Fig. S1. Correlation between 20 RNA-Seq samples based on a spearman correlation analysis. Deeper blue indicates a higher correlation. The sample names are indicated on the left and top of the heatmap. Each cell shows the correlation between two samples. The correlation coefficients of two technical replicates are highlighted by red squares. The sample abbreviation is listed in Table 1. The suffix indicates the replicate number (rp1 or rp2).