Salicylic acid and potassium nitrate promote flowering through modulating the hormonal levels and protein pattern of date palm *Phoenix dactylifera* 'Sayer' offshoot

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Abstract: Salicylic acid enhances the flowering process in the plant by creating new proteins under salinity stress. The study was to determine the role of salicylic acid (500 ppm) and potassium nitrate (1500 ppm), on flowering of date palm 'Sayer' offshoots under salinity effect. Application of salicylic acid increased the number of clusters, the number of new leaves, the content of carbohydrates, ascorbic acid, indoleacetic acid, zeatin, gibberellin, and abscisic acid significantly under salinity compared with control. Although the measured parameters were the highest in plants treated with salicylic acid, there was no distinction among potassium nitrate treatment under saltwater, and salicylic acid treatment with saltwater. Salicylic acid and potassium nitrate treatment demonstrated some amazing contrasts in protein patterns in light of gel electrophoresis. Plants treated with salicylic acid with fresh water and with saltwater showed five and six protein bands, respectively, that differed in the molecular mass of one polypeptide compared to control with freshwater. However, there was a difference in the molecular mass of two polypeptides compared to control with salt water, which showed six bands. In contrast, potassium nitrate application showed five protein bands, whether with freshwater or with saltwater. The findings could facilitate to elucidate the flowering mechanisms in date palm.

Key words: abscisic acid; clusters number; zeatin; electrophoresis; gibberellin; indoleacetic acid Salicilna kislina in kalijev nitrat pospešujeta cvetenje stranskih poganjkov dateljeve palme (*Phoenix dactylifera* 'Sayer') z moduliranjem ravni hormonov in vzorca proteinov

Izvleček: Salicilna kislina pospešuje cvetenje preko tvorbe novih proteinov v razmerah slanostnega stresa. Namen raziskave je bil določiti vlogo salicilne kisline (500 ppm) in kalijevega nitrata (1500 ppm) na cvetenje stranskih poganjkov dateljeve palme ('Sayer') v razmerah slanosti. Uporaba salicilne kisline je značilno povečala število stranskih poganjkov, število novih listov, vsebnost ogljikovih hidratov, askorbinske kisline, indolocetne kisline, zeatina, giberelina in abscizinske kisline v razmerah slanosti v primerjavi s kontrolo. Čeprav so imeli vsi merjeni parametri največje vrednosti pri obravnavanju s salicilno kislino, ni bilo razlike v obravnavanjih s kalijevim nitratom in salicilno kislino v istih razmerah. Obravnavanja s salicilno kislino in kalijevim nitratom so imela velike razlike v vzorcu proteinov, določenem z gelsko elektroforezo. Rastline, ki so bile tretirane s salicilno kislino, sladko in slano vodo so imele pet oziroma šest proteinskih trakov, ki so se razlikovali v molekulski masi enega izmed polipeptidov v primerjavi s kontrolo, kjer je bilo samo obravnavanje s sladko vodo. Kakorkoli, v primerjavi s kontrolo, kjer je bilo obravnavanje samo s slano vodo in je bilo šest proteinskih trakov, je bila razlika v molekulski masi dveh polipeptidov. Obravnavanje samo s kalijevim nitratom je pokazalo samo pet proteinov, ne glede na obravnavanje s slano ali sladko vodo. Ti izsledki bi lahko pomagali razjasniti mehanizem cvetenja pri dateljevi palmi.

Ključne besede: abscizinska kislina; število poganjkov; zeatin; elektroforeza; giberelin; indolocetna kislina

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1 INTRODUCTION

Date palm (Phoenix dactylifera L.) is usually exposed to abiotic stresses in an arid and semi-arid region. Notably, high temperatures, the shortage of irrigation water, and soil salinity are the real constraining variables to productivity (Allbed et al., 2017). The production of fruits is restricted in the extreme conditions. In any case, their development and yield profitability are reduced (Moustafa et al., 2018), where the yield reduced to half at 18 dS m⁻¹ and 100 % at 32 dS m⁻¹ (Elsadig et al., 2017). The production of date palm fruits relies upon the number of clusters and the success of the pollination process. Flowering in date palm is delayed when presented to unforgiving natural conditions. Date palm flowering is a complex process in which numerous natural factors and the physiological material of the plant interact. However, the transformation of the axillary buds to floral buds in the date palm relies upon a high sugar level and a low nitrogen C/N ratio. For this situation, the role of the ecological conditions includes a substantial impact on the regulation of this quantitative relation, particularly the impact of salinity, whether the salinity of the soil or water system conditions (Shareef, 2016). We have observed earlier that foliar application of salicylic acid on offshoots induced flowering in date palm under salt stress, especially at the early stage (Shareef et al., 2017).

Salicylic acid (2-hydroxybenzoic acid) (SA), is thought to activate general plant resistance ordinarily related to activation of defense genes (Tamaoki et al., 2013). Moreover, SA has a specific role in plant growth, induction of flowering, and uptake of ions. Thus, it has an essential role in the regulation of flowering (Wada & Takeno, 2013). The formation of the buds as a result of the SA application is unusual as SA changes the synthesis and signaling pathways of plant hormones, like jasmonic acid, ethylene, and auxin (IAA) (Vlot et al., 2009). Auxin, abscisic acid, and zeatin gradually decrease during flowering in date palm leaves in early varieties and then rise after flowering while the gibberellin (GA3) increased (Cheruth et al., 2015). Gibberellins accordingly are hormones responsible for dynamics in the survival and persistence of plants (Lymperopoulos et al., 2018). The effects on the phase of plant development and blossoming at the suitable season increased plant production. The interaction of environmental signals with endogenous biological process signals the main mechanisms which control flowering time in plant and are stimulated by rising SA (Zhang et al., 2018). Khayyat et al. (2018) found that the stimulation of the flowering of the saffron plant was successful using 2 mM SA or 1000 ppm potassium.

Under the water shortage conditions, leaf stomata do not function actively in plants with potassium deficiency and, therefore, cause excessive water loss (Farooq et al., 2015). Potassium (K) plays a vitally role in the physiological processes of photosynthesis, in the formation of carbohydrates and proteins, the transfer of water and nutrients, in use of nitrogen (N), and the stimulation of early plant growth (Lakudzala, 2013). In plant tissues, the transport of water, nutrients, and carbohydrates is enhanced by potassium application (Safar-Noori et al., 2018). Potassium ions induce gene expression in light of environmental changing (Zhang et al., 2018a). Gene expression drives the biological process and stress properties. It is, therefore responsible for subsequent translation modifications of the pure nucleus proteins and sometimes methylation of deoxyribonucleic acid (Zhang et al., 2018b). The process of flowering is controlled by a group of genes made up of gene expression (Yan et al., 2019).

This study aimed to check whether the exogenous use of salicylic acid (SA) or potassium nitrate (KNO_3) will positively influence the growth and formation of flower buds in the late flowering date palm.

2 MATERIALS AND METHODS

2.1 EXPERIMENT FIELD

The experiment was equipped at the General Authority of Palm station, Burjysia, Basrah, Iraq (latitude 30 ° 22 ' 6.294 "N and longitude 47 ° 36 ' 39.639 ") at 26 km of Basrah center, in 2017 and 2018 growing season. In the experiment, 30 plants used, Sayer date palm cultivar was five years old, planted on 5 x 5 m in sandy loam. The drip irrigation system was used. The normal EC value for soil was 12 dS m⁻¹. The average temperature of the experiment months was in October 26.2 °C, November 20.4 °C, December 14.4 °C, January12.2 °C, February 14.1 °C, March 18.1 °C, and April 23.8 °C. Plants were treated with saltwater and freshwater on April 1st in season 2017, according to the block design during the six months before the treatments, and continued until the samples were taken on April 1st in the season of 2018 after the flowers were completed in this variety. The plants were treated with foliar treatment once, with one plant for each replicate on October 1st in the season of 2017 as follows: only spray water (0 dS m⁻¹) + irrigation with freshwater (EC water 1.5 dS m⁻¹) (control); spray only with water + irrigate with salt water (EC 8 dS m⁻¹) (control); foliar spray of 500 ppm salicylic acid (SA) + irrigation with fresh water; foliar spray of SA at 500 ppm + irrigated with saltwater; foliar spray with potassium nitrate (KNO₃) 1500 ppm + irrigated with freshwater; foliar spray with KNO₃ 1500 ppm + irrigated with saltwater. On the 1st of April in the season of 2018, the data on plant parameters described in the following subchapters were recorded.

2.2 NUMBER OF CLUSTERS

The date palm starts to flower usually in February until April. The flowering is completed at the Hillawi cultivar, usually in April. The number of clusters was determined by calculating the number of clusters per plant to different treatments in April one time.

2.3 THE NUMBER OF NEW LEAVES

At the beginning of October, the number of leaves was assessed before the treatment time for each plant. The number of new leaves is completed in date palm in the spring. Therefore, at the beginning of April, the number of newly formed leaves was determined according to the following formula:

New leaves = Total of leaves on April – Total of leaves before treatment.

2.4 CARBOHYDRATE ANALYSIS

Soluble carbohydrates were determined after Yemm & Willis (1954). Samples of fresh pinnae were weighed (0.2 g) and homogenized using 70 % ethanol. Then they were filtered, and the use of benzene removed pigments. An aliquot of 0.2 ml of leaf extract was added to 1.0 ml of 5 % phenol + 5 ml H_2SO_4 95 % to react in a water bath for 10 min at 100 °C. Soon after, the test tube was cooled in an ice bath, and then the absorbance was read at 620 nm.

2.5 ASCORBIC ACID (ASA) CONTENT

AsA content was determined by employing a slightly changed methodology of Luwe et al. (1993) in which 0.5 g of green leaf pieces were ground in liquid nitrogen and afterward homogenized in 1 % cold trichloroacetic acid. The homogenate was then centrifuged at 12,000 x g for 20 minutes at 4 °C, and the supernatant (50 μ l) blended with100 mM potassium phosphate buffer, and ascorbate estimated at 265 nm.

2.6 HORMONES ANALYSIS

Five g of a fresh leaf tissue sample, which was homogenized in 70 % methanol was stirred overnight at 4 °C. The extract was filtered through Whatman filter paper (No. 1) and evaporated under vacuum. The pH of the aqueous phase was adjusted to 8.5 using 0.1 M phosphate buffer. Later the aqueous phase was partitioned using methanol twice. The methanol phase was removed by a rotary evaporator. The aqueous phase pH was adjusted to 2.5, using 1 N hydrochloric acid (HCI). Phytohormones were determined by the injection of the concentrate into a reversed-phase HPLC, C18 column in an isocratic elution mode utilizing a portable stage comprising of acetone: water (26:74) with 30 mM phosphoric acid as per Tang et al. (2011). The pH was kept up at 4, utilizing 1 N sodium hydroxide. A temperature was kept upt at 25 °C. The flux rate was 0.8 ml min⁻¹, and the elution of the phytohormones was observed at 208, 265, 270, and 280 nm for indoleacetic acid, abscisic acid, gibberellins, and zeatin, respectively.

2.7 EXTRACTION OF PROTEINS AND GEL ELECTROPHORESIS

Proteins were extracted by homogenizing the 333 mg of solidified dried leaf to 1 ml of extraction cradle [0.2 M, tris-hydroxymethyl aminomethane (Tris) + 0.001 M ethylene diamine tetra acetic acid + $(Na_2 + EDTA) + 12$ % glycerol + 0.01 M dithio threitol (DTT) + 0.05 mM phenyl methyl sulfonyl fluoride (PMSF)] by utilizing the mortar and pestle. At that point the samples were centrifuged at 15,000 × g for 15 min. Splitting buffer consisted of 0.125 M Tris HCl (pH 6.8) + 4 % SDS + 20 %, glycerol + 10 % b-mercapto ethanol + 0.01 % bromo phenol blue. Protein samples were denaturized by bubbling in the water bath at 90 °C for 3 min. Protein electrophoresis was performed in an irregular SDS polyacrylamide gel, as indicated by a strategy depicted by Laemmli (1970).

2.8 STATISTICAL ANALYSIS

Randomized completely block design of six treatments of salicylic acid and potassium nitrate replicated five times were utilized. Experimental data on all factors were analyzed by ANOVA. SPSS variant 19.0 (SPSS, Chicago, IL), and Duncan test were used for different correlations treatments considered at the $p \le$ 0.05 levels.

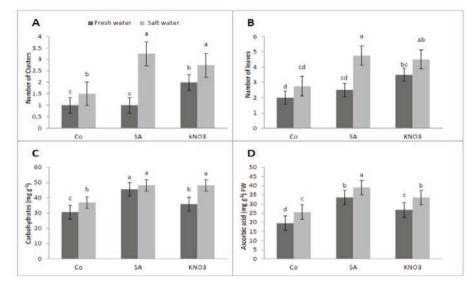


Figure 1: Salicylic acid and potassium nitrate effects on the number of clusters (A), number of new leaves (B), carbohydrates (C), and ascorbic acid (D)content in leaves of date palm offshoots after irrigated with salt water or with fresh water. The means of five replicates \pm SE are presented. Bars with different letters are significantly different at $p \le 0.05$ after a Duncan correction.

3 RESULTS

3.1 EXOGENOUS UTILIZATION OF SA AND KNO₃ PROMOTES FLOWERING, NUMBER OF NEW LEAVES, CARBOHYDRATES AND ASCORBIC ACID

The highest flowering rate was observed through SA treatment in salinity conditions followed by KNO, under the same conditions, whereas the lowest clusters number was found in control irrigated with fresh water, compared with saltwater (Fig. 1 A). The number of new leaves was fundamentally influenced by treatments; however, there was no difference between control, freshwater, and saltwater. Although the highest value of this variable was seen in plants treated with SA, there was no distinction among KNO₃ treatment under saltwater and SA treatment with salt water (Fig. 1 B). Application of both KNO₃ and SA treatments significantly increased carbohydrates in leaves under salt stress; there was no difference among treatments in this trait (Fig. 1 C). Ascorbic acid content under salt stress was higher than with freshwater in control; however, SA increased significantly ascorbic acid content under salt stress compared with other treatments (Fig. 1 C).

3.2 SA AND KNO₃ MODULATE IAA, CK, ABA, AND GA3 LEVELS

The application of treatments significantly in-

creased IAA, CK, and GA3 levels, either under saltwater or freshwater, compared with controls (Fig. 2). Whereas the content of ABA increased under salt stress only, and no significant effect of treatment was observed under normal conditions. SA increased ABA and GA3 content significantly under salt stress (Fig. 2 C, D).

3.3 SA AND KNO₃ MODULATE THE PROTEIN PATTERN OF LEAVES

The protein pattern of date palm leaves was determined according to the molecular mass (KD) of protein bands affected by SA and KNO, under freshwater and saltwater conditions by SDS-PAGE gel electrophoresis (Fig. 3). The leaf proteins in both controls were separated into five different polypeptides through acrylamide gels (Fig. 3). The control treatment with freshwater has a molecular mass of polypeptides 236.000, 132.072, 69.333, 40.642, and 34.389 KD, whereas control with salt water, had different molecular mass to three polypeptides 177.500, 44.179, and 30.090 KD. The application of SA showed five and six-band of proteins with fresh water and with saltwater, respectively, that differed in the molecular mass of one polypeptide compared to control with fresh water 28.232 KD. Whereas, SA with saltwater changed the molecular mass of two polypeptides compared to control with salt water 17.090 and 13.980 KD and showed six bands (Fig. 3). However, potassium application showed five bands of protein, whether with freshwater or with saltwater. One of the polypeptides

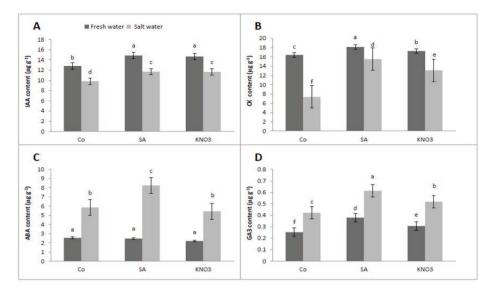


Figure 2: Salicylic acid and potassium nitrate effects on IAA (A), CK (B), ABA (C), and GA3 content (D) in leaves of date palm offshoots under irrigation with salt water or with fresh water. The means of five replicates \pm SE are presented. Bars with different letters are significantly different at $p \le 0.05$ after a Duncan correction.

differed in the molecular mass compared to the treatment of potassium with fresh water 33.188 KD. In contrast, two polypeptides differed in the molecular mass compared to potassium with saltwater treatment 33.600 and 14.646 KD (Fig. 3).

4 DISCUSSION

Date palm offshoots give a few numbers of floral buds that turn into a cluster at the beginning of their transformation from the vegetative to the flowering phase under extreme conditions. The salicylic acidtreated plant accelerated flowering initiation in plants under the salinity conditions (Wada & Takeno, 2013). An increase in the number of clusters observed after the application of SA (Fig. 1 A) was due to its effect on increase of carbohydrates to higher levels than the impact of salinity alone (Fig. 1 C). SA acts as an internal signal to regulate the physiological processes (Appu & Muthukrishnan, 2014).

Desoky and Merwad (2015) reported that spraying the leaves of wheat plants with SA increased the soluble carbohydrate content. Carbohydrate content in the plant increased the chances of the axillary buds transformation into a floral bud (Dierck, 2016). The number of increased leaves reflected the increase in the number of clusters (Fig. 1B, A). Inducing flowering, when connected to salicylic acid, might be the consequence of the positive impact on plant development. It can be indirect as SA adjusts the synthesis and signaling pathways of other plant hormones, including gibberellin, auxin, and abscisic acid (Fig. 2). SA and K⁺ ion have a role in cell division and stimulate secondary metabolism to produce antioxidants such as ascorbic acid. However, SA with saltwater increased AsA content (Fig. D). AsA is a primary co-factor in the synthesis of ethylene, gibberellic acid, and abscisic acid. Therefore, the endogenous level of AsA can affect the signaling of synthesis of those molecules (Anwar et al., 2018). Also, AsA is considered as an organic acid that stimulates the process of flowering (Akram et al., 2017)ascorbic acid (AsA. The high AsA content in plants supports a particular balance in preventing pigment damage and membrane injury (Costa et al., 2018). The antioxidant activity of ascorbate peroxidase and superoxide dismutase, and synthesis of special protein groups were increased by SA and AsA, together with salt stress (Al-Mayahi, 2016).

SA increased ABA and GA3 significantly under salt stress (Fig. 2 C, D). Salicylic acid rose IAA, CK and ABA levels, enhanced cell division in the apical meristem, and improved flowering of the plant (Sytar et al., 2019). In contrast, Alonso-Ramírez et al. (2009) reported that there was a cross-talk between SA and GA3 in *Arabidopsis thaliana* during abiotic stress conditions. GA3 induced flowering in early varieties of date palm (Cheruth et al., 2015). ABA has molecular effects on the downstream states of the autonomous biological pathway and, as such, improves the plant's capacity to encounter the change to flowering (Duncan et al., 2018).

The proteins are the final product of genetic pathways inside the plant cells that are created in light of cell

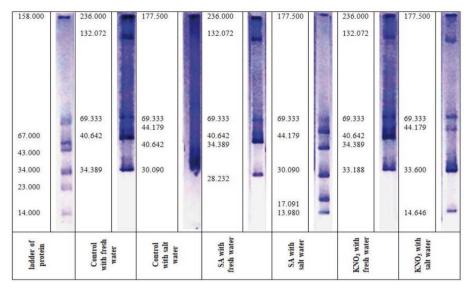


Figure 3: hematic diagrams of the electrophoretic protein pattern of date palm leaves with the molecular mass (KD) of protein bands affected by salicylic acid and potassium nitrate under freshwater and saltwater conditions.

needs and moved imbalances in some areas throughout entirely different stages of life and stress conditions (Razavizadeh, 2015)Brassica napus L. Applications of SA and K⁺ showed new protein bands with low molecular mass and the disappearance of other groups (Fig. 3). Salinity and SA induced de novo caused compilation of particular polypeptides and regulated the expression of salt-stress-tolerant proteins (Amirbakhtiar et al., 2019) which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited. Salt stress is one of the major adverse environmental factors limiting crop productivity. Considering Iran as one of the bread wheat origins, we sequenced root transcriptome of an Iranian salt tolerant cultivar, Arg, under salt stress to extend our knowledge of the molecular basis of salinity tolerance in Triticum aestivum. RNA sequencing resulted in more than 113 million reads and about 104013 genes were obtained, among which 26171 novel transcripts were identified. A comparison of abundances showed that 5128 genes were differentially expressed due to salt stress. The differentially expressed genes (DEGs. In Arabidopsis thaliana and different other flowering plants, it was suggested that molecular mechanisms which include gibberellic acid regulate the LFY promoter (Blazquez, 1998). However, ABA-induced RNAbinding proteins SNF5 and FCA control flowering time and stress responses (Fahraji et al., 2014) kinetin and salicylic acid may increase yield of different crops due to reduction in stress induced inhibition of plant growth. Salicylic acid (SA. In this respect, SA and salinity conditions together induced high levels of GA3 and ABA which caused synthesis of new isozymes of low molecular mass to form the new floral buds. These findings could facilitate understanding the mechanisms of flowering and salt-tolerance of the date palm.

5 CONCLUSION

Exogenous application of SA and KNO₃ enhanced the biochemical mechanisms of flowering in the date palm through conferring adaptation to salinity stress and creating new isozyme. IAA, ABA, CK, and GA3 successfully increased the number of the cluster. Furthermore, these hormones in the plant under salinity conditions contribute to the process of flowering by the formation of specific proteins through the epigenetic pathway to promote the transformation of the offshoots into an adult plant. This study is the first, which highlighted the stimulative effect of SA and KNO₃ on the flowering of the date palm.

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