RESEARCH ARTICLE

**Investigations on Growth, Biochemical Compounds and Antioxidant activities in Hybrid Maize Genotypes as influenced by Silicon Fertilization**

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|  | ABSTRACTA pot culture study was conducted to investigate the effect of silicon fertilization on the growth and physiological traits of ten different hybrid maize genotypes. The treatment structure comprises three levels of silicon (0, 75 and 150 kg ha-1) as calcium silicate and ten hybrid maize genotypes in a completely randomized design. The genotypes were imposed with recommended NPK (250:75:75kg ha-1) and grown-up to 40 days. At the end of 40 days, the dry matter production, SPAD, biochemical compounds like soluble proteins, phenols, proline and antioxidants such as peroxidase and superoxide dismutase (SOD) were recorded. Results showed a linear increase in the overall development of hybrid maize genotypes with increasing silicon fertilization upto 150 kg Si ha-1. The genotype COH(M)8 was the most responsive genotype to Si application among all the genotypes, followed by CMH12-586 and VaMH12014. In the case of biochemical attributes, genotypic variation played a significant role in expressing the impact of Si. Lesser response to Si fertilization and its accumulation was noticed with NK6240 and COH(M)9 on the growth and biochemical attributes. Silicon fertilization improved the biochemical and antioxidant activity of the maize genotypes and the response varied with levels of Si. A positive and significant response to Si fertilization was observed with soluble protein (9.82 mg g-1), phenols (0.70 mg g-1), peroxidase (0.83 units min-1 mg-1) and SOD activity (8.93 U g-1) whereas an inverse significant relationship was noticed with the proline content (4.43 µmoles g-1) of the plants which confirmed the role of Si in plant defense mechanism. The relationship study also confirmed the positive and significant correlation between growth parameters and dry matter production of the genotypes for silicon fertilization.  |

Keywords: Antioxidants, Biochemical compounds, Chlorophyll, Dry matter production, Maize genotypes,

 Silicon levels

## INTRODUCTION

Silicon (Si), the second most abundant element on the earth surface, is rapidly gaining attention in agriculture because of its many beneficial effects for plants. Current efforts to resolve this enigma involve comprehensive analyses of the effect of Si supplementation under various abiotic and biotic stresses. At the same time, research activities are focused on improving Si fertilization and Si sources for better crop cultivation to understand the involvement of Si in improving the growth and yield of crops. (Frew *et al.,* 2018). The ability of Si to alleviate stress is a fundamental component of plant life-processes, and it enhances biochemical tolerance in plants, which is mainly associated with increased activity of defense-related enzymes, such as poly-phenoloxidase, glucanase, peroxidase, and phenylalanine ammonia-lyase (PAL), production of antimicrobial compounds such as phenolic, flavonoids, phytoalexins and pathogenesis-related proteins in plants besides regulating systemic signals, such as salicylic acid, jasmonic acid, and ethylene (Fortunato *et al*., 2012; Van *et al*., 2013). It plays a crucial role in plant growth by conferring mechanical strength to crops and resistance against pathogens and herbivory. Silicon additionally influences plant chlorophyll content and photosynthetic capabilities of crops by improving their light receiving posture (Tubana *et al.,* 2016).

Grasses are well-known high accumulators of Si, and, therefore, serve as an excellent model to study the role of Si in plant physiology. Maize (*Zea mays* L.) is also known as Si-accumulator grown widely as an important cereal in India. It is known to take up actively and accumulate Si in its organs(Liang *et al.,* 2007; Coskun *et al.,* 2019) and is highly responsive to Si fertilization (Liu *et al.,* 2011).

Silicon is an agronomically important fertilizer element that enhances plant tolerance to biotic and abiotic stress. It plays a dual role in soil-plant system by reinforcing plant protection measures and by optimizing soil fertility through improved water, physical and chemical soil properties, and maintenance of nutrients in plant-available forms (Meena *et al.,* 2014). The benefits of Si on maize are manifold like improving the population, quality, photosynthetic efficiency, increasing leaf area, yield, etc (Covshoff and Hibberd, 2012). Thus, a short term experiment was carried out with different genotypes of hybrid maize to assess the effectiveness of various levels of silicon applied as calcium silicate on dry matter production and physiological aspects of hybrid maize.

## MATERIAL AND METHODS

***Experimental details***

A pot culture study was conducted in the glasshouse of the Department of Soil Science and Agricultural Chemistry, Tamil Nadu Agricultural University, Coimbatore, to know the effect of silicon on growth, biochemical constituents and enzyme activity in ten maize genotypes. The soil chosen for the study was sandy loam in texture having deficient Si content (36.5 mg kg-1) with neutral soil reaction (7.50), non-saline (0.09 dS m-1), non-calcareous and having low organic carbon content (1.18 g kg-1). The available N (86.8 kg ha-1) and K (123 kg ha-1) status were low but high in available P (23.1 kg ha-1). Each pot was filled with 4.50 kg of processed soil and three seeds per pot were sown. The soil test-based basal recommended dose of N, P2O5 and K2O was applied and the silicon was applied as calcium silicate at three levels 0, 75 and 150 kg ha-1. Ten different hybrid maize genotypes were chosen for the study, out of which eight were TNAU hybrids (CO 6, COH(M) 8, COH(M)9, VaMH2014) and cultures (CMH12- 586, CMH2-686, CMH14-716, CMH15-005) and two private hybrids (NK6240 and 900M Gold). All the plants were grown and harvested at 40 days after sowing for recording SPAD, root volume, and total dry matter production. The biochemical constituents (protein, proline and phenols) and enzyme activities (peroxidase and superoxide dismutase) in the fresh leaves were also analyzed.

***Dry matter production:*** For determining dry matter production, all maize genotypes were harvested, washed with water and separated into shoot and root samples. The samples were then dried in a hot air oven at 70 0 C for 24 h till they got completely dried following which they were weighed and expressed as g plant-1.

***SPAD:*** A chlorophyll meter (SPAD 502) designed by the soil plant analysis development (SPAD 502), Minolta Camera Co. Ltd., Japan, was used to record the chlorophyll index. Measurements were taken on the middle of a fully expanded leaf (3rd or 4th leaf from the apex), and five readings were taken in each replicate to represent the mean chlorophyll index for each treatment.

***Root volume:*** The root volume was measured using water displacement technique by immersing the root in water-filled measuring cylinder. The water displaced is measured using a graduated cylinder and is expressed in cm3.

***Biochemical compounds***

*Soluble protein:* Estimation of soluble protein was carried out using Lowry’s method (1951). Two fifty mg of fresh leaf samples was taken and macerated with 10 mL of 0.02M phosphate buffer (pH 7.8). One mL of this extract was taken in a test tube, to which 5 mL of ACT (Alkaline Copper Tartarate) and 0.5 mL of Folin-Ciocalteau reagent were added. The solution was kept as such for 30 min for color development in the dark at room temperature and the absorbance was measured at 660 nm. Fifty mg of Bovine serum albumin (BSA) dissolved in a hundred mL of distilled water (500 ppm) was used as the standard. The soluble protein was calculated and expressed in mg g-1.

*Proline:* Proline content of fresh leave samples was analyzed by the method outlined by Bates *et al.,* (1973). About 0.5 g of leaf sample was macerated with 10 mL of 3% sulpho salicyclic acid and centrifuged at 3000 rpm for 10 minutes. From this, 2 mL of extract was taken in a test tube and added with 2 mL of acid ninhydrin, 2 mL of glacial acetic acid and 2 mL of ortho phosphoric acid. The test tubes were kept in a hot water bath for 1 h wherein the solution color changes from yellow to red. Then, the solution was transferred to a separating funnel and 4 mL of toluene was added. The whole mixture was vigorously shaken for 20-30 sec and then allowed to settle. The lower colorless layer was discarded and the upper pink layer was used for measuring the absorbance at 520 nm. The proline content was calculated and expressed as µmoles g-1 fresh weight.

*Total phenols:* Estimation of total phenols was carried out as per Bray and Thorpe’s method (1954) in which 500 mg of fresh leaf sample was cut into small bits and transferred to test tubes to which 5 mL of 80% ethanol was added. The test tubes are heated for 10 mins and cooled off. The leaf samples are macerated with another 5 mL of 80% ethanol and centrifuged at 5000 rpm for 10 mins, after which the supernatant is collected and the volume is made up to 25 mL. One mL of supernatant is taken to which2 mL of 20% sodium carbonate (Na2CO3) and 0.5 mL of Folin’s reagent were added. After the development of blue color, absorbance was taken at 660 nm. Standards are prepared using catechol at different concentrations (10, 20…..100 ppm). Total phenols was calculated and expressed in mg g-1.

***Enzyme activities***

*Peroxidase:* Leaf samples were homogenized using a phosphate buffer of which one mL of supernatant was taken and 3 mL of 0.05M pyrogallol and 0.5 mL of 30% H2O2 were added. The change in absorbance was measured at 430 nm for every 30 seconds up to 180 seconds and the enzyme activity was calculated and expressed as Units min-1 mg-1 of sample. (Sadasivam and Manickam, 1992).

*Super oxide dismutase:* Super**-**oxide dismutase was estimated by Nitro blue tetrazolium (NBT) method as given byBeau-Champ and Fridovich (1971). Five hundred milligram of leaf sample was macerated using 10 ml HEPES-KOH buffer containing 0.1 mM EDTA and centrifuged at 15000 rpm for 15 min. The supernatant was collected and made up to 50 mL volume. One mL of the enzyme extract was mixed with 3 mL of the reaction mixture and the absorbance was recorded at 560 nm after the reaction tubes were illuminated for 15 min. Non-illuminated and illuminated reactions without supernatant were used as controls. One unit of SOD activity was defined as the amount of enzyme required to result in 50% inhibition of NBT reduction at 560 nm. The result was expressed as units per gram of fresh weight.

***Statistical analysis***

The experiment was laid out in a completely randomized design with three replications. The data obtained from the investigations were analyzed using OPSTAT software and were subjected to analysis of variance to find out the significance (Panse and Sukhatme, 1978). Wherever the treatment differences were found, significant critical differences (CD) were worked out at 5% level with a mean separation by the least significant difference and denoted by symbol \* ,\*\* for 5% and 1%, respectively. Non-significant comparisons were indicated as NS. A simple correlation was worked out between different parameters to know the positive or negative relationships between them.

## RESULTS AND DISCUSSION

***Dry matter production***

The data on total dry matter production of various maize genotypes were recorded and given in table 1. The application of silicon fertilizers significantly influenced the dry matter production of all the maize hybrids. The highest dry matter production (DMP) of all the maize hybrids was reported with the application of 150 kg Si ha-1as calcium silicate (13.3 g pot -1). Among the genotypes, COH (M)8 recorded the highest DMP (13.3g pot -1) followed by CMH12-586 (13.1g pot -1). The lowest DMP was registered with no silicon applied pots (control) in NK6240 (9.60g pot -1) and CMH15-005 (9.84g pot -1). The increased DMP might be due to the beneficial effect of Si, which is known to promote the growth of maize seedlings and eventually contributed to increase in dry matter of maize hybrids, which was also reported by Yang *et al*. (2008) for its essentiality in plant growth and development. Further, Si is also known to make the plants to stand erect, which helps in effective utilization of sunlight. This further helped in improved translocation of nutrients, which ultimately increased the dry matter production (Patil, 2017).

An attempt was made to understand the relationship between dry matter production and root volume, which revealed that DMP is positively correlated with root volume (r = 0.973\*\*, Fig. 1). Also, the relationship between DMP with a relative water content of the genotypes were also assessed and found a linear relationship between them as well (r = 0.935\*\*, Fig. 1).

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**Figure 1: Relationship of dry matter production with relative water content and root volume**

***Chlorophyll index***

Chlorophyll content was measured using the SPAD meter for all the maize genotypes, which also increased with increasing levels of Si application. The highest SPAD value was observed in the genotype COH (M)8 (40.2) followed by CMH12-586 (37.6) whereas lesser SPAD values was observed in the genotypes NK6240 (35.0) and CMH5-005 (Table 1). Chlorophylls play an important role not only in the capacity but also in the efficiency of plants’ photosynthesis activity. Photosynthetic capacities of crops’ applied with Si are improved by the enlarged size of chloroplasts and the increased number of grana in leaves (Xie *et al.,* 2014). SPAD values enhanced with Si fertilization might be due to the fact that Si regulates the stomatal activity and increased the photosynthesis and water use efficiency of plants which eventually resulted in better vegetative growth (Shedeed, 2018). A significant relationship was noticed between DMP and chlorophyll index for all the maize genotypes (r = 0.922\*\*).

***Biochemical constituents***

The effect of silicon fertilization was studied on soluble proteins, proline and phenolic content in the leaves of all the maize genotypes. It was observed that increasing levels of silicon increased the soluble protein and phenol content of the plants but decreased the proline content of the leaves. The highest soluble protein content was measured with the genotype COH (M)8, which varied from 8.73 mg g-1 in no silica control to 9.82 mg g-1 in 150 kg Si ha-1. This was closely followed by CMH12-586 with 7.84 mg g-1 in control and 8.58 mg g-1 in 150 kg Si ha-1 (Table 2). This is in good accordance with the findings of Gong (2005), who observed that silicon application prevented oxidation of proteins in wheat and helped in enhancing the protein content of the plant.

Proline has been considered as a carbon and nitrogen source for rapid recovery from stress and growth, a stabilizer for membranes and a free radical scavenger (Shahnaz *et al.,* 2011). A decrease in levels of proline was observed with an increasing dose of silicon (Table 2). Among the maize genotypes, COH(M)8 was observed to contain high proline content in the NPK treatment (9.87 µmoles g-1) followed by VaMH1204 (9.65 µmoles g-1) and CO6 (9.44 µmoles g-1). With the application of calcium silicate at 150 kg Si ha-1 , the proline content was lowered and the lowest content was recorded in COH(M) 9 (2.60 µmoles g-1) and 900M Gold (2.67 µmoles g-1). Increasing dosage of silicon might have allowed the plants to resist abiotic or biotic stress with the production of silaproline molecules just as in the case of humans. To date, no further studies have been done on this aspect (Kaya *et al.,* 2009).

The total phenol content was observed to increase slightly with the application of silicon (Table 2). Application of silicon at 150 kg ha-1 increased the phenol content of all maize genotypes and the highest phenol content was found in COH(M)8 (0.70 mg g-1) followed by CMH12-586 (0.68 mg g-1) and VaMH1204 (0.65 mg g-1 ). The lowest phenol content was observed in NPK control in CMH12-686 (0.43 mg g-1) and 900M Gold (0.47 mg g-1). This is in good corroborations with the works done by Malcovska and his co-workers (2014). Silicon is known to regulate the metabolism and utilization of phenols in plant leaves by producing various Si polyphenol complexes (Shi *et al.,* 2014).

***Antioxidant enzymes***

The application of silicon at different levels increased both the peroxidase and the superoxide dismutase activity in hybrid maize regardless of all genotypes (Fig. 3).The highest peroxidase activity was exhibited by COH(M) 8 in 150 kg Si ha-1 treatment (0.83 units min-1 mg-1) followed by CMH12-586 (0.74 units min-1 mg-1) whereas the lowest activity was recorded in control treatment in CMH14-716 (0.20 units min-1 mg-1) and NK6240 (0.25 units min-1 mg-1). Peroxidase has a protective effect also involved in chlorophyll degradation, ROS generation and membrane lipid peroxidation, which are response to injury and a product of senescence (Cao *et al.,* 2015).

As expected, increasing silicon levels increased the superoxide dismutase (SOD) activity in the leaves of maize genotypes and the mean values varied from 5.27 to 8.18 U g-1. The highest value was recorded with the application of 150 kg Si ha-1 in the genotype COH(M)8 (8.93 U g-1). Lesser enzyme activity was observed with COH (M) 9 (3.82 U g-1) in control treatment (Fig. 3). ). Generation of reactive oxygen species such as superoxide radicals, hydroxyl radicals and hydrogen peroxide causes oxidative damage to plants, but is inevitable in all the aerobic organisms. Plants with high levels of antioxidants, either constitutive or induced, have been reported to have greater resistance to oxidative damage. Plants have evolved specific protective mechanisms, involving antioxidant molecules and enzymes in order to defend themselves against oxidants (Devi *et al.*, 2017). It is evident that silicon is known to influence the anatomical, morphological, biochemical and physiological reactions in the plants under stress by decreasing the permeability as well as the lipid peroxidation of the membrane, thus maintaining membrane stability to the crops as mentioned by Balakhina and Borkowska (2013).





**Figure 3. Effect of different levels of silicon on antioxidant enzymes activity in maize genotypes**

**Table 1: Effect of silicon on dry matter production and SPAD of maize genotypes**

|  |  |  |
| --- | --- | --- |
|  Si levels (kgha-1) Genotypes | Total Dry matter production (g plant-1) | Chlorophyll index |
| 0 | 75 | 150 | Mean | 0 | 75 | 150 | Mean |
| CO 6 | 10.7 | 11.7 | 12.2 | 11.5 | 32.2 | 34.8 | 36.7 | 34.9 |
| COH (M)8 | 12.2 | 12.9 | 13.3 | 12.8 | 36.8 | 38.3 | 40.2 | 38.7 |
| COH (M)9 | 10.0 | 11.2 | 11.7 | 11.0 | 30.7 | 34.2 | 36.5 | 34.2 |
| CMH12-586 | 11.8 | 12.6 | 13.1 | 12.5 | 34.0 | 35.9 | 37.6 | 36.1 |
| CMH12-686 | 11.2 | 12.2 | 12.6 | 12.0 | 32.9 | 35.0 | 36.4 | 35.0 |
| CMH14-716 | 10.4 | 11.5 | 11.9 | 11.3 | 31.0 | 34.0 | 35.6 | 33.8 |
| CMH15-005 | 9.84 | 11.2 | 11.5 | 10.8 | 31.9 | 34.2 | 36.0 | 34.4 |
| NK6240 | 9.60 | 11.0 | 11.4 | 10.7 | 30.9 | 33.6 | 35.3 | 33.6 |
| 900M Gold | 10.2 | 11.3 | 11.8 | 11.1 | 30.5 | 32.9 | 35.0 | 33.1 |
| VaMH12014 | 11.6 | 12.5 | 13.0 | 12.3 | 33.1 | 35.1 | 37.0 | 35.2 |
| Mean | 10.8 | 11.8 | 12.3 | 11.6 | 32.4 | 34.8 | 36.6 | 34.9 |
|  | G | L | G×L |  | G | L | G×L |  |
| SEd | 0.028 | 0.015 | 0.048 |  | 0.315 | 0.172 | 0.545 |  |
| CD (P=0.05) | 0.055 | 0.030 | 0.096 |  | 0.631 | 0.346 | NS |  |

G – Maize genotypes L – Levels of silica NS – Non significant

**Table 2: Effect of different levels of silicon on various biochemical compounds in maize genotypes**

|  |  |  |  |
| --- | --- | --- | --- |
|  Si levels (kgha-1) Genotypes  | Soluble protein content (mg g-1) | Total phenol content (mg g-1) | Proline( μmoles g-1) |
| 0 | 75 | 150 | Mean | 0 | 75 | 150 | Mean | 0 | 75 | 150 | Mean |
| CO 6 | 6.75 | 7.30 | 7.55 | 7.20 | 0.54 | 0.59 | 0.64 | 0.59 | 9.44 | 7.28 | 4.04 | 6.92 |
| COH (M)8 | 8.73 | 9.33 | 9.82 | 9.29 | 0.61 | 0.66 | 0.70 | 0.66 | 9.87 | 7.74 | 4.43 | 7.35 |
| COH (M)9 | 7.37 | 7.76 | 7.95 | 7.69 | 0.48 | 0.54 | 0.60 | 0.54 | 9.39 | 7.02 | 2.60 | 6.33 |
| CMH12-586 | 7.84 | 8.28 | 8.58 | 8.24 | 0.58 | 0.64 | 0.68 | 0.63 | 9.05 | 6.47 | 2.99 | 6.17 |
| CMH12-686 | 5.85 | 6.14 | 6.36 | 6.12 | 0.43 | 0.50 | 0.54 | 0.49 | 9.26 | 7.17 | 3.66 | 6.70 |
| CMH14-716 | 5.68 | 5.88 | 6.21 | 5.93 | 0.56 | 0.63 | 0.67 | 0.62 | 8.79 | 6.48 | 3.05 | 6.11 |
| CMH15-005 | 6.72 | 6.96 | 7.17 | 6.95 | 0.53 | 0.59 | 0.60 | 0.58 | 8.89 | 6.74 | 3.68 | 6.43 |
| NK6240 | 5.40 | 5.64 | 5.88 | 5.64 | 0.51 | 0.54 | 0.59 | 0.55 | 9.09 | 6.98 | 3.47 | 6.51 |
| 900M Gold | 6.67 | 6.88 | 7.10 | 6.89 | 0.47 | 0.50 | 0.55 | 0.51 | 9.34 | 6.73 | 2.67 | 6.25 |
| VaMH12014 | 8.36 | 8.60 | 8.85 | 8.60 | 0.50 | 0.62 | 0.65 | 0.59 | 9.65 | 7.13 | 3.88 | 6.89 |
| Mean | 6.94 | 7.28 | 7.55 | 8.20 | 0.52 | 0.58 | 0.62 | 0.58 | 9.28 | 6.97 | 3.45 | 6.57 |
|  | G | L | G×L |  | G | L | G×L |  | G | L | G×L |  |
| SEd | 0.051 | 0.028 | 0.089 |  | 0.011 | 0.006 | 0.019 |  | 0.050 | 0.027 | 0.086 |  |
| CD (P=0.05) | 0.103 | 0.056 | 0.178 |  | 0.022 | 0.012 | NS |  | 0.100 | 0.055 | 0.172 |  |

G – Maize genotypes L – Levels of silica NS – Non significant

## CONCLUSION

This study was a small attempt to know the effect of silicon fertilization on crop growth and their physiological characters. From the results, it is evident that Si does have a role in plant development and growth. Out of all the ten maize genotypes, COH(M)8 was found to be Si efficient as it was able to utilize Si better than others to produce more DMP and better growth attributes. Different physiological parameters and their relationship with growth due to silicon fertilization revealed that Si as calcium silicate improved the growth, physiology and antioxidant enzymes of hybrid maize genotypes. Also, relationship studies between dry matter production and growth attributes like root volume and relative water content showed a linear relationship, which also proved that Si increases the plant strength, promotes root and shoot growth, improves photosynthetic activity and gives the plants resistance against stress as well. Hence, Si fertilization is not only involved in stress tolerance mechanisms of the plants but also increased the plant establishment and growth.

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