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Characterization of Minor Millets (Panicum sumatrense and Eleusine coracana) for Trait Related to Moisture Stress Tolerance

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Abstract

Drought stress is one of the abiotic stresses which may alter plant growth, metabolism and yield. Water stress limits the growth, productivity and quality of agricultural crops in the world. Water Stress is not only due to the scarcity of water but also due other factors such as salinity, high temperatures and severe cold that make plants not able to absorb enough water from soil to grow well and this is called physiological drought that leads to a series of disorders in physiological and biochemical processes. Millets are resilient to extreme environmental conditions especially to inadequate moisture and are rich in nutrients. The aim of this research was to analyze the effects of water stress on relative water content, proline, soluble carbohydrates and chlorophyll content of minor millet genotypes under control conditions. The photosynthetic pigments like chlorophyll a, chlorophyll b and total chlorophyll decreased and the biochemical components like, proline, leaf protein and carbohydrate increased under water stress. In this study little millet genotypes showed minimum decrease in chlorophyll content and maximum increase in proline, protein and carbohydrate content when compared to the previously reported tolerant millet genotype. This study suggested the little millet (Panicum sumatrense) genotype RLM 37 having drought tolerant adaptive mechanism and better performance under water stress.

Keywords: Minor millet, water stress, relative water content, proline, protein

1. Introduction

Plant growth and development as well as crop production are highly influenced and sometimes limited by environmental conditions, such as drought, salinity and temperature stresses. Among these, drought stress is the most important environmental constrains to world agricultural production (Bray et al., 2000). Thus, an understanding of drought stress and water use in relation to plant growth is of importance for sustainable agriculture. Water stress not only affects the morphology but also severely affects the physiological metabolism of the plant. In response to various environmental stresses, plants have developed different physiological and biochemical mechanisms to adapt or tolerate stress (Rahnama and Ebrahimzadeh, 2005; Faical et al., 2009). Osmoregulation is one of the important biochemical phenomena in plants to cope up adverse environmental conditions. Osmotic adjustment in plants subjected to drought stress occurs by the accumulation of high concentrations of osmotically active compounds known as osmolytes such as proline, glycine betaine, soluble sugars, polyamines etc., in order to lower the osmotic potential (Rontein et al., 2002; Jouve et al., 2004).

Proline accumulates in many plant species under a broad range of stress conditions such as water shortage, salinity, extreme temperatures, and high light intensity. Proline is considered to be a compatible solute. Photosynthetic pigments are important to plants mainly for harvesting light and production of reducing power such as ATP and NADPH. Both the chlorophyll a (CHLa) and (CHLb) are prone to soil drying damages. Drought stress induced changes in the ratio of CHLa and CHLb and carotenoids (Faroog et al., 2009). The chlorophyll content decreased to a significant level at higher water deficits in plants (Kiani et al., 2008). Millets are examples of less-utilized crops with adaptation to marginal lands where they can withstand various stress conditions and contribute to sustainable low-input food production. Minor millets also called small millet are a group of grassy plants with short slender culm and small grains. They are categorized as coarse cereals and mainly form staple food for the tribal people where cultivation of major cereals like rice, wheat and maize is either not popular or fail to produce substantial yield (Chopra and Neelam, 2004). In India, small millets are cultivated in the semi-arid and hilly regions inhabited by traditional farmers. Finger millet is the principal crop amidst the small

millets, occupying 60–70% of the total area under small millet production globally, followed by kodo millet, foxtail millet, little millet, proso millet, and barnyard millet. Grains of small millets are extremely resistant to storage pests and can be stored for indefinite periods (Yenagi et al., 2010). Nutrition wise, grains of small millets are rich in micronutrients, particularly calcium and iron. They have high dietary fiber content, rich essential amino acids, and low glycemic index (Devi et al., 2011; Shobana et al., 2013). Minor millet is known for its great level of tolerance against drought, salinity and diseases. Water deficit affects the development, growth and yield in plant crop, but the tolerance crops to this stress varies remarkably. Changes in morphological, physiological, biochemical and molecular aspects are generally noted in response to drought stress. Understanding these responses to drought is important for screening tolerance of genotypes to water-limited conditions. The purpose of this research was to study water stress effect on some biochemical processes of minor millet cultivars, so that responses of these cultivars can be evaluated in resistance to drought stress.

2. Materials and Methods

2.1. Plant material

The experimental materials of the present investigation comprised of minor millet were planted in pots separately and maintained in green house at 28±2 °C. Pots were watered normally (once per day) until the plants attain 21day period. The 21-day-old seedlings were subjected to water stress. Water was withdrawn from pot on 21st up to 29th days. Samples were harvested on 30th days from both control and water stressed plants in liquid nitrogen and stored at -80 °C.

2.2. Relative water content of leaf (RWC %)

RWC was measures for both stress and control plant leaves. Measurements of RWC were made between 11:00 to 13:00 hours after the dew had dried. One fully extended next to flag leave per plot was sampled in pre-weighed centrifuge tubes. Immediately after cutting at the base of lamina, leaves were sealed in pre-weighed centrifuge tubes and quickly transferred to the laboratory. Fresh weights (FW) of leaves were determined. Turgid weight (TW) were obtained after soaking leaves in distilled water in test tubes for 4 to 6 hr at room temperature (about 20 °C) and under the low light conditions of laboratory. After soaking, leaves were quickly and carefully blotted dry with tissue paper in preparation for determining turgid weight. Dry weights (DW) were obtained after oven drying the leaf samples for 48 hr at 70 °C. The RWC was calculated based on the formula suggested by Barr and Weatherley (1962) as follows:

RWC%=(FW-DW/TW-DW)×100

Where

FW=fresh weights of leaf taken immediately after excision TW=Turgid weight of leaf

DW=Dry weight of leaf drying at 70 °C for 48 h.



Proline content was estimated following the method described by Bates et al. (1973).

Express the proline content on fresh weight basis as follows:

$$μ$$
 moles per g tissue= $\frac{μg \text{ proline ml}^{-1} \times \text{ml toluene}}{115.5} \times \frac{5}{\text{sample-1}}$

2.4. Estimation of total carbohydrates

Total carbohydrates in the leaves and seeds were determined by phenol sulphuric acid method proposed by Krishnaveni et al. (1984). For the estimation, 100 mg of tissue control and stress plants was weighed. The tissue was hydrolyzed by adding 5 ml 2.5N HCl and boiling in hot water bath for three hours. After cooling, it was neutralized using solid sodium carbonate until effervescence ceases. The final volume was made to 100 ml and centrifuged. From this sample, 0.1 and 0.2 ml was pipette out in two separate test tubes. The volume of the test tube was made to 1 ml by using distilled water. To the test tubes, 1 ml phenol and 5 ml 96% H₂SO₄ was added. The tubes were kept for 10 minutes and shaken well. These test tubes were kept in hot water bath at 20-30 °C for 20 minutes. The absorbance was taken at 490 nm after cooling by using the mixture of 1 ml water, 1 ml phenol and 96% H₂SO₄.

2.5. Estimation of protein content

The total leaf protein content of eighteen rice genotypes was estimated as per the method given by Lowry et al. (1951).

2.6. Chlorophyll estimation

Chlorophyll were extracted from the leaves and estimated by the method of Arnon (1949).

Chlorophyll content was calculated using the formula of Arnon as follows:

Total chlorophyll $(mg/ml)=(0.0202)\times(A.645)+(0.00802)\times(A.663)$

Chlorophyll 'a' (mg/ml)= $(0.0127)\times(A.663)$ – $(0.00269)\times(A.645)$ Chlorophyll 'b' (mg/ml)= $(0.0229)\times(A.645)$ – $(0.00468)\times(A.663)$.

The values were expressed in milligram per gram fresh weight.

2.7. Estimation of chlorophyll stability index

The chlorophyll stability index (CSI) of eighteen rice genotypes was determined according to Sairam et al. (1997) and calculated as follows:

2.8. Statistical analysis

The OPSTAT software developed at BHU was applied for statistical analysis.

3. Results and Discussion

3.1. Effect of water stress on relative water content of leaf (RWC %)

Water stress treatment produced a significant decline in

relative water content of leaf (RWC) among seven millet genotypes. The mean RWC was recorded as 84.711% (ranged from 81.709-87.040%) at control condition and 32.502% (ranged from 20.939–49.289%) under stress condition (Table 1, 2). Among the seven genotypes, higher RWC was recorded in little millet genotype BL 4 (87.040%) and lowest RWC was observed in OLM 203 (81.709%) under control condition and higher RWC was recorded in little millet genotype RLM 37 (49.289%) and lowest RWC was observed in PR 10 14 (20.939%) under stress condition. Minimum decrease in RWC content over control were recorded in RLM 37 (1.699 fold), and GPU 67 (1.841 fold) under water stress condition. Schonfeld et al. (1988) expressed with increase of drought stress of wheat, RWC decrease and usually but not always,

Table 1: Mean and range for biochemical traits of seven millet genotypes under control condition									
Genotype	RWC Control	Proline	Carbohydrate	Protein	Chlorophyll	Chlorophyll b	Total chloro-		
		control	control	control	a Control	control	phyll control		
	Mean±S.E.	Mean±S.E.	Mean±S.E.	Mean±S.E.	Mean±S.E.	Mean±S.E.	Mean±S.E.		
RLM 37	83.743±0.000	0.029±0.007	38.016±0.299	0.011±0.001	0.916±0.002	0.435±0.003	1.350±0.006		
BL 4	87.040±0.260	0.686±0.006	53.819±0.209	0.092±0.000	0.938±0.003	0.641±0.004	1.580±0.007		
MM 10	86.928±0.470	0.591±0.006	38.359±0.238	0.095±0.001	0.856±0.002	0.417±0.003	1.273±0.005		
OLM 203	81.709±0.337	0.262±0.008	26.018±0.214	0.096±0.000	0.928±0.002	0.459±0.004	1.387±0.007		
GPU 67	84.186±0.233	0.100±0.006	18.717±0.297	0.109±0.002	2.607±0.003	1.546±0.000	4.153±0.003		
BR 36	84.309±0.507	0.063±0.007	27.492±0.305	0.010±0.001	1.379±0.002	1.176±0.003	2.555±0.004		
PR 10 14	85.065±0.047	0.539±0.008	21.802±0.178	0.271±0.002	1.087±0.002	0.435±0.004	1.522±0.006		
Mean	84.711	0.324	32.032	0.098	1.244	0.730	1.974		
Minimum	81.709	0.029	18.717	0.010	0.856	0.417	1.273		
Maximum	87.040	0.686	53.819	0.271	2.607	1.546	4.153		
CD (p=0.05)	0.980	0.022	0.775	0.005	0.007	0.010	0.017		
SEm±	0.320	0.007	0.253	0.001	0.002	0.003	0.006		
SEd±	0.452	0.010	0.358	0.002	0.003	0.005	0.008		
C.V.	0.654	3.749	1.368	2.624	0.334	0.780	0.485		

Table 2: Mea	n and range for	biochemical tra	its of seven mille	et genotypes ur	nder water stre	ss condition	
Genotype	RWC Stress	Proline	Carbohydrate	Protein	Chlorophyll	Chlorophyll b	Total chloro-
		stress	stress	stress	a stress	stress	phyll stress
	Mean±S.E.	Mean ± S.E.	Mean ± S.E.	Mean ± S.E.	Mean ± S.E.	Mean ± S.E.	Mean ± S.E.
RLM 37	49.289±0.297	0.496±0.009	97.731±0.298	0.246±0.005	0.452±0.003	0.310±0.003	0.762±0.006
BL 4	28.452±0.227	0.723±0.005	61.155±0.299	0.241±0.004	0.123±0.003	0.143±0.006	0.266±0.003
MM 10	31.673±0.375	2.083±0.006	60.537±0.240	0.320±0.002	0.229±0.002	0.173±0.004	0.401±0.006
OLM 203	21.468±0.574	0.822±0.005	39.867±0.299	0.313±0.001	0.436±0.002	0.462±0.004	0.898±0.005
GPU 67	45.739±0.286	0.249±0.007	30.029±0.297	0.257±0.003	2.309±0.001	1.233±0.003	3.540±0.004
BR 36	29.956±0.039	1.067±0.008	38.839±0.280	0.177±0.018	1.09±0.003	0.406±0.003	1.495±0.006
PR 10 14	20.939±0.306	2.719±0.005	44.358±0.240	0.352±0.003	0.751±0.003	0.306±0.004	1.057±0.007
Mean	32.502	1.166	53.217	0.272	0.770	0.433	1.203
Minimum	20.939	0.249	30.029	0.177	0.123	0.143	0.266
Maximum	49.289	2.719	97.731	0.352	2.309	1.233	3.540
CD (p=0.05)	1.027	0.021	0.858	0.022	0.007	0.012	0.017
SEm±	0.335	0.007	0.280	0.007	0.002	0.004	0.005
SEd±	0.474	0.01	0.396	0.01	0.003	0.006	0.008
C.V.	1.787	1.003	0.912	4.653	0.516	1.628	0.78

in drought stress conditions, the cultivars that are resistant to drought have more RWC. Relative water content (RWC) of leaves has been reported as direct indicator of plant water contents under water deficit conditions (Lugojan and Ciulca, 2011). Under water stress condition decrease in water status and osmotic potential in plants is the ultimate outcome of lower relative water content. Osmoregulation mechanism plays a phenomenal role in preserving turgor pressure which helps in soil water absorption and continue plant metabolic activities for its survival. In this study, little millet genotype RLM 37 showed minimum decrease in relation water content which indicates their comparable potential for drought tolerance with respect to known drought tolerant finger millet genotypes GPU 67.

3.2. Effect of drought stress on leaf proline content

Accumulation of proline as an osmolyte under water stress was observed in seven different millet genotypes. Mean leaf proline content was found to be 0.324 μ mol g⁻¹ fresh weight of leaf tissue (ranged from 0.029 μ mol g⁻¹ f.wt.- 0.686 μ mol g-1 f.wt.) at control condition, which increased to 1.166 μ mol g-1 fresh weight under water stress (ranged from 0.249 μ mol g⁻¹ f.wt.- 2.719 μ mol g⁻¹ f.wt.), (Table 1, 2). Highest increase in leaf proline content over control was recorded in little millet genotype RLM 37 (17.103 fold) followed by finger millet genotype BR 37 (16.937 fold) under water stress. In general, proline content of leaves increased with the decline in irrigation water, suggesting that the production of proline is probably a common response of millet under drought conditions. The role of proline in adaptation and survival of plants under has been well documented by Watanabe et al. (2000) and Saruhan et al. (2006). Proline is one of the most studied solutes and high proline content in plants under water stress is frequently observed in several species (Clifford et al., 1998; Bajji et al., 2001) and may act as a regulatory or signaling molecule to activate multiple responses that are part of the adaptation process (Maggio et al., 2002; Claussen, 2005). Osmotic adjustment through the accumulation of cellular solutes, such as proline, has been suggested as one of the possible means for overcoming osmotic stress caused by the loss of water (Caballero et al., 2005). Lobato et al. (2011) reported that the accumulation of proline and free amino acids in soybean (Glycine max cv. Sambaiba) leaves were increased under water deficit 67 and 388.1%, respectively. Teixeira and Pereira (2006) indicated that proline content significantly increased in all potato organs in response to the stress conditions. This increase was more remarkable in roots and tubers than in the leaves. High levels of proline enable the plant to maintain low water potentials causing the accumulation of compatible osmolytes that allows additional water to be taken up from the environment, thus buffering the immediate effect of water deficit within the organism (Mousa and Abdel-Aziz, 2008). In this study on the basis of accumulation of leaf proline content under water stress condition little millet genotype RLM 37 and finger millet

genotype BR 36 was found to be maximum compared to tolerant genotype GPU 67 hence were identified as potential drought tolerant genotypes.

3.3. Effect of drought stress on carbohydrate

The observation of carbohydrate content under control and stress condition showed that the carbohydrate increases significantly with prolongation of water stress. Mean carbohydrate content of the minor millet genotype was found to be 32.032 mg g-1 f.wt (ranged from 18.717 mg g-1 f.wt.-53.819 mg g-1 f.wt.) at control condition, which increased further to 53.217 mg g⁻¹ f.wt under water stress (ranged from 30.029 mg g⁻¹ f.wt. –97.731 mg g⁻¹ f.wt.). (Table 1, 2). Water stress induced highest increase in carbohydrate content was obtained in little millet genotype RLM 37 (2.571 fold) followed by finger millet genotype PR 10 14 (2.035 fold) when compared with tolerant genotype GPU 67. The accumulation of sugars in response to drought is quite well documented (Izanloo et al., 2008; Watanabe et al., 2000). Soluble sugars may function as a typical osmoprotectant, stabilizing cellular membranes and maintaining turgor pressure. Gene ontology attributes such as proline and soluble sugar accumulations were highly enriched in the drought-up-regulated genes, suggesting that those metabolic pathways are important in responses to drought stress. Indeed, the importance of many of these pathways to drought tolerance has been empirically supported by transgenic experiments (Umezava et al., 2006). Proline and total soluble sugar content in the leaves was significantly (p<0.05) increased due to the increase in the level of drought stress. The differences in the responses to drought stress among the nine selected corn cultivars suggested that each cultivar has different ability to synthesis proline and total soluble sugar with an increase in drought stress treatment (Hermalina et al., 2014).

3.4. Effect of drought stress on protein content

The protein content of seven millet genotypes was recorded under control and water stress conditions. The mean soluble protein content was found to be 0.098 mg g-1 f.wt. (ranged from 0.010-0.271 mg g-1 f.wt.) in control condition, which increased to 0.271 mg g⁻¹ f.wt. (ranged from 0.177–0.352 mg g⁻¹ f.wt.) under stress condition. (Table 1,2). Water stress induced highest increase in protein content was obtained in little millet genotype RLM 37 (22.36 fold) followed by finger millet genotype BR 36 (17.70 fold) when compared with tolerant genotype GPU 67. Chinoy et al. (1974) who also reported high protein content in drought stressed rice plant. Ashraf and Foolad (2005) had reported that higher protein content in tolerant genotypes under water stress condition is due to higher DNA and RNA content, which stimulate synthesis and inhibit protein decomposition. According to Ahmad et al soluble sugars and the protein content of the leaves from water stressed plants was significantly higher than the irrigated control. Comparison within each stress period showed that the protein levels were significantly higher at all stress periods than the controls. In this study little millet genotype RLM 37 was found to be maximum protein content compared to tolerant genotype GPU 67. Increases in protein suggest that under water stress conditions synthetic activity is enhanced to accommodate increased metabolic activity for maintaining the osmotic balance.

3.5. Effect of drought stress on chlorophyll content

The chlorophyll content of seven millet genotypes was recorded under control and water stress conditions. A significant decrease in chlorophyll content (chlorophyll a, b and total chlorophyll) was observed in seven millet genotypes under water stress conditions (Table 1, 2).

3.5.1. Chlorophyll a content

The mean chlorophyll a content was recorded to be 1.244 mg g⁻¹ leaf tissue. It ranged from 0.856 mg g⁻¹ in little millet genotype MM 10 to 2.607 mg g-1 leaf tissue in finger millet genotype GPU 67 under control condition. While mean chlorophyll content under stress was found to be 0.770 mg g⁻¹ leaf tissue which ranged from 0.123 in little millet genotype BL 4 to 2.309 mg g⁻¹ leaf tissue in finger millet genotype GPU 67. (Table 1, 2). Minimum fold decrease in chlorophyll a content was recorded in finger millet genotype GPU 67 (1.129 Fold) followed by BR 36 (1.265 fold).

3.5.2. Chlorophyll b content

The mean chlorophyll b content was recorded to be 0.730 mg g⁻¹ leaf tissue. It ranged from 0.417 in little millet genotype MM 10 to 1.546 mg g⁻¹ leaf tissue in finger millet genotype GPU 67 under control condition. Under stress, the mean chlorophyll b content was found to be 0.433 mg g⁻¹ leaf tissue. This was in the range of 0.143 in little millet genotype BL 4 to 1.223 mg g⁻¹ leaf tissue in finger millet genotype GPU 67 under water stress condition (Table 1, 2). Out of seven genotypes under study, minimum decrease was recorded in little millet genotype OLM 203 (0.994 fold) followed by finger millet genotype GPU 67 (1.254 fold).

3.5.3. Total chlorophyll content

At control condition the mean total chlorophyll content was found to be 1.974 mg g⁻¹ leaf tissue. It ranged from 1.273 (MM 10) to 4.153 mg g⁻¹ leaf tissue (GPU 67) which decreased to 1.203 mg g⁻¹ leaf tissue (ranged from 0.266–3.540 mg g⁻¹ leaf tissue) under water stress condition (Table 1, 2). Out of seven genotypes, minimum decrease were recorded in finger millet GPU 67 (1.173 fold) followed by PR 10 14 (1.440 fold) and BR 36 (1.709 fold) hence were considered as potential drought tolerant genotypes. The chlorophyll was declined from well watered (control) condition to severe drought stress (13 DID) and plant pigment was increased by the age of maturity in plant (Paul et al., 2013). Manirannan et al. (2007) detected a depression in CHL a and b and TC in Helianthus annuus L. under water stress. According to Poljakoff and Gale (1975), the ability to synthesize more chlorophyll under water stress is a good criterion for the species tolerant to drought. In this

study finger millet genotype showed minimum decrease in chlorophyll content that suggested their tolerance mechanism under water stress.

3.6. Effect of drought stress on chlorophyll stability index (CSI %)

Water stress induced decreasing trend in chlorophyll stability index (%) was observed in seven millet genotypes. A significant difference (p<0.05) for chlorophyll stability index (CSI) was observed among seven millet genotypes under water stress condition. The mean chlorophyll stability index was found to be 54.686% (ranged from 16.840-85.258%) under water stress. Among seven millet genotypes, lower decrease in CSI (%) was obtained in BL 4 (16.840%). The higher CSI% obtained in tolerant finger millet genotype GPU 67 followed by PR 10 14 (69.453%) indicated enhanced ability of identified millet genotypes to withstand water stress condition.

4. Conclusion

Plants in drought stress time make changes in some of their physiological and biochemical features. Accumulations of soluble carbohydrates, proline and protein increased under water stress and the genotypes that have more RWC and chlorophyll content are more resistant to drought stress. Our results represented that the little millet genotype RLM 37 showed minimum decrease in RWC and maximum increase in proline, carbohydrate and protein which indicates their comparable potential for drought tolerance with respect to tolerant genotype.

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