**Additional file 8:**

**Figure S5**

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**Figure S5** RNA-seq analysis of WT and *wsl9* seedlings at 30℃. mRNA was enriched from total RNA isolated from 10-d-old WT and *wsl9* mutant seedlings using oligo-(dT) fragmented and reverse-transcribed using random hexamer primers. The library was thenconstructed and sequenced using an Illumina HiSEquation 2000. a Frequencies of detected genes sorted according to expression level. b Read numbers of WT and *wsl9* mutant sequences. c Volcano plot showing the overall alterations in geneexpression in WT and *wsl9* mutant b- Y-AXIS ‘Read number’*.*