



Extended Data Figure 4 | Growth of *C. campestris* on *A. thaliana* *sgs2-1* and *dcl4-2t* mutants with varying methodologies, as indicated.

a–d, *P* values (Wilcoxon rank-sum tests, unpaired, two-tailed) from comparison of mutant to wild-type (Col-0) are shown. Box plots show the median, box edges represent the first and third quartiles, the whiskers extend to $1.5 \times$ interquartile range, and all data are shown as dots. **a**, $n = 16$ and 9 biologically independent samples for Col-0 and *sgs2-1*, respectively. 95% confidence interval (Col-0 minus *sgs2-1*), 0.120 to 0.000. **b**, $n = 10$ and 8 biologically independent samples for Col-0 and *dcl4-2t*, respectively.

95% confidence interval (Col-0 minus *dcl4-2t*), -0.052 to 0.012 .

c, $n = 11$ and 10 biologically independent samples for Col-0 and *dcl4-2t*, respectively. 95% confidence interval (Col-0 minus *dcl4-2t*), -0.014 to 0.018 . **d**, $n = 8$ and 9 biologically independent samples for Col-0 and *dcl4-2t*, respectively. 95% confidence interval (Col-0 minus *dcl4-2t*), -0.184 to 0.008 . **e**, $n = 14$, 14 and 12 biologically independent samples for Col-0, *sgs2-1*, and *dcl4-2t*, respectively. 95% confidence interval (Col-0 minus *sgs2-1*), -0.020 to 0.090 . 95% confidence interval (Col-0 minus *dcl4-2t*), -0.010 to 0.110 .