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JIP-test as a tool to identify salinity tolerance in sweet sorghum genotypes

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Abstract

The effect of salinity on primary photochemical reactions (using JIP-test) in six sweet sorghum genotypes was tested. An increase in salt concentrations induced significantly the accumulation of proline and caused a decline in leaf osmotic potential. Except for 100 mM NaCl concentration, salinity significantly decreased chlorophyll content and photosynthetic efficiency of plants. Increasing salinity led to a higher accumulation of Q_B-nonreducing PSII reaction centers. K-step in OJIP fluorescence transient was observed for the most sensitive genotypes under the high NaCl concentration. The studied sorghum genotypes responded differently to salinity stress. Thus, the study helps understand the plant tolerance mechanisms of different sweet sorghum genotypes to increasing salinity stress. The study also confirmed that the use of JIP-test is suitable for the identification of sorghum genotypes according to their growth under salinity stress.

Additional key words: abiotic stress; chlorophyll fluorescence; photosynthesis.

Introduction

With an increase of the Earth's population, a significant proportion of grasslands or wetlands are being exploited through intensive cultivation and management (Knox *et al.* 2015, Yamaura *et al.* 2019). The exploitation of different ecosystem leads to damage of the original vegetation and soil structure, which results in soil salinization. The

combination of salinity with drought further adds to the nutritional constraints for agriculture productivity (Kumar et al. 2017). Indeed, the development of agriculture, involving systematic use of irrigation, is the main factor behind the extension of this abiotic constraint, which severely limits crop yields and contributes to the degradation of vegetal cover. With the increase in population, the freshwater availability is becoming limited, whereas the

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Abbreviations: ABS/CS_m – absorbtion per excited cross-section; DI_0/CS_m – effective dissipation of energy per excited cross section; ET_0/CS_m – electron transport flux from reduced Q_A to Q_B per excited cross-section; F_0 – minimal intensity of chlorophyll a fluorescence, measured when all PSII reaction centers are assumed to be open (at 50 μ s); F_m – maximum fluorescence intensity; F_t – fluorescence intensity at time t; F_v/F_m – photosynthetic efficiency of PSII under dark adaptation; LA – leaf area; Lp – plant length; Lr – root length; Ls – shoot length; OEC – oxygen-evolving complex; PI_{ABS} – index of overall photochemical performance; RC/ABS – fraction of active PSII reaction centers; TR_0/CS_m – maximal trapping rate of absorbed photons per excited cross section; V_j – variable fluorescence at time 2 ms; W_k – the amplitude of the K step of OJIP fluorescence transient; ψ_0 – electron transport at PSII acceptor side.

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groundwater salinity is increasing (Kabeel and El-Said 2015). Several factors, such as intensive use of saline water in irrigation, evaporation, and condensation due to climate change, low precipitation, etc., play an important role in rapidly increasing saline land throughout the world. Maas and Hoffman (1977) have shown that the crop yield does not get affected until a salinity threshold is exceeded. Research has been performed to show that low salinity does not impact crop productivity, whereas when salt is present in a high concentration, it may severely affect the crop yield (Machado and Serralheiro 2017, Mbarki et al. 2018, Arif et al. 2019). The mechanism behind the plant response can be related to osmotic stress, ionic toxicity, and ionic imbalances caused by salinity in plants (Kim et al. 2007, Munns and Tester 2008). The inhibition of leaf expansion by the salinity was reported to be associated with changes in several factors, such as leaf turgor, net CO₂ assimilation, accumulation of toxic ions, and/or disturbance in mineral nutrition (Bernstein et al. 1993). A number of studies have been performed to understand the salinity interaction with photosynthetic machinery in different crops (Sudhir and Murthy 2004, Kalaji et al. 2018, Wungrampha et al. 2018, Betzen et al. 2019).

Among agricultural crops, sorghum is one of the major cereal, which is widely grown in arid, subtropical, semiarid tropical, and temperate regions. In several regions of the world, it is one of the most important sources of food. Sorghum is used as a raw material for brewing and is considered to be a bioenergy crop. As an important crop species, it feeds more than 500 million people in 98 countries (Pennisi 2009). The effects of salt stress and high temperature on sorghum have been widely reported and include the inhibition of photosynthesis and PSII activity (Azhar and McNeilly 1987, 1989; Esechie 1994, Marambe and Ando 1995, Kader and Jutzi 2004, Netondo et al. 2004a,b; Prasad et al. 2008, Jain et al. 2010). A large number of genotypic variation has been reported for the tolerance to salinity in sorghum (Igartua et al. 1994, Maiti et al. 1994). As the salinity problem is increasing, there is a need for quick identification of salinity tolerant variety in order to improve crop productivity in saline soil.

Photosynthesis is closely related to crop yield but is affected by salinity. The decline of photosynthetic capacity commonly occurred in plants under salt stress, but the underlying mechanisms are complex and not clear (Isayenkov and Maathuis 2019). Chlorophyll (Chl) fluorescence is often used to predict crop yields under various environmental conditions (Baker and Rosenqvist 2004, Kalaji and Pietkiewicz 2004). In recent years, Chl fluorescence parameters were used as the criteria for selecting plants in the breeding programs (Kalaji and Guo 2008, Kalaji et al. 2017). It can be used to analyze detailed information about the energy transfer in photosynthesis (Maxwell and Johnson 2000, Kalaji et al. 2017). After a dark adaptation, when plants are exposed to actinic light, a rapid increase in Chl fluorescence can be observed and explored to provide information about the status of PSII through a JIP-test (Strasser 1978, 1981). JIP-test uses several mathematical formulae for the purpose to decipher the energy flow in the thylakoids membrane, and is able

to describe primary photosynthetic reactions (Strasser 1978, 1981). In recent years, JIP-test has been successfully applied to understand the mechanism of action of several stress factors in plants (Bussotti *et al.* 2011, Bayat *et al.* 2018, Rastogi *et al.* 2019a). Therefore in this study, we used the JIP-test to understand the salinity tolerance mechanism in sorghum in order to select the best genotype (among studied six genotypes), which can be suitable for growing under salinity conditions.

Materials and methods

Plant material and cultivation conditions: Six sorghum [Sorghum bicolor (L.) Moench.] genotypes (five originated from China - Tian Xuan 35, Tian Xuan 90, Tian Xuan 97, Dalishi, and Sutian-1 and one from USA – Rio) were grown under hydroponic cultivation under light-, humidity-, and temperature-controlled conditions. Seeds were surface-disinfected for 30 min using 10% SAVO solution (Biochemie, Bohumín, Czech Republic), which contains 5% NaClO, and washed three times in distilled water. Seeds were germinated on moisture filter paper (Whatman 3) at Petri dishes under 14/10-h day/night regime with light intensity of 150 μmol(photon) m⁻² s⁻¹ and temperature of 24/18°C. The 7-d-old seedlings were selected for a similar size and transferred to 7-L plastic trays with aerated (regularly six times a day for two hours) Hoagland nutrient solution (pH 6.5). The nutrient solution was changed every four days. Plants were cultivated under 14-h photoperiod, at temperature of 24/18°C, relative humidity of 55–60%, and light intensity of 150 µmol(photon) m⁻² s⁻¹. Salt treatment was applied at the stage of 21 d after germination by adding NaCl into Hoagland solution to final concentration 50, 100, 150, 200, and 250 mM, respectively. The time duration of salt treatment was ten days.

Morphological analysis: The length of the aboveground part of a plant (Lp), root (Lr), and shoot (Ls) were measured using caliper on randomly selected plants (7 plants per treatment). Leaf area (LA) was measured for 3rd mature leaf of the plant by drawing the leaf structure on graph paper; further the leaf drawn was cut and measured through leaf area meter *CI-202* (*CID-Bioscience*, USA), with 10 repetitions.

Chl a fluorescence: Chl a fluorescence transients were measured with Handy-PEA (Hansatech Instruments Ltd., King's Lynn, Norfolk, UK). All fluorescence measurements were taken in the middle part of the 3rd mature leaf (out of the main leaf vein). The leaf samples after 30 min of dark adaptation (using leaf clips) were illuminated with continuous red light (peak in 650 nm wavelength; the spectral line half-width of 22 nm) provided by an array of three light-emitting diodes. The light pulse intensity used was 3,500 µmol(photon) m⁻² s⁻¹ and the duration of the light pulse was 1 s. The measured data were used for calculation of the JIP-test parameters using Biolyzer v. 3.06 HP software (Strasser et al. 2010, Kalaji et al. 2016).

Osmotic potential, proline content, and Chl index: Samples of leaves (two circle segments, diameter of 6.0 mm) for measurement of osmotic potential (ψ_s [MPa]) were frozen in liquid nitrogen immediately after the cut and stored at -80° C. Before ψ_s determination, the segment was thawed and inserted into psychrometric chamber *C-52* connected to *PSYPRO* datalogger (*Wescor*, Logan, Utah, USA). Time interval of temperature and humidity equilibration was set at 60 min and the cooling interval was 15 s.

Free proline content in leaves was determined by the ninhydrin method (Bates *et al.* 1973) using spectrophotometer *Jenway 6405 UV/VIS (Jenway*, Stone, Staffordshire, UK).

Chl (SPAD) index was measured on the adaxial surface of the 3rd leaf using *SPAD-502* chlorophyll meter (*Minolta*, Japan) immediately after the measurement of Chl *a* fluorescence. The SPAD index values correspond to the amount of Chl present in the leaf and are calculated based on the amount of light transmitted by the leaf in two wavelength regions (650 and 940 nm). SPAD index was an average of ten measurements on the middle part of the leaf (except base and apex).

Sodium and potassium ion concentration: The root and leaves were sampled for dry mass for the purpose to determine Na⁺ and K⁺ content. Samples were ovendried at 70°C for 72 h. Na⁺ and K⁺ content in plant tissues (root and leaves) were determined according to Allen *et al.* (1986) using a flame photometer (*FP6410*, *INESA*, China). Dry plant tissues were ground using a mortar and pestle and samples (0.1 g) were digested in 2 ml of sulfuric acid and hydrogen peroxide mixture (3:1). Because the tissue mass of plants grown in few conditions was very low, we pooled the sample together and took only one measurement for the given condition. We selected 150 mM NaCl concentration for the measurement as under this concentration the plants clearly showed the stress response.

Statistical analysis: Each data point represented the mean of seven replicates for morphological and SPAD measurements, four replicates for osmotic potential, and proline analysis, and 3–5 replicates for Chl *a* fluorescence measurements. All data were evaluated by one-way analysis of variance (*ANOVA*), and the mean differences were compared by post hoc test at *P* level of 0.05. The normal distribution and homogeneity of experimental results were tested by *Kolmogorov-Smirnov* and *Lavene*'s test. In all figures, the spread of values is shown as error bars representing standard error (SE) of the means. All analyses were performed using the *Statistica v. 10* software (*StatSoft Inc.*, Tulsa, Oklahoma, USA).

Results

Morphology of sorghum genotypes: Different sorghum genotypes treated with different concentrations of salt were analyzed for its plant length (Lp), root length (Lr), shoot length (Ls), and the leaf area (LA) (Table 1). The response was dependent on salt concentration in different

genotypes. With NaCl treatment of 250 mM, Tian Xuan 90 showed the least sensitivity; we observed around 5, 44, 12, and 40% decrease in Lp, Lr, Ls, and LA, respectively, whereas cv. Tian Xuan 97 was very sensitive to 250 mM NaCl with around 61, 58, 59, and 76% decrease in Lp, Lr, Ls, and LA, respectively. Interestingly, the salt treatment of 200 mM was observed to be more harmful to Tian Xuan 90 than the impact of 250 mM salt concentration, especially for plant length. LA was the least in genotype Rio under 250 mM concentration. The shortest root was observed in Tian Xuan 97 under 250 mM salt treatment.

Transient fluorescence curves: Transient fluorescence curves, also known as OJIP curves, were analyzed for six sorghum genotypes exposed to different concentrations of salt (0, 100, 150, 200, and 250 mM). For better visualization, the curve was normalized for F₀ (fluorescence at time 0) and presented as F_t/F₀ (where F_t is fluorescence at time t) against the time (Fig. 1). The figure shows the variable response to salinity stress on OJIP curves for different sorghum genotypes. Apart from O, J, I, and P steps in the OJIP curve, some other peaks may appear, which indicates various phenomena of photosynthetic mechanism (Kalaji et al. 2014). Among them, K step is an important step which indicates a decrease in oxygenevolving complex (OEC) capacity due to the impairment of electron transport between the OEC and reaction center of PSII (Srivastava et al. 1997). Generally, these additional effects were masked under moderate stress conditions in the graph with absolute value of Chl a fluorescence transient curve. Therefore, the mathematically double-normalized values of OJ phase, i.e., the time plots of relative variable fluorescence V_t/V_j from 0.05 to 2 ms, were used in the analysis (Fig. 2). The graph indicates a different response of salinity stress at K step for sorghum genotypes.

Different photosynthetic parameters: Among various photosynthetic parameters, maximal photochemical efficiency of PSII (F_v/F_m) and performance index (PI_{ABS}) were selected to be represented in this manuscript due to its proven sensitivity to identify the abiotic stress response in different plants (Kalaji et al. 2017, Rastogi et al. 2019a,b). F_v/F_m indicates the maximum quantum yield of PSII photochemistry, whereas PIABS is a photosynthetic parameter that takes into consideration a number of different phenomena related to PSII photochemistry (Zivcak et al. 2008). With an increase in the salt concentration, the impact of salinity on different sorghum genotypes was different (Fig. 3). Other important photosynthetic parameters obtained through JIP-test analysis, which represent the activity of acceptor and donor sites of PSII, were analyzed and presented in Fig. 4. Relative variable fluorescence at 2 ms (V_i), the probability that a trapped exciton transfers as electron into the electron transport chain beyond Q_A (ψ_0), the ratio of the K phase to the J phase (W_k) and the fraction of OEC in relation to control plants were observed and presented for its differences in different sorghum genotypes under different salinity conditions.

Table 1. Morphological characterization of sorghum genotypes treated by different concentrations of NaCl in hydroponic solution. Lp – plant length, Lr – root length, Ls – shoot length, LA – leaf area. The data represent means \pm SE, n = 7. Uppercase letters and lowercase letters denote significant difference (Duncan's post hoc test) between genotypes at a given growth NaCl concentration level and all treatments for given sorghum genotype at p < 0.05, respectively.

Genotype	NaCl [mM]	Lp [mm]	Lr [mm]	Ls [mm]	LA [cm ²]
Tian Xuan 35	0	$243.6 \pm 19.7^{\mathrm{Ba}}$	249.4 ± 15.0^{Aa}	$59.2 \pm 5.1^{\mathrm{Ba}}$	$29.4\pm3.2^{\mathrm{Ba}}$
	100	$235.0 \pm 23.4^{\rm Aa}$	$175.7\pm3.8^{\mathrm{Aa}}$	$54.0 \pm 5.3^{\mathrm{Ba}}$	$28.2\pm5.5^{\mathrm{Aa}}$
	150	$178.3\pm15.0^{\mathrm{Bb}}$	$228.0\pm28.8^{\mathrm{Ab}}$	$52.7\pm7.1^{\rm Ba}$	$15.4 \pm 4.7^{\mathrm{Bb}}$
	200	$201.7\pm14.5^{\mathrm{Ab}}$	$162.7\pm10.5^{\mathrm{Ab}}$	$59.0\pm3.0^{\rm Ba}$	$18.2\pm2.6^{\mathrm{Bb}}$
	250	$200.0\pm19.8^{\mathrm{ABb}}$	$178.5\pm6.4^{\mathrm{Ab}}$	$54.5\pm2.1^{\rm Ba}$	$15.7\pm3.6^{\mathrm{ABb}}$
Tian Xuan 90	0	$188.0 \pm 8.0^{\text{CDa}}$	$215.4 \pm 31.0^{\mathrm{ABCa}}$	$57.0 \pm 3.9^{\mathrm{Bab}}$	$18.0 \pm 2.5^{\mathrm{CDa}}$
	100	$176.0\pm32.8^{\mathrm{Ba}}b$	$164.0\pm10.1^{\mathrm{Ab}}$	$57.7 \pm 6.1^{\mathrm{ABa}}$	$17.3\pm6.5^{\rm Ba}$
	150	$175.7\pm20.2^{\rm Bab}$	$170.0\pm36.3^{\rm BCb}$	$52.3 \pm 3.1^{\mathrm{Bab}}$	$8.6 \pm 3.0^{\text{BCb}}$
	200	$149.7\pm23.5^{\mathrm{Bb}}$	$161.7\pm24.4^{\mathrm{Ab}}$	$52.7 \pm 2.5^{\mathrm{Bab}}$	$11.2\pm1.4^{\text{Cab}}$
	250	$179.7\pm14.8^{\text{Bab}}$	$144.3\pm10.8^{\mathrm{Bb}}$	$50.0\pm2.6^{\mathrm{Bb}}$	$10.9 \pm 1.1^{\text{BCab}}$
Tian Xuan 97	0	$159.6\pm34.8^{\mathrm{Da}}$	$185.8\pm16.7^{\text{Ca}}$	$52.4\pm8.4^{\rm Ba}$	$11.2 \pm 5.6^{\mathrm{Dab}}$
	100	$173.3\pm22.5^{\mathrm{Ba}}$	$163.0\pm11.0^{\mathrm{Aa}}$	$51.3\pm7.8^{\rm Ba}$	$13.8 \pm 4.4^{\mathrm{Ba}}$
	150	$150.3\pm10.6^{\mathrm{Bab}}$	$127.7\pm12.7^{\text{Cb}}$	$45.7\pm8.5^{\rm Ba}$	$7.4 \pm 4.7^{\text{Cabc}}$
	200	$106.7\pm13.9^{\mathrm{Bbc}}$	$100.7\pm27.3^{\text{Cbc}}$	$31.3\pm1.2^{\text{Cb}}$	$4.2\pm1.6^{\rm Dbc}$
	250	$62.0\pm7.0^{\rm Dc}$	$79.3\pm18.5^{\rm Dc}$	$21.7\pm1.5^{\text{Eb}}$	$2.7 \pm 1.1^{\rm Dc}$
Dalishi	0	$293.0\pm39.0^{\mathrm{Aa}}$	$201.0\pm36.5^{\mathrm{BCa}}$	$81.3 \pm 4.6^{\mathrm{Aa}}$	$54.7\pm7.7^{\rm Aa}$
	100	$197.3\pm9.9^{\mathrm{Bb}}$	$159.3\pm15.2^{\mathrm{Ab}}$	$66.3\pm3.1^{\mathrm{Ab}}$	$33.4 \pm 4.4^{\text{Ab}}$
	150	$232.3\pm21.7^{\mathrm{Ab}}$	$184.3\pm17.5^{\mathrm{ABb}}$	$68.0 \pm 6.6^{\mathrm{Ab}}$	$30.4\pm1.7^{\mathrm{Ab}}$
	200	$241.3\pm15.9^{\mathrm{Ab}}$	$148.0 \pm 8.9^{\mathrm{ABb}}$	$69.3\pm6.5^{\mathrm{Ab}}$	$31.8 \pm 4.6^{\text{Ab}}$
	250	$219.0\pm21.3^{\mathrm{Ab}}$	$153.3 \pm 12.7^{\mathrm{Bb}}$	$60.3\pm4.5^{\mathrm{Ab}}$	$21.1 \pm 5.3^{\mathrm{Ac}}$
Rio	0	$212.6\pm13.3^{\mathrm{BCa}}$	$237.0\pm14.2^{\mathrm{ABa}}$	$58.7 \pm 3.5^{\mathrm{Ba}}$	$21.8\pm2.9^{\text{Ca}}$
	100	$186.3\pm8.4^{\mathrm{Bab}}$	$176.0 \pm 4.2^{\mathrm{Ab}}$	$52.7 \pm 2.1^{\mathrm{Bb}}$	$17.0 \pm 2.5^{\rm Bab}$
	150	$167.3\pm8.0^{\mathrm{Bb}}$	$156.3\pm13.8^{\mathrm{BCb}}$	$49.0 \pm 4.4^{\mathrm{Bb}}$	$14.1 \pm 3.4^{\rm BCbc}$
	200	$128.7\pm28.4^{\mathrm{Bc}}$	$112.7\pm26.6^{\rm BCc}$	$35.0 \pm 2.8^{\text{Cc}}$	$9.1 \pm 3.6^{\text{CDcd}}$
	250	$129.7\pm3.2^{\text{Cc}}$	$103.3\pm7.6^{\text{Cc}}$	$29.0 \pm 1.0^{\mathrm{Dd}}$	$5.6 \pm 1.0^{\text{CDd}}$
Sutian-1	0	$181.0\pm16.7C^{\text{Db}}$	$121.3\pm7.0^{\mathrm{Da}}$	$60.7\pm1.5^{\rm Bb}$	$15.9 \pm 2.6^{\text{CDb}}$
	100	$236.7\pm15.7^{\mathrm{Ba}}$	$151.0\pm10.8^{\mathrm{Ba}}$	$71.3 \pm 5.7^{\mathrm{ABa}}$	$26.6\pm6.7^{\rm Ba}$
	150	$163.0\pm27.9^{\rm Bbc}$	$124.0\pm31.2^{\rm Ca}$	$49.0\pm10.6^{\mathrm{Bc}}$	$13.1 \pm 4.1^{\rm BCbc}$
	200	$137.3\pm15.0^{\rm Bc}$	$125.3\pm17.0^{\mathrm{ABCa}}$	$32.3\pm2.5^{\rm Cd}$	$8.4 \pm 3.1^{\rm CDbc}$
	250	$143.3\pm7.1^{\text{Cc}}$	$87.0 \pm 4.6^{\text{CDb}}$	$39.7 \pm 3.1^{\rm Ccd}$	$7.2 \pm 1.2^{\text{CDc}}$

Leaf model for energy flux: For a better understanding of photosystem functioning, the *Biolyzer* software-generated leaf model was presented in Fig. 5. The figure shows the proportion of phenomenological energy fluxes within a leaf, calculated based on per excited cross section (CS_m).

Osmotic potential, proline concentration, and Chl concentration: Salt is known to interfere with solute movement, therefore, the osmotic potential of plants was measured and observed to be increasing with the increase in salt concentration (Table 2). Proline is the amino acid, which is elevated under stress conditions (Hayat *et al.* 2012); in this study, the concentration of proline increased with the increase in salt concentrations for different sorghum genotypes (Table 2). Chl concentration is the factor that is generally decreasing under plant stress conditions; the SPAD index indicated a decrease in Chl concentration due to salinity in studied sorghum genotypes.

 $\mathbf{Na^+}$ and $\mathbf{K^+}$ concentration: The ratio of Na^+ and K^+ increased significantly in salt-treated samples (Table 3) for studied sorghum genotypes, which was due to a significant increase in Na^+ due to the salt treatment, which also replaced K^+ ; as a result, the concentration of K^+ decreased in the treated samples.

Discussion

Different stress factors interact differently with different plants, but we start recognizing it only when we see their impacts on plant growth parameters, such as plant size, leaf shape, yield, *etc*. Therefore, for this study, we first focused on morphological parameters and observed a significant decrease in plant length, root length, shoot length, and leaf area with an increasing concentration of salt. Morphological variation showed that Tian Xuan 97 was the most sensitive genotype to salt stress, whereas Tian Xuan 90 was the least

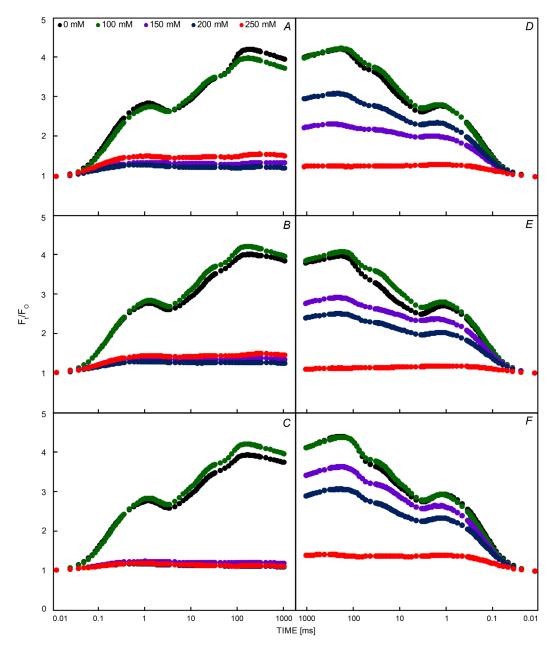


Fig. 1. Chlorophyll a normalized fluorescence intensity (F_1) relative to F_0 (A–F) in sorghum leaves of different genotypes affected by different NaCl concentration. Genotypes: Tian Xuan 35 (A), Tian Xuan 90 (B), Tian Xuan 97 (C), Dalishi (D), Rio (E), and Sutian-1 (F). Data represent the mean (n = 3–5).

sensitive genotype of sorghum. But the morphological changes are not sufficient to indicate why the deviation in growth parameters happened, and the morphological deviations are identified mostly at a later stage of the plant life cycle. Therefore, different methods have been developed for the purpose to identify the stress response at an early stage. Some of the methods are destructive and some are noninvasive. The noninvasive and quick methods are the most useful for farmers as they can avoid the loss of the crop yield by early detection of the stress factor. Recently, the JIP-test has established itself as a quick and reliable technique for field identification of different stress

factors at the early stage (Bussotti et al. 2011, Kalaji et al. 2014, Rastogi et al. 2019a). Chl induction curve is an indicator of different phenomena of photosynthetic light reactions. Significant suppression in Chl induction curve clearly indicated that Tian Xuan 35, Tian Xuan 90, and Tian Xuan 97 are salt-sensitive genotypes, whereas Dalishi, Rio, and Sutian-1 are relatively salt-resistant sorghum genotypes, where photosynthetic machinery was functioning even at salt concentration of 200 mM. The studied genotypes of sorghum were resistant to 100 mM salt except Tian Xuan 97, which was highly sensitive to salt stress. The sensitivity of Tian Xuan 97 was clear with a

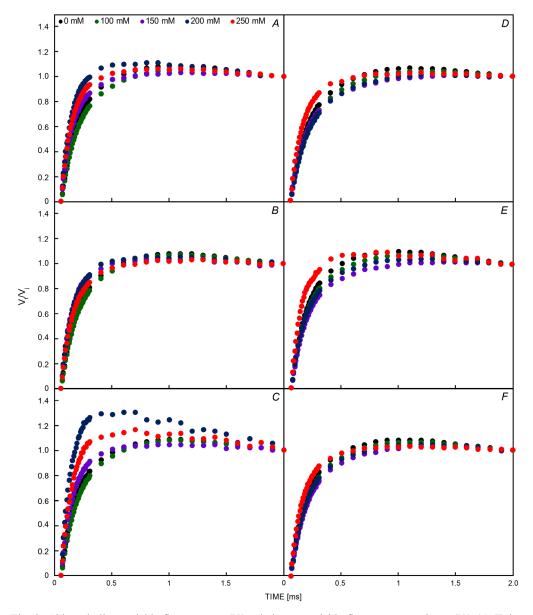


Fig. 2. Chlorophyll a variable fluorescence (V₁) relative to variable fluorescence at 2 ms (V₁) (A–F) in sorghum leaves of different genotypes affected by different NaCl concentration. Genotypes: Tian Xuan 35 (A), Tian Xuan 90 (B), Tian Xuan 97 (C), Dalishi (D), Rio (E), and Sutian-1 (F). Data represent the mean (n = 3–5).

clear deviation observed for K step, indicating dissociation of the OEC (Oukarroum *et al.* 2007), thus showing significant damage to the photosystems. The observation of photosynthetic parameters, such as F_{ν}/F_{m} and PI_{ABS} , clearly indicated the lowest value for Tian Xuan 97, which further confirmed the claim that Tian Xuan 97 was the most sensitive genotype. Data also indicated that the plants were able to tolerate the salt concentration up to 100 mM, but 150 mM was already lethal for salt-sensitive varieties, *i.e.*, Tian Xuan 35, Tian Xuan 90, and Tian Xuan 97. As the salt-resistant varieties also showed a significant reduction in F_{ν}/F_{m} and PI_{ABS} at 150 mM, this concentration was identified as the threshold level, harmful to all studied genotypes. The best photosynthetically performing geno-

type was Sutain-1 which showed a significantly better Chl induction curve, F_v/F_m , and PI_{ABS} when compared to other salt-resistant genotypes. The same was visible in other photosynthetic parameters studied in Fig. 4, where a number of closed reaction centers was higher for sensitive varieties as seen by V_j . The probability of electron to pass through Q_A^- was also low for sensitive genotypes, in the form of lower value of ψ_0 (Kalaji *et al.* 2017). The ratio of K phase to the J phase (W_k) was high for Tian Xuan 97. The value of W_k indicates the activity of the OECs on the donor side of PSII where an increase in W_k indicates damage to the complexes. The observation was confirmed by the values of the fraction of active oxygen-evolving complex which fell down to zero at 150 mM in Tian Xuan

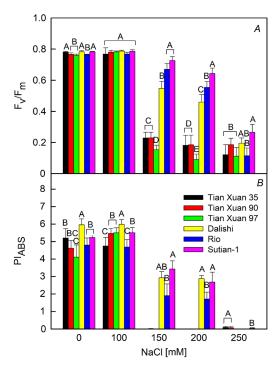


Fig. 3. Maximal photochemical efficiency of PSII (F_{ν}/F_{m}) and performance index (PI_{ABS}) in sorghum genotypes affected by different NaCl concentration in hydroponic solution. Data represent the mean \pm SE (n=3-5). Different letters indicate the significant differences between genotypes (Duncan's post hoc test, p<0.05).

97 genotype, whereas, the fraction of active OEC was high up to 200 mM in the resistant genotypes. In the leaf pipeline model presented, we can see that the number of inactive reaction centers was high (shown by dark cycle) for plants treated with 150 mM and higher salt concentration except for genotype Sutian-1 (which was the most salt-tolerant one). Similar to a number of reaction centers, the lightuse efficiency of different plants also decreased which can be seen by smaller sizes of ABS/CS_m and TR_0/CS_m , where ABS/CS_m indicates light absorbed by per cross section and TR_0/CS_m indicates trapped light energy per cross section. Electron flux per cross section of leaf (ET_0/CS_m) was negligible for salt-sensitive genotypes, whereas it was significantly higher for the salt-resistant genotypes.

To see further the impact of salinity on some other physiological processes, we measured osmotic potential (ψ_S) , proline content (Pro), and SPAD index. Different studies have shown that the accumulation of salts in the root zone may cause a decrease in ψ_s (Franco et al. 2011), which was lower with the increase of NaCl concentration in different sorghum genotypes in our study. Tian Xuan 70, being the most sensitive, did not show any measurable ψ_S for 150, 200, and 250 mM NaCl. An overproduction of proline is a common phenomenon in plants under stress. The increase in the proline concentration has a positive effects on stress tolerance by maintaining cell turgor, preventing electrolyte leakage, and preventing plants from oxidative burst (Hayat et al. 2012). The increase in the proline content under salinity stress was observed in different sorghum genotypes, which is in agreement

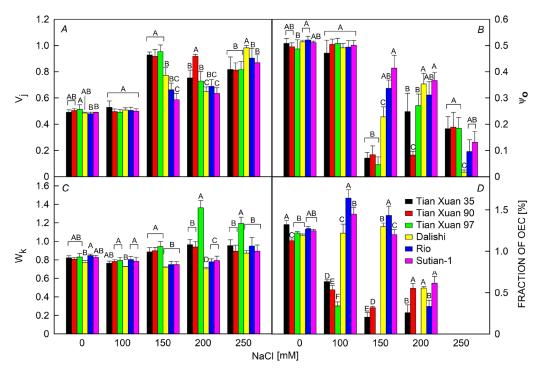


Fig. 4. Parameters reflected the activity of acceptor (A,B) and donor (C,D) side of PSII in sorghum leaves affected by different NaCl concentration. (A) Relative variable fluorescence at 2 ms (V_j) . (B) Probability that a trapped exciton transfers as electron into the electron transport chain beyond Q_A (ψ_0) . (C) Ratio of the K phase to the J phase (W_k) . (D) The fraction of oxygen-evolving complex relative to control plants. Data represent the mean \pm SE (n = 3-5). Different letters indicate the significant differences between genotypes (Duncan's post hoc test, p<0.05).

Table 2. Osmotic potential (ψ_s), proline content (Pro), and SPAD index in sorghum genotypes treated by different concentrations of NaCl in hydroponic solution. The data represent means \pm SE, n=7 for SPAD and n=4 for ψ_s and Pro, respectively. n.m. – not measurable, FM – fresh mass. *Uppercase letters* and *lowercase letters* denote significant differences (*Duncan*'s post hoc test) between genotypes at different NaCl concentration levels (p<0.05), respectively.

Genotype	NaCl [mM]	$\psi_S [MPa]$	$Pro \ [\mu mol \ g^{\text{l}}(FM)]$	SPAD index
Tian Xuan 35	0	-1.28 ± 0.25^{Aa}	$0.60\pm0.02^{\mathrm{Ba}}$	$35.5 \pm 1.6^{\text{Ba}}$
	100	$-1.11\pm0.23^{\mathrm{Aa}}$	$0.99\pm0.07^{\rm Ca}$	$33.4 \pm 1.3^{\mathrm{ABa}}$
	150	$-3.69\pm0.29^{\mathrm{Cb}}$	$52.27 \pm 7.80^{\rm Cb}$	$25.9 \pm 0.8^{\mathrm{ABb}}$
	200	$-3.84\pm0.62^{\mathrm{Ab}}$	$54.23\pm4.36^{\mathrm{Ab}}$	$24.0 \pm 1.7^{\mathrm{Bbc}}$
	250	n.m.	$65.50 \pm 3.15^{\rm Ba}$	$24.2\pm1.3^{\rm Ac}$
Tian Xuan 90	0	$-1.42\pm0.51^{\mathrm{Aa}}$	$0.92 \pm 0.53^{\mathrm{Bc}}$	$35.5\pm1.6^{\mathrm{Ba}}$
	100	$-1.35\pm0.06^{\mathrm{Aa}}$	$1.34\pm0.25^{\text{Cc}}$	$35.6\pm0.9^{\rm Aa}$
	150	$-2.00\pm0.48^{\mathrm{Aa}}$	$73.47 \pm 12.98^{\rm Ab}$	$24.2 \pm 2.8^{\mathrm{BCb}}$
	200	n.m.	n.m.	$24.0 \pm 0.7^{\mathrm{Bb}}$
	250	n.m.	$143.19 \pm 15.21^{\mathrm{Aa}}$	$24.8\pm1.2^{\rm Ab}$
Tian Xuan 97	0	$-1.26\pm0.13^{\mathrm{Aa}}$	$2.39 \pm 0.51^{\mathrm{Ac}}$	$32.6\pm0.9^{\rm Ca}$
	100	$-1.28\pm0.07^{\mathrm{Aa}}$	$6.73\pm2.00^{\rm Bc}$	$32.6\pm1.3^{\rm Ca}$
	150	n.m.	48.92 ± 3.41^{Cb}	$23.3\pm1.2^{\text{Cb}}$
	200	n.m.	n.m.	$23.3\pm2.2^{\rm Bb}$
	250	n.m.	$89.09\pm7.58^{\mathrm{Ba}}$	$21.3\pm2.8^{\mathrm{Ab}}$
Dalishi	0	$-0.96\pm0.27^{\mathrm{Aa}}$	$0.73\pm0.16^{\rm Bc}$	$38.6 \pm 1.7^{\mathrm{Aa}}$
	100	$-1.76\pm0.19^{\mathrm{Bb}}$	$9.69 \pm 3.02^{\mathrm{Bb}}$	$32.7\pm1.4^{\text{Cb}}$
	150	$-2.41\pm0.37^{\mathrm{ABc}}$	$25.88\pm3.37^{\mathrm{Da}}$	$26.8 \pm 1.4^{\rm Acd}$
	200	$-2.73\pm0.41^{\mathrm{Ac}}$	$29.78\pm4.47^{\mathrm{Ba}}$	$26.9 \pm 2.6^{\mathrm{ABcd}}$
	250	$-4.65 \pm 0.36^{\rm d}$	n.m.	$24.0 \pm 3.2^{\mathrm{Ad}}$
Rio	0	$-0.98\pm0.23^{\mathrm{Aa}}$	$1.23\pm0.27^{\rm Be}$	$35.7 \pm 1.1^{\mathrm{Ba}}$
	100	$-2.23\pm0.12^{\mathrm{Cb}}$	$16.57\pm2.34^{\mathrm{Ad}}$	$33.3\pm3.2^{\mathrm{ABa}}$
	150	$-2.07\pm0.39^{\mathrm{Ab}}$	$32.71\pm6.68^{\mathrm{Dc}}$	$26.2 \pm 0.9^{\mathrm{ABb}}$
	200	$-3.52\pm0.41^{\mathrm{Ac}}$	$57.16 \pm 12.84^{\mathrm{Ab}}$	$26.2 \pm 3.8^{\mathrm{ABb}}$
	250	n.m.	$79.42\pm7.34^{\mathrm{Ba}}$	$24.4 \pm 3.3^{\mathrm{Ab}}$
Sutian-1	0	$-1.03\pm0.17^{\mathrm{Aa}}$	$1.05\pm0.21^{\rm Bd}$	$34.1 \pm 2.3^{\mathrm{BCa}}$
	100	$-2.11\pm0.12^{\mathrm{Cb}}$	$1.41\pm0.28^{\mathrm{Cd}}$	$32.6\pm2.0^{\rm Ca}$
	150	$-2.77\pm0.18^{\mathrm{Bb}}$	$10.25 \pm 1.99^{\rm Bc}$	$26.1\pm1.1^{\rm ABb}$
	200	$-2.87\pm0.72^{\mathrm{Ac}}$	$25.62\pm6.91^{\mathrm{Bb}}$	$27.8 \pm 2.9^{\mathrm{Ab}}$
	250	n.m.	$78.57 \pm 14.85^{\rm Ba}$	$22.4\pm1.6^{\rm Ac}$

Table 3. Accumulation of Na⁺ and K⁺ in sorghum genotypes treated by different concentration of NaCl in hydroponic solution.

Genotype	NaCl [mM]	Root content Na ⁺ [mg g ⁻¹]	Leaves content K ⁺ [mg g ⁻¹]	Na+/K+	Na ⁺ [mg g ⁻¹]	K^+ [mg g ⁻¹]	Na+/K+
Tian Xuan 35	0	5.94	38.40	0.15	5.57	44.96	0.12
	150	22.86	36.52	0.63	36.57	26.26	1.39
Tian Xuan 90	0	3.01	30.22	0.10	2.79	27.59	0.10
	150	26.99	21.69	1.24	39.94	24.90	1.60
Tian Xuan 97	0	9.05	28.11	0.32	9.75	24.71	0.39
	150	38.84	22.20	1.75	43.94	32.40	1.36
Dalishi	0	1.51	45.92	0.03	0.49	49.82	0.01
	150	33.04	32.11	1.03	27.56	24.93	1.11
Rio	0	6.43	41.41	0.16	2.83	28.49	0.10
	150	23.05	20.73	1.11	27.93	30.82	0.91
Sutian-1	0	2.28	46.29	0.05	8.43	24.24	0.35
	150	28.60	33.15	0.86	37.62	27.77	1.35

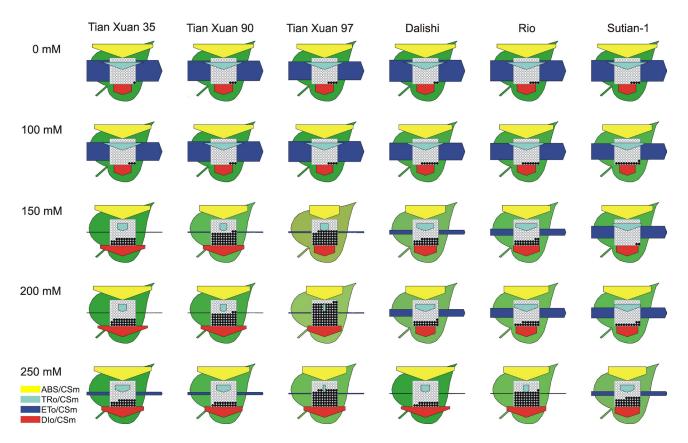


Fig. 5. Leaf pipeline model showing the proportion of phenomenological energy fluxes within a leaf calculated per excited cross section (CS_m). Models are drawn with *Biolyzer* software. The value of each parameter can be seen in relative changes in width of each arrow (see the color legend). Active RCs are shown as open circles and inactive RCs are closed circles. ABS/ CS_m – absorbtion per excited cross section; DI_0/CS_m – effective dissipation of energy per excited cross section; ET_0/CS_m – electron transport flux from reduced Q_A to Q_B per excited cross section; TR_0/CS_m – maximal trapping rate of absorbed photons per excited cross section.

with previous studies on sorghum and other plants (Al-Karaki *et al.* 1996, Iqbal *et al.* 2014, Hmidi *et al.* 2018). Interestingly, Tian Xuan 97 at 250 mM NaCl showed the maximum amount of proline, which might indicate the hyperactive protective mechanism of the plant under such a high salt concentration. SPAD index varied slightly for different genotypes under the same salt concentration indicating its lesser sensitivity for the identification of salt stress, while the variation in proline and osmotic potential were significant between genotypes, which indicated that these parameters are more reliable for the given purpose, *i.e.*, to assess the salt stress in comparison to SPAD index.

The concentration of Na^+ and K^+ ion was measured for 150 mM salt concentration; Na^+ concentration was higher in leaves of salt-sensitive genotypes, whereas comparatively lower in leaves of salt-resistant genotypes. The concentration of Na^+ was similar in roots of salt-resistant and salt-sensitive sorghum genotypes, indicating that the transportation of Na^+ in salt-resistant genotypes was significantly reduced, thus indicating the resistant mechanism for salt-resistant genotypes.

The study clearly indicated how JIP-test can be used to identify the salt-resistant genotypes of sorghum, with an indication of the sensitive photosynthetic parameters. Compared to chlorophyll fluorescence, the chlorophyll concentration was found to be a less sensitive parameter for identifying the susceptible individuals. In addition, our results suggest that the regulation of ion transport may have an important role for salt resistance in sorghum genotypes.

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