

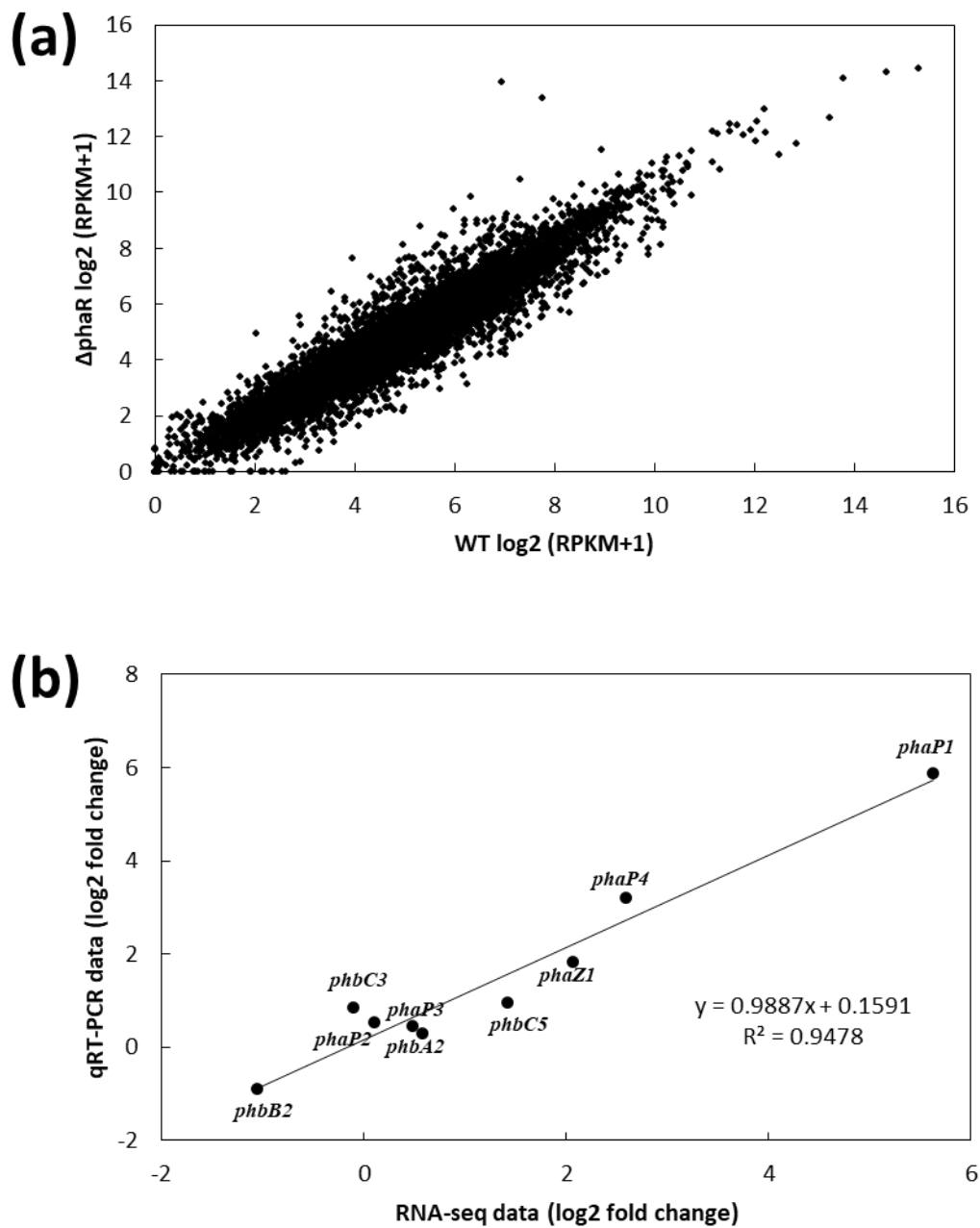


Fig. S3. Overview of RNA sequencing analysis data quality. (a) Comparison of USDA110 (WT) and ΔphaR RPKM values. (b) Comparison of log2 transformed data of quantitative RT-PCR and RNA sequencing. Fold change refers to the relative expression in ΔphaR to USDA110 (WT).