



Fig. S2. Comparison of three different estimates of the number of ORFs affected by endophyte infection. Shown are the number of ORFs in different tissue types for which expression differed between infected and uninfected plants, as estimated by the statistical procedure described in the manuscript, and the number of ORFs for which the software DEGseq v. 1.16.0 indicated significant (false discovery rate $q < 0.05$) effects. For comparison also included are the numbers of ORFs for which another software, DESeq v. 1.14.0 indicated significant effects. Tissue types are labeled as described in Fig. 1.