

SUPPLEMENTAL MATERIAL

Table S1: Genes with detectable expression during PG and PTG.

Table S2: Genes with altered transcription (up- or down-regulated more than 1.6-fold) during PG and PTG.

Table S3: Cluster analysis results of the expressed genes during PG and PTG.

Table S4: Results of functional category and gene family analysis of genes with altered expression levels during PG and PTG.

Table S5: MAPMAN analysis of genes with altered expression levels during PG and PTG.

Table S6: Expressed genes in the CaM/CML, CHX and Hsp gene families.

Table S7: Transcriptionally up-regulated genes of the CaM/CML, CHX and Hsp families during PG and PTG.

Table S8: Sequences of the specific primers for the genes tested in real-time PCR experiments.

Figure S1: Cluster analysis of the expressed genes during PG and PTG.

Figure S2: MAPMAN analysis of the difference in transcription between PG and PTG.