

Poster Presentation

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Expression of trehalose-6-phosphate synthase gene from *Arabidopsis thaliana* in transgenic tobacco: a strategy to increase temperature stress tolerance

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Background

Genetic engineering of plants towards osmoprotectant accumulation is gaining increased importance within the broad context of abiotic stress tolerance [1]. An enzyme, trehalose-6-phosphate synthase, is believed to play a key role in the synthesis of the disaccharide trehalose and hence on the improvement of abiotic stress tolerance [2]. We used *Agrobacterium* to transform tobacco plants to express the trehalose-6-phosphate synthase gene from *Arabidopsis thaliana*, under the control of CaMV 35S promoter and using the vector pGreen 0229 [3]. Transgenic T2 plants were evaluated for gene expression by northern and western blots. Seeds were sown in media germinated at: 15, 25 and 35°C for evaluating germination rates under high and low temperatures.

Results

Three of the transgenic lines obtained (B5A, B5H and B1F) have distinct levels of gene expression: B5H and B5A are high expressing lines while B1F is a low expressing one. In non-transgenic controls no expression was detected (Figure 1).

Transgenic lines were shown to have significantly higher germination rates under low and high temperatures (respectively, 15 and 35°C) than wild type plants (Table 1).

Conclusion

Our results demonstrate that transgenic plants accumulating trehalose-6-phosphate synthase have an altered phenotype that includes temperature stress tolerance upon germination. We suggest that AtTPS1 can be used to engi-

Table 1: Germination rates (Number of seeds germinated per 100 seeds placed on germination medium) of three transgenic lines at three different temperatures.

Temperature	WT	B5A	B5H	B1F
24°C	99.0 ^a (1.0)	99.0 ^a (1.50)	99.0 ^a (2.00)	100.0 ^a (0.00)
15°C	32.3 ^a (4.1)	95.0 ^b (3.97)	97.4 ^b (2.95)	97.0 ^b (2.80)
35°C	18.0 ^a (3.8)	88.0 ^b (7.9)	85.2 ^b (6.35)	83.2 ^b (5.9)

^{a, b}Percentages with different superscripts indicate statistical significance ($p < 0.05$); Standard deviations are shown between parenthesis; WT – Wild Type plants; B5A, B5H and B1F – Transgenic lines

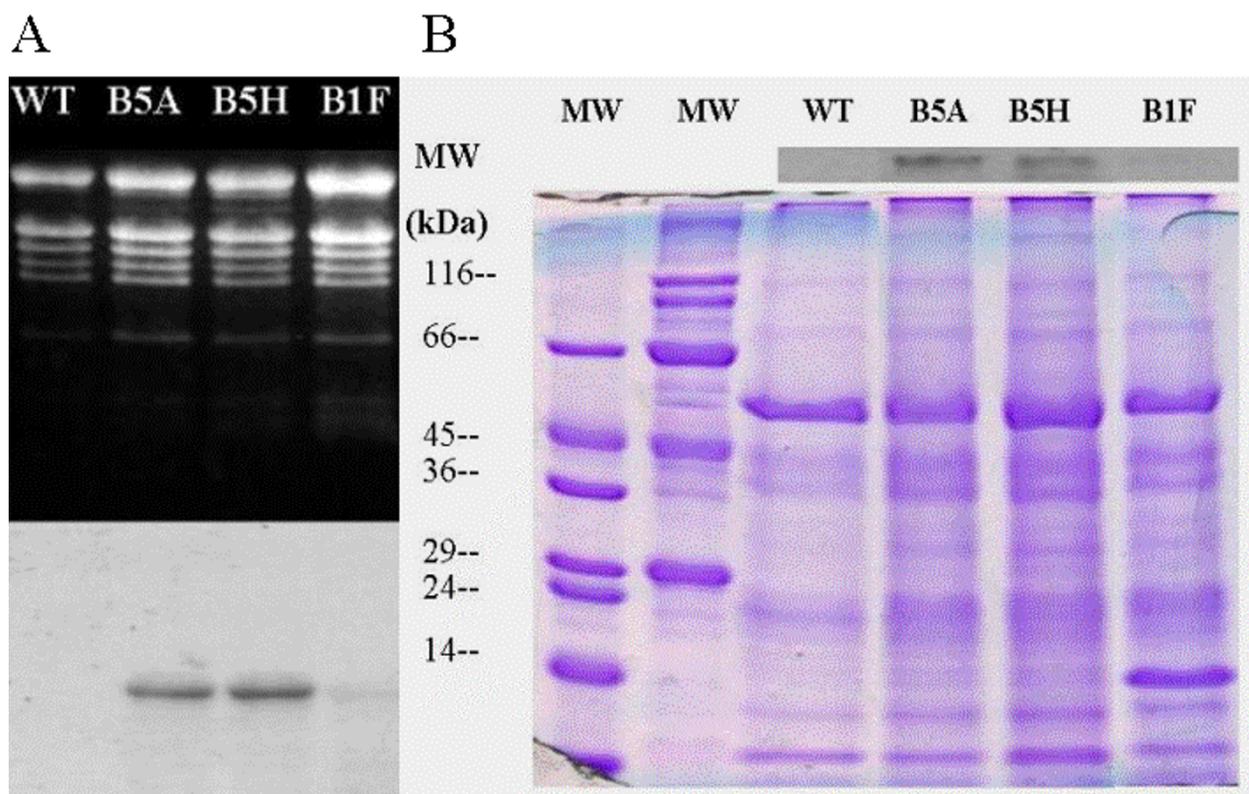


Figure 1

AtTPS1 gene expression analysis in WT and transgenic tobacco lines. A – Northern blot B – Western blot. In both cases, no AtTPS1 protein production was detected in control wild type plants while transgenic lines showed accumulation of AtTPS1 transcripts and enzyme. **WT** – Wild Type; **B5A**, **B5H** and **B1F** – Transgenic line. MW – Molecular weight markers (kDa).

near important crop plants such as maize, wheat or rice to withstand different environmental stresses.

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