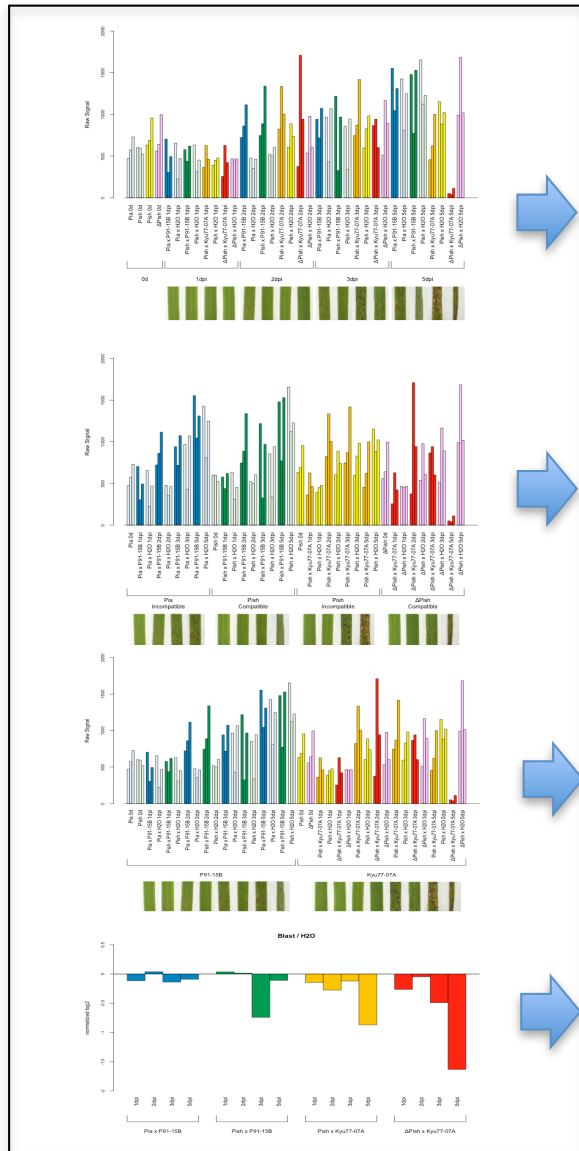


Expression profile in graphical format

The expression profile for each gene is presented in graphical format with the raw signal intensity data arranged according to time course experiments (graph 1), susceptible / resistant lines of rice (graph 2), and strains of the *M. oryzae* used for inoculation (graph 3), and the normalized signal intensity (graph 4) for expression data .



Graph 1. Raw signal intensity data arranged according to number of days post incubation (dpi). The figures at the bottom show the corresponding lesion in the leaf samples.

Graph 2. Raw signal intensity data arranged according to lines of rice inoculated with *M. oryzae*. The figures at the bottom show the corresponding lesion in the leaf samples.

Graph 3. Raw signal intensity data arranged according to the strains of *M. oryzae* used for inoculation. The figures at the bottom show the corresponding lesion in the leaf samples.

Graph 4. Signal intensity data based on 75 percentile normalization and log2 transformation for the average relative value to H₂O treatment (blast / H₂O) of 3 replicates among susceptible and resistant lines.