INTERACTIVE EFFECT OF SALINITY AND POTASSIUM ON GROWTH, BIOCHEMICAL PARAMETERS, PROTEIN AND OIL QUALITY OF SOYBEAN GENOTYPES

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Potassium (K⁺) nutrition ensures growth and quality of crops both under normal and saline soils. In view of that a study was conducted to assess the impact of various doses of K⁺ on growth, bio-chemicals, protein and oil contents of four soybean (*Glycine max* L.) genotypes under saline condition. Soybean genotypes were grown in pots having treatment combination of salinity (control, 6 and 12 dS m⁻¹) and K⁺ doses (control, 50 and 75 kg ha⁻¹). The results showed that sufficient K⁺ nutrition enhanced plant growth, biomass production, protein and oil percentage of soybean genotypes both in normal and saline condition. Water relation in plant as well as chlorophyll contents and leaf area improved with the application of potassium under salinity stress. Across K⁺ application rates, all soybean genotypes significantly varied from each other. K⁺ accumulation was positively correlated with growth and other parameters of soybean genotypes, but these relationships were stronger in case of genotypes No.62 and No 13 as compared to Ajmeri and William-82 genotypes. Soybean genotypes No. 62 and No. 13 classified as potassium efficient genotypes while Ajmeri and William-82 categorized as potassium inefficient genotypes on the basis of growth and K uptake.

Keywords: Salinity, Potassium, Protein, antioxidant enzymes, Oil, Soybean

INTRODUCTION

Soybean is one of the most important oil seed crop that is multi-dimensional in its uses however it contains the best quality protein due to which it is being called as "the meat that is grown on plant". The seeds of soybean contain 50% good quality protein, 17-24% highly palatable oil containing zero cholesterol level, 6% ash, 29% sugars (Essa and Al-Ani, 2001; Agarwal, 2007; Shi and Cai, 2010). Except above all benefits, it is also a good source of polyunsaturated fat, fibers, vitamins and contains a small amount of minerals and energy (Krishnan, 2001). It contains about 85% of the world oil seed and placed in group of the oil seed crop. Soybean seed is processed into soybean meal and vegetable oil. Soybean may be a significant source for biofuel production (Anonymous, 2009). In 2009-2010 the total demand of food oil in Pakistan was 4.125 million tons and about 65% of the country requirement of edible oil was met through imports (Akinori et al., 2000; Balasubramaniyan et al., 2001; Anonymous, 2012).

Agriculture sector is facing various threats today, one of the major stresses among all is salinity which is caused due to high temperature, less rainfall, poor quality water and soil management and eventually high evapotranspiration (Neto *et al.*, 2006; Flowers and Colmer, 2008). The affected area due to salinity in all over the world is about 45 m ha of irrigated regions and about 1.5 m ha cannot give productions due to

high salinity (Munns and Tester, 2008). In Pakistan the cultivated land of 6.67 m ha is saline among which Punjab province covers an area of about 2.67 m ha (Ghafoor *et al.*, 2004). In world over, salinity has damaged about 7% of agricultural area, out of which 3% land is considered highly saline and eventually this area is expected to increase up to 20% in the future.

Continuous reduction of cultivable area due to salinity and increasing population of the world is causing immense pressure on agriculture to ensure food security. Therefore, the current decade's explorations are going to emphasize the response of plants in salt stress (Zora, 2006). Plant species show different behavior in salt stress in order to exclude salts from their cells or to endure their presence within the cells by involving many kinds of physiological and biochemical changes. Generally, there are two approaches for increasing the production of crops from salt affected soils: first reclamation of these soils and second by introduction of salt tolerant genotypes (Blumwald et al., 2004; Yilmaz et al., 2004). The former approach is not practicable due to wide salt-affected areas, insufficient availability of good quality water, soil permeability and high cost of amendments (Akhtar et al., 2010). Therefore, we are left with choice of introducing salt tolerant genotypes in these problematic soils and also finding the options to improve salt tolerance to certain levels that can minimize all these problems (Iqbal et al., 2007). It is observed that the salt sensitive plants does not maintain a higher K⁺/Na⁺ ratio, as it is needed while salt tolerant plants conserve a high K⁺ content (Rejili *et al.*, 2007). With the increasing concentration of salts in the plants (stem tissues) of *Spina Christi Z*.an increased amount of N and P were observed. However, salinity caused by NaCl had hardly been the source of reducing K⁺/Na⁺ ratio in different parts of plants (Sohail *et al.*, 2010).

Potassium is one of the major components of plant tissue which makes about 1-10% of dry matter (Epstein and Bloom, 2005). Potassium (K⁺) plays a fundamental role in balancing turgor potential, membrane potential, stoma movement, tropisms and enzymes activation (Cherel, 2004). Both elements potassium and sodium are found together in the soil solution. Both of these possess positive charge on them and hence during reciprocal absorption and translocation, they employ antagonistic or synergistic effects under salinity stress (Hussain *et al.*, 2013). Eventually, maintained amount of K⁺/Na⁺ ratio is precisely critical for the working stomatal function, for the synthesis of protein, cell osmoregulation, activation of enzymes, and turgor maintenance (Shabala *et al.*, 2003).

Application of K^+ under salinity lowers the creation of reactive oxygen species while activities of antioxidative enzymes containing catalase, superoxide dismutase and peroxidase enhanced during stress (Liang *et al.*, 2007; Abbasi, *et al.*, 2015). Increased accumulation of K^+ in salinity stressed plants is important for maintenance of increased cellular K^+/Na^+ ratio. Supplementary K^+ can reduce the adverse effects of increasing salinity (Kavitha *et al.*, 2012). Soybean responds positively to potassium application in regulation of oil and protein contents in seeds (Tiwari *et al.*, 2002).

In view of above discussions, the current research work was conducted with objectives to investigate the behavior of K^+ application on the performance, antioxidant enzymes, protein and oil quality of soybean genotypes during salinity stress.

MATERIALS AND METHODS

Experimental conditions: A pot study was conducted in University of Agriculture, Faisalabad-Pakistan by using four soybean genotypes (*Glycine max* L.) including; No. 62, No. 13, William and Ajmeri. In blank earthen pots filled with pulverized 12 kg soil (EC=1.5 dS m⁻¹, pH=8.23, SAR=6.74 (mmol L⁻¹)^{1/2}) soil texture=sandy clay loam) after mixing with required fertilizer and potassium dose, the soybean seeds were sown at 2 cm depth and their response was determined under different levels of NaCl (control, 6 and 12 dS m⁻¹) alone and in combination with K₂SO₄.Seven treatments were applied to soybean viz. T1: Control, T2: EC 6 dS m⁻¹, T3: EC 12 dS m⁻¹, T4: EC 6 dS m⁻¹ + 50 kg ha⁻¹ K⁺, T5: EC 6 dS m⁻¹ + 75 kg ha⁻¹ K⁺, T6: EC 12 dS m⁻¹ + 50

kg ha⁻¹ K⁺, T7: EC 12 dS m⁻¹ +75 kg ha⁻¹ K⁺. The experimental design was CRD using factorial arrangement with four replications. The total period of the experiment was 4 months. The tap water was used for irrigation (EC= 0.88 dS m^{-1} , RSC= $0.75 \text{ me } \text{L}^{-1}$, SAR= 2.5) to the pots whenever irrigation necessary.

Following parameters were determined: Relative water contents (RWC) (Sairam *et al.*, 2002), membrane stability index (MSI) (Sairam *et al.*, 2002), gas exchange parameters (Infrared gas analyzer), chlorophyll contents (SPAD chlorophyll meter), leaf area (leaf area meter), antioxidant enzymes (Giannopolitis and Ries, 1977; Chance and Maehly, 1955) and oil and protein contents (Hymowitz *et al.*, 1974) and plants were harvested for further analysis.

Statistical Analysis: All information presented in this experiment was means of four replications. Results were interpreted by using a statistical package, statistix 8.1®.

RESULTS

Growth parameters: Salinity stress reduced fresh and dry biomass in all soybean genotypes, while application of potassium was helpful to survive during stress (Fig. 1). In all soybean genotypes salinity stress (12 dS m⁻¹) caused reduction in shoot fresh (LSD=1.57) and dry weight (LSD=0.62), root fresh (LSD=0.93) and dry weights (LSD=0.10) at 5% probability level. Among all treatments the treatment with maximum K^+ dose (9 mM) with 6 dSm⁻¹ salinity stress (NaCl) performed better after control and showed maximum shoot fresh weight (31.84 g), shoot dry weight (7.86 g), root fresh weight (7.86 g) and root dry weight (1.56 g) in No. 62 followed by No.13 while minimum shoot fresh weight (8.77 g), shoot dry weight (2.58 g), root fresh weight (2.58 g) and root dry weight (0.42 g) was observed in Ajmeri followed by William-82. Among all treatments the treatment with K⁺ dose (9 mM) at salinity 6 dSm⁻¹ performed better after control and showed maximum shoot fresh and dry weights. However, K⁺ application improved all plant growth attributed in all soybean genotypes while best response towards K⁺ application was detected in soybean genotype No. 62. Moreover, in all soybean genotypes 75 kg K⁺ ha⁻¹ dose was found better at 6 dS m⁻¹ salinity in reducing the adverse effects of salinity. Effect of salinity and potassium application on shoot and root length (LSD= 3.70 and 1.76 respectively) at 5% probability level in four soybean genotypes shown in (Figure 1). Accumulation of NaCl caused significant reduction in shoot length and root length in all soybean genotypes than control. The greater reductions in these factors were observed in salt sensitive soybean genotypes than salt tolerant soybean genotypes on the basis of potassium use efficiency. But, potassium application considerably minimized the negative effects of salinity both in salt tolerant and salt sensitive soybean genotypes especially at higher 75 kg ha⁻¹ K⁺ level.



Figure 1. Effect of salinity and potassium application on growth of soybean genotypes. (T1: Control, T2: EC 6 dS m⁻¹, T3: EC 12 dS m⁻¹, T4: EC 6 dS m⁻¹ + 50 Kg ha⁻¹ K⁺, T5: EC 6 dS m⁻¹ + 75 Kg ha⁻¹ K⁺, T6: EC 12 dS m⁻¹ + 50 Kg ha⁻¹ K⁺, T7: EC 12 dS m⁻¹ + 75 Kg ha⁻¹ K⁺)

Leaf area, chlorophyll contents, MSI and RWC: Leaf area, chlorophyll contents, relative water contents and membrane stability index expressed in Fig. 2. Reduction in leaf area (LSD= 7.13), chlorophyll contents (LSD=2.89), membrane stability index (LSD=4.46), and relative water contents (LSD=3.89) parameters at 5% probability was recorded in all soybean genotypes with induced salinity (6 and 12 dS m⁻¹) levels. Among all the treatments which were used, reduction was maximum in pots which were supplied with combinations of maximum dose of K⁺ and NaCl (K⁺75 kg

ha⁻¹ +12 dS m⁻¹ NaCl). In response of treatment 6 dS m⁻¹NaCl induced salinity with 9mM K⁺ application No. 62 exhibited percent of control 94. 99 and 97% followed by No. 13 with 91, 95 and 96% while Ajmeri showed 63, 81, 80% followed by William-82 with 67, 83 and 83% in leaf area, relative water contents and membrane stability index respectively. Reduction was maximum in potassium inefficient soybean genotypes (Ajmeri and William-82) than potassium efficient genotypes (No. 62 and No. 13). Plants showed positive improvement in leaf area, chlorophyll



Figure 2. Effect of salinity and potassium application on leaf area, chlorophyll contents and leaf water relations. (T1: Control, T2: EC 6 dS m⁻¹, T3: EC 12 dS m⁻¹, T4: EC 6 dS m⁻¹ + 50 Kg ha⁻¹ K⁺, T5: EC 6 dS m⁻¹ + 75 Kg ha⁻¹ K⁺, T6: EC 12 dS m⁻¹ + 50 Kg ha⁻¹ K⁺, T7: EC 12 dS m⁻¹ + 75 Kg ha⁻¹ K⁺)

contents, relative water contents and membrane stability index at 5% probability level in response to K⁺ application and the level of enhancement was better in salt tolerant soybean genotypes than salt sensitive. The reaction of maximum dose of K⁺ application with 12 dS m⁻¹ NaCl was either negligible or bad in all soybean genotypes.

Gas exchange parameters: Gas exchange parameters were determined in response to salinity and potassium application. Reduction was greater at maximum salinity in photosynthetic rate (LSD=1.48), transpiration rate (LSD=0.89), stomatatal conductance (LSD=0.04), and substomatal conductance (LSD=5.41) at 5% probability level while potassium application was useful in improving salinity tolerance (Fig. 3.). The improvement of potassium was relatively less with 12 dS m⁻¹ salinity stress (NaCl) than

with 6 dS m⁻¹ salinity stress (NaCl). Soybean genotype No. 62 showed percent of control 87, 90, 98 and 94% followed by No. 13 with 82, 88, 97 and 93% while Ajmeri showed 56, 61, 74 and 71% followed by William-82 with 59, 63, 77 and 70% in photosynthesis rate, transpiration rate, stomatal conductance and intercellular CO₂ Conc. respectively in combined application of 6 dS m⁻¹ salt stress with 75 kg ha⁻¹ K.However maximum reduction in all soybean genotypes was measured at 75 kg ha⁻¹ K⁺ and 12 dS m⁻¹ (NaCl) combine application. The reaction of all the soybean genotypes was dissimilar due to their diverse genetic potential of salt tolerance. No. 62 followed by No. 13 presented fewer declines in gas exchange parameters in response to salt stress than with Ajmeri and William-82.



Figure 3. Effect of salinity and potassium application on gas exchange parameters of soybean genotypes. (T1: Control, T2: EC 6 dS m⁻¹, T3: EC 12 dS m⁻¹, T4: EC 6 dS m⁻¹ + 50 Kg ha⁻¹ K⁺, T5: EC 6 dS m⁻¹ + 75 Kg ha⁻¹ K⁺, T6: EC 12 dS m⁻¹ + 50 Kg ha⁻¹ K⁺, T7: EC 12 dS m⁻¹ + 75 Kg ha⁻¹ K⁺)

Improvement of gas exchange parameters was more at higher dose of K at 6 dS m⁻¹ salinity stress.

Antioxidants enzymes: Antioxidant enzymes were measured in all soybean genotypes under induced salt stress (NaCl) and potassium application. The activity of SOD was positively enhanced in salt tolerant soybean genotypes (No. 62 and No.13) but no improvement was measured in salt sensitive soybean genotypes (Ajmeri and William-82) at 6 dS m⁻¹ salt stress alone and also with application of K (Table 1.) At maximum salinity level, SOD activity reduced in all Ajmeri and William-82 than control. The activities of POD and CAT were reduced at low salinity level than control in all soybean genotypes. But at low salinity level, CAT and POD activities reduced but reduction was less in salt tolerant soybean genotypes than salt sensitive soybean genotypes at high salinity level. Application of K^+ improved CAT activity in salt tolerant soybean genotypes in both salinity and K^+ levels. Obvious increment in activities of SOD and POD were recorded at low salinity level at high potassium level (75 kg ha⁻¹ K) in soybean genotypes. However, no major effect of application of K⁺ was perceived on SOD, POD and CAT activities in salt sensitive soybean genotypes.

Soybean yield, protein and oil: Soybean yield, Protein and oil concentration of soybean genotypes increased with succeeding seed development under saline and non-saline conditions. Under control or normal conditions higher

Varieties	Treatments	SOD (unit mg ⁻¹ of protein)	POD (unit mg ⁻¹ of protein)	CAT (unit mg ⁻¹ of protein)
No. 62	T1	51.43±1.04hi	70.75±0.85e	10.23±0.73c
	T2	62.11±1.06 (121)d	82.57± (117)c	9.82±0.70 (96)cd
	Т3	54.75±0.87 (106)fg	66.71± (94)e	8.38±0.60 (82)fgh
	T4	67.31±0.71 (131)b	89.63± (127)ab	11.45±0.82 (112)ab
	T5	71.52±0.72 (139)a	91.39± (129)a	12.17±0.87 (119)a
	T6	55.81±0.90 (109)ef	69.26± (98)e	8.69±0.62 (85)efg
	Τ7	49.87±1.25 (97)hij	61.44± (87)f	8.03±0.31 (79)fghijk
No. 13	T1	49.27±1.18ij	67.75±1.11e	9.58±0.22cde
	T2	58.51±0.26 (119)e	75.64±0.60 (112)d	9.00±0.20 (94)def
	Т3	51.38±0.90 (104)hi	59.58±1.42 (88)fg	7.85±0.18 (82)ghijkl
	T4	63.52±1.72 (129)cd	86.78±0.89 (128)bc	10.53±0.24 (110)bc
	T5	65.64±1.16 (133)bc	91.07±2.28 (134)ab	11.49±0.26 (120)ab
	T6	52.50±0.89 (107)gh	62.18±1.05 (92)f	8.23±0.19 (86)fghi
	Τ7	47.31±0.85 (96)jk	56.75±1.38 (84)gh	7.21±0.14 (75)jklm
William 82	T1	40.17±0.81mn	45.25±1.11jk	7.45±0.15hijklm
	T2	45.04±0.68 (112)kl	49.67±1.66 (110)ij	7.65±0.10 (103)hijkl
	T3	39.15±0.52 (97)nop	37.62±1.24 (83)1	5.07±0.10 (68)no
	T4	47.62±0.73 (119)jk	47.62±2.46 (105)jk	8.12±0.16 (109)fghij
	T5	50.82±1.09 (127)hi	52.86±0.80 (117)hi	8.42±0.17 (113)fgh
	T6	39.78±1.70 (99)no	37.84±1.50 (84)1	5.14±0.10 (69)n
	Τ7	36.24±1.03 (90)q	34.67±1.11 (77)lm	4.77±0.12 (64)no
Ajmeri	T1	37.09±0.880pq	44.25±1.11k	6.55±0.13m
-	T2	40.97±1.08 (110)mn	46.82±1.30 (106)jk	6.87±0.12 (105)lm
	Т3	35.51±0.87 (96)qr	30.91±0.99 (70)mn	4.39±0.09 (67)no
	T4	42.62±1.03 (115)lm	46.23±2.07 (104)jk	7.07±0.14 (108)klm
	T5	44.29±0.89 (119)1	47.26±1.41 (107)jk	7.34±0.15 (112)ijklm
	T6	36.92±0.64 (100)pq	34.76±1.53 (79)lm	4.52±0.09 (69)no
	T7	33.38±0.73 (90)r	28.78±0.65 (65)n	4.13±0.08 (63)o

Table 1. Effect of salt treatment and potassium application on superoxide dismutase (SOD), peroxidase (POD) and catalase (CAT) of soybean genotypes

T1: Control, T2: EC 6 dS m^{-1} , T3: EC 12 dS m^{-1} , T4: EC 6 dS m^{-1} + 50 Kg ha⁻¹ K⁺, T5: EC 6 dS m^{-1} + 75 Kg ha⁻¹ K⁺, T6: EC 12 dS m^{-1} + 50 Kg ha⁻¹ K⁺, T7: EC 12 dS m^{-1} + 75 Kg ha⁻¹ K⁺.

Each value is an average of 4 replicates \pm S.E., and values in parenthesis are the percent of control.

contents of protein and oil contents were measured while under saline condition reduction was higher. In improving the protein (LSD= 2.20) and oil concentration (LSD=0.57) at 5% probability level the potassium application was helpful in all soybean genotypes but smallest enhancement was measured at maximum dose of K with highest salinity level. Though both protein and oil concentration reduced with increasing salinity (NaCl) extent (Table 2.).

DISCUSSION

Genetic differences in plants provide a valued tool in assortment of genotypes with required traits (Misra and Dwivedi, 2004). It is noticeable that salinity stress adversely affects all the morphological, ionic, biochemical and physiological parameters (Sairam *et al.*, 2002; Zhu, 2002;

Hajer et al., 2006; Chen et al., 2013; Abbasi et al., 2015). Potassium application significantly enhanced plant growth, gas exchange factors, K⁺/Na⁺ ratio and antioxidant enzymes activities in all soybean genotypes and minimized the adverse effects of salt stress (Chen et al., 2007; Zheng et al., 2008; Abbasi et al., 2014). The results also showed that soybean genotype No. 62 and No. 13 are comparatively more salt tolerant and potassium efficient than soybean genotypes Ajmeri and William-82. There was significant reduction of plant biomass, leaf area, chlorophyll contents, MSI, RWC, antioxidant enzymes, protein and oil contents in all soybean genotypes at all the levels of salinity. These findings are also comparable with previous results which confirmed that salinity caused reduction in plant growth due to osmotic stress, ionic toxicity and less cell wall extensibility (Hajer et al., 2006; Abbasi, et al., 2015). The

Varieties	Treatments	Pods per plant (Yield)	Protein (% DM)	Oil (% DM)
No. 62	T1	17.25±1.10a	45.56±0.74a	19.01±0.62a
	T2	11.75±0.85 (68)bc	40.21±0.75 (88)de	18.06±0.59 (95)bcd
	T3	7.88±0.43 (46)fghi	34.74±0.65 (76)hi	17.11±0.56 (90)gh
	T4	12.25±0.63 (71)bc	42.10±0.64 (92)cd	18.25±0.60 (96)bc
	T5	13.00±0.41 (75)b	43.34±0.39 (95)bc	18.44±0.60 (97)ab
	T6	8.75±0.75 (51)fgh	39.37±0.57 (86)ef	17.49±0.57 (92)defg
	T7	7.75±0.63 (45)fghi	32.06±0.87 (70)klmn	16.92±0.55 (89)ghi
No. 13	T1	16.00±0.91a	44.56±1.28ab	18.93±0.26a
	T2	11.00±1.08 (69)cde	38.32±1.10 (86)ef	17.79±0.24 (94)cdef
	T3	7.13±0.52 (45)ghij	32.97±0.94 (74)ijkl	16.84±0.23 (89)hi
	T4	11.50±0.65 (72)bcd	40.10±1.15 (90)de	17.98±0.24 (95)bcde
	T5	11.88±0.43 (74)bc	42.19±1.06 (95)cd	18.17±0.25 (96)bc
	T6	8.00±0.71 (50)fgh	35.93±0.50 (81)gh	17.22±0.23 (91)fgh
	T7	7.00±0.41 (44)hijk	30.30±0.87 (68)mno	16.65±0.23 (88)hij
William 82	T1	15.75±0.85a	44.90±0.77ef	17.83±0.35cde
	T2	8.75±0.48 (56)fgh	35.02±0.60 (78)lmno	16.04±0.31 (90)kl
	Т3	5.68±0.71 (36)jk	30.09±0.52 (67)q	15.15±0.30 (85)mn
	T4	9.25±0.63 (59)ef	37.27±0.64 (83)jklm	16.22±0.32 (91)jk
	T5	9.63±0.62 (61)def	39.51±0.68 (88)hij	16.40±0.32 (92)ijk
	T6	6.00±0.41 (38)ijk	33.68±0.58 (75)op	15.33±0.30 (86)m
	T7	5.43±0.59 (34)jk	27.39±0.47 (61)r	14.97±0.29 (84)mno
Ajmeri	T1	15.75±0.85a	40.21±0.46ab	17.47±0.35efg
	T2	8.50±0.50 (54)fgh	30.84±0.85 (77)hi	15.37±0.31 (88)m
	T3	5.43±0.83 (34)jk	26.22±0.72 (65)no	14.50±0.29 (83)op
	T4	9.00±0.41 (57)fg	32.39±0.89 (81)fg	15.37±0.31 (88)m
	T5	9.38±0.47 (60)ef	34.32±0.94 (85)e	15.55±0.31 (89)lm
	T6	5.75±0.48 (37)jk	29.30±0.81 (73)ijk	14.67±0.30 (84)nop
	T7	5.18±0.69 (33)k	23.90±0.66 (59)pq	14.32±0.29 (82)p

Table 2. Effect of salt treatment and potassium application on yield, protein and oil contents of soybean genotypes.

T1: Control, T2: EC 6 dS m^{-1} , T3: EC 12 dS m^{-1} , T4: EC 6 dS m^{-1} + 50 Kg ha⁻¹ K⁺, T5: EC 6 dS m^{-1} + 75 Kg ha⁻¹ K⁺, T6: EC 12 dS m^{-1} + 50 Kg ha⁻¹ K⁺, T7: EC 12 dS m^{-1} + 75 Kg ha⁻¹ K⁺.

Each value is an average of 4 replicates ±S.E., and values in parenthesis are the percent of control.

adverse effects of NaCl stress was significantly reduced with potassium application also plant growth and development in soybean genotypes improved. That was accomplished to incompatible effects of K⁺ with Na⁺(Lynch and Lauchli, 1985; Abbasi *et al.*, 2014). Similarly, improvement in plant growth and dry mass assembly with potassium application was stated in different crops e.g. rice (Bohra and Doerffling, 1993), maize (Abbasi *et al.*, 2014) in saline condition.

Relative water contents and membrane stability index are the chief factors which measure the extent of tolerance in plant and with increasing salt stress that were reduced significantly (Noreen and Ashraf, 2010). Under salinity stress, reduction in RWC in all soybean genotypes proved the previously reported consequences (Sairam *et al.*, 2002; Abbasi *et al.*, 2014). Potassium application improved relative water contents and membrane stability index under

NaCl stress because water uptake improved with potassium application for turgidity maintenance in different crops (Subbarao *et al.*, 2000a). Salt stress significantly reduced leaf area while application of potassium increased leaf area in soybean genotypes as also stated by Ayub *et al.* (2012) in cluster beans.

It is well documented that the key factor for growth and final biomass production is photosynthetic rate of plants. The adverse effects of salinity minimize photosynthetic rate significantly by reduction in photosynthetic rate (A), transpiration rate (E), stomatal conductance (Gs), internal CO_2 concentration (Ci) and chlorophyll contents (Chen *et al.*, 2013). Significant reduction in gas exchange attributes resulted with salinity stress in all soybean genotypes but application of potassium improved all gas exchange parameters and the best performance was documented at

maximum dose of K with 6 dS m⁻¹ salinity level. Reduction in gas exchange parameters was greater in potassium inefficient soybean genotypes (Ajmeri and William-82) under saline condition while potassium efficient soybean genotypes showed minimum reduction in gas exchange parameters.

Salinity stress produces reactive oxygen species that cause damage in intera cellular structure which results reduction in photosynthetic rate (Chen and Heuer, 2013). Antioxidant enzymes provide protection to cell structures and organelles against ROS under salt stress condition (Reddy et al., 2004). In the present study, significant increase in the activities of SOD, POD, and CAT were recorded in all soybean genotypes which formed as a result of oxidative stress but higher activities were recorded with higher dose of potassium (9 mM) with EC 6 dS m⁻¹. In scavenging reactive oxygen species SOD is the main enzyme deliberated to be the first line of defense against ROS (Hamilton and Heckathorn, 2001) which change O^{2-} to H_2O_2 the less noxious form of oxygen (Costa et al., 2005). The CAT and POD abolish the H₂O₂ produced by SOD into other less toxic formulae (Foyer et al., 1994). Comparatively greater activities of antioxidant enzymes have been recorded in soybean genotypes No. 62 and No. 13 than Ajmeri and William-82. Consequently, soybean genotypes react contrariwise as a result of differences in their antioxidant systems, salt stress and application of potassium (Mohammadkhani and Heidari, 2007; Nawaz and Ashraf, 2007). Nevertheless, potassium application under salt stress further upgraded the activities SOD, POD and CAT in all soybean genotypes. These result also comparable to earlier argument that under salinity stress potassium application significantly improved antioxidant activities in plants (Zheng et al., 2008; Soleimanzadeh et al., 2010; Abbasi et al., 2015).

Oil percentage and protein contents per unit area, increased with adequate potassium nutrition also cost of production decreased and yield and profit increased (Khan et al., 2010). Substantial effect was verified in growth and yield increment by application of potassium in soybean genotypes and similar findings were examined in sunflower and safflower (Bakht et al., 2006; Gerendás et al., 2008; Asadia, 2010). Availability of nitrogen to plants is upgraded by phosphorus and potassium application with increasing the biological nitrogen fixation that is used for the production of crude protein in cluster bean (Ayub et al., 2012). Similar consequences were found in previous studies in several crop species e.g. lucerne (Patel and Kotecha, 2006), mungbean (Hussainand Arshad, 2011). Reducing protein with higher salinity and this might be due to disruption in nitrogen metabolism or reticence of nitrate uptake due to decline of absorbed water and fewer root permeability (Medhat, 2002). In present study all the soybean seeds under control conditions showed higher concentration of protein and oil as

compared to NaCl salt stress and this was associated with production of larger grain size of soybeans as compared to saline soils and similar findings were verified by (Ghassemi-Golezani *et al.*, 2009). Decrease in protein and oil yield per plant with higher salinity usually related with reduction in the interval of protein and oil accumulation and grain yield per plant under saline condition (Table 2.) but the amount of protein and oil accumulation of Ajmeri and william-82 was slightly less than No. 62 and No. 13. Higher grain, oil and protein per plant of No. 62 and No. 13 soybean genotypes were due to production of more grains per plant than other soybean genotypes as also stated by (Ghassemi-Golezani *et al.*, 2009).

Conclusion: Salt stress strictly reduced plant growth and yield by disturbing plant morphological, physiological, protein and oil contents of all soybean genotypes. Effects of salinity stress were more distinct on soybean genotypes Ajmeri and William-82 than No. 62 and No. 13 soybean genotypes. Though, potassium application significantly reduced adverse effects of salinity by humanizing the plant growth, gas exchange attributes, improving antioxidant enzyme activities, protein and oil quality of all soybean genotypes. Potassium dose (75 kg K⁺ ha⁻¹) was found more effective at EC 6 dS m⁻¹salt stress level in easing damaging effects of salinity. Potassium efficient soybean genotypes No. 62 and No. 13 produced high biomass, chlorophyll contents, antioxidant enzymes, gas exchange parameters under salt stress condition as compared to salt sensitive soybean genotypes Ajmeri and William-82.

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