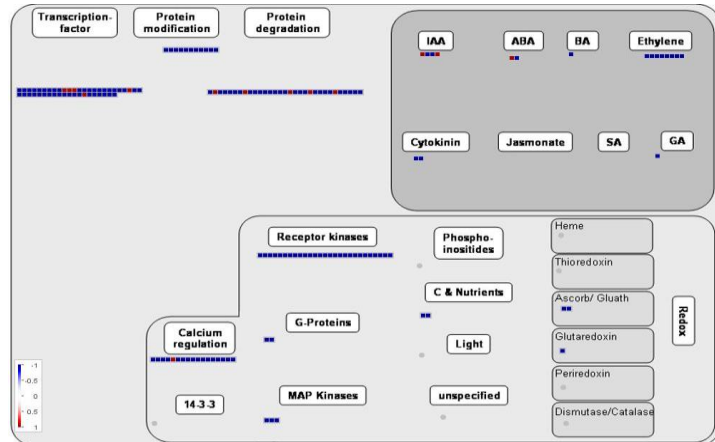
8R6-PGPB



35S::acdS

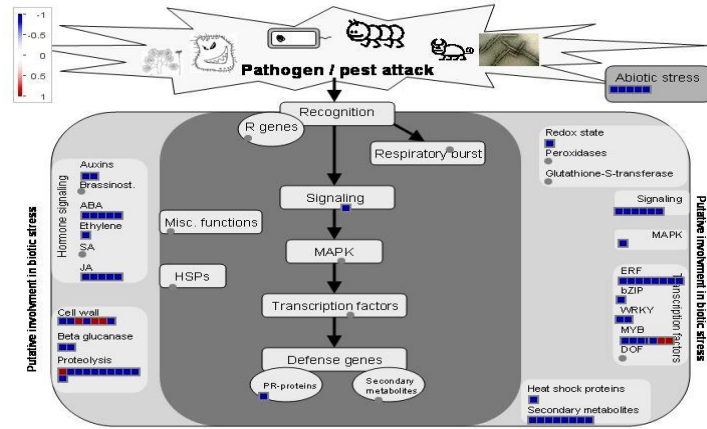


rolD::acdS

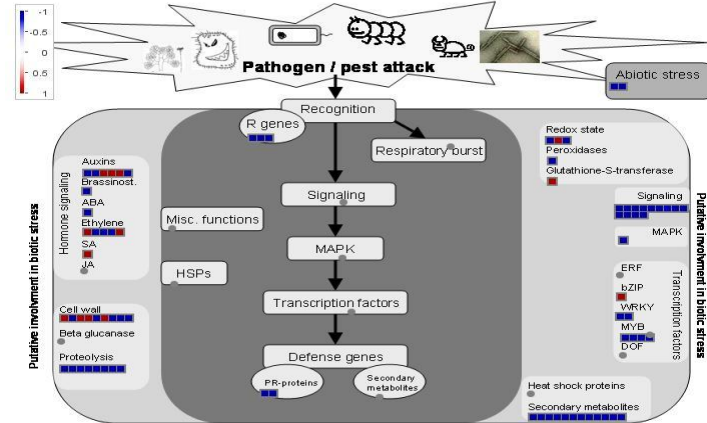


Biotic Stress Response

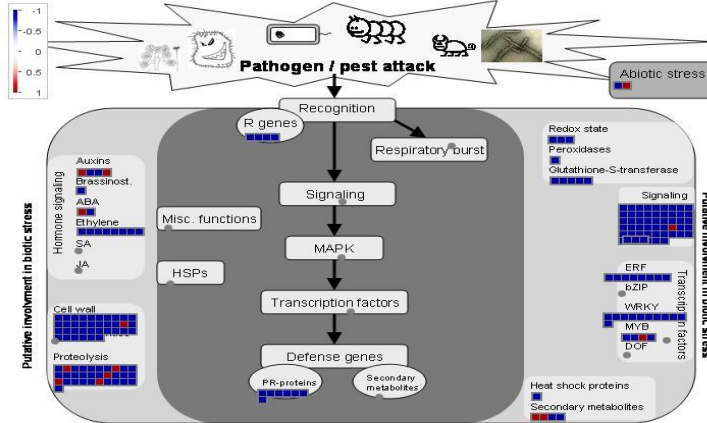
8R6-PGPB



35S::acdS

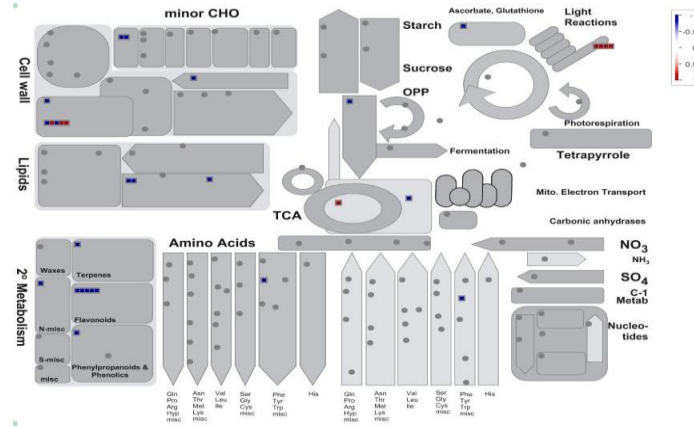


rolD::acdS

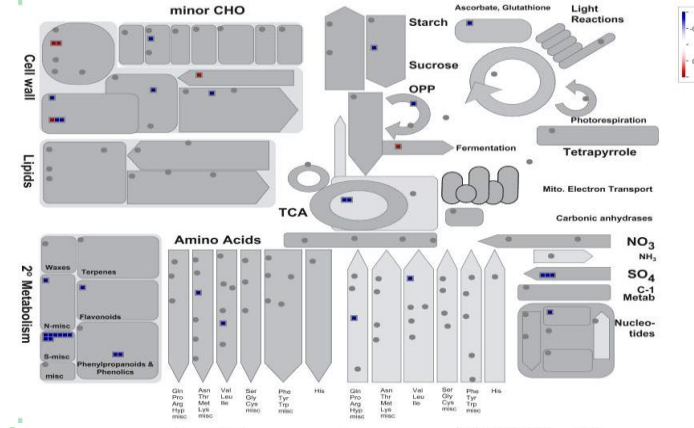


Metabolism

8R6-PGPB



35S::acdS



rolD::acdS

