

## Role of abscisic acid in regulating the expression of EcMyb gene for drought stress tolerance in *Eleusine coracana*

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ORIGINAL RESEARCH ARTICLE

### ABSTRACT

Abscisic acid (ABA) plays many important roles in plants, including seed germination, dormancy, stomata closure, and adaptation to water stress. ABA has been implicated in responses to environmental stresses such as drought and salinity. Drought is a major abiotic factor that limits agricultural crop productivity. The study was performed to investigate the effect of drought on expression of Ecmyb gene on drought tolerant (PRM 6107) and sensitive genotypes (PES 400) of *Eleusine coracana* in seedling stage and investigate whether it followed the ABA dependent pathway. Different experiments were conducted wherein drought was induced by treatment with PEG (polyethylene glycol) and ABA (abscisic acid) was applied in three different concentrations i.e. 25, 50 and 100  $\mu$ M. Expression of EcMyb gene was observed only in tolerant variety PRM 6107 under drought stress. It could be thought that the expression of EcMyb transcription factor gene may be upregulating the expression of many drought responsive genes which is providing drought tolerance to PRM 6107. In response to ABA, EcMyb gene was expressed in both the genotypes. However, expression was strong and early i.e. after 6 h of the treatment in PRM 6107 than in PES 400. As drought is a complex phenomenon, therefore, to understand how ABA accumulates in response to drought in both tolerant and sensitive varieties, primers were designed for NCED (9-cis epoxy carotenoid dioxygenase) gene, an important gene in ABA synthesis pathway and co-expression of NCED and EcMyb gene was recorded. The co-expression of NCED gene and EcMyb gene was observed only in PRM 6107 in drought conditions since first day whereas in PES 400 a basal level expression of NCED gene was observed after 5<sup>th</sup> day. There was no expression of EcMyb gene. It was concluded that EcMyb gene expressed only in that genotype which can synthesise a certain level of ABA. Morphological and physiological parameters were also recorded in both the genotypes after drought and ABA treatment. Root length increased, whereas shoot length, relative water content (RWC) and chlorophyll content decreased with exposure of drought as well as ABA treatment. The study suggests that EcMyb gene expresses only when ABA accumulates in plant cell and may follow ABA dependent signalling pathway.

### KEYWORDS

abscisic acid; drought; EcMyb gene; *Eleusine coracana*; NCED gene

## 1. INTRODUCTION

Global warming possesses a serious threat to contemporary agriculture. It is speculated that average global temperature will rise by 1 °C till 2050 and by 2-4 °C by 2100 (Nelson and Gerald, 2010). This increase in temperature may adversely affect agricultural production and productivity. Conditions that adversely

affect the crop growth and yield is called as stress. Stress may be biotic or abiotic. Abiotic stress such as drought and salinity causes loss up to 50 % of the potential yield in major crop plants. On the other hand the world population is estimated to increase from 6.8 billion in 2011 to 9.3 billion by 2050 (United Nations, 2011). So it is a great challenge to feed the growing population with shrinking agricultural land. In such

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situation, it is imperative to search for an alternative source either in the form of modified cultural practices or genetic modifications per se of crops that could better adapt to such stress situations and bridge the gap between demand and supply of food grains. C4 plants like millets, maize, sorghum perform well under stress condition. In general, abiotic stress often causes a series of morphological, physiological, biochemical and molecular changes that affect plant growth, development and productivity. Drought, salinity, extreme temperatures (cold and heat) and oxidative stress are often interrelated. These conditions either alone or in combination induce cellular damage. Abiotic stress such as drought leads to accumulation of ABA endogenously. Accumulation of ABA triggers ABA dependent signalling pathway. Accumulation of ABA causes up regulation and down regulation of ABA responsive gene. This up regulation and down regulation occurs due to a number of factors; among these transcription factors (TFs) and RNA binding proteins are of great importance.

Finger millet (*Eleusine coracana*) is one such crop which is well known for its hardiness and cultivation on marginal land deprived of inputs like irrigation. Therefore it can be used for fishing genes responsible for drought tolerance. Once such genes are identified, these can be isolated and integrated into the genome of other crops like wheat and rice using recombinant DNA technology to tailor abiotic stress resilient crop. Therefore, the molecular exploration of finger millet, an underutilized but drought tolerant and nutritionally rich crop would hold a high promise in the present global warming scenario.

Drought responsive Myb proteins are TFs (Transcription factors) composed of one, two, or three imperfect helix-turn helix repeats that recognize the major groove of DNA and are thought to play an important role in imparting drought tolerance to plants (Yanhui et al., 2006). Therefore, investigating the pathway underlying drought tolerance in plants is important and finger millet can be taken as a model plant for this purpose. EcMyb, homologue of a drought responsive Myb gene has been shown to express in *Eleusine coracana* and its expression increased with the severity of drought (Salvi et al., 2012). The present study was carried out to investigate the expression pathway of EcMyb gene in seedling stage of *Eleusine coracana* under drought stress and to determine whether it followed the ABA dependent pathway.

## 2. MATERIALS AND METHODS

### 2.1. Plant material

Two contrasting genotypes of finger millet viz. PES-400 & PRM-6107 were used. The former was drought susceptible and the latter was drought tolerant (Rai et al., 2009).

Seeds were sown in pots filled with garden soil, peat moss and vermicompost in 3:1:1 ratio and plants were grown in a poly house. Uniform germinated seedlings were transplanted into plastic foam frames suspended in a plastic tray containing an aerated nutrient solution (Sonoda et al., 2003). The pH of the solution was maintained at 5.8. The seedlings were grown in nutrient solution for 4 days before subjecting them to experimental conditions.

### 2.2. ABA treatment

Both the contrasting genotypes were subjected to ABA treatments of different concentrations (25, 50 and 100  $\mu\text{M}$ ) by mixing with the nutrient solution. The pH of the ABA solution was adjusted to 6.5 by adding HCl or NaOH. The treatment was given through roots. No ABA treatment was given to control.

### 2.3. PEG treatment

In a separate experiment, drought was imposed to the seedlings using PEG (15 % polyethylene glycol having a molecular weight 8000 MW) in the nutrient solution. Plantlets were subjected to drought by immersing the roots of both the genotypes in the nutrient solution containing PEG. Control was not subjected to PEG treatment.

### 2.4. Primer designing

Primers were designed against two genes, EcMyb and NCED. The target specific internal primer was designed for EcMyb1 using NCBI data base (accession no. JN107890.1) and online PRIMER3 tool. Since no information was available regarding NCED genes of finger millets at NCBI, homologous sequences from other species like *Zea mays*, *Oryza sativa*, *Sorghum bicolor*, *Setaria italica*, *Triticum aestivum* and *Hordeum vulgare* present in NCBI data base were downloaded. These sequences were aligned using on-line bioinformatics tool CLUSTALW. Based on homology and conserved region, degenerate primers

were designed. Tubulin primers were used as control. Primers were custom synthesised by IDT (Integrated DNA Technologies).

### **2.5. RNA isolation and semi quantitative expression analysis**

An RNase free environment was created prior to RNA isolation. Tips and tubes of different sizes were thoroughly treated with 0.1% DEPC (Diethyl pyrocarbonate) water and then completely dried in oven and autoclaved. Aluminium foil and tissue paper to be used were also autoclaved. The pestle and mortar were dipped in 0.1% DEPC water overnight, wiped with chloroform, autoclaved and subsequently kept at 0 °C to cool it before grinding the samples. Workbench was also cleaned with DEPC water.

In order to study the expression of genes (EcMyb and NCED), total RNA was isolated from leaves using RNA isolation solution (Invitrogen) and digested with RNase-free DNase I to eliminate genomic DNA contamination of the desired gene. Isolated and purified RNA was quantified by nano-drop instrument. From quantified RNA, 2 µg RNA was reverse transcribed using cDNA synthesis kit of Fermentas according to the manufacturer's instructions. The transcript level of each gene was measured using semi quantitative PCR in a total volume of 25 µL reactions. The amplification program was as follows: Initial denaturation at 94 °C for 3 min, and then 32 cycle of sequential step at 94 °C for 1 min, 55 °C for 30 sec, 72 °C for 30 sec and the last reaction cycle was at 72 °C for 7 min.

The expression levels of EcMyb gene and NCED gene were analysed by densitometry analysis on gel using gel doc system known as Semi quantitative expression analysis.

## **3. RESULTS AND DISCUSSION**

In PEG treated genotype PRM 6107, the expression of EcMyb gene was observed and it increased with increase in duration of exposure of PEG. In other words, the expression of Ecmyb gene increased in drought tolerant genotype with increase in severity of drought. There was no expression of EcMyb gene in drought sensitive genotype PES on exposure of PEG. PEG was applied to mimic the drought stress and as PRM 6107 is drought tolerant genotype so its tolerant phenotype was positively correlated with the expression of EcMyb while PES 400, a drought sensitive genotype failed to

accumulate an optimum level of EcMyb gene product to combat drought.

In order to investigate their ABA dependency, the expression of EcMyb gene was observed in response to ABA under unstressed condition and it was found that the gene was expressed in both the genotypes even without any drought stress. The expression of EcMyb gene was found to increase with increase in concentration as well as duration of exposure to ABA in both the genotypes. It indicates that expression of EcMyb gene is ABA dependent and it may be possible that in sensitive genotype ABA did not accumulate to such a level so as to induce the expression EcMyb gene.

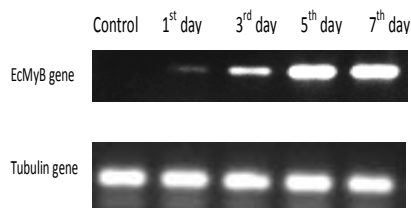
The expression of genes encoding the first two enzymes of the ABA biosynthetic pathway, zeaxanthin epoxidase (ZEP) and 9-cis-epoxycarotenoid dioxygenase (NCED), is significantly induced by stress conditions in order to increase drought tolerance (Tan et al., 2003; Moumeni et al., 2011; Ye et al., 2011). The expression of ZEP, MCSU and AAO3 in Arabidopsis (Xiong et al., 2002) and of AhNCED1 in peanut (Guo et al., 2009) was significantly up-regulated by exogenous ABA.

The expression analysis of NCED gene showed that there was a slight increase in expression from 1<sup>st</sup> to 7<sup>th</sup> day in drought tolerant genotype, PRM 6107 after PEG treatment while, only slight expression was observed in drought sensitive genotype, PES 400 after 5<sup>th</sup> and 7<sup>th</sup> day. It means that ABA accumulation in sensitive genotype was less than tolerant genotype, which may not be sufficient to increase the expression of EcMyb gene. Co-expression of NCED gene and EcMyb gene in PRM 6107 indicates that EcMyb gene which is drought responsive is expressed when optimum amount of ABA accumulates in the plant.

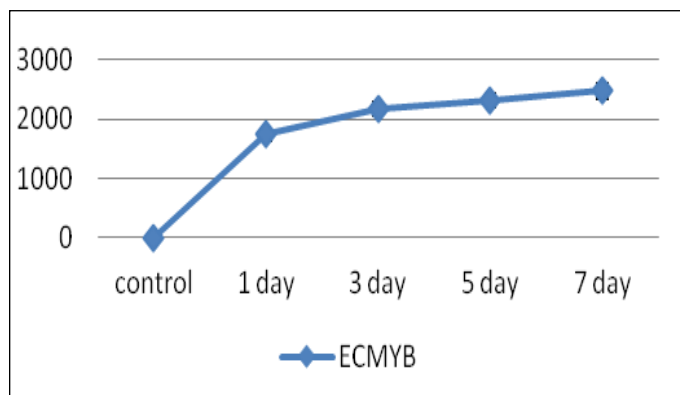
### **3.1. Effect of PEG treatment on EcMyb and NCED gene expression**

A gradual increment in the expression level of EcMyb was recorded in PRM 6107 right from the first day of PEG treatment that created a physiological drought condition in the plant. The expression reached its plateau at 5<sup>th</sup> day and was constant up to 7<sup>th</sup> day of PEG treatment (Figures 1 and 2). However no expression of EcMyb gene was observed in PES 400 under the same experimental condition. To further investigate whether regulation of EcMyb is ABA dependent or not, the level of NCED transcript accumulation was tracked which indicated the endogenous ABA accumulation. The expression of NCED gene increased from 1<sup>st</sup> day to 7<sup>th</sup> day in the tolerant genotype. In sensitive genotype

expression of NCED was observed at 5<sup>th</sup> and 7<sup>th</sup> days of PEG treatment (Figures 3 and 4). The expression of NCED and EcMyb gene was positively co-related with each other (Figure 5). This result indicates that accumulation of EcMyb gene product is ABA dependent.



**Figure 1.** EcMyb gene expression after PEG treatment in PRM 6107 genotype.



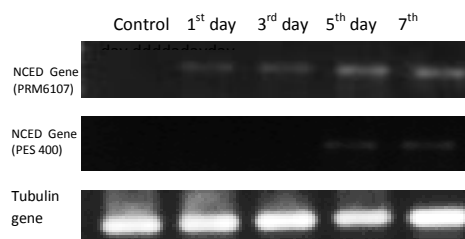
**Figure 2.** Graphical representation of EcMyb gene expression after densitometry analysis.

### 3.2. Effect of ABA treatment on EcMyb and NCED gene expression

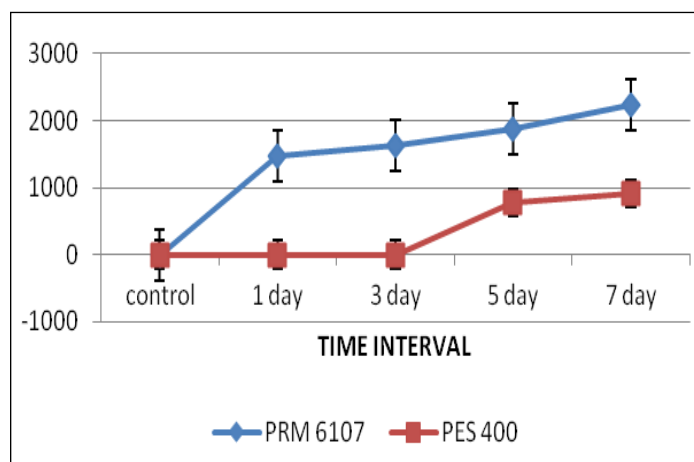
The results (Figures 1, 2, 3 and 4) showed that the expression of EcMyb gene took place only in PRM6107 genotype which also showed optimum expression of NCED gene. Therefore, it could be thought expression of EcMyb gene follow ABA dependent signalling pathway. However, it is not conclusive in nature as drought results in manifestation of a concerted set of reactions within plant system and genes other than NCED might have contributed in endogenous accumulation of EcMyb. Co-expression based assay is not sufficient to confirm such results. Therefore second set of experiment was designed where tolerant and susceptible genotypes were treated exogenously with ABA under optimal moisture condition (unstressed condition).

Plantlets subjected to varying concentration (25, 50 and 100  $\mu$ M) of exogenous ABA under unstressed condition showed the expression of EcMyb gene in

both the tolerant as well as sensitive genotypes. The expression of EcMyb was observed in drought tolerant genotype PRM 6107 for all the concentrations of ABA at an early stage ie 6 h after ABA treatment (Figures 6 and 7). However the expression of the same was deferred until 12 h of treatment imposition with low level of expression at 25  $\mu$ M and 50  $\mu$ M concentration of exogenous ABA treatment in PES 400 (Figure: 8 and 9). However, a strong and quick (6 h) expression was reported at higher dose (100  $\mu$ M) of ABA treatment in PES 400.



**Figure 3.** NCED gene expression after PEG treatment in PRM 6107 and PES 400 genotype.



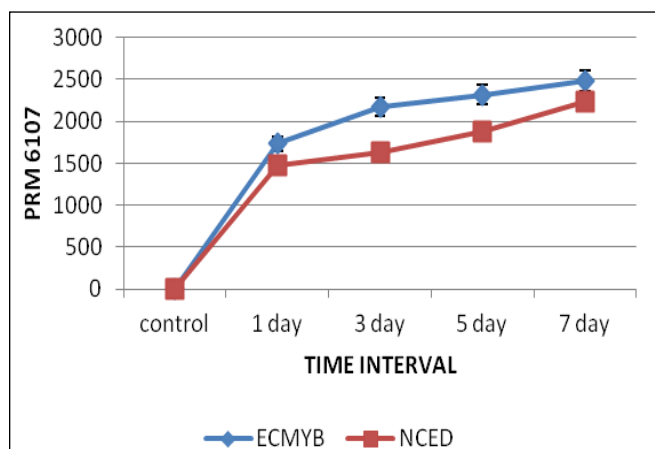
**Figure 4.** NCED gene expression after PEG treatment in PRM 6107 genotype.

NCED (9-cis epoxy carotenoid dioxygenase) is a key gene in ABA biosynthesis pathway and expressed towards the early stages of the pathway. To increase drought tolerance, the expression of genes encoding the first two enzymes of the ABA biosynthetic pathway i.e. zeaxanthin epoxidase (ZEP) and 9-cis epoxy carotenoid dioxygenase (NCED) is significantly induced by stress conditions (Tan et al., 2003; Moumeni et al., 2011; Ye et al., 2011).

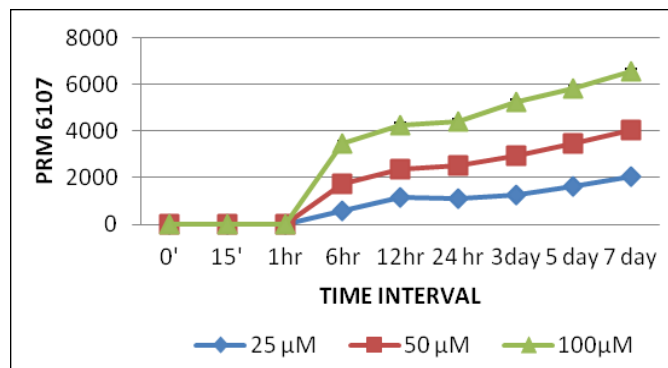
In response to exogenous ABA, no expression of NCED was found in any genotype at any concentration and duration of exposure. This might have taken place because of feedback inhibition. We applied ABA under

unstressed condition; therefore only basal level of expression was observed which was not sufficient to be tracked under gel. Also the exogenous ABA which was provided in nutrient media through root might have led to suppression of endogenous NCED gene. Similar results were reported for NCED gene expression in response to exogenous ABA in tomato (Thompson et al., 2000) and cowpea plants (Iuchi et al., 2000). However response of ABA biosynthetic genes to exogenous ABA may vary among plant species (Xiong and Zhu, 2003).

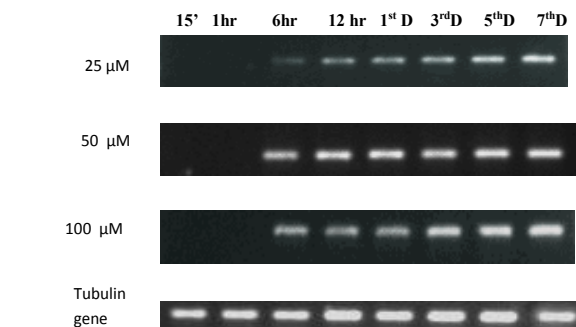
by ethylene accumulation via ABA. Arabidopsis root growth was found to be inhibited by ABA due to ethylene accumulation (Xingju et al., 2014).



**Figure 5.** Correlation between expression of NCED and EcMyb gene in PRM 6107.



**Figure 7.** Densitometric analysis for EcMyb gene expression in PRM 6107 under different concentration of ABA.



**Figure 6.** EcMyb gene expression in PRM 6107 under different concentration of ABA.

**Table 1.** Effect of drought and ABA treatment on root length of seedlings of *Eleusine coracana*.

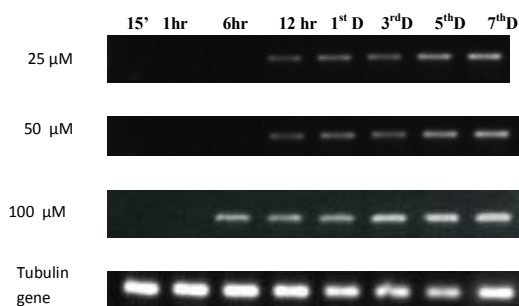
		Variety	0 day	1 <sup>st</sup> day	3 <sup>rd</sup> day	5 <sup>th</sup> day	7 <sup>th</sup> day
PEG		PRM 6107	7.5	7.5	8.2	8.2	8.3
		control	6.5	7.3	7.7	8	8.9
		PES 400	7	7.3	7.3	7.3	6.2
		control	7.5	7.8	8	8.3	8.5
ABA	25 μM	PRM 6107	18	18	18.2	18.3	18.5
		PES 400	19.5	19.5	19.6	19.7	19.9
	50 μM	PRM 6107	20.5	20.5	21	21	21.3
		PES 400	23.1	23.1	23.3	23.5	23.6
100 μM	PRM 6107	19.5	19.5	20	20.5	21	
	PES 400	14.5	14.5	15	15.2	15.4	

### 3.3. Root length

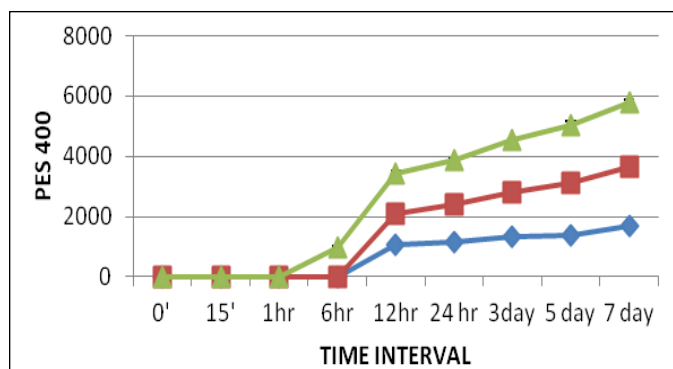
Drought is a complex trait that can be avoided by alteration in morphological and physiological characters. Root and shoot length (root: shoot ratio) plays an important role in avoiding drought stress (Hoogenboom et al., 1987; Guoxiong et al., 2002). Root plays an important role in drought avoidance by uptake of water from soil. Therefore plant genotypes with higher root growth have higher ability to avoid drought (Leishman and Westoby, 1994). An increased level of osmotic stress led to higher root to shoot ratio (Bajji et al., 2000). Root and shoot growth is regulated

In the present study, increase in root length in both the drought tolerant and sensitive genotypes was normal in controlled condition. In PRM 6107, root length increased up to 5<sup>th</sup> day of exposure to PEG and after that it did not increase further (Table 1). In PES 400, root length was maintained up to 3<sup>rd</sup> day of PEG treatment after which it also did not increase. This might be possibly due to wilt induced by PEG (8000) as the experiment was performed in hydroponic condition and PEG created physiological drought. Increase in root length up to 5<sup>th</sup> day might be due to the avoidance mechanism of the plant after which the accumulation of stress component overtook

the avoidance mechanism resulting in no increase in root length after 5<sup>th</sup> day. PEG induces lignification of roots which causes wilting of roots and hence leads to reduced root length (Hongtao et al., 2014).



**Figure 8.** EcMyb gene expression in PES 400 under different concentration of ABA.



**Figure 9.** Densitometric analysis for EcMyb gene expression in PES 400 under different concentration of ABA.

**Table 2.** Effect of drought and ABA treatment on shoot length of seedlings of *Eleusine coracana*.

		Variety	0 day	1 <sup>ST</sup> day	3 <sup>rd</sup> day	5 <sup>th</sup> day	7 <sup>th</sup> day
PEG		PRM 6107	13.3	13.5	13.0	13.0	13.0
		control	15	15.2	15.5	16	16.3
		PES 400	15	15.0	15.0	14.9	14.9
		control	14	14.1	15	15.4	16
ABA	25 μM	PRM 6107	20	20	20.2	20	19.9
		PES 400	21.5	21.5	21.6	21.5	21.5
	50 μM	PRM 6107	20.5	20.6	20.7	20.7	20.6
		PES 400	19.5	19.5	19.5	19.5	19.5
	100 μM	PRM 6107	19.5	19.5	19.8	19.7	19.6
		PES 400	21.3	21.3	21.4	21.3	21.1

Both the genotypes had similar variation in root length at various concentration of ABA applied under unstressed condition and no adverse effect on root length was recorded. Similar results were reported in wheat and maize (Moosavi et al., 2005). However the correlation of ABA concentration and root length varied in different studies. It was found to increase with increase in concentration of ABA applied in maize (Mulkey et al., 1983; Pilet and Saugy, 1987) whereas a reverse trend was reported in wheat (Jones et al., 1987) and maize (Pilet and Chanson, 1981). Effect of ABA on soybean seedling was found to be similar to drought (Creelman et al., 1990).

### 3.4. Shoot length

Shoot length for both genotypes was found to decrease in response to the PEG treatment since the first day. However, the decrease was more prominent for the sensitive one (Table 2). Treatment with ABA led to a decrease in overall shoot length in comparison to control in both the genotypes at all the concentrations, but this decrease was noticed only after the 3<sup>rd</sup> day of treatment. However, shoot length decreased in response to both the treatments, but treatment with PEG caused loss of turgidity in the whole plant while exposure to different concentrations of ABA led to wilting of the tip only. Wilting in the case of PEG treatment occurred due to loss of turgidity as seedlings failed to take water from the solution in the presence of PEG. Similar results were reported in tomato where ABA was applied exogenously at the seedling stage (Basak et al., 2012). In another study, it was reported that ABA accumulated during water stress inhibits shoot growth to reduce water loss (Saab et al., 1990; Sharp and LeNoble, 2002). However, some studies showed a positive correlation between ABA applied through soil and shoot growth as ABA restricts the ethylene production responsible for shoot growth inhibition (Abeles et al., 1992; Hussain et al., 2000).

### 3.5. Relative water content

Relative water content (RWC) was found to decrease in both the genotypes upon drought treatment; however, this decrease was more prominent for the sensitive genotype than the tolerant one. Drought is sensed by the roots in the soil and ABA gets accumulated in the root from where it is transported to the leaf through xylem (Davies and Zhang, 1991) where it regulates stomatal conductance so that water deficit in the shoot region can be avoided by transpiration. Therefore, endogenous accumulation

of ABA initially plays main role in avoiding water stress. Simultaneously water in nutrient solution is also limited to be taken up by the plant because of PEG which ultimately leads to low water availability in leaves. Production of osmolytes like proline is also induced under drought which reduces relative water content of plant under drought.

**Table 3.** Effect of drought and ABA treatment on relative water content of seedlings of *Eleusine coracana*

		Variety	0 day	1 <sup>st</sup> day	3 <sup>rd</sup> day	5 <sup>th</sup> day	7 <sup>th</sup> day
PEG		PRM 6107	101.5	97.2	90.5	79.3	65.1
		control	97.1	97.6	98.1	98.2	98.9
		PES 400	97.2	91.0	82.2	72.1	53.4
		control	95.0	95.3	95.8	95.5	95.6
ABA	25 $\mu$ M	PRM 6107	95.0	95.4	98.8	103.9	84.3
		PES 400	89.9	90.3	97.0	99.4	86.7
	50 $\mu$ M	PRM 6107	99.3	100.5	102.2	107.6	79.1
		PES 400	94.9	95.0	96.5	98.3	88.6
	100 $\mu$ M	PRM 6107	87.9	90.2	99.0	104.6	66.9
		PES 400	82.8	85.0	98.0	99.5	93.6

Relative water content (RWC) of ABA treated plants increased initially slightly but with increasing duration of exposure after 5<sup>th</sup> day of treatment it decreased drastically (Table 3). ABA might have caused closure of stomata and thus reduced transpirational loss of water initially resulting in an increase in relative water content (Hussain et al., 2012). Later on the reduction of RWC might occur due to reduced water transport through the roots to the leaves due to presence of ABA in nutrient solution (Manuel and Marc, 2011). Similar results were reported by (Xing et al. 2002). This negative correlation of ABA concentration and relative water content showed that it mimics the water stress condition.

On the other hand RWC may increase in response to ABA as there are reports that ABA transiently increases the opening of water channels like Aquaporins (Hose et al., 2000; Mahdieh and Mostajeran, 2009). But the expression is only transient and prolonged exposure and high concentration of ABA limits the root conductance which reduces the water

uptake (Fiscus, 1981). One of the possible reasons for temporary increase in RWC may be increased water uptake through its impact on aquaporins and then limitation by morphological changes to the roots i.e. suberization as a means of protecting the plant from drought stress. RWC of tolerant genotype was found to be higher than sensitive one but the reduction in tolerant genotype was also found to be higher and RWC of tolerant genotype at 7<sup>th</sup> day was lower than that of sensitive one. The possible reason behind this may be relatively higher sensitivity of tolerant genotype to ABA.

### 3.6. Chlorophyll content

Chlorophyll is a light absorbing pigment that plays a critical role in photosynthesis and therefore is an important component to be considered during stress, especially drought. Chlorophyll content was found to decrease with increased exposure to PEG in both the genotype (Table 4). However the extent of decrease in chlorophyll was more in sensitive genotype. Water deficit may trigger the production of reactive oxygen species (ROS) which in turn may lead to lipid peroxidation and consequently reduction in chlorophyll content (Kumar et al., 2011).

**Table 4.** Effect of drought and ABA treatment on chlorophyll content of seedlings of *Eleusine coracana*.

		Variety	0 day	1 <sup>st</sup> day	3 <sup>rd</sup> day	5 <sup>th</sup> day	7 <sup>th</sup> day
PEG		PRM 6107	0.217	0.207	0.193	0.185	0.177
		control	0.198	0.198	0.198	0.200	0.201
		PES 400	0.182	0.180	0.139	0.108	0.054
		control	0.272	0.274	0.278	0.279	0.301
ABA	25 $\mu$ M	PRM 6107	0.151	0.156	0.161	0.162	0.151
		PES 400	0.130	0.131	0.146	0.146	0.140
	50 $\mu$ M	PRM 6107	0.145	0.146	0.158	0.161	0.157
		PES 400	0.136	0.136	0.148	0.148	0.147
	100 $\mu$ M	PRM 6107	0.160	0.161	0.198	0.193	0.190
		PES 400	0.152	0.152	0.148	0.147	0.146

Cytokinin is a phytohormone that plays role in chlorophyll accumulation and keeps plant green and ABA being antagonist of cytokinin (Kulaeva, 1982) may interfere with chlorophyll accumulation on prolonged exposure (Fletcher and McCullach, 1971; Fletcher et al., 1973; Lew and Tsuji, 1982; Reiss and Beale, 1995). This may cause chlorosis and leaf abscission when high concentrations of ABA are applied (Blanchard et al., 2007; Kim and Iersel, 2011; Waterland et al., 2010). ABA drenches may induce wilting, even if the plants are well watered (Barrett et al., 2009).

## 4. CONCLUSIONS

ABA plays an important role in regulating drought response by inducing drought responsive genes like EcMyb whose product accumulate under drought condition in tolerant genotype PRM 6107 whereas no expression was observed in sensitive genotype PES 400. This gene was expressed in both the genotypes in presence of ABA which indicates that ABA regulates the expression of EcMyb in both the genotypes. Thus, the expression of EcMyb gene followed ABA dependent signalling pathway under drought condition. Difference in the responses of two genotypes under drought condition occurred due to differential accumulation of ABA. In sensitive cultivar, lower accumulation of ABA occurred which was basal and not sufficient to induce stress tolerance response under drought condition. Therefore, ABA may act as biochemical marker for varietal characterisation of tolerant and resistant genotypes under drought condition and only cloning of EcMyb gene is not sufficient to obtain drought tolerance traits. Sometimes gene may be present but due to lack of proper signalling, cloned gene may not be expressed. In future, for transgenic approach involving transfer of EcMyb gene from underutilised crop like finger millet to cereals grain like rice, wheat and maize, we will have to focus on cloning and also over expression of ABA for proper expression of stress related EcMyb gene.

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