

Supplemental Figure 1. Expression of select *C. sativa* genes after salt stress (15 dSm-1) as determined by RNA-Seq and droplet digital PCR (dd PCR) analysis. Plots show expression (fold change) in control (wild type DH55) plus salt, 35S::*acdS* and *rolD*::*acdS* transgenic lines or after treatment with the plant growth promoting bacterium *P. migulae* 8R6 relative to the control without salt (set at 1.0) and are reported as means and standard errors of three biological replicates.

