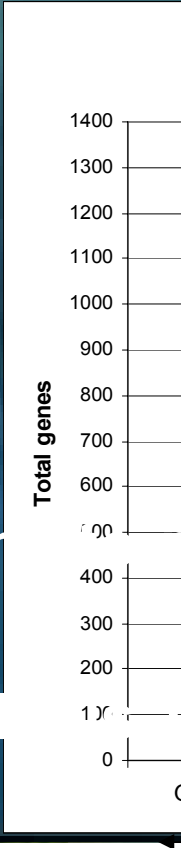


of the four infection times tested were considered a biological replicate. Each infection point together with the corresponding control were analyzed. The data displayed changes in gene expression. Several transcriptional changes were observed. 1.5% of the induced or repressed genes were involved in gall formation (Fig.2). The early gall formation (1-3 dpi, 1366-1314 genes). The transcriptional profile of the compatible interaction is especially at 1 and 3 dpi. The number of up-regulated genes is 1366 and the number of down-regulated genes is 1314. In contrast, the number of



figuras

Te he camabiado el desarrollo no he usado el para que tuviera pero convendria