Identification of Discriminant Factors after Exposure of Maize and Common Bean Plantlets to Abiotic Stresses

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Abstract

Adverse environmental conditions limit crop yield and better understanding of plant response to stress will assist the development of more tolerant cultivars. Maize and common bean plantlets were evaluated under salinity, high temperature, drought and waterlogging conditions to identify biochemical markers which could be useful for rapid identification of putative stress tolerant plants. The levels of phenolics (free, cell wall-linked, total), aldehydes including malondialdehyde and chlorophylls (a, b, total) were measured on stressed plantlets. Only two indicators were statistically non-significant: chlorophyll b in maize plantlets stressed with sodium chloride and malondialdehyde content in drought stressed maize. The most remarkable effects of abiotic stresses can be summarized as follows: (i) salinity increased levels of free phenolics in maize plantlets and chlorophylls (a, b, total) in common bean; (ii) high temperature (40 °C) elevated levels of chlorophylls (a, b, total) in maize but decreased chlorophylls (a, b, total) and free phenolics in common bean; (iii) drought increased phenolics and decreased chlorophylls (a, b, total) in maize and increased chlorophyll pigments (a, b, total) in common bean; (iv) waterlogging increased free phenolics and decreased chlorophylls (a, b, total) in maize and increased chlorophyll (a, total) in common bean. Free phenolics and chlorophylls, especially a, were the most responsive indicators to stress and can, therefore, be considered putative biochemical markers for abiotic stress tolerance in maize and common bean. The use of Fisher’s linear discriminant analysis to differentiate non-stressed and stressed plants in breeding programs is also a novel aspect of this report. Fisher’s linear discriminant functions classified correctly 100% of non-stressed or stressed originally grouped plants.

Keywords: abiotic stress, biochemical markers, genetic improvement, Phaseolus vulgaris, Zea mays

Introduction

Adverse environmental conditions, such as salinity, high temperature, drought and waterlogging, limit the geographical distribution of plant species and crop yield (Osmond et al., 1987). Predicted climatic change, population growth and the importance of sustainable food production makes the development of stress tolerant crop cultivars a high-priority globally (Zhu, 2001). Maize is the second most important agricultural crop globally. It is a human and livestock food and also used in the processing of industrial goods (Qing et al., 2009). Global maize production in 2011 exceeded 700 million tons (FAOSTAT, 2013). However, legumes also play a critical role in human and animal diets and contribute to sustainability by maintaining soil fertility (Tilman et al., 2002). The protein content of grain legumes can be three times that of cereal grains, thus a significant proportion of human protein and nutritional requirements can be supplied by legumes (Gepts et al., 2005). Common bean (Phaseolus vulgaris L.), one of the world’s most important grain legumes, is consumed as a dietary staple worldwide, particularly in Latin America and Africa (FAOSTAT, 2013).

Efficient and effective genetic improvement of stress tolerance of crops such as maize and common bean requires easy to measure markers that have a higher heritability than...
the targeted abiotic stress trait (William et al., 2007). A number of biochemical markers have been reported for abiotic stresses. For example, salinity is associated with increases in abscisic acid (Shaﬁ et al., 2011), proline (Benhassaini et al., 2012), glycine-betaine (Quan et al., 2004), polyols, sugar alcohols and soluble sugar concentrations (Gurmani et al., 2007). Salinity stress also decreases plant growth (Munn, 2005), nutrient uptake (Abdelgadir et al., 2005), K+ : Na+ ratio (Diaz-Lopez et al., 2012a), stomatal aperture and density (Huang et al., 2009), hexoses, sucrose and starch (Arbona et al., 2005) and chlorophyll contents (Rivelli et al., 2012).

Moreover, high temperature stress is associated with increased lipid peroxidation (Silva et al., 2010b) and decreased photosynthesis (Ribeiro et al., 2009), CO2 : O2 ratio in chloroplasts (Foyer and Noctor, 2000) and stomatal aperture (Ribeiro et al., 2004). Whereas drought stress is linked with increased abscisic acid (Gurmadi et al., 2007), myo-inositol (Diaz-Lopez et al., 2012b) and glycine-betaine levels (Quan et al., 2004); and decreased CO2 assimilation (Gindaba et al., 2004), relative water content (Galle et al., 2007), leaf turgor pressure (Schachtman and Goodger, 2008), osmotic potential (Silva et al., 2010a), starch content (Chao et al., 2006) and sugars and oligosaccharides (Anderson and Kohorn, 2001). Likewise, waterlogging is associated with increased free amino acids (Medina et al., 2009), abscisic acid (Xu et al., 2007), and Na+ and Cl− concentrations (Wetson and Flowers, 2010), and decreased total biomass (Colmer and Voeseke, 2009), relative growth rate (Mielke et al., 2003), stomatal conductance and photosynthesis (Lopez and Kursar, 2003), CO2 assimilation (Gimeno et al., 2012), soluble sugars and starch concentration (Gimeno et al., 2012).

In a study of maize under salinity stress, Omoto et al. (2012) found that the activities of pyruvate orthophosphate dikinase, phosphoenolpyruvate carboxylase, NADP-dependent malate dehydrogenase and NAD-dependent malate dehydrogenase, which are derived mainly from mesophyll cells, increased, whereas those of NADP-malic enzyme and ribulose-1,5-biphosphate carboxylase / oxygenase, which are derived mainly from bundle sheath cells, decreased. In salt-treated plants, the photosynthetic metabolites malate, pyruvate and starch decreased by 40, 89 and 81%, respectively. Gas-exchange analysis revealed that the net photosynthetic rate, the transpiration rate, stomatal conductance and the intercellular CO2 concentration decreased strongly in salt-treated plants. Moreover, maize net photosynthesis was inhibited at leaf temperatures above 38°C, transpiration rate increased progressively while nonphotochemical fluorescence quenching increased (Crafts-Brandner and Salvucci, 2002). However, under drought stress a substantial decrease in gas exchange attributes (net photosynthetic rate, transpiration rate, stomatal conductance, water use efficiency, instantaneous water use efficiency and intercellular CO2) was observed in maize (Anjum et al., 2011).

An aerobic treatment dramatically altered the patterns of gene expression in maize seedlings (Subbajah and Sachs, 2002). During anaerobiosis pre-existing protein synthesis is immediately repressed, with the concurrent initiation of selective synthesis of approximately 20 proteins. Among these anaerobic proteins were enzymes involved in glycolysis and related processes. However, inducible genes that have different functions were also found; these may function in other, perhaps more long-term, processes of adaptations to flooding, such as aerenchyma formation and root-tip death (Subbajah and Sachs, 2002).

In common bean, salinity had adverse effects not only on biomass yield and relative growth rate, but also on other morphological parameters such as plant height, number of leaves, root length and shoot/root weight ratio. Photosynthesis, transpiration rate and stomatal conductance were also adversely affected (Gama et al., 2007). In contrast, high temperature that exceeds optimal growth conditions tends to decrease both NO3− uptake and N2 fixation (Hungria and Kaschuk, 2014). Drought stress reduces leaf water potential and gas-exchange characteristics (CO2 assimilation, stomatal conductance) (Fenta et al., 2014). It has been suggested that nodule characteristics and symbiotic nitrogen ﬁxation ability should be included with above- and below-ground traits as phenotypic markers in germplasm evaluation and breeding programs aimed at improving drought tolerance in common bean (Fenta et al., 2014). Flooding tends to reduce root dry weight, leaf area and total chlorophyll content in common bean (Celik and Turhan, 2011).

This work focuses on two of the most important grain crops in Cuba and many other countries: maize and common bean. Our aim was to identify previously unreported biochemical markers for tolerance to salinity, high temperature, drought and waterlogging, which could be used for the rapid identiﬁcation of putative stress tolerant maize and common bean plants in crop breeding programs. Stress screening was conducted on young plantlets; a method that allows large numbers of plants to be inexpensively screened. These protocols will therefore be attractive to crop breeding programs. We measured phenolics, aldehydes and chlorophylls to examine their expression in maize and common bean under abiotic stress. Data collected were used to generate Fisher’s linear discriminant functions and differentiate non-stressed and stressed plants.

Materials and Methods

Survival response to different levels of stress

After harvesting in Ciego de Avila, Cuba (2012), maize (cv. ‘Tuzón’) and common bean (cv. ‘Milagro Villalacreño’) seeds were stored at 4°C in the dark in hermetically closed containers. Seeds at 12% moisture content based on fresh weight (ISTA, 2005) were stored. Seeds of maize and common bean were sown in plant containers (200 cm3) of ferralic red soil collected in Ciego de Avila, Cuba, pH 6.8, conductivity: 0.88 S cm−1, (3 seeds per container) and allowed to germinate and grow in a growth chamber at 28°C before the imposition of stress treatments. The photosynthetic photon flux density was 800 µmol m−2 s−1. Chemical fertilizers were not used and each plant container was irrigated with 25 ml water daily for 10 d. After 10 d the plantlets were subjected to different stress treatments using five containers per treatment.

Salt stress was imposed by irrigating each pot daily with 25 ml of NaCl solution at increasing concentrations (200, 400, 600 and 800 mM) – or with water in the non-stressed controls – and
Fig. 1. Effect of sodium chloride on maize and common bean plantlets. Seeds were allowed to germinate and grow without salt stress during 10 days, then plantlets were stressed for 72 hours. Each plant container was irrigated every day with 25 ml water (without or with NaCl). In each photograph, black vertical bars represent 10 cm. Pot volume = 200 cm$^3$. Substrate: Ferralic red soil. In C, D and E, OCV means Overall Coefficient of Variation = (Standard deviation/Average)$^2$ 100. To calculate this coefficient, average values of each treatment were considered. The higher difference between the treatments compared, the higher the OCV.

Fig. 2. Effect of exposure to high temperature (40 °C) on maize and common bean plantlets. Seeds were allowed to germinate and grow without high temperature stress (28 °C) during 10 days, then plantlets were exposed to 40 °C during 12 hours. In each photograph, black vertical bars represent 10 cm. Pot volume = 200 cm$^3$. Substrate: Ferralic red soil. In C, D and E, OCV means Overall Coefficient of Variation = (Standard deviation/Average)$^2$ 100. To calculate this coefficient, average values of each treatment were considered. The higher difference between the treatments compared, the higher the OCV.
percentage plantlet survival was recorded after 72 h. Heat stress was generated by exposing plantlets to 40 °C for 12 h and survival was assessed every 3 h of treatment. Drought stress was imposed by suspending watering for 96 h; during this period, plantlet survival was registered every 24 h. Finally, to assess the effect of waterlogging, pots were immersed in 350 ml water for an additional 10 d and survival rates were determined every 24 h during this period.

Biochemical changes induced by stresses

Phenolics, aldehydes and chlorophylls were assessed in stress-treated maize and common bean plantlets ten days after sowing. Plantlets otherwise maintained under the conditions described above, were either treated with 567 mM NaCl for 72 h (salt stress), exposed to 40 °C for 9.3 h (heat stress), kept without irrigation for 51.8 h (water stress), or immersed in water (350 ml per pot) for 51.8 h. After treatment, middle-aged leaves were collected from the three plantlets of each container, pooled and ground in liquid nitrogen to a fine powder. Leaf material was similarly collected from the corresponding non-stressed controls. Three independent samples (1 g powder each) per treatment were used for all biochemical assays.

Chlorophylls (a, b, total) were quantified following Porra (2002), phenolics (free, cell wall-linked, total) by the method of Gurr et al. (1992), and malondialdehyde and other aldehydes as described.
in Heath and Packer (1968). To determine the levels of chlorophyll pigments, extraction was carried out with 5.0 ml acetone (80%, v:v). The samples were centrifuged (12,000 rpm, 4 °C, 15 min) and supernatants collected and absorbances at 647 and 664 nm recorded.

Phenolic compounds were extracted and quantified using a spectrophotometer by a colorimetric method based on reaction with thiobarbituric acid. Malondialdehyde and other aldehydes were quantified by a Folin Ciocalteu reagent (mg gallic acid equivalents per g fresh mass) using a spectrophotometer by a colorimetric method based on reaction with thiobarbituric acid. The samples were centrifuged (12,000 rpm, 4 °C, 15 min) and supernatants collected and absorbances at 647 and 664 nm recorded.

### Results

**Survival response to different levels of stress**

The effect of salt stress on maize and common bean is shown in Fig. 1. Both crop species are susceptible to dosages higher than 400 mM NaCl (Fig. 1C, E). Although plant survival at 600 mM NaCl was lower in common bean than in maize this was not significant (Fig. 1D). In contrast, common bean plantlets showed higher tolerance to heat stress than maize (Fig. 2). In maize, plant survival decreased significantly with exposure to 40 °C for more than 3 h; whereas this reduction was very slight in common bean plantlets up to 9 h of treatment (Fig. 2E). Addition of 50% survival was observed after 9.3 h of treatment (Fig. 2E). When the time of exposure to heat stress is considered an independent factor; that is plantlets of both species are pooled, 50% survival was observed after 9.3 h of treatment (Fig. 1D). In contrast, common bean plantlets showed higher tolerance to heat stress than maize (Fig. 2). In maize, plant survival decreased significantly with exposure to 40 °C for more than 3 h; whereas this reduction was very slight in common bean plantlets up to 9 h of treatment (Fig. 2E). When the time of exposure to heat stress is considered an independent factor; that is plantlets of both species are pooled, 50% survival was observed after 9.3 h of treatment (Fig. 1D). In contrast, common bean plantlets showed higher tolerance to heat stress than maize (Fig. 2). In maize, plant survival decreased significantly with exposure to 40 °C for more than 3 h; whereas this reduction was very slight in common bean plantlets up to 9 h of treatment (Fig. 2E). When the time of exposure to heat stress is considered an independent factor; that is plantlets of both species are pooled, 50% survival was observed after 9.3 h of treatment (Fig. 1D). In contrast, common bean plantlets showed higher tolerance to heat stress than maize (Fig. 2). In maize, plant survival decreased significantly with exposure to 40 °C for more than 3 h; whereas this reduction was very slight in common bean plantlets up to 9 h of treatment (Fig. 2E). When the time of exposure to heat stress is considered an independent factor; that is plantlets of both species are pooled, 50% survival was observed after 9.3 h of treatment (Fig. 1D). In contrast, common bean plantlets showed higher tolerance to heat stress than maize (Fig. 2).

### Biochemical changes produced by stresses

Phenolic compounds, aldehydes and chlorophylls were assessed in middle-aged leaves from plants surviving 72 hours of treatment with 567 mM NaCl conditions which caused 50%
Table 4. Effect of waterlogging on maize and common bean plantlets at 51.8 h of stress

<table>
<thead>
<tr>
<th>Indicators evaluated in middle-aged leaves*</th>
<th>Maize</th>
<th>Common bean</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>25 ml water/day</td>
<td>Flooded</td>
</tr>
<tr>
<td>Free phenolics (mg gallic acid equivalents/g fresh mass)</td>
<td>1.80b</td>
<td>12.14a</td>
</tr>
<tr>
<td>Cell wall-linked phenolics (mg gallic acid equivalents/g fresh mass)</td>
<td>86.03a</td>
<td>43.54b</td>
</tr>
<tr>
<td>Total content of phenolics (mg gallic acid equivalents/g fresh mass)</td>
<td>87.82a</td>
<td>55.06b</td>
</tr>
<tr>
<td>Malondialdehyde (µM/g fresh mass)</td>
<td>22.72a</td>
<td>20.67b</td>
</tr>
<tr>
<td>Other aldehydes (µM/g fresh mass)</td>
<td>81.14b</td>
<td>83.52a</td>
</tr>
<tr>
<td>Chlorophyll a (mg/g fresh mass)</td>
<td>2.96a</td>
<td>0.34b</td>
</tr>
<tr>
<td>Chlorophyll b (mg/g fresh mass)</td>
<td>1.78a</td>
<td>0.22b</td>
</tr>
<tr>
<td>Total content of chlorophyll (mg/g fresh mass)</td>
<td>4.67a</td>
<td>0.52b</td>
</tr>
</tbody>
</table>

*In each crop, results with the same letter are not statistically different (t-test, p > 0.05).

Table 5. Classification as non-stressed or stressed made by Fisher’s discriminant functions

<table>
<thead>
<tr>
<th>Type of stress evaluated</th>
<th>Plant</th>
<th>Results of discriminant functions</th>
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</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Function for non-stressed</td>
</tr>
<tr>
<td>Salinity</td>
<td>Maize</td>
<td>0 mM NaCl</td>
</tr>
<tr>
<td></td>
<td></td>
<td>22.72a</td>
</tr>
<tr>
<td></td>
<td></td>
<td>25 ml water/day</td>
</tr>
<tr>
<td></td>
<td>Common bean</td>
<td>0 mM NaCl</td>
</tr>
<tr>
<td>High temperature</td>
<td>Maize</td>
<td>28 °C</td>
</tr>
<tr>
<td></td>
<td>Common bean</td>
<td>28 °C</td>
</tr>
<tr>
<td></td>
<td></td>
<td>40 °C</td>
</tr>
<tr>
<td></td>
<td></td>
<td>40 °C</td>
</tr>
<tr>
<td></td>
<td></td>
<td>25 ml water/day</td>
</tr>
<tr>
<td>Drought</td>
<td>Maize</td>
<td>20 °C</td>
</tr>
<tr>
<td></td>
<td>Common bean</td>
<td>20 °C</td>
</tr>
<tr>
<td>Waterlogging</td>
<td>Maize</td>
<td>28 °C</td>
</tr>
<tr>
<td></td>
<td>Common bean</td>
<td>28 °C</td>
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<tr>
<td></td>
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<td>40 °C</td>
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plantlet death as calculated from Fig. 1E and compared to levels in the corresponding non-stressed controls. In general, salt stress induced statistically significant changes in the levels of all indicators with the exception of chlorophyll b contents in maize, although in most cases these changes were relatively small. The most remarkable effects of abiotic stresses can be summarized as follows. The salinity stress treatment produced an 8-fold increase in free phenolics in maize and between 3 and 4-fold increases in chlorophyll content (a, b, and total) in common bean (Table 1). On the other hand, maize and common bean responded differently to heat stress in the levels of biochemical markers (Table 2). In common bean, when biochemical marker expression was assessed after 40°C treatment for 9.3 h a 6-fold decrease in free phenolics and a 20-fold reduction of chlorophylls (a, b, and total) compared to the control was observed. In contrast, chlorophyll levels increased in maize between 5 and 10-fold under the same conditions. Moreover, fifty percent of all plantlets died 51.8 h after irrigation was suspended (Fig. 3E). Under these conditions, the most relevant observed changes in biochemical marker levels were a 7-fold increase in free phenolic compounds and a 5-fold reduction in chlorophyll contents (a, b, and total) in maize, and an increased in chlorophyll levels in common bean (Table 2). In the waterlogging stress experiment, biochemical evaluations were made 51.8 h post water immersion; a time that coincided with 50% plant death under this condition (Fig. 5). Total contents of phenolics and chlorophylls, especially a, were the most affected indicators and therefore can be regarded as potential abiotic stress biochemical markers. Leaf chlorophyll content was affected by salinity in tetraploid wheat (Munns and James, 2003), rice (Sultana et al., 1999), Brassica oleracea (Bhattacharya et al., 2004), Brassica juncea (Qasim, 1998) and Brassica napus (Pak et al., 2009). Salinity can affect chlorophyll content through inhibition of chlorophyll synthesis or an acceleration of its degradation (Zhao et al., 2007). Thiayonong et al. (2004) found that the chlorophyll losses due to salinity stress is consistent with possible differences in reactive oxygen species (ROS) production among the genotypes and suggested that in salt sensitive genotypes, ROS scavenging systems were unable to destroy ROS generated. Our results do not support these findings as common bean chlorophyll levels increased under salinity (Table 1).

According to Baker (1993), changes in the photochemical efficiency of plants under drought may be assessed by the analysis of chlorophyll a fluorescence efficiency associated with photosystem II. Under stress, a decrease in the ratio of variable fluorescence / maximum fluorescence has been attributed to the inactivity of the photosystem II reaction centers due to the degradation of the D1 and D2 proteins responsible for the transfer of water electrons to chlorophyll a associated with the photosystem II reaction center (Hao et al., 1999; Lazár, 1999). Chlorophyll content could therefore be correlated to chlorophyll fluorescence thus indicating its suitability as a future biochemical marker. Abiotic stresses decrease photosynthesis, mainly by limiting CO2 entrance to leaves through stomatal closure. Moreover, membrane systems containing chlorophylls are destabilized affecting the luminous phase thus leading to increased synthesis of chlorophylls that are unable to fix more CO2 (Hörtensteiner, 2006; Hörtensteiner and Kräutler, 2011).

A consequence of the abiotic stress-induced limitation of photosynthesis is the exposure of plants to excess energy, which, if not safely dissipated, may be harmful to photosystem II because of over reduction of the reaction centers (Demmig-Adams and Adams, 1992) and increased production of ROS in the chloroplasts (Simmoff, 1993). On the other hand excess energy could be used to synthesize secondary metabolites as suggested by Selmar and Kleinwächter (2013).

Phenolic compounds and flavonoids are among the most influential and widely distributed secondary products in the plant kingdom (Ali and Abbas, 2003). Many play important physiological and ecological roles and are involved in resistance to different types of stress (Ayaz et al., 2000). These metabolites have several defense functions and, therefore, their biosynthesis in plants is generally induced in response to biotic and abiotic stimuli such as UV-B radiation, drought, chilling ozone, heavy metals, and attacks by pathogens, wounding, or nutrient deficiency (Bettsaib et al., 2011; Dixon and Paiva, 1995; Grace, 2005).

Our results indicated that free phenolics and chlorophylls, especially a, were the most responsive indicators. The differences recorded between maize and common bean is to some extent a
function of their differences in photosynthetic efficiency, however there may also be a genotype within species effect. We have hypothesized that, in stressed plants, levels of free phenolics and chlorophylls first increase and subsequently decrease. However, such changes take place in different time frames depending on the plant species and genotype. In the experiments shown here, moderate and severe stress conditions were applied that did not necessarily represent any specific natural environment, but were used for selection purposes only. In follow-up experiments, mild to moderate stress conditions may enable plant metabolism to respond properly to the respective stress conditions. Further studies are in progress to determine the practical use of free phenolics, chlorophylls and other biochemical markers for stress tolerance in breeding programs.

The Fisher’s linear discriminant functions shown in this paper (Fig. 5) are important tools for those breeding programs focused on the production of abiotic-stress-tolerant plants. Maize and common bean seeds of new genotypes can be grown for 10 days and then stressed or not as described here. Levels of phenolics (free and cell-wall linked), malondialdehyde, other aldehydes and chlorophylls ($a$, $b$) are determined. Such new data are evaluated in both discriminant functions. If the resulting value of the stressed discriminant function is similar to that of the non-stressed discriminant function, the new genotype can be regarded as putatively tolerant, as it shows similar physiology under either non-stressing or stressing conditions. Although the new genotype tolerance still requires additional confirmation under a field environment, the results described here allow some research cost reductions because there is no inclusion of a large number of susceptible cultivars in expensive field trials. At present, this research group is carrying out further experiments to know if these discriminant functions can be used in other plant species.

Discriminant analysis is useful for situations where the building of a predictive model of group membership based on observed characteristics of each case is desirable. The procedure generates discriminant functions based on linear combinations of the predictor variables, which provide the best discrimination between groups. The functions are generated from a sample of cases for which group membership is known. The functions can then be applied to new cases with measurements for the predictor variables but unknown group membership (Bunette and Prasanna, 2003; Cardi, 1998; Daouy and Lawes, 2000; Fighvollo et al., 2001; Somersalo, 1998; Teshome et al., 1997). The use of this kind of analysis for differentiation of non-stressed or stressed plants is a novel aspect of this report that can be applied for early selection of plant tolerance to abiotic factors.

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