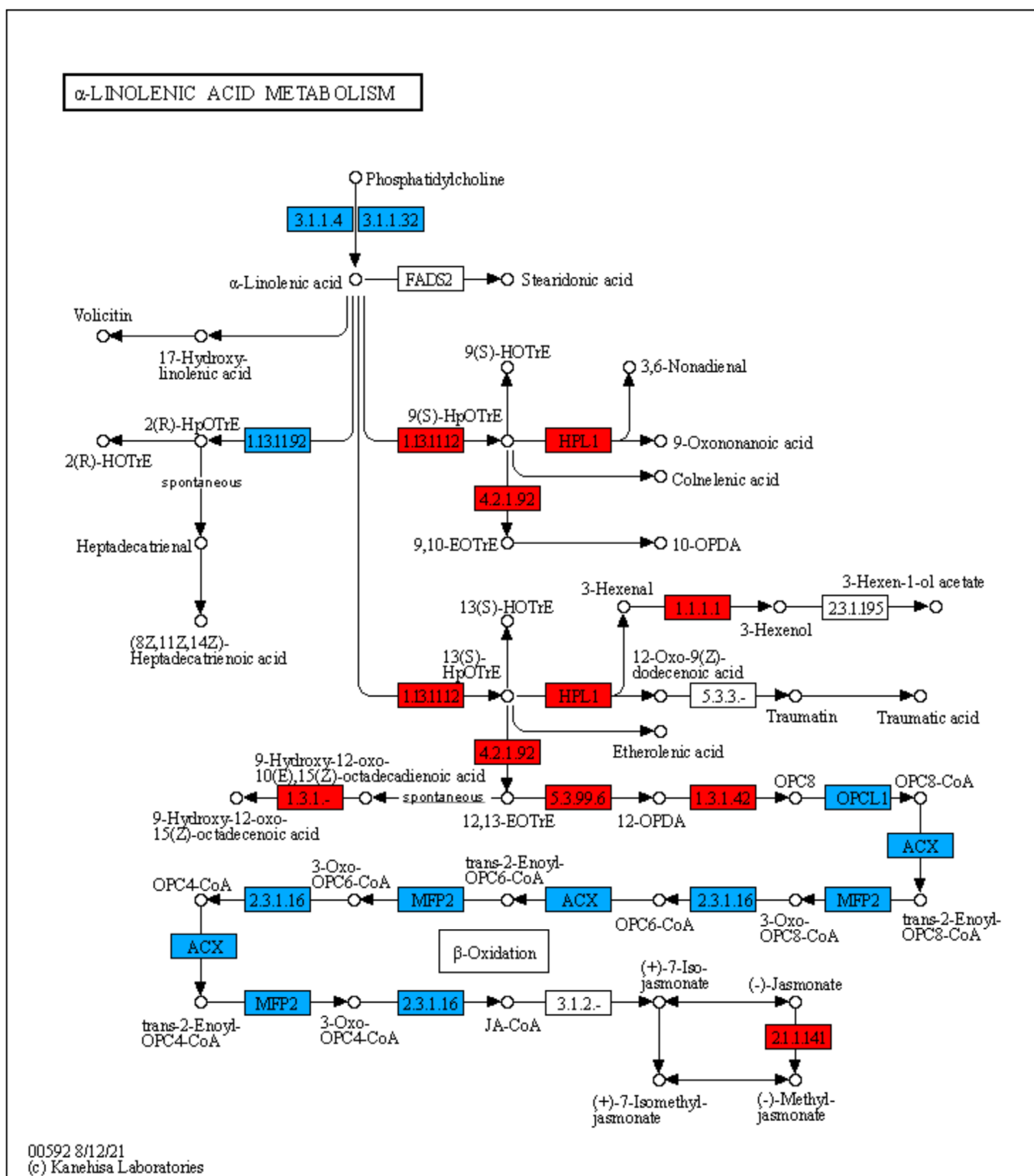
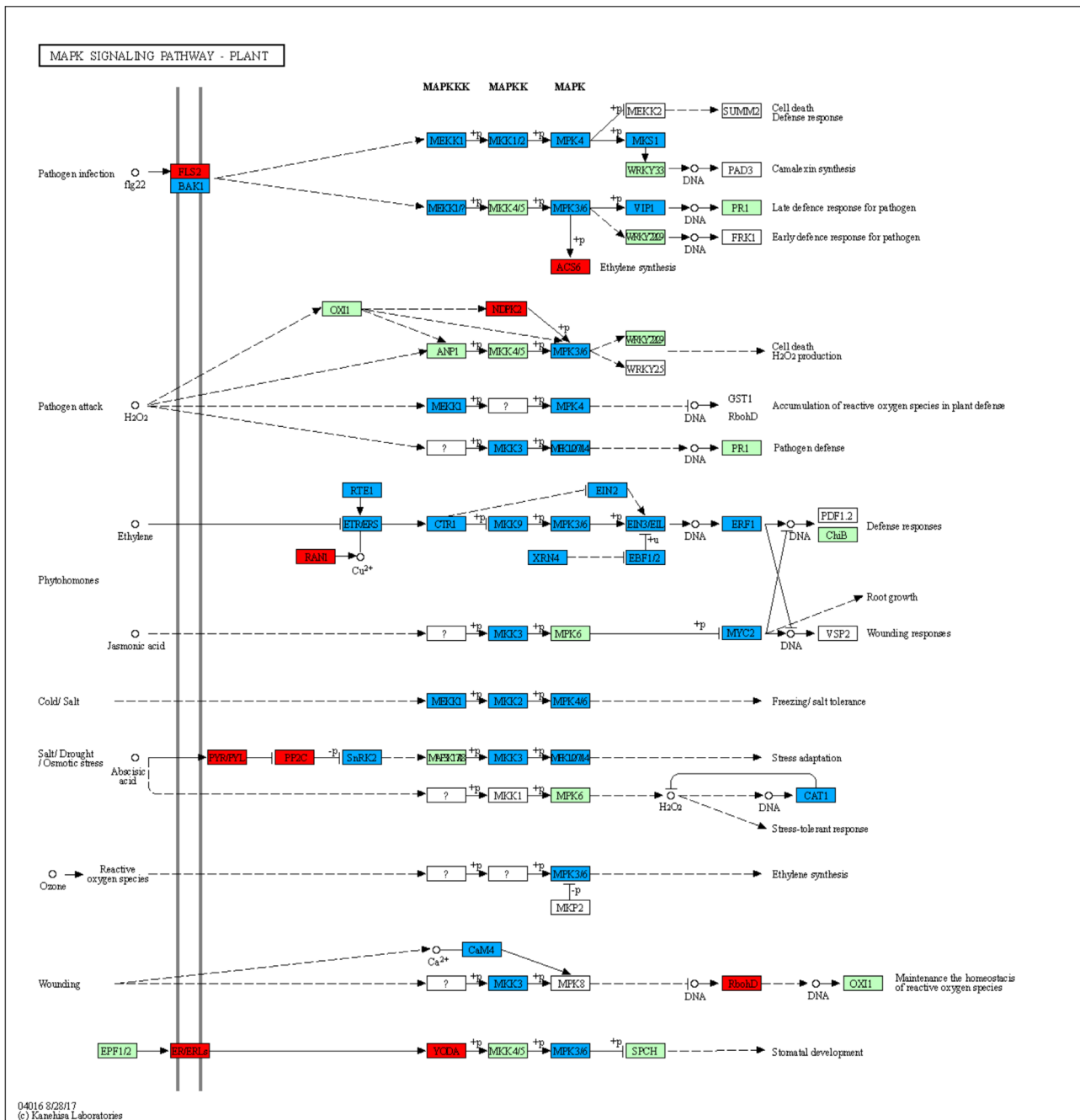


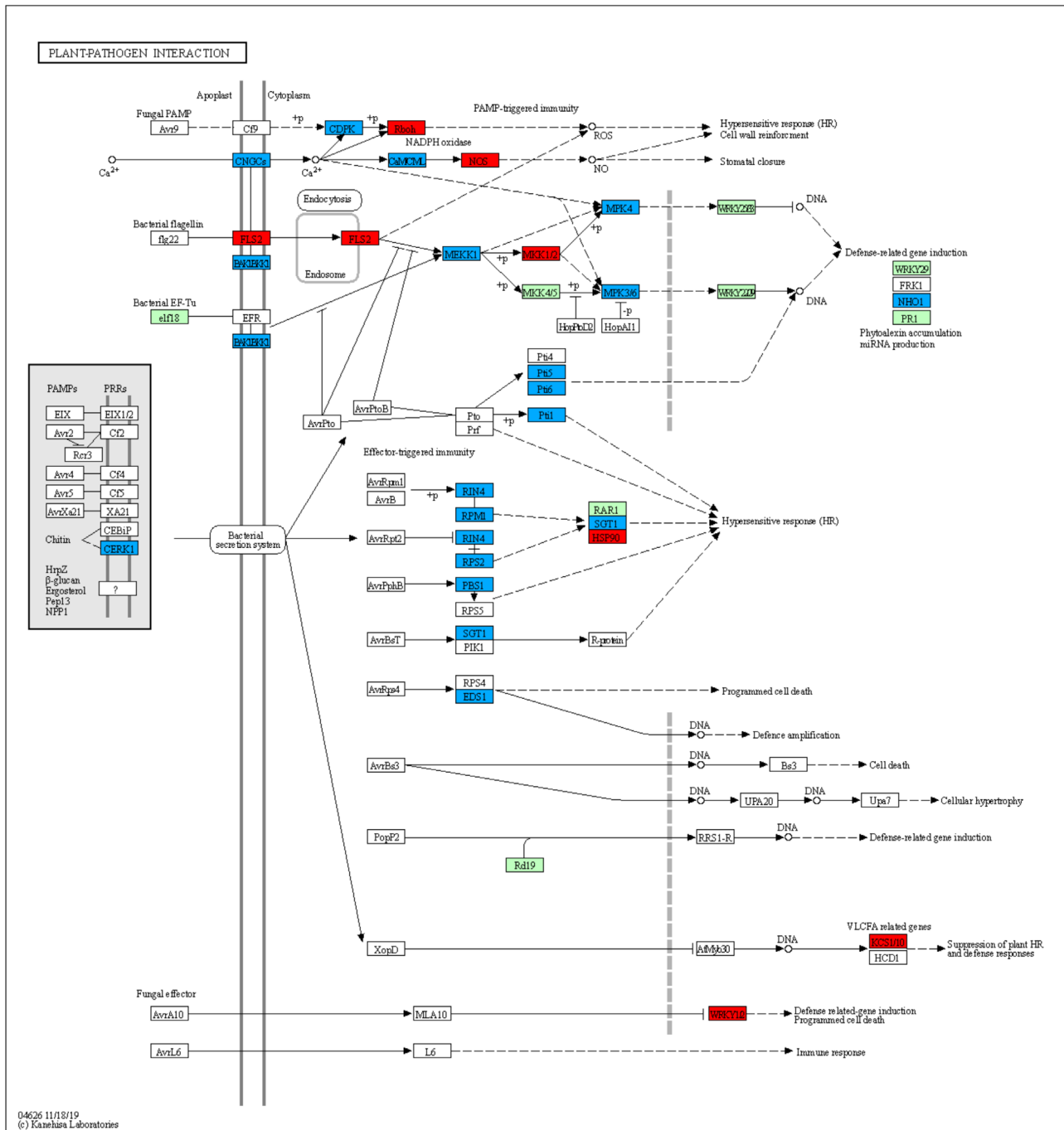
Supplementary Fig. 1. Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway showing plant hormone signal transduction in soybean. KEGG orthologue (KO) genes whose expressions are upregulated (red) and down-regulated (blue) in infected plants were presented in this. White boxes mean there are no identified functional orthologs in soybean. Green ones indicate that the gene expression was not changed by pathogen infection or the gene was not detected in this transcriptome analysis.



Supplementary Fig. 2. Kyoto Encyclopedia of Genes and Genomes pathway showing α -linolenic acid metabolism in soybean. Color codes are the same as those described in the legend of Supplementary Fig. 1.



Supplementary Fig. 3. Kyoto Encyclopedia of Genes and Genomes pathway showing MAPK signaling pathway-plant in soybean. Color codes are the same as those described in the legend of Supplementary Fig. 1.



Supplementary Fig. 4. Kyoto Encyclopedia of Genes and Genomes pathway showing plant-pathogen interaction in soybean. Color codes are the same as those described in the legend of Supplementary Fig. 1.