Supplemental Figure 1 (Giegé *et al.*)

Overall dispersion of gene expression in the individual microarray experiments.
Boxplots for the individual microarray hybridisations S1 (1 to 3), S2 (1 to 3) and RS (1 to 3) before normalisation (**A**) and after quantile method normalisation (**B**). Log-ratio values are given on the y axis. The horizontal line in the middle of the boxes shows the log-ratio medians and the horizontal bars indicate the maximum and minimum log ratios.
The median absolute deviation (MAD) values are shown above the plots for the individual experiments in **A** and for the group of biological replicates in **B**.