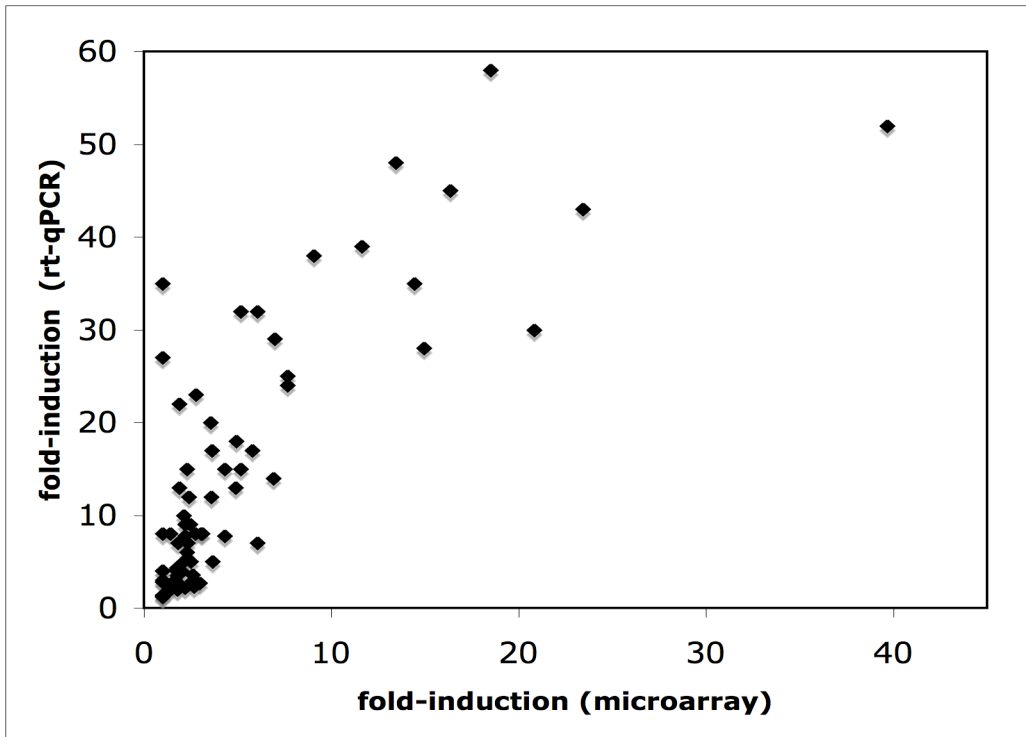


Figure S4. Data validation by rt-qPCR.

(A) Fold-induction by rt-PCR vs fold induction by microarrays [2 exp(ratio)]. The expression levels of 51 genes listed in Table S6 were examined. (B) Time-course of the expression of three upregulated genes in *atm* and WT.

A



B

