



Fig. S5 Probability of detection of significant Lp19 effects on expression of ORFs significantly affected by FI1 as a function of reads mapping to the ORF. Shown is the frequency of significant (FDR $q < 0.05$) ≥ 2 -fold effects of Lp19 infection on ORFs reported as significantly affected by FI1 infection (Dupont et al., 2015), as a function of sequencing depth (number of all reads mapping to the ORFs in all tissues in all sample biological repeats in this work). In the x axis sequencing depth intervals are shown, followed, in brackets, by the number of ORFs affected by FI1 infection in each read depth category.