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RESPONSE OF SOME FORAGE PEA GENOTYPES TO SALT STRESS DURING THE SEEDLING STAGE

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Abstract

Salinity stress is a common problem under dry climatic condition all over the world. The determination of salt tolerant genotypes has a crucial importance to alleviate this problem. Pea known as the most salt tolerant plant among legumes and seedling stage is the most sensitive to salt stress than the other growth stages. The experiment was carried out in the growth chamber to observe tolerance of the examined genotypes to different salt doses. Therefore germination rates, mean germination time (mgt), root/shoot lengths and fresh/dry seedling weights of some

forage pea genotypes (Local population, Crackerjack, Golyazi, Ozkaynak, Rose, Taskent, Tore, Ulubatlı) under different salt concentrations (Control, 5, 10, 15 and 20 dS m⁻¹) were determined. The results showed that salinity x genotype interaction was significant among genotypes with respect to all investigated parameters. Crackerjack had the highest germination rate and root length with the increasing salt levels and it was followed by Ozkaynak, Rose, Taskent and Tore. Ozkaynak was the fastest germinated genotype with respect to mgt with 1,92 days and the genotype had the longest shoot length. Fresh seedling weight of Crackerjack was the higher than all other genotypes but there was not a significant difference between Rose and Crackerjack. Results indicated that Taskent, Tore, Ozkaynak, cultivars could be recommended upto moderate saline areas while Crackerjack and Rose could be recommended for slightly saline areas.

Keywords

Forage pea, Germination, Salinity, Seedling vigor

1. Introduction

Soil salinity is a widely known problem of plant growth especially in arid and semi-arid areas of the world. The problem may occur due to improper usage of irrigation and fertilizers or it can be also originated from parent materials or weathering of salt minerals (Matternicht et al., 2008; Barbouchi et al., 2014; Gorji et al., 2017). All over the world, more than 20% of the cultivation lands suffer from salinization (Moud & Maghsoudi, 2008). Fundamentally, salt stress is an osmotic problem that reduces water uptake and thereby plant growth from germination to blooming stage affects negatively but it also causes significant problems in enzyme activities and other metabolic processes (Abdul-Jaleel et al., 2007; Wang et al., 2009). The severity of adverse effect of salinity on plants differs among the genotypes (Carpici et al., 2009) but most of the legumes are showed sensitivity to salt stress (Bonilla et al., 2004). Therefore, salt tolerant genotypes should be determined for salinity problematic areas.

Forage pea is an annual forage legume cultivated for both forage and grain feed for livestock industry and the plant was also known that the most salt tolerant genus among the legumes (Noreen & Ashraf, 2009). Both forage and grains of the plant are rich in minerals and amino acids (Acikgoz, 2001; Tan et al, 2011). High protein content of the forage (about 20%) and the grain (about 30%) and about 70-80% forage digestibility make it a valuable forage plant (Uzun et al., 2005; Avcioglu et al., 2009). It can be cultivated in the Central Anatolia conditions

as winter and spring sowing and also can be grown sole or mixtured with cereals. The plant is also used widely for multiple purposes as forage, hay, grain, feed or green manure.

Seed germination and seedling growth were early stages of the plant growth that were considered very critical for crop production (Almansouri et al., 2001; Okcu et al., 2005). Various studies indicate that responses of the genotype to salt stress vary in different growth stages (Almansouri et al., 2001; Jamil et al., 2006). However germination and seedling stages are considered as an early indicator to salt tolerance (Carpici et al., 2009; Zhu & Banuelos, 2016). Noreen et al. (2007) observed that germination of field pea seeds was severely reduced with increasing salinity levels (0, 60 120, 180 and 249 mM NaCl) and Steppuhn et al. (2001) reported that seed production decreased in line with increasing salinity stress and did not produced any grain under severe salinity condition but there were significant differences among pea genotypes (Olmos & Hellin, 1996; Noreen & Ashraf, 2009).

This research was carried out to determine seedling growth performance of some forage pea genotypes under different salt levels because seedling response is considered as an early indicator of tolerance to salt stress.

2. Materials and Methods

The experiment was carried out in the growth chamber of Eskisehir Osmangazi University, Faculty of Agriculture, and Department of Field Crops in 2016. Eight forage pea (*Pisum sativum* L.) genotypes (one of them was local population and the other 7 were Crackerjack, Golyazi, Ozkaynak, Rose, Taskent, Tore, Ulubatli cultivars) were used as material.

Solutions were prepared in 4 levels using NaCl and adjusted in 5, 10, 15 and 20 dS m⁻¹ EC values and distilled water was used as control. With 4 replications and in every replication, 25 seeds of each genotypes were germinated in triple rolled filter paper that moistened using 14 ml of test solutions per paper. Seeds were left to germinate at 20 °C in the dark for 10 days and then seedling measurements were performed (Okcu et al., 2005). The paper was replaced every two days to prevent accumulation of salt (Rehman et al., 1996). Seeds were considered as germinated when the radicle length reached to 1 mm and germinations were recorded in every 24 h (Okcu et al., 2005). Mean germination time was calculated considering Ellis & Roberts (1980)'s suggestion. Ten seedlings from every replicates were chosen randomly than, shoot and



root lengths were measured separately from these seedlings. Roots and shoots were weighed to calculate fresh seedlings weight and oven dried at 70 °C for 48 h (Uzun et al., 2005).

All data were subjected to analysis of variance based on general linear models for factorial arrangement (5 solutions x 8 genotypes) of treatments in a completely randomized design using SAS 9.3 software. Means were separated using TUKEY Multiple Range Test.

3. Results and Discussion

Genotypes showed different response with respect to investigated parameters to salt concentration, thus salt levels x genotype interactions were significant for all investigated characteristics. About all genotypes showed highest germination ratio at 5 dS m⁻¹ salt concentration but the response to salt levels changed at the other levels. For example, the lowest germination ratio recorded at control treatment for Golyazi cultivar while Rose and Taskent had the lowest germination ratio at 20 dS m⁻¹ salt level.

Salt level was significantly affect mgt, especially mgt was delayed after 10 dS m⁻¹ salt level. Genotypes also showed significant differences with respect to mgt. While mgt did not change in Taskent and Tore cultivars, the other genotypes showed different responses to salt levels. It was suprising that the highest salt content did not adversely affect mgt in the genotypes except for Ulubatli and Rose cultivars.

Root length was affected differently by salt content and it changed significantly among the genotypes. While there were great decreases at Crackerjack, Rose, Taskent and Tore in 10 dS m⁻¹, Ozkaynak was not significantly affected upto 15 dS m⁻¹ with respect to root length. However higher and similar root lengths were recorded at 5 dS m⁻¹ among the genotypes.

Genotypes showed different responses to changing salt concentrations with respect to shoot length. Shoot length of the genotypes were higher at 5 dS m⁻¹ salt level than the other levels but Population, Crackerjack, Golyazi, Taskent and Tore showed great decreasing trend after 5 dS m⁻¹ salt level, Ozkaynak was not greatly affected upto 10 dS m⁻¹. Crackerjack, Ozkaynak, Rose, Taskent and Tore cultivars had slightly higher but statistically insignificant shoot length.

Table 1: Germination rates of genotypes x salt interactions

Cultivar/Dosage	Control	5 ds m ⁻¹	10 ds m ⁻¹	15 ds m ⁻¹	20 ds m ⁻¹	Mean
Population	91,5 ^{abc}	90,0 ^{abc}	91,0 ^{abc}	90,0 ^{abc}	87,0 ^c	89,9 ^d

Crackerjack	100 ^a	100 ^a	99,5 ^a	99,0 ^{ab}	98,0 ^{abc}	99,3 ^a
Golyazi	88,0 ^{bc}	99,0 ^{ab}	97,0 ^{abc}	94,5 ^{abc}	94,5 ^{abc}	94,4 ^c
Ozkaynak	100 ^a	99,0 ^{ab}	98,5 ^{ab}	99,5 ^a	95,5 ^{abc}	98,5 ^{ab}
Rose	99,0 ^{ab}	100 ^a	99,0 ^{ab}	99,0 ^{ab}	92,0 ^{abc}	97,8 ^{abc}
Taskent	100 ^a	99,5 ^a	97,5 ^{abc}	98,0 ^{abc}	87,0 ^c	96,4 ^{abc}
Tore	97,0 ^{abc}	97,5 ^{abc}	95,5 ^{abc}	95,5 ^{abc}	90,5 ^{abc}	95,2 ^{bc}
Ulubatlı	35,5 ^e	50,0 ^d	36,0 ^e	48,5 ^d	32,0 ^e	40,4 ^e
Mean	88,9 ^b	91,7 ^a	89,2 ^{ab}	90,5 ^{ab}	84,6 ^c	

P<0.1

Table 2: Mean germination time values of genotypes x salt interactions

Cultivar / Salt	Control	5 ds m ⁻¹	10 ds m ⁻¹	15 ds m ⁻¹	20 ds m ⁻¹	Mean
Population	2,73	2,51	2,27	2,45	2,88	2,57 ^d
Crackerjack	2,66	2,30	2,16	2,18	2,50	2,36 ^c
Golyazi	3,53	2,66	2,31	2,83	3,42	2,95 ^f
Ozkaynak	1,97	2,00	1,61	1,86	2,17	1,92 ^a
Rose	2,53	2,61	2,40	2,70	3,43	2,74 ^e
Taskent	2,08	2,02	1,86	2,04	2,06	2,01 ^{ab}
Tore	2,01	2,06	2,04	2,02	2,08	2,04 ^b
Ulubatlı	2,96	2,86	2,60	2,99	3,47	2,98 ^f
Mean	2,56 ^c	2,38 ^b	2,16 ^a	2,38 ^b	2,75 ^d	

P<0.1

Fresh seedling weights were always higher in all genotypes at 5 dS m⁻¹ salt level compared to the other treatments but the response of fresh seedling weight of the genotypes was fluxional at the other treatments. Crackerjack, Rose, Taşkent and Tore cultivars were affected negatively after 5 dS m⁻¹ salt content but Ozkaynak was not affected seriously upto 10 dS m⁻¹. Similar responses to salt levels were also recorded for dry seedling weight for genotypes.

Table 3: Root lengths of genotypes x salt interactions

Cultivar / Salt	Control	5 ds m ⁻¹	10 ds m ⁻¹	15 ds m ⁻¹	20 ds m ⁻¹	Mean
Population	4,65	4,85	4,72	3,77	2,30	4,06 ^d
Crackerjack	9,35	9,57	6,15	6,32	4,05	7,09 ^a
Golyazi	2,47	8,52	4,17	3,35	2,97	4,30 ^{cd}
Ozkaynak	7,92	7,70	6,97	6,05	2,17	6,16 ^b

Rose	9,10	8,32	4,60	6,32	3,10	6,29 ^b
Taskent	6,82	6,95	4,27	4,50	2,10	4,93 ^c
Tore	7,77	6,17	3,75	4,30	2,37	4,87 ^c
Ulubath	0,32	6,22	3,70	2,05	0,18	2,49 ^e
Mean	6,05 ^b	7,29 ^a	4,79 ^c	4,58 ^c	2,40 ^d	

P<0.1

Table 4: Shoot lengths of genotypes x salt interactions

Cultivar / Salt	Control	5 ds m ⁻¹	10 ds m ⁻¹	15 ds m ⁻¹	20 ds m ⁻¹	Mean
Population	6,42	8,27	5,00	2,40	1,02	4,62 ^c
Crackerjack	6,55	4,97	3,00	2,32	2,27	3,82 ^{de}
Golyazi	1,00	6,75	4,27	2,57	2,00	3,32 ^e
Ozkaynak	12,05	10,27	9,67	6,77	2,45	8,24 ^a
Rose	8,05	6,32	3,17	2,40	1,60	4,31 ^{cd}
Taskent	10,00	9,32	6,60	6,65	3,12	7,14 ^b
Tore	11,75	10,45	7,30	4,27	1,50	7,05 ^b
Ulubath	2,85	4,90	3,70	0,47	0,30	2,44 ^f
Mean	7,33 ^a	7,65 ^a	5,34 ^b	3,48 ^c	1,78 ^d	

P<0.1

The plants are the most sensitive to salt stress during seedling stage and considerable differences occur among genotypes belonging to species are originated from genetical differences (Cerda et al., 1982; Jayasundara et al., 1998; Noreen & Ashraf, 2009). In this study, investigated genotypes showed different response to salt stress with respect to germination percentage and it was decreased at high salt levels. Similar results were also recorded by Dua et al. (1989).

Table 5: Fresh seedling weights of genotypes x salt interactions

Cultivar / Salt	Control	5 ds m ⁻¹	10 ds m ⁻¹	15 ds m ⁻¹	20 ds m ⁻¹	Mean
Population	1,59	2,16	1,37	0,66	0,48	1,25 ^c
Crackerjack	4,05	4,17	2,61	1,65	1,39	2,77 ^a
Golyazi	0,22	2,37	1,00	1,10	0,91	1,12 ^c
Ozkaynak	2,25	2,36	2,02	1,23	0,91	1,75 ^b
Rose	3,89	4,40	2,21	1,15	1,40	2,61 ^a

Taskent	1,71	2,04	1,43	1,06	0,78	1,40 ^{bc}
Tore	1,84	2,14	1,62	1,03	0,70	1,47 ^{bc}
Ulubath	0,21	1,29	0,46	0,29	0,11	0,47 ^d
Mean	1,97 ^b	2,62 ^a	1,59 ^c	1,02 ^d	0,83 ^d	

P<0.1

Table 6: Dry seedling weights of genotypes x salt interactions

Cultivar / Salt	Control	5 ds m ⁻¹	10 ds m ⁻¹	15 ds m ⁻¹	20 ds m ⁻¹	Mean
Population	0,20	0,20	0,16	0,12	0,09	0,15 ^d
Crackerjack	0,42	0,39	0,31	0,27	0,22	0,32 ^a
Golyazi	0,06	0,24	0,19	0,14	0,13	0,15 ^d
Ozkaynak	0,26	0,25	0,21	0,17	0,12	0,21 ^b
Rose	0,48	0,42	0,28	0,21	0,17	0,31 ^a
Taskent	0,25	0,23	0,15	0,15	0,12	0,18 ^c
Tore	0,25	0,24	0,18	0,14	0,10	0,18 ^c
Ulubath	0,05	0,16	0,07	0,04	0,02	0,07 ^e
Mean	0,25 ^b	0,27 ^a	0,20 ^c	0,16 ^d	0,12 ^e	

P<0.1

Mean germination time increased upto 10 dS m⁻¹ salt treatment in most of the investigated genotypes in the experiment. But this increasing ratio showed differences among genotypes. These results mainly originated from salt stress tolerance differences of the genotypes as mentioned by Noreen & Ashraf (2009).

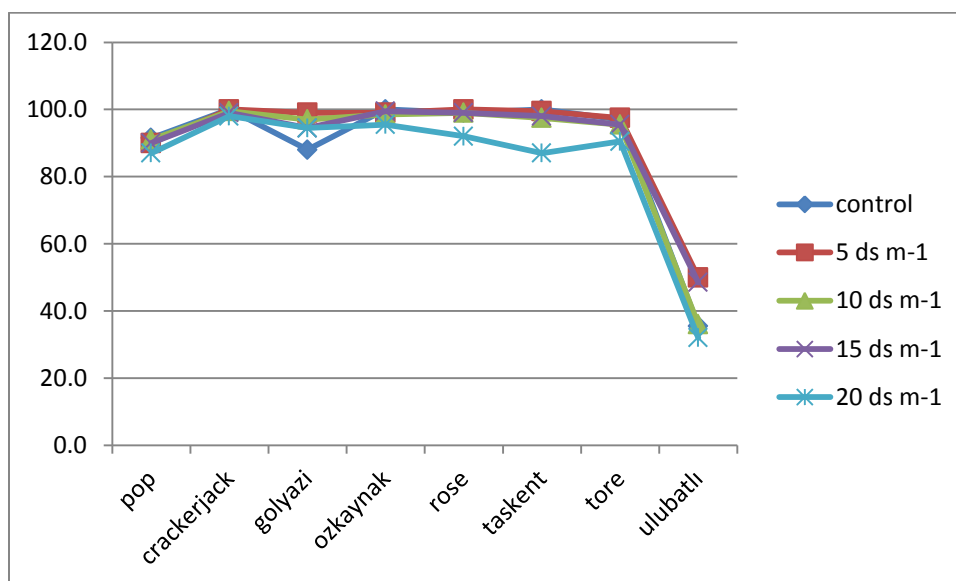


Figure 1: Germination rates of genotypes under different salinity levels

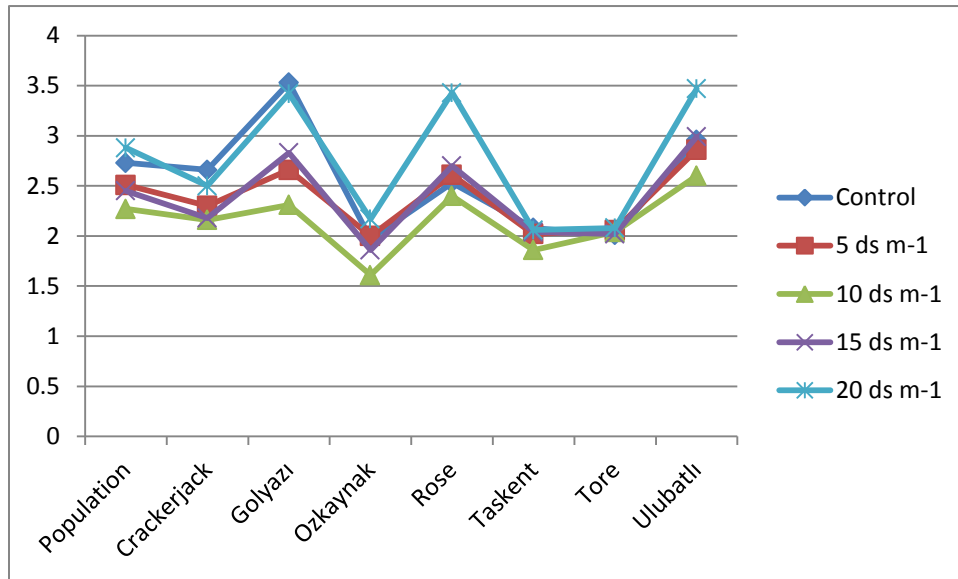


Figure 2: Mean germination time of genotypes under different salinity levels

Root and shoot lengths showed considerable changes among the genotypes. In general, root and shoot lengths were decreased as salt level increase but decreasing trend was different among the genotypes. Generally, root growth was less affected by salt stress than shoot growth (Table 1). Similar results were also reported by Jayasundara et al. (1998) and Noreen & Ashraf (2009).

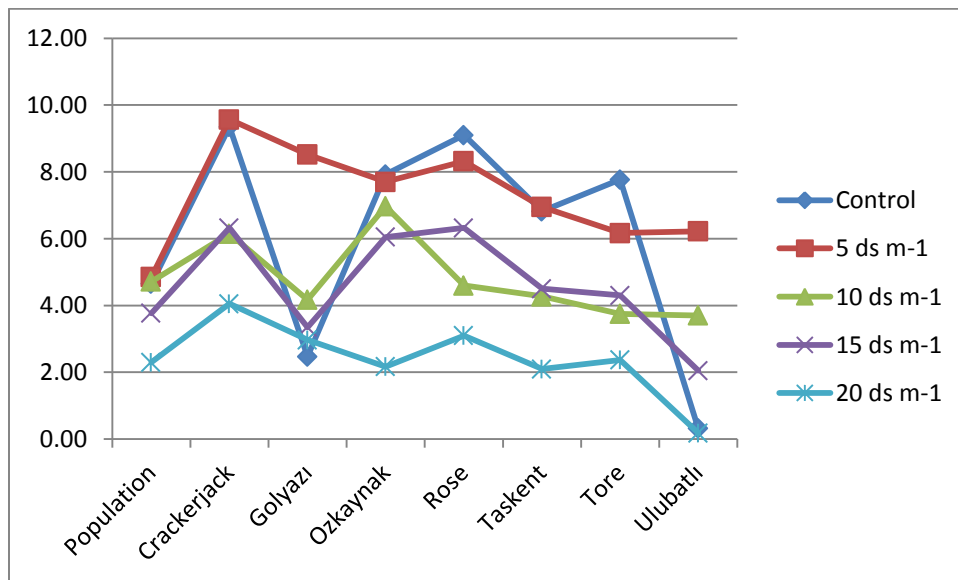


Figure 3: Root lengths of genotypes under different salinity levels

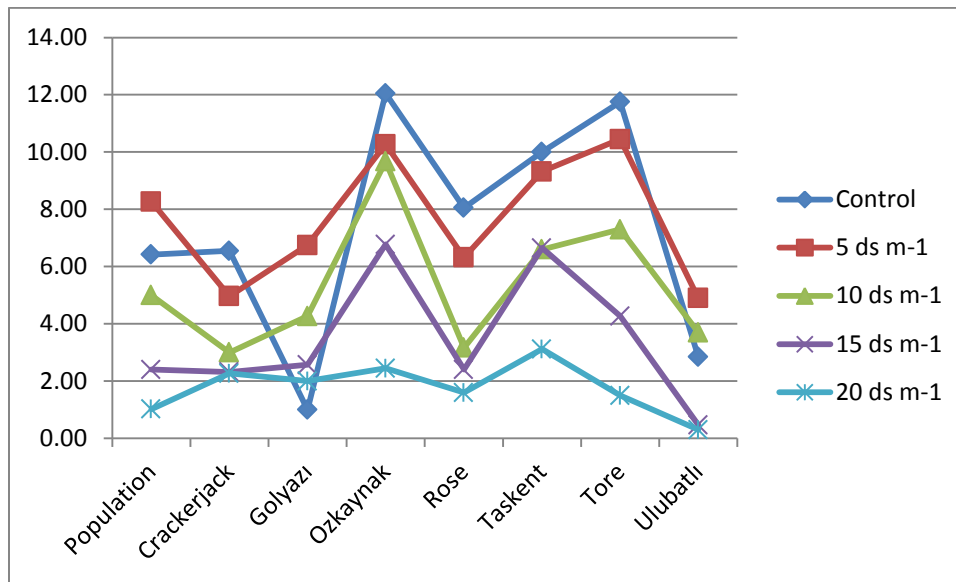


Figure 4: Shoot lengths of genotypes under different salinity levels

As salt level increase, seedling fresh and dry weights were decreased especially it was more pronounced after 5 dS m⁻¹ salt content (Table 1). This was mainly originated from negative effect of salt stress on plant growth (Parida & Das, 2005). The genotypes showed different response to salt levels with respect to fresh and dry seedling weight (Figure 5,6). Reduction in fresh and dry seedling weight were low in Taşkent, Ozkaynak and Tore cultivars upto 10 dS m⁻¹ salt level and in Rose and Crackerjack upto 5 dS m⁻¹. These results clearly pointed out those salt tolerances of the investigated genotypes were different, as reported in the other studies (Najafi et al., 2007; Noreen & Ashraf, 2009).

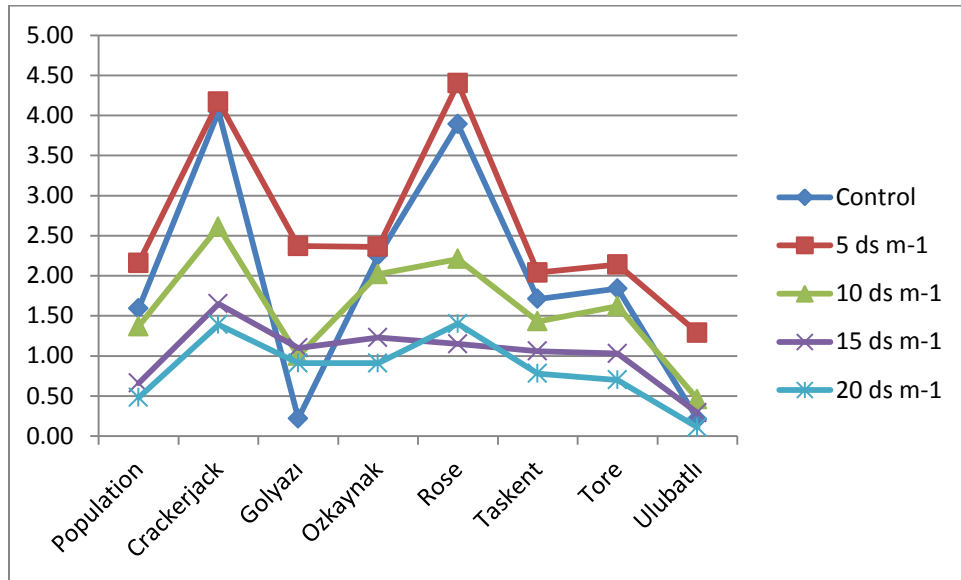


Figure 5: Fresh seedling weights of genotypes under different salinity levels

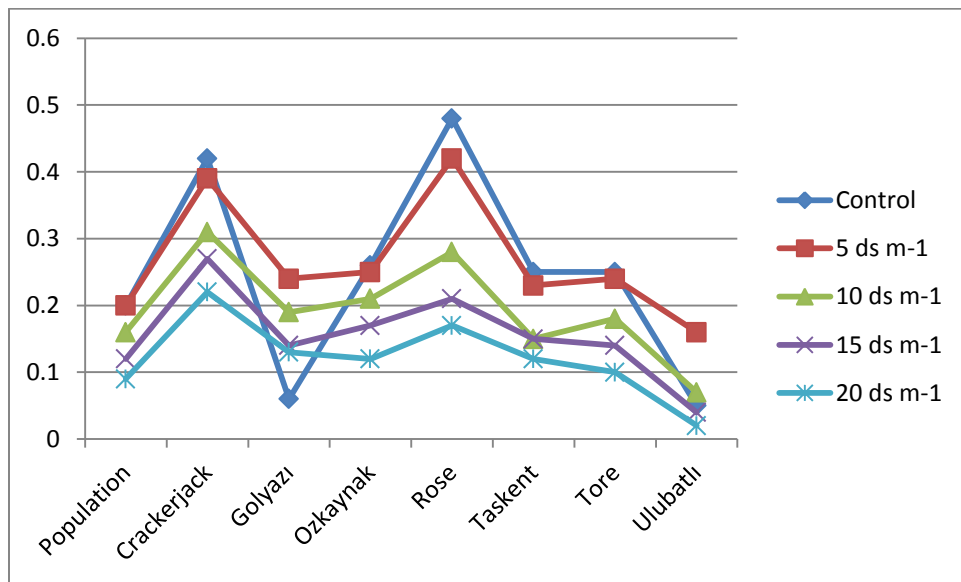


Figure 6: Dry seedling weights of genotypes under different salinity levels

In the experiment, some investigated parameters showed slight increase in lower salt content compared to the control. This situation can be related to characteristics of the plant because the pea genus has been categorized as one of the most tolerant legume to salinity (Noreen & Ashraf, 2009). Similarly, Najafi et al (2007) were also reported positive result under lower salt levels in pea under greenhouse conditions.

4. Conclusion

Salination is an inevitable and a serious problem under dry environment, especially irrigated areas. It is important to select suitable crops species to alleviate negative effect of salinization under salt affected areas.

Germination and seedling growth in the investigated genotypes showed different response to salinity levels. Among the genotypes, Taskent, Tore and Ozkaynak were seen to be more resistant to salt stress because their growing performances were not affected seriously upto 10 dS m⁻¹ salt level. On the other hand, Crackerjack and Rose cultivars performance was not affected seriously by 5 dS m⁻¹ salt level. According to the result; these 5 cultivars can be suggested for slightly salt affected areas and firstly mentioned 3 cultivars (Taskent, Tore, Ozkaynak) can be suggested for moderately salt affected areas. However, further researches are needed with respect to investigate field performance of the suggested cultivars.

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