

# Evaluation of wheat events transformed with the *p5cs* gene under conditions of water stress

## Avaliação de eventos de trigo transformados com o gene *p5cs* e submetidos à seca

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### ABSTRACT

The enzyme  $\Delta^1$ -pyrroline-5-carboxylate synthetase (P5CS), encoded by the *p5cs* gene, is the rate-limiting step for proline biosynthesis, which acts as an osmoprotectant. The launch of transgenic commercial varieties depends on physiological and phenotypic characterization of events. The objective of this study was to evaluate five *p5cs* gene events (591, 14, 326, 164, 312), through physiological analyses and production rates during droughts. Plants of events and irrigated (IC) and stressed controls (SC) in flowering stage were submitted to 16 days of complete water restriction. The following elements were evaluated: relative water content (RWC); malondialdehyde concentration (MDA%); membrane stability index (MSI%); proline content; and production rates. As a result, the effect of water stress was only perceived after 12 days of water restriction with a decrease in the RWC values. Transgenic plants showed higher proline levels on the 16th day, compared to the SC (33 to 62%). The levels of MDA were lower on the 12<sup>th</sup> and 16<sup>th</sup> days for all the plants evaluated, but the MSI were similar to the IC for the events 591, 326, and 164, demonstrating the osmoprotectant effect. However, higher levels of proline did not result in higher production rates in the transgenic plants.

**Keywords:** Poaceae, Abiotic stresses, Proline, Physiological Index, Production

### RESUMO

A enzima  $\Delta^1$ -pirrolina-5-carboxilato sintetase (P5CS), codificada pelo gene *p5cs*, é o passo limitante da biossíntese de prolina, que age como um osmoprotetor. O lançamento de variedades comerciais transgênicas depende da caracterização fisiológica e fenotípica de eventos. O objetivo deste estudo foi avaliar cinco eventos (591, 14, 326, 164, 312) contendo o gene *p5cs*, através de análises fisiológicas e taxas de produção sob estresse hídrico. Plantas de eventos e controles irrigados e estressados (CI e CS) em estágio de florescimento foram submetidos a 16 dias de restrição hídrica. Foram avaliados: teor relativo de água (RWC); concentração de malondialdeído (MDA%); índice de estabilidade da membrana (MSI%); conteúdo de prolina; e taxas de produção. Como resultado, o efeito do estresse hídrico só foi percebido após 12 dias de restrição hídrica com uma diminuição nos valores de RWC. Plantas transgênicas apresentaram maiores níveis de prolina no 16<sup>o</sup> dia, comparado ao SC (33 a 62%). Os teores de MDA foram menores no 12<sup>o</sup> e 16<sup>o</sup> dias para todas as plantas avaliadas, mas os MSI foram semelhantes ao IC para os eventos 591, 326 e 164, demonstrando o efeito osmoprotetor. Porém, níveis altos de prolina não resultaram em maiores taxas de produção de grãos.

**Palavras-Chave:** Poaceae, estresse abiótico, prolina, índice fisiológico, produção

## INTRODUCTION

Wheat is cultivated in approximately 17% of the world's agricultural land, which corresponds to 200 million hectares, and is one of the major sources of calories and protein in the human diet (Jones, 2005). Due to its significant economic importance, wheat is produced in various countries and climates. Drought and salinity are thought to be the most important abiotic stresses that damage this crop worldwide (Dolferus *et al.*, 2011; Saad *et al.*, 2013).

Water stress promotes loss of cellular water or osmotic stress, and the tolerance capacity is related to the maintenance of cellular integrity, targeting a functioning metabolism, which favors the plants' recovery after the end of the water deficit period (Ozden *et al.*, 2009).

One of the most important biochemical modifications in plants subjected to abiotic stresses is the production of reactive oxygen species (ROS) (Ahmad *et al.*, 2008). Under water restriction, there is a reduction in the rate of photosynthesis due to the closing of stomata and, along with the water shortage and constant capture of light, the photosynthetic fixation of CO<sub>2</sub> is limited – resulting in an accumulation of electrons (Ashraf *et al.*, 2008; Munns & Tester, 2008). The ROS thus produced damage cell membrane through the peroxidation and re-esterification of fatty acids.

Recent studies have shown that genetic manipulation of genes responsible for the biosynthesis of low-molecular-weight metabolites has granted the plants a better tolerance to drought and salinity (Dolferus *et al.*, 2011; Dunwell, 2014). Proline is an osmoprotectant with a fundamental role in the protection against oxidative damage, due to a more efficient antioxidant system, in which the proline seems to be more involved in the sequestration and inactivation of free radicals and in the protection of antioxidant enzymes (Ozden *et al.*, 2009; Bhas-kara *et al.*, 2015). Specifically, the enzyme  $\Delta^1$ -pyrroline-5-carboxylate synthetase (P5CS), encoded by the *p5cs* gene, is considered to be the rate-limiting step in the synthesis route of proline.

The comparatively high proline accumulation in transgenic plants is responsible to a better growth,

more chlorophyll and relative water content and lower levels of lipid peroxidation of plants under osmotic stress (Khan *et al.*, 2015).

Increasing tolerance to water deficit stress, by *p5cs* genetic transformation, has been obtained for wheat (Vendruscolo *et al.*, 2007; Pavei *et al.*, 2016). Generally, after the transformation, the positive events for the transgenesis are selected according to the presence of a single copy, fertility, and normal phenotypic appearance (Xiao *et al.*, 2009). Several genetic factors can affect transgene expression, including epigenetic factors (Dietz-Pfeilstetter, 2010). The stable expression throughout generations and the non-occurrence of gene silencing seem to be the major challenges for plant improvement in polyploid genomes (Anand *et al.*, 2003; Rooke *et al.*, 2003; Meng *et al.*, 2006; Yao *et al.*, 2006).

In addition, the pre-selection of high productivity events is fundamental for the launch of commercial transgenic varieties, especially when the system of genetic transformation used is the gene gun, which consists of the insertion of gene fragments without the number and position control of insertion in the genome (Mlynorowa *et al.*, 1996; Ulker *et al.*, 1999; Basri, 2005).

This study was carried out with the objective of evaluating the behavior of five transgenic events of wheat containing the exogenous *p5cs* gene in the 3<sup>rd</sup> generation (T3), under severe water stress, in order to establish a physiological characterization and an evaluation of the efficiency of transformation events in terms of wheat plants' tolerance to water stress.

## MATERIAL AND METHODS

The experiment for the phenotypic evaluation was conducted in a greenhouse, at Coodetec's Biotechnology Laboratory, in Cascavel-PR, Brazil. The transformation procedure used in this experiment in wheat (*Triticum aestivum* L. cv CD200126) was the gene gun. It was executed according to the protocol proposed by Bohorova *et al.* (1999). The pJS107 plasmid, which contains the cDNA of the *p5cs* gene ( $\Delta^1$ -pyrroline-5-carboxylate synthetase) originated from *Vigna aconitifolia* (Jacq.) Maréchal

was used, driven by the stress-inducible promoter *AIPC* (Zhu *et al.*, 1998). The *pJS107* also contains the gene bar merged into the 35S promoter, which was used for the selection of transformants in the presence of phosphinothricin (PPT) – confirmed by analyses made by Southern Blot (Vendruscolo *et al.*, 2007).

Wheat seeds derived from five transformation events  $T_1$  from 12 originally obtained (events 1(164); 2(591); 3(326); 4(312) and 6(14) (Vendruscolo *et al.*, 2007-Figure 1) were planted in greenhouses under controlled conditions of temperature and irrigation (average temperature  $24 \pm 2$  °C, Relative Humidity 55-75%, watering every day). Seeds of the same genotype (CD200126) were used as a control group, although they were not transformed. The transgenic plants were multiplied and cultivated by self-fertilization until they originated the  $T_3$  descent. They were considered homozygous lines for the transgenic insertion through the obtained inheritance pattern of 3:1 in  $T_1$  (Chi-squared 1.1 to 3.2).

$T_3$  seeds were placed to germinate in Germitest Paper. After germinating for 3 days, seedlings were transferred into pots (four seedlings per pot), each containing 4kg of previously sifted and fertilized soil (pH 6.4; P = 60 mg.dm<sup>-3</sup>; K = 1.14 cmol.dm<sup>-3</sup>; Ca = 6.69 cmol.dm<sup>-3</sup>; Mg = 3.03 cmol.dm<sup>-3</sup>; H+Al = 3.18 cmol.dm<sup>-3</sup>; Cu = 14.45 mg.dm<sup>-3</sup>; Mn = 400 mg.dm<sup>-3</sup>; Fe = 21mg.dm<sup>-3</sup>; Zn = 32.49 mg.dm<sup>-3</sup>; SB = 10.86 cmol.dm<sup>-3</sup>; 41.6 g.dm<sup>-3</sup> organic matter). Thinning was applied after 15 days of sowing, remaining only one plant per pot. The experiment involved five transformation events (591, 14, 326, 164, and 312) and the control groups (with and without water stress), in completely randomized design with five repetitions, totaling 35 pots.

When the plants reached the booting stage, around 60 days after germination, corresponding to the stage 45 in the Zadocks scale (Zadocks *et al.*, 1974), water stress was applied and measured by Relative Water Content (RWC%) (Schonfeld *et al.*, 1988). The treatments underwent 16 days of complete water restriction (except for the control group without water stress). For the evaluation of the physiological parameters, the second leaves of all the plants were collected and, for the quantification of proline and malondialdehyde (MDA), flag leaves were

sampled every 4 days. After application of water shortage, the pots were irrigated again. The last collection was made four days after the irrigation was restored, totaling 6 collections. The collections of leaves for physiological analyses were always performed in the morning, between 8:30 a.m. and 10 a.m., at an average temperature of 24°C.

The following parameters were evaluated: proline content (Bates *et al.*, 1973); Membrane Stability Index (MSI) (Babu *et al.*, 2004), and lipids peroxidation level was determined in terms of MDA content (Sairam *et al.*, 1998).

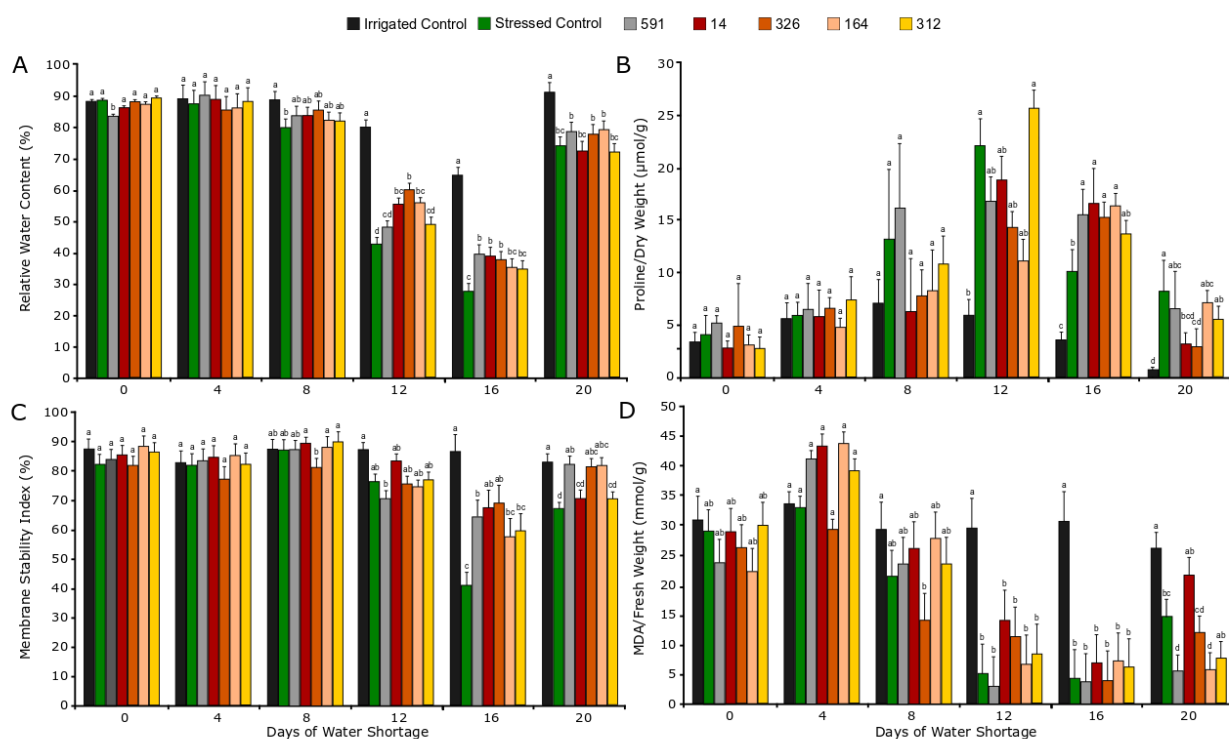
In order to evaluate the production capacity of both the plants of events subjected to water stress and the control group plants, the wheat ears were collected and counted, and the total number of seeds, the weight of 100 seeds, and the total weight of seeds per plant were determined at the end of the cycle.

The results were submitted to analysis of variance (ANOVA) and to the Turkey test to 5% of probability, with the aid of the GENES program (Cruz, 2006).

## RESULTS AND DISCUSSION

Although the events used in this study show the insertion of the transgene – confirmed by the Southern Blot in the  $T_1$  generation, the number of inserted copies of the gene *bar* and *p5cs* has not been determined yet. However, some authors recommend the verification of performance of transgenic events containing multiple copies to confirm the action and effects of the transgene position in drought conditions (Xiao *et al.*, 2009). Yao *et al.* (2006), for example, reported that the occurrence of transformants with a small number of copies in wheat is relatively rare, especially when the transformation technology used is the gene gun.

In normal water supply conditions before water stress was applied, the relative water contents (RWC) in the control group plants were not different from those in the transgenic plants in the different events evaluated (Figure 1A). Under water stress conditions, the values of RWC were very similar until the 8<sup>th</sup> day of stress application.



**Figure 1** - Effect of water stress on: A-Relative water content (%);B-Proline content( $\mu\text{mol/g}$ ); C- Membrane stability index(%) and D-Malondialdehyde (MDA)content (mmol/g). Values are the means of  $n=5 \pm S D$ . The same letters in the figures do not show statistical differences by the Tukey's test at a probability of 5%.

A different behavior among the transgenic plants and the control group plants was observed from the 12<sup>th</sup> day of water stress on. The highest numbers of RWC were found in the events 14, 326, and 164, in the order of 29%, 40%, and 29%, respectively, when compared to the plants under stress of the control group. The RWC numbers decreased even more as the stress was prolonged (16<sup>th</sup> day), and the events 591, 14, and 326 continued to present a RWC higher than the control group under water stress. The decrease in the RWC demonstrates that the plants experienced water stress as of the 12<sup>th</sup> day of water restriction. After the end of the water restriction period and plant rehydration (20<sup>th</sup> day), the plants recovered the RWC numbers to 70% and 80%, including the stressed control group.

Regarding proline concentration, we noted a natural fluctuation in the irrigated control values, which varied from 1.5 to 7  $\mu\text{mol} \cdot \text{g}^{-1}$  of dry weight over the different periods we assessed (Figure 1B). Control and transgenic plants presented proline

contents varying from 2.9 to 25  $\mu\text{mol} \cdot \text{g}^{-1}$  of dry weight under water stress in the different events we evaluated. It proves the importance of this osmo-protectant during stress. The events 591 (8<sup>th</sup> day) and 312 (12<sup>th</sup> day) presented the highest increase in proline indexes compared to the irrigated controls (2.2 and 4.3 fold, respectively). In control plants submitted to stressed, that increase was 3.7 fold higher on the 12<sup>th</sup> day compared to the irrigation control. However, we did not register events displaying higher proline contents that could indicate an over-expression due to the presence of *p5cs* multi copies. In this regard, Rooke *et al.* (2003) observed that there is no direct correlation between the number of transgene insertions and the expression level.

We observed a decrease in proline levels on the 16<sup>th</sup> day of water restriction. However, when compared to the control group submitted to water stress, transgenic plants in the different events we evaluated maintained high levels of proline due to

the induced-stress expression, as the transgene has the *aipc* promoter (Vendruscolo *et al.*, 2007). In the assessment administered on the 16<sup>th</sup> day, the percentages of proline in the transgenic plants in comparison to the control group (under stress) were as follows: 52% (event 591); 63% (event 14); 49% (event 326); 61% (event 164); and 33% (event 312).

The results of proline levels that were different from those of the control groups (both under stress and irrigated) demonstrate the differential expression of the insert holding the gene *p5cs* and the *aipc* promoter, which shows that there was no gene silencing or transgene co-suppression due to a possible presence of multicopies in the different aspects we evaluated. Rooke *et al.* (2003) observed that polyploidy species, such as wheat, are less prone to gene silencing. This is justified by the large buffering capacity of the inserted genes. The variation we noted between the events, regarding the different physiological parameters observed, can be explained by considering experimental and environmental factors that are not controlled and are not directly associated with the transgene (James *et al.*, 2004; Saint Pierre *et al.*, 2012). After rehydration (Day 20), proline levels decreased, which proved once again the important action of osmolyte during the water stress process (Figure 1B).

Figure 1C shows values obtained for the parameter MSI, under water stress conditions. Under normal conditions of water supply, control and transgenic plants displayed similar results (83-87%). We observed cell membrane injury due to water stress from the 12<sup>th</sup> and 16<sup>th</sup> days on after applying water restriction. The lowest values for MSI were found in plants in the control group under stress (40%) when compared to transgenic events: 66% (event 591), 83% (event 14), 78% (event 326), 76% (event 164), and 78% (event 312). After irrigation was carried out again, in the assessment of the 20<sup>th</sup> day, only the events 591, 326, and 164 presented an MSI close to the irrigated control value (83%). This suggests the maintenance of the integrity of cell membranes. Plants in the control group under water stress and the events 14 and 312 had their MSI near 68%.

MDA content estimated for control and transgenic plants in the different events was similar

before water stress started (Figure 1D). After the 12<sup>th</sup> day, transgenic plants and the stressed control group displayed decreased values of MDA, which were also different from the irrigated control group. The assessment of the 20<sup>th</sup> day, after rehydration, showed that MDA values had increased again, which can be justified by metabolism acceleration and the onset of senescence, since only plants in the event 591 presented lower and significant MDA levels when compared to the control group. That increase might have happened due to a lack of proline, as it might have moved to reproductive tissues. Therefore, plant cells (control and transgenic) would suffer more damage to oxidative stress because of water deficit (Ozden *et al.*, 2009).

Low MDA content represents low lipid peroxidation. It is, therefore, inversely proportional to the membrane stability (Bhatnagar-Mathur *et al.*, 2009). Data obtained regarding lipid peroxidation, estimated through MDA content, allowed us to conclude that transgenic plants have a better ability to preserve the plant cell intact for a longer period. We were able to prove it on the 20<sup>th</sup> day of assessment, possibly due to the high levels of proline (observed on the 16<sup>th</sup> day of evaluation) (Figures 1B and 1D). Several studies have showed that high proline levels in cells might prevent oxidative stress caused by the presence of reactive oxygen species (ROS), which is a result of water stress (Hong-Bo *et al.*, 2006; Hoque *et al.*, 2007; Ahmad *et al.*, 2008; Bhatnagar-Mathur *et al.*, 2009). This condition resulted in a higher maintenance of cellular integrity in the events we evaluated compared to the control group.

Similar to our experiment, Pavei *et al.* (2016) evaluated 16 different transgenic plants ( $T_2$ ) transformed to *p5cs* using constitutive promoter (*Ubi*) and stress induced promoter (*aipc*) subjected to 12 days of water stress. As results, transgenic plants independently of the promoter used produced approximately 1.85 times more proline than non-transgenic plants, but MDA values did not differ between transgenic plants and control. However, transgenic plants showed a better RWC at the end of evaluation period (84% – transgenic and 57%–control). We can conclude that proline is essential to copy the oxidative stress damage, perhaps the MDA measurement was not sufficient to quantify the antioxidant effect of proline on cells.



Productive indexes indicate the decrease of production in water deficit conditions (Table 1). The irrigated control group presented: a mean of 3 cobs per plant; 138 seeds per plant; 475 mg of weight per 100 seeds; and 3.3 g of seeds total weight per plant. The stressed control group showed a decrease of 33% in the number of cobs, 75% in the mean number of seeds per plant, 34% in the weight of 100 seeds (310.46 mg), and 5.5 times the total weight of seeds (0.6 g). Transgenic plants in the event 14 showed the best productive indexes compared to the stressed control group. They had an increase of 106% in the number of seeds, 1.3 times in the number of cobs, 26% in the weight of 100 seeds, and 116% in the total weight of seeds. Event 312 showed an increase of 1.5 times in the number of seeds, 51% in the number of cobs, 50% in the weight of 100 seeds, and 100% in the total weight of seeds.

Lemos *et al.* (2008) observed that the dry period applied in the booting stage of CD200126 plants resulted in a decreased weight of 51.8% of in 100 seeds and of 80% in COB sprouting. After that, severe water stress was applied for 8 days. Saint Pierre *et al.* (2012), on the other hand, stated that the decrease in plant productivity might be because plants submitted to water stress display a sharp delay in growth. When evaluating transgenic wheat's (*DREB1A*) productive performance, they observed that transgenic plants displayed a more conservative profile for plant growth when exposed to reduced water consumption but showed higher survival rates when compared to

plants in the control group, with no delay in the culture phenological cycle.

Finally, the results obtained in our study show that proline would act as an osmotic agent, maintaining the membrane integrity, which might be helpful to plants' survival. However, productivity is considered a polygenic trait in which different genes and environmental factors are involved. Obtaining drought-resistant wheat elite genotypes has been a challenge, as it is a multigenic trait highly affected by environmental conditions (Saad *et al.*, 2013).

Due to the fact that timeline for commercialization of an transgenic event is long (~14 years for the first commercial launch) (Fraley, 2015) and the investments are quite high (Rudelsheim *et al.*, 2018), biotech trait developers aim to create and identify events that optimally meets the biotech trait product goal in terms of expression, stability, safety, and utility to support commercial release (Mumm, 2013).

## CONCLUSIONS

We noticed an increase in proline levels in transgenic plants from the 12<sup>th</sup> day on, and the highest index was observed in the event 312 and in all transgenic events on the 16<sup>th</sup> day, resulting in higher stability of transgenic plants' membranes compared to the stressed control group. These results show the osmoprotectant effect of proline on cell membranes. We also observed differential phenotypical characteristics in events containing the gene *p5cs* and the *aipc* promoter in the T<sub>3</sub> generation, in different events of the 16<sup>th</sup> day. However, higher levels of proline did not result in higher productivity indexes in transgenic plants.

**Table 1** - Productive indexes obtained in transgenic events that were evaluated and submitted to a dry period

	Mean of cobs/plant	Mean of seeds/plant	Weight of 100 seeds (mg)	Total weight of seeds/plant (g)
IC	3	138.08 *	475.93	3.3 *
SC	2.04	33.8	310.46	0.6
591	2.28	36.36	447.81	0.8
14	4.64 *	69.92	391.61	1.3
326	2.08	28.68	339.62	0.7
164	2.68	44.32	412.05	0.8
312	3.08	51.08	466.06	1.2

IC – Irrigated Control, SC – Stressed Control. Values are the mean of n = 5 ± SD  
\*Significant values according to the Tukey's test (5%)

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