Chapter 4 Role of Micronutrients in Biochemical Responses of Crops Under Abiotic Stresses

Shyam Narain Pandey

Abstract Essential micronutrients regulate and change the functioning of plants. The plants show tolerance in diverse ways under abiotic stress conditions, like drought, mineral deficiency, elevated salts concentrations in soil, etc., greatly contributed by the cellular biochemical reactions under suitable availability of micronutrients. Micronutrients such as zinc, copper, manganese, iron, molybdenum, boron and chloride support physiology and biochemical constituents in plants to neutralize the abiotic stresses. Under abiotic stresses, the micronutrients in plants, being constituents of biomolecules, function as an activator of many enzymes, electron carriers, etc. Also, their functions in metabolism regulations, reproduction, and protection against abiotic and biotic stress conditions are very significant. During diverse stress conditions, a large number of reactive oxygen species (ROS) are generated, and they damage the cellular metabolism of plants. The sufficient micronutrient status protects the plants against ROS by activating the enzymes and biomolecules to become a constituent or activator and directly favour the growth and cellular metabolism. This chapter describes the better interactions of plants and micronutrients for maintaining growth and metabolism and providing tolerance against abiotic stresses through changing the biochemical constituents in plants.

Keywords Abiotic stresses · Cellular metabolism · Micronutrients · Oxidative stress

1 Introduction

Today, global food demand is increasing due to the continuously growing population. Despite the development of agricultural technology for high production of food grain, abiotic stresses cause havoc on crop production. The major abiotic stresses in the world are drought, salinity, temperature (including global climatic change), mineral deficiency, heavy metals toxicity, etc. (Hussain and Shi [2014;](#page-16-0)

S. N. Pandey (\boxtimes)

Department of Botany, University of Lucknow, Lucknow, Uttar Pradesh, India

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Gautam et al. [2017](#page-16-1)). All plants require essential elements for their structure and functions. In recent decades, environmental stress conditions increased, and in this situation, the role of nutrients, particularly micronutrients, is very significant in the survival of plants (Pandey [2018\)](#page-17-0). The micronutrients such as zinc, iron, copper, manganese, molybdenum, boron, chlorine, etc. along with other macronutrients play a direct role in the growth and metabolism of plants (Sharma [2006\)](#page-18-0). Plants cannot survive in the absence of these micronutrients (Arnon and Stout [1939](#page-14-0)) even in normal growing conditions. The major abiotic stress conditions such as drought, salt and nutrients deficiency/toxicity are increasing day by day due to various anthropogenic activities. A large number of enzymes and organic molecules in cellular metabolism supported the defence system of plants against various stresses (Pandey et al. [2018](#page-17-1)). A large number of enzymes and biomolecules involved in plant defence system require micronutrient elements as constituents or activators (Pandey [2014\)](#page-17-2). In food chains, plant nutrition effects both on plants growth and survival of other living organisms (Karthika et al. [2018](#page-16-2)). Micronutrients play their role in various biotic and abiotic stresses, because they are electron carriers, regulate cellular metabolism and play a role in osmotic regulation. Micronutrients maintain the structural integrity of the plasma membrane during abiotic stress conditions (Brown et al. [2002\)](#page-15-0). Micronutrient such as iron is a cofactor of choline monooxygenase, a catalyst that helps in the biosynthesis of an osmoprotectant glycine betaine, which maintains the integrity of plasma membrane and stabilize the structure of proteins during salt and drought stress conditions (Gorham [1995\)](#page-16-3).

Micronutrients play their protective role in plants during stress conditions such as drought, salt and minerals deficiency/toxicity, heavy metals, etc. Under such stress conditions, production of reactive oxygen species (ROS) creates oxidative stress damage to cellular structure and functions (Pandey [2018](#page-17-0)). Micronutrients such as iron, manganese, copper and zinc are participants in detoxification of ROS. Also, they play a role in signal transduction (Apel and Hirt [2004\)](#page-14-1). In the antioxidant system of enzymes, many micronutrients are a cofactor for protecting cellular structure and functions from the excessive generation of ROS (Pandey and Gautam [2009](#page-17-3)). Under abiotic stresses, during electron transport system (in chloroplast and mitochondria), excessive ROS are produced. An important enzyme superoxide dismutase (SOD), which has micronutrient cofactors such as Cu-Zn SOD, Mn-SOD and Fe-SOD, supports the defence system against the ROS (Sharma [2006;](#page-18-0) Pandey [2014](#page-17-2)). These micronutrients containing SOD show high activity in mitochondria (Mn-SOD), chloroplast (Fe-SOD and Cu-Zn SOD) and cytosol (the activity of Cu-Zn SOD). The activity of SOD converts superoxide ions (O_2^-) into hydrogen peroxide (H₂O₂). The high concentrations of H₂O₂ are inducing lipid peroxidation and damage the cellular metabolic systems (Pandey 2018). So H_2O_2 concentration is prevented by its conversion into water (H₂O). This conversion of H₂O₂ to H2O is catalysed by Fe-containing enzymes catalase (CAT) and ascorbate peroxidase (APX). Ascorbate peroxidase scavenges H_2O_2 in the chloroplast in combination with the ascorbate-glutathione cycle (Asada [1997\)](#page-14-2). The enzyme catalase breaking down the H_2O_2 is present in peroxisomes. In peroxisomes, the conversion of glycolate to glyoxylate also contributes to accumulation of H_2O_2 , thus becoming a cofactor of the above antioxidative enzymes (catalase, APX and SOD), the micronutrients involved in the cellular defence system against oxidative stress (Van Wiren et al. [1996\)](#page-19-0). In cellular defence system, antioxidative enzymes decrease their activity due to limited availability of metal cofactors by various stresses in soil, which weakens the antioxidant defence system (Pandey [2014](#page-17-2)). Micronutrients are involved in various biosynthetic pathways (Pandey et al. [2018\)](#page-17-1). Biosynthesis of secondary metabolites is formed with catalytic actions of enzymes. These enzymes contain micronutrient as cofactors. Also, they play a role in lignifications of the cell wall, genes regulation, hormonal balance, ionic homeostasis and cell signalling (Stiles [2013\)](#page-18-1). A large number of enzymes present in the various compartment of the cell contain various micronutrients for performing their normal functions (Tej and Zeiger [2002](#page-18-2)). Even normal growth plants require a sufficient level of micronutrients. But availability of nutrients is greatly affected by soil physical, chemical and biological properties which determine the dynamics of the equilibrium of micronutrients in soil and its labile pool, from which micronutrients are absorbed by plant roots. From the labile pool, the acquisition of micronutrients is affected by the physical factors of the environment such as light intensity, pH, temperature, moisture, etc. and soil cultural practices. In addition, the genetic composition and stress disposition of plants influence the acquisition of micronutrients (Brady and Weil [1999;](#page-15-1) Karthika et al. [2018](#page-16-2)).

Salt stress conditions (salinity and sodicity) are a major threat to crops responsible to yield less by 4–6.5% annually across the world. In salt-affected soils, high concentration of sodium and chloride ions impairs the availability of essential nutrients to plant roots for absorption (Chakraborty et al. [2016](#page-15-2)), and these results in a reduction of crop productivity (Pandey [2014\)](#page-17-2). Similarly, the stress in soil due to excessive heavy metals activity affects plant growth and contaminates our delicate food chain (Naaz and Pandey [2010](#page-17-4); Roychowdhury and Tah [2011](#page-18-3); Basu et al. [2012;](#page-15-3) Roychowdhury et al. [2018,](#page-18-4) [2019\)](#page-18-5). Soil salinity causes nutrient loss, poor soil characteristics and essential nutrients imbalance. High concentration of salts in saline soils affects osmotic potential. Mostly, micronutrients such as Zn, Fe, Cu and Mn are less available for plants under high soil pH (alkaline) range, low soil moisture, high temperature and low organic matter content conditions (Brady and Weil [1999;](#page-15-1) Pandey et al. [2018\)](#page-17-1). The status of some micronutrients influences plants-water relations. The role of chlorine in osmoregulation involves cell osmotic relations and turgor-dependent extension growth of cells (Sharma [2006](#page-18-0)).

2 Zinc (Zn)

Zinc is an essential micronutrient for the structural and functional role of all living organisms (Pandey et al. 2002). Mostly, plants absorb zinc from the soil as Zn^{++} ion. The uptake of zinc from the soil and translocation to plant shoot portion are concentration-dependent, and some workers have reported carrier-mediated transport of Zn^{2+} (Norvell and Welch [1993\)](#page-17-6). In adverse soil conditions such as calcareousness, low temperature, salinity and alkalinity, zinc is less available in the soil for plants root absorption (Gautam et al. [2017\)](#page-16-1). The zinc-deficient soil conditions induce the plants for the secretion of phytosiderophores which mobilize zinc (Treeby et al. [1989\)](#page-19-1) and help their uptake in plants. Plant genotypes are variable in the secretion of phytosiderophores and efficiency of zinc uptake (Van Wiren et al. [1996\)](#page-19-0). Several identified transporters such as *ZIP* (Guerinot [2000\)](#page-16-4) and CDF (cation diffusion facilitator) family (Williams et al. [2000](#page-19-2)) facilitated zinc transport in plants. Zinc transporter genes (ZIP 1, 2, 3 and 4) have been isolated from *Arabidopsis,* and these genes are expressed in plant roots due to zinc deficiency (Guerinot [2000](#page-16-4)). A large number of proteins contain zinc as their structural component for performing their catalytic functions of more than 250 enzymes. These metal chelates with polypeptides are synthesized during heavy metals stress conditions in plants (Gautam et al. [2017\)](#page-16-1). Zinc provides stability of various proteins involved in regulatory functions, and such proteins are zinc fingers, RING finger domains and zinc clusters. Under excessive heavy metals concentrations in plant tissues, metallo-polypeptides help tolerance in plants against heavy metals in tissues (Pandey et al. [2002](#page-17-5)).

2.1 Enzymes

Zinc does not function during oxidation-reduction reactions due to its single oxidation state (Zn^{2+}) . Some important enzymes playing a vital role in cellular metabolism contain zinc as cofactor or activator, and these are DNA-dependent RNA polymerases, carbonic anhydrases, superoxide dismutase, carboxypeptidase A, etc. Zinc is used as a substrate by the enzyme H+ ATPase and functions as Zn-ATPase. Most enzymes contain zinc as constituents catalyse hydrolysis reactions (Sharma [2006\)](#page-18-0). The carbonic anhydrase catalyses the reversible conversion of carbon dioxide to bicarbonate $(HCO₃⁻)$ and is therefore critical to photosynthesis in $C₄$ plants growing in drought and high-temperature conditions. In C_4 plants in carboxylation reaction, bicarbonate is used as substrate and enzyme is PEPCase. Carbonic anhydrase enzyme is localized in mesophyll chloroplast in C_3 plants, while cytosol of mesophyll cell in C_4 plants. The cytosol is also the site for phosphoenolpyruvate carboxylase (PEPCase) using bicarbonate as substrate (Hatch and Burnell [1990\)](#page-16-5). The active sites of SOD contain copper and zinc. SOD (Cu-Zn SOD) protects plant cells against ROS. Zinc is also a constituent of some DNA-dependent RNA polymerases involved in transcription (Petranyl et al. [1978\)](#page-17-7). A large number of enzymes required zinc as a cofactor in animals, bacteria and fungi. The availability of zinc is influenced under various stress conditions or zinc deficiency, and the activity of all zinc-related enzymes is adversely affected. The decrease in the activity of carbonic anhydrase in leaves is a strong indicator of zinc deficiency (Rengel [1995\)](#page-18-6). Zincdeficient plants also show decreased activity of SOD (Pandey et al. [2002](#page-17-5)). The low activity of catalase and ascorbate peroxidase in plants due to zinc deficiency has been reported (Yu et al. [1998;](#page-19-3) Pandey et al. [2002](#page-17-5)). The role of glutathione reductase (GR) is increased in response to environmental stresses and xenobiotics which

provide tolerance to the plants against these stresses. The decrease in GR activity in response to zinc deficiency in bean plant has been reported by Cakmak and Marschner ([1993\)](#page-15-4). Under stress conditions, zinc protects plants by an inhibitory effect on plasma membrane-bound NADPH oxidase (catalyses the production of superoxide ions).

2.2 Physiological Role

Plants show zinc deficiency grown under soil salinity, sodic or calcareousness stresses performed poor photosynthetic functions (Karthika et al. [2018](#page-16-2)). Low zinc in cells changes chloroplast structure, electron transport and carbon dioxide fixation (Sharma et al. [1994\)](#page-18-7). The production of ROS increases due to zinc deficiency in plant tissues, and ROS damage in the thylakoid membranes consequently decreases the photosynthetic active area in leaf (Cakmak and Engels [1999](#page-15-5); Henriques [2001\)](#page-16-6). Carbonic anhydrase catalyses the production of the substrate $(HCO₃⁻)$ in $C₄$ plants and maintains $CO₂$ concentration in $C₃$ plants at the site of carboxylation by the conversion of HCO_3^- to CO_2 under the role of RuBP carboxylase (Hatch and Burnell [1990\)](#page-16-5). The activity of carbonic anhydrase is inhibited in plants under the influence of zinc deficiency (Badger and Price [1994\)](#page-15-6). In zinc-deficient plants, decrease in catalase and peroxidase activity, photosynthetic rate and accumulation of essential oil in leaves have been studied (Pandey [2018](#page-17-0); Srivastava et al. [1997\)](#page-18-8). Marschner et al. ([1996\)](#page-17-8) observed increased accumulation of carbohydrates in leaves of zincdeficient plants. The increased accumulation of starch in zinc-deficient condition correlates the role of zinc in sucrose biosynthesis (Shrotri et al. [1980](#page-18-9)) and starch metabolism. Zinc is also involved in auxin metabolism by the help of tryptophandependent auxin biosynthesis. Tryptophan is a significant precursor of auxin which is a growth-promoting plant hormone. Zinc deficiency reduces the protein content, while increases the non-protein nitrogen in plants (Bisht et al. [2002](#page-15-7)) indicates the role of zinc in protein synthesis. The metabolism of nucleic acids and ribosome and zinc as a constituent of a transcription factor may also affect zinc deficiency which leads to decrease in protein synthesis. Low zinc status in plants suppressed the various stages of reproductive development (Pandey and Gautam [2009\)](#page-17-3) such as flowering, male and female gametogenesis, seed development, development of anther, etc. (Sharma [2006](#page-18-0); Pandey [2014](#page-17-2)). Zinc is a critical nutrient element for microsporogenesis and pollen fertility. Pollen tube accumulated zinc at their part that participated in fertilization. Zinc finger protein (TFIIIA type) controls the development of the flower. The zinc finger proteins (anther-specific) play a role in microsporogenesis (Kobayashi et al. [1998\)](#page-16-7). Zinc-deficient plants accumulate high proline content which is indicative of water stress (Sharma and Sharma [1987](#page-18-10); Pandey [2018\)](#page-17-0).

2.3 Oxidative Stress

Plants produce high reactive oxygen species when influenced with zinc deficiency. Also, the power of defence system decreases which neutralizes the toxic effects of ROS (Cakmak [2000;](#page-15-8) Pandey and Gautam [2009;](#page-17-3) Singh and Pandey [2011\)](#page-18-11). The activity of copper-Zn SOD, ascorbate peroxidase and catalase antioxidant enzymes is reduced in plants due to zinc deficiency (Yu and Rengel [1999](#page-19-4); Naaz and Pandey [2009;](#page-17-9) Gautam et al. [2017\)](#page-16-1). High production of free oxygen radicals oxidizes lipid in membrane causing leakage of cell sap due to structural changes in the plasma membrane. The toxic effect of ROS also leads damage to the photosynthetic apparatus (Cakmak and Engels [1999;](#page-15-5) Henriques [2001](#page-16-6)).

3 Copper (Cu)

Copper exists in $Cu⁺$ (cuprous) and $Cu⁺⁺$ (cupric) oxidation states, particularly in oxidation-reduction reactions. In cellular metabolism, many copper-containing proteins act as enzymes and electron carriers. It is an important constituent of cytochrome C oxidase. In the mitochondrial electron transport system, copper as constituents of cytochrome C oxidase is the terminal component. The copper protein (plastocyanin) is an electron carrier in the light reaction phase of photosynthesis. Copper is a component of many proteins detoxifying the effect of reactive oxygen species highly produced during abiotic stress conditions. Copper is taken by plants roots through active transport proteins across the plasma membrane (Fox and Guerinot [1998\)](#page-15-9). Copper transporter proteins COPT1, COPT2, COPT3, COPT4 and COPT5 have been identified from *Arabidopsis* (Kampfenkel et al. [1995](#page-16-8); Sancenon et al. [2003](#page-18-12)). These copper transporters have been detected in leaf, stem, flowers and roots, but their expression was different in different parts of the plant. The cupric ions long-distance transports are facilitated by Cu^{++} -amino acid complex formation, and the amino acids involved in the transport process are asparagine, histidine and glutamic acid (Loneragan [1981](#page-16-9)). Under copper deficiency, transport from leaves to grains is poor (Loneragan [1981](#page-16-9)), particularly at the time of seed-filling stage.

3.1 Physiological Functions

Copper causes reduction of one atom of O_2 to water and another atom of O_2 to hydroxylation of substrates such as phenolase complex (Sharma [2006](#page-18-0)). It is a cofactor of a large number of enzymes in plants such as Cu-Zn SOD and cytochrome C oxidase. The metal receptor protein (metallochaperones) delivers copper metal Cu-proteins (Lippard [1999\)](#page-16-10). Some important copper-containing enzymes are ascorbate oxidizer, SOD, cytochrome C oxidase, diamine oxidase, etc. Catechol oxidase

(copper proteins) or polyphenol oxidases are important for secondary metabolism in plants. They play a role in the biosynthesis of lignin and alkaloids. Also, in the formation of phytoalexins, they have antifungal activity. Therefore, phenolases involved in metabolism provide strength to the cell wall for the protection of pathogens (Pandey et al. [2018](#page-17-1)). Laccase or p-diphenols oxidase contains four copper ions (both $Cu⁺$ and $Cu⁺⁺$) acting as a catalyst in the oxidation of p-diphenols to p-quinones. Laccase is involved in the synthesis of plastoquinones, useful in PS-II (Ayala et al. [1992\)](#page-14-3). Tyrosinase enzyme was identified as mushrooms involved in the conversion of tyrosine to dopa-quinone. The Cu-Zn SOD enzyme involved in the conversion of ROS to H_2O_2 and oxygen, therefore, provides protection of cells from ROS. The enzymes SOD and ascorbate peroxidase are attached to the chloroplastic thylakoids in the vicinity of the PS-I. Ascorbate peroxidase is involved in the reduction of H_2O_2 to $H₂O$ and protects cellular lipid peroxidation (Pandey [2018\)](#page-17-0). Copper is a constituent of single-electron carriers protein plastocyanin found in chloroplasts. Plastocyanin is an electron carrier linking two pigment systems (PS-I to PS-II). Copper element that binds to polypeptides forms copper-chelating biomolecules (Maksymiec [1997](#page-17-10)). Copper deficiency causes a reduction in photosynthesis (Ayala et al. [1992;](#page-14-3) Pandey [2018\)](#page-17-0). In some blue-green algae, plastocyanin synthesis is with the availability of copper. Under salinity and alkali soil conditions, low availability of Cu inhibits the Cu-Zn SOD activity and plastocyanin-mediated transport of electrons within the chloroplasts (Shikanai et al. [2003](#page-18-13)). Low copper in plants causes ultrastructural changes in chloroplast thylakoids in grana and decrease in chlorophyll content (Casimiro et al. [1990](#page-15-10)).

3.2 Stress Neutralization

Copper involves in the disproportion of superoxide ions (O_2^-) produced due to abiotic stresses because it is constituent of the enzyme Cu-Zn SOD for the conversion of H_2O_2 and molecular oxygen. Therefore, low copper weakens the oxidative defence system in plants. Copper plays a protective role in plants against pathogens through their involvement in the synthesis of lignin (Graham [1983](#page-16-11)). Copperdeficient plants are more vulnerable to pathogenic infections. During stress conditions, copper induces the formation of ROS and activates copper diamine oxidases that trigger defence mechanisms in the plant (Sharma [2006\)](#page-18-0).

4 Iron (Fe)

Iron exists in ferric (Fe^{***}) and ferrous (Fe^{**}) oxidation states and participates in oxidation-reduction reactions. It participates in the electron transport system, and it is a cofactor of several enzymes. They play important roles in fatty acids metabolism and biosynthesis of terpenoids, signalling molecules and growth hormones.

The role of iron is known for induction to produce and detoxification of reactive oxygen species. The biochemical reactions related to iron are very complex. Iron also plays a very important role in signal transduction and biosynthesis of Jasmonic acid (Sharma [2006\)](#page-18-0). Under Fe deficiency conditions, mostly plants uptake reductionbased iron (Bienfait et al. [1983\)](#page-15-11). The ferric ion chelate reduced into ferrous ion chelate by the catalytic activity of an enzyme known as Fe (III) chelate reductase. This reductase enzyme is induced in the plasma membranes of root epidermal cells under Fe deficiency (Sharma [2006\)](#page-18-0). Yi and Guerinot ([1996\)](#page-19-5) described that the *Arabidopsis* mutants ferric reductase defective 1, 2 and 3 suppress the activity of Fe+++ chelate in *Arabidopsis* as reported by Robinson et al. ([1999\)](#page-17-11) and named it ferric reductase oxidase 2 (FRO2). The various genes involved in uptake in roots, as well as their distribution in shoot portions, are characterized (Robinson et al. [1999;](#page-17-11) Waters et al. 2002). Extracellular pH influences the solubility of $Fe⁺⁺$ in the root zone area, which is observed maximum at pH 5.0. High pH reduces the Fe^{+++} reduction (Manthey et al. [1996](#page-17-12); Zheng et al. [2003](#page-19-7)). Plant hormone ethylene helps in the uptake of iron in roots (Zaid et al. [2003](#page-19-8)) by changing the root morphology of irondeficient plants (Schmidt et al. [2000\)](#page-18-14). Several Fe transports proteins in plasma membranes, and several transporter genes of different gene families have been identified and characterized such as iron-regulated transporter (IRTI) belonging to ZIP family identified in *Arabidopsis* (Eide et al. [1996](#page-15-12)). Nramp family transporters are also involved in the uptake of iron (Thomine et al. [2000\)](#page-18-15). The strategy of the plant to uptake the iron with the reduction of Fe^{3+} to Fe^{2+} by the catalytic activity of Fe chelate reductase is established only under iron deficiency condition. The second strategy is established for iron uptake by the formation of Fe complex of phytosiderophores. The mugineic acid forms complex with $Fe³⁺$ that is prerequisite for its recognition by the specific Fe⁺⁺⁺ transport system in roots (Ma and Nomoto [1996;](#page-16-12) Ma et al. [1993\)](#page-17-13). Mugineic acid synthesizes from methionine chelate $Fe³$ + with their carboxyl groups and amino groups. The graminaceous plant species secrete mugineic acid (phytosiderophores, when soil is deficient in iron), but the secretion of specific mugineic acids (non-proteinogenic amino acids) is different (Sharma [2006\)](#page-18-0). The mugineic acids (MA) are secreted by rice, corn and wheat. Soybean genotypes are 2-DMA, 3-HMA (3-hydroxyl mutagenic acid) and 3-epihydroxy mugineic acid (epi-HMA). These MAs provide tolerance to plants against iron chlorosis. The transport of iron to the shoot portion takes place through xylem as ferric citrate organic complex (Cataldo et al. [1996\)](#page-15-13).

4.1 Physiological Roles

A large number of iron (Fe)-containing enzymes and carrier proteins are present in electron transport systems. These enzymes are essential for the biosynthesis of secondary metabolites, plant hormones and some other biomolecules significant as signalling molecules and developmental activities. Some enzymes are di-iron enzymes in which two iron atoms forming an Oxbridge are bound to two histidine and four carboxylate residues such as mitochondrial alternative oxidase (Sharma [2006](#page-18-0)). In non-haem iron enzymes, Fe cofactor is present as Fe-sulphur cluster which enables iron-sulphur proteins to be as donors or acceptors of electrons. The iron clusters are Fe-S, 2Fe-2S (two Fe ions are coordinated to two inorganic sulphide ions) and 4Fe-4S (contains four Fe ions and four sulphide ions). The enzymes containing Fe as activator are catalase, cytochrome C oxidase and peroxidases, and they are iron bound to the apoprotein as iron porphyrin. There, Fe ion is complexed to the four nitrogen atoms of the four pyrrole groups. Some non-haem iron enzymes are superoxide dismutase (SOD), lipoxygenase, alternative oxidase and 2-oxoglutaratedependent enzymes. Some important non-haem iron enzymes with Fe-S cluster cofactor are nitrite reductase aconitase, sulphite reductase, formate dehydrogenase, succinate dehydrogenase, NADH-Q oxidoreductase and FDX-thioredoxin reductase. The various groups of enzymes containing Fe involved in detoxification of reactive oxygen species are SOD, catalase, peroxidase, alternative oxidase, etc. A large number of Fe proteins are electron carriers, a significant part of mitochondrial and chloroplastic electron transport system including ferredoxins and cytochromes. Leghaemoglobin in leguminous plants has a high affinity for O_2 molecule. The phytoferritin serves as a store of cell's excess iron in plants (Sharma [2006](#page-18-0)). Haem protein cytochromes are localized in chloroplasts, in mitochondrial inner membrane and in the endoplasmic reticulum, and cytochromes are involved in reversible reactions of two iron oxidation states ($Fe^{++} = Fe^{++}$). Ferredoxins (2Fe-2S)-type ironsulphur clusters function in electron transport in between iron oxidation states, mainly localized in chloroplasts and root plastids. Chloroplastic ferredoxin is reduced by taking electrons from P700, and it is an electron donor for several reductive reactions such as glutamine to glutamate and reduction of nitrite and sulphite. The phytoferritins in plants constitute a major sink of iron accommodating about 4500 iron atoms (Smith [1984\)](#page-18-16). Iron plays a significant role in photophosphorylation and, as a part of ferredoxin and ferredoxin-thioredoxin reductase, functions in the photoreduction of thioredoxin. In the case of some plants, the ferredoxin-thioredoxin regulates the activity of Rubisco (Zhang and Portis [1999\)](#page-19-9) and also functions in the Calvin cycle. The iron proteins NADH-cytochrome b_5 reductase, cytochrome b_5 and a desaturase are membrane-bound proteins involved in fatty acid metabolism (Sharma [2006](#page-18-0)).

4.2 Free Radical Production

The reduction of O_2 and H_2O_2 generates superoxide ions which are dissimulated by SOD and make two ROS causing lipid peroxidation damaging cellular components (Pandey [2018\)](#page-17-0). Superoxide ions react to H_2O_2 to produce hydroxyl radicals

$$
\text{Fe}^{2+} + \text{H}_2\text{O}_2 \rightarrow \text{Fe}^{3+} + \text{OH}^{\circ} + \text{OH}^{-}
$$

These ROS damage proteins, lipid and DNA and induce mutations. Ferrous ions also react with oxygen molecules to generate $Fe^{2+}O$ (ferryl) and $Fe^{2+}O_2$ (per ferryl), which are more toxic compounds. The activity of lipoxygenase (iron enzyme) in the hydroxylation of linoleic acid catalyses the production of singlet oxygen species $(^1O_2)$.

4.3 Protective Role

In plants, iron also participates in the defence system against salt stress, oxidative stress and various pathogens attack. Although iron ions promote the production of ROS, iron-containing enzymes are participating in detoxification of ROS (Roychowdhury et al. [2018,](#page-18-4) [2019\)](#page-18-5). The iron-SOD detoxify the superoxide ion (O2−) conversion into H_2O_2 . Other enzymes catalase and ascorbate peroxidase (haem enzyme) detoxify the H_2O_2 by converting them into H_2O . In the iron-deficient plant, decrease in the activity of ascorbate peroxidase and consequently increase in H_2O_2 concentration in sunflower have been reported (Ranieri et al. [2001](#page-17-14)). Quinol reacting with oxygen generates ROS, and iron enzyme alternative oxidase reduces quinol by providing an alternate pathway in the mitochondrial electron transport system, thus preventing the production of ROS. Iron plays a role as an osmoprotectant through biosynthesis of glycine betaine against high temperature and high salinity (Gorham [1995\)](#page-16-3). Also, iron is involved in the biosynthesis of lignin, which provides mechanical strength to cell wall against pathogens (Caruso et al. [2001\)](#page-15-14).

5 Manganese (Mn)

Manganese (Mn) exists in various oxidation states such as Mn^{2+} , Mn^{3+} , Mn^{4+} and Mn^{5+} , out of which the most prominent is Mn^{2+} . It forms a rapid bond with oxygencontaining species. Mn behaves as an activator of enzymes, plays a catalytic role in cellular metabolism and also functions in $CO₂$ fixation in CAM and $C₄$ plants. It also plays a significant role in scavenging oxygen free radicals. Mn2+ions are absorbed by plant roots through facilitated diffusion. The acidification outside the plasma membrane by H⁺ efflux increases uptake of Mn (Yan et al. [1992](#page-19-10)). Several genes have been identified and characterized, and they are encoding Mn transporter proteins (Hall and Williams [2003\)](#page-16-13). The P-type ATPase is involved in Mn transport. The transporter ECA1 in *Arabidopsis* functions as an endoplasmic reticulum-bound $Ca²⁺/Mn$ pump, which also provides tolerance to the plants to excess manganese (Wu et al. [2002](#page-19-11)). Mn anti-portal Sh MTP1 confers tolerance to manganese in plants through internal sequestration (Delhaize et al. [2003\)](#page-15-15). Nramp family transporter participates in manganese transport, and Nramp 3 plays a role in long-distance transport of manganese (Thomine et al. [2003](#page-18-17)). The translocation of manganese to developing grains is slowly observed in wheat.

5.1 Physiological Role

Over 30 enzymes contain Mn as constituents. Some important enzymes with Mn are Mn-SOD (manganese superoxide dismutase), PEPCK (phosphoenolpyruvate carboxyl kinase), NAD + malic enzyme, IDH (isocitrate dehydrogenase), NADP + malate enzyme, PEPCase (phosphoenolpyruvate carboxylase), glutamine synthetase, enolases, etc. PEPCK catalyzes the decarboxylation of oxaloacetate to PEP in the chloroplast of bundle sheath in C_4 plant. The released carbon dioxide is the starting point of the Calvin cycle.

Oxaloacetate + $ATP \rightarrow Phosphoenolpyruvate + ADP + CO₂$

The significant role of Mn is the oxidation of water in PS II (Ono and Onone [1991\)](#page-17-15). Many enzymes in C_4 plants containing Mn²⁺ as cofactor help in photosynthesis in C_4 plants. Manganese is also involved in the decarboxylation of C_4 acids to generate carbon dioxide in bundle sheath cells. Such produced $CO₂$ is again fixed in a $C₃$ compound by Rubisco (Sharma [2006](#page-18-0)). Manganese-activated enzymes catalyse the biosynthesis of secondary metabolites in plants. It is also involved in biosynthesis pathway systems of pigments as an activator of many enzymes. The manganese plays a significant role in the biosynthesis of aromatic amino acids (tyrosine, tryptophan and phenylalanine). These amino acids are precursors of flavonoids, indole and lignin (Burnell [1988\)](#page-15-16). Manganese is the activator of arginase and plays a role in the synthesis of polyamines. Polyamines are important for plant growth and development. Deficiency of Mn decreases photosynthesis in plants because Mn is an important constituent of PS II (Yachandra et al. [1993\)](#page-19-12), and leaf ultrastructure changes are induced by Mn deficiency (Polle et al. [1992\)](#page-17-16). Plants show an increase in soluble carbohydrate and a decrease in fatty oil content under manganese deficiency (Campbell and Nable [1988\)](#page-15-17). Increase in soluble nitrogenous compounds due to Mn deficiency has been reported in plants because Mn is an activator of arginase and allantoate amidohydrolase (Winkler et al. [1985](#page-19-13)). Also, Mn deficiency is related to the accumulation of arginine and uriedes and reduction in nucleic acids (Chatterjee et al. [1994\)](#page-15-18).

5.2 Protective Role

Manganese-SOD (superoxide dismutase) localized in mitochondria is an important enzyme of the antioxidant system; thus, Mn is involved in the defence system against reactive oxygen species. It prevents the accumulation of superoxide ions by dismutation of them and converts it to hydrogen peroxide (Sharma [2006\)](#page-18-0). Low Mn in plants shows oxidative stress. Overexpression of Mn-SOD in transgenic tobacco increases tolerance in plant against oxidative stress as reported by Slooten et al. [\(1995](#page-18-18)). The Mn-SOD also contributes tolerance to plants against drought stress.

Increased Mn-SOD expression induced under drought stress has been observed in wheat seedlings as reported by Wu et al. [\(1999](#page-19-14)). The concentrations of tissue manganese are related to disease in plants, and it supports the resistance in plants to pathogenic diseases (Graham [1983](#page-16-11)). Manganese contributes protection and strength in plants to fungal infection through its involvement in the biosynthesis of phenolics and lignin (Morab et al. [2003](#page-17-17)).

6 Molybdenum (Mo)

Molybdenum (Mo) is a second transition series metal that exists in several oxidation states, and the most stable is the Mo VI (hexavalent) form. Molybdenum has an affinity for sulphur-containing groups. It participates in oxidation-reduction reactions due to its variable and easy convertibility oxidation states. Molybdenum is a cofactor of several enzymes. Mo plays a role in dinitrogen fixation and assimilation of nitrate in plants, and it is also essential for free-living and symbiotic bacterial nitrogen fixation processes. Plants take Mo as Mo O_4^2 (molybdate ion). Molybdenum follows some transporter path as transport of phosphate. A higher concentration of sulphate decreases the uptake of molybdenum (Marschner [1995\)](#page-16-14). For redox reactions in plants and microbes, Mo is a cofactor of several enzymes (Sharma [2006\)](#page-18-0). Xanthine oxidase and nitrate reductase are Mo-containing enzymes. These enzymes involve in catalysing the reactions for nitrogen and nitrogenous compounds. Aldehyde oxidase contains iron and MoCo as prosthetic groups catalyse the terminal reaction of the biosynthetic pathway of abscisic acid and auxins (Taylor [1991\)](#page-19-15). Molybdenum-containing enzymes also contain additional cofactors such as Fe-S, iron, haem, etc. (Koshiba et al. [1996](#page-16-15)).

6.1 Physiological Roles

The molybdoenzyme nitrate reductase is very significant to catalyse the conversion of nitrate to nitrite, and the enzyme plays a role in the synthesis of organic nitrogenous compounds. The bacterial and eukaryotic nitrate reductase enzymes belong to two different families of molybdoenzymes (Sharma [2006](#page-18-0)). The enzyme nitrate reductase distribution differs in different plant species but is localized in the cytosol in all plant species. The molybdoprotein xanthine dehydrogenase has two similar and catalytically independent subunits. Each subunit contains a molybdoprotein, two Fe-S clusters and one flavin cofactor which catalyses oxidative degradation of xanthine to uric acid with the use of NAD+ as an electron acceptor (Sharma [2006\)](#page-18-0). Sulphite oxidase works in plants and human beings, and it is an essential liver enzyme. Sulphite oxidase is a molybdoprotein but lacks the redox-active centres (Mendel and Hansch [2002](#page-17-18)). Aldehyde oxidase contains FAD (flavin), iron and MoCo as prosthetic groups, it catalyses the oxidation of the abscisic aldehyde to abscisic acid (Seo and Koshiba [2002\)](#page-18-19) and indole-3-acetic acetaldehyde to indole-3 acetic acid, and it regulates growth in plants. Dinitrogenase (Mo-Fe protein) and dinitrogenase reductase (Fe protein) are nitrogenase (prokaryotic enzyme) that catalyses the nitrogen fixation process. Several other non-molybdoenzymes such as catalase (Agarwala et al. [1986](#page-14-4)), cytochrome oxidase (Chatterjee et al. [1985](#page-15-19)) and succinate dehydrogenase (Agarwala et al. [1986\)](#page-14-4) are influenced by Mo status in plants. Deficiency of Mo induces changes in several organic compounds (Sharma [2006](#page-18-0)).

7 Boron (B)

Boron (B) is metalloid that exists in three-valence status with high affinity to oxygen. Boron-diol complexes are formed between as-diol furanoid groups of sugars (apiose and fucose) and boric acid. Therefore, boron plays an important role in the structural and functional process for plant cell walls and membranes (Roychowdhury et al. [2019\)](#page-18-5). Boron bridging the hydroxyl groups gives a large number of boroncontaining compounds in plants (Dembitsky et al. [2002\)](#page-15-20). The important role of boron is cross-linking of cell wall polysaccharides which has been established (O' Neill et al. [2004](#page-17-19)). A most significant form of boron in soils is boric acid (B [OH] 3) that exists mostly at pH 7.5 in the cytoplasm (Woods [1996\)](#page-19-16). The uptake of boron from the soil is a passive process with formation of boron complexes which create concentration gradient. A investigation has also been reported for the uptake of boron is active or passive or both (Pfeffer et al. [1998\)](#page-17-20). The transport of boron for long distances is facilitated by transpiration stream through the xylem, whereas retranslocation of boron to sites that do not lose water such as fruits, inflorescence, etc. involves phloem (Brown et al. [2002](#page-15-0)).

7.1 Physiological Role

Boron plays an important structural role in the cell wall as it binds with dial groups of polysaccharides. It is also highly accumulated in the cell wall as an integral component of polysaccharide complexes (Moth et al. [1993\)](#page-17-21). Boron influences the uptake of ions from roots. Its effect on membrane potential (Em) is demonstrated by Schon et al. (1990) (1990) . Boron-deficient plants show high efflux of $K⁺$, sucrose and amino acids (Cakmak et al. [1995\)](#page-15-21) due to structural changes in the cell wall. Short-term boron deficiency increases the accumulation of actin and tubulin involved in alteration in the polymerization pattern of the cytoskeletal assemblies as investigated in maize root apices by Yu et al. [\(2002](#page-19-17)). Boron increases photosynthetic oxygen evolution and photosynthetic efficiency in leaves. The supply of boron in plants affects the metabolism of carbohydrate, affecting the activity of related enzymes. The pentose phosphate pathway (alternative to glycolysis) is influenced by boron

concentration in the plant (Sharma [2006](#page-18-0)). The activity of glucose-6-phosphate dehydrogenase and 6-phosphogluconate dehydrogenase increases in response to boron deficiency in plants (Lee and Aronoff [1967\)](#page-16-16). In boron-deficient plants, the activity of acid phosphatase increases (Agarwala et al. [1991](#page-14-5)).

8 Chlorine (Cl)

Chlorine is a halogen element (group VII) with a single oxidation state as Cl− ions. These ions are mostly bound to organic molecules or exchange sites. Among chlorine-containing organic compounds, 4-chloroindole acetic acid shows high auxin activity (Flowers [1988](#page-15-22)). The absorption of the free anion of chloride is by plant roots and also by shoot portion of plants in gaseous form or as chloride ions (White and Broadley [2001](#page-19-18)). Chloride uptake through plasma membrane occurs either by active transport (by H^+ ATPases) or by facilitated diffusion. The active transport with ATP hydrolysis involves a Cl[−]/nH⁺ symporter (Sharma [2006](#page-18-0)).

8.1 Physiological Roles

In manganese-linked oxygen evolution complex (of photosystem II), chlorine is a structural constituent. Chlorine deficiency in plants decreases photosynthesis and evolution of $O₂$ by chloroplasts of higher plants (Kelley and Izawa [1978](#page-16-17)). Chlorine deficiency inhibits extension growth of cells and cell division, and consequently, leaf area for photosynthesis is reduced (Terry [1977](#page-18-21)). Stomatal functioning indirectly affects photosynthesis process because proton pumping across vacuolar membrane induces an influx of chloride simultaneous with K^+ , followed by water movement in guard cells, increase turgor and the opening of stomata, and affects the exchange of carbon dioxide (Schnabl [1980](#page-18-22)). Chlorine does not function as a catalyst, directly. It stimulates asparagine synthetase, which is important to catalyse the glutamate-dependent synthesis of asparagine. Asparagine synthetase also catalyses the amination of aspartate to asparagine. The synthesis of asparagine resulted in the storage of nitrogen and its transportation from source to sink (Rognes [1980\)](#page-17-22).

8.2 Osmoregulation

Chlorine functions as osmoregulator in plants and therefore affects plant-water relations. The osmoregulatory functions such as cell division in shoot and root apices, stomatal functioning and development of stigma require chlorine. It causes turgorinduced extension cell growth in the apical portion of root and shoot. The rapid growth of stigma induces extension growth of cells which increases cell turgidity

followed by quick mobilization of Cl− and K+ ions from the neighbouring cells as reported by Heslop-Harrison and Roger [\(1986](#page-16-18)). Such osmoregulatory functions of chlorine are very significant in reproductive biology. A carbon dioxide-induced chlorine efflux from guard cells into the apoplastic fluid is linked to the activation of the anion of protein channels in the plasma membrane of guard cells (Hanstein and Felle [2002\)](#page-16-19) observed in faba bean. The concentrations of chlorine for osmoregulation process are variable from plant to plant, and osmotic responses are also variable under abiotic stress conditions (Sharma [2006](#page-18-0); Roychowdhury et al. [2013;](#page-18-23) Ganie et al. [2014](#page-16-20); Roychowdhury [2014](#page-18-24); Hasanuzzaman et al. [2015;](#page-16-21) Pandey [2018](#page-17-0)).

9 Conclusions

The role of micronutrients in plant's biochemistry is very significant, particularly during biotic and abiotic stress conditions, because they play a significant role in the regulation of enzymatic reactions and biosynthesis of organic molecules. During abiotic stress conditions, the strength of the defence system, new synthesis and induction of biomolecules, induction of signalling, etc. are facilitated by appropriate concentration of essential micronutrients in plant tissues. The critical normal range of micronutrients not only promotes the growth but also protects cellular metabolism against abiotic stresses such as drought, mineral deficiency, salinity, adverse temperature, heavy metals and chemical toxicity.

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