

Supporting Information

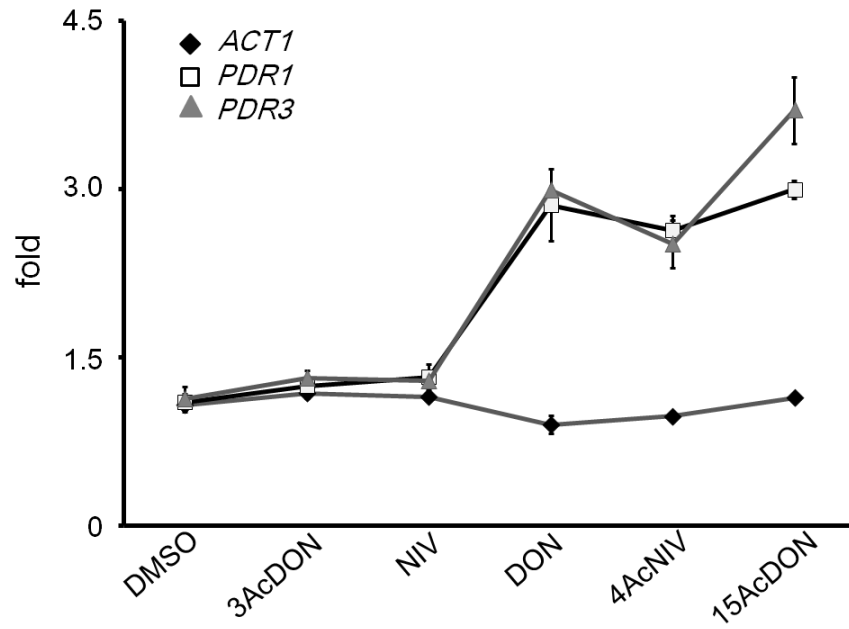


Figure S1. Confirmation of the DNA microarray by quantitative PCR. Real-time PCR-based quantitative estimation was executed using *ACT1* expression as an internal control. (◆), (□), and (▲) indicate *ACT1*, *PDR1*, and *PDR3* gene expression changes, respectively. The data set of each gene expression corresponded with the expression patterns of the DNA microarray (see Figure 6). Bars = SE; $n = 3$.