Supplemental Figure 3.
Cluster analysis of common differentially-expression genes in vegetative tissue from <i>C. sativa</i> lines expressing <i>acdS</i> under the control of the <i>rolD</i> promoter or the constitutive <i>CaMV 35S</i> promoter, or from plants grown in soil treated with <i>P. migulae 8R6</i> in the presence of salt (NaCl, 15 dSm ⁻¹). Cluster analysis done using JMP software version 15.

