



Fig. S1 Correlation between RT-qPCR and RNAseq results. Each gene id indicates *B. napus* genes that were differentially expressed in RNAseq and qRT-PCR. Relative log₂ fold changes at 1-, 4-, and 7-DPI were calculated between inoculated clubroot resistant (CR) and inoculated clubroot susceptible (CS) genotypes. 1-, 4-, and 7- DPI= indicates, days post inoculation. Error bar shows the standard error of mean (n=3).