

Organic acids of crop plants in aluminium detoxification

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Phytotoxicity of aluminium is one of the most serious problems in limiting plant growth in acidic soils. A number of plant species exhibit inheritable aluminium tolerance by the secretion of organic acids which is highly specific to aluminium stress and localized to the root apices. Organic acids have been considered to play an important role in the detoxification of aluminium, both externally and internally. Some plants detoxify aluminium by the secretion of organic acids from the roots. Other plant species that accumulate aluminium in their leaves, detoxify aluminium internally by forming complexes with organic acids. The kind of organic acids and the secretion pattern depends on plant species. This review summarizes current understanding of the mechanism and regulation of the secretion of organic acids from roots under aluminium stress. The advantages that plants get from the presence of organic acids in the rhizosphere are described and the biotechnological approaches to increase the secretion of organic acids are highlighted.

Keywords: Aluminium phytotoxicity, crop plants, detoxification, organic acids.

PHYTOTOXICITY of aluminium ion (Al^{3+}) is a serious problem limiting crop production in acidic soils in many parts of the India. This is particularly important in the northeastern region, where more than 90% of the soils are acidic, creating the potential for Al toxicity in surface and subsurface layers¹. The initial response to Al toxicity is inhibition of root elongation by destroying the root apex², resulting in inefficient uptake of water and nutrients. Several strategies have been adopted to manage acid soils. The primary method has been the application of large amount of lime to raise soil pH. However, liming is not a remedy for sub-soil acidity and it is not always economically feasible. An alternative to liming is to select and breed aluminium-tolerant cultivars on the acid soil. Several studies provided strong evidence that Al-tolerant genotypes have developed strategies to adapt to Al toxicity, and one of these strategies involves the efflux of organic acids^{3,4}. The Al-dependent stimulation of organic acid efflux from roots has now been reported in many species, and this response has been associated with an

increase in Al resistance. The anions of organic acids secreted by the roots are thought to chelate the toxic Al cations, and thus prevent them from interacting with the root apices. However, some researchers observed that the secretion of organic acids is not the only mechanisms for aluminium resistance in plants. The understanding of the mechanisms and regulation of organic anion secretion from the roots under aluminium stress are not yet fully understood. The biochemical, physiological and genetic bases of the mechanisms can produce and export organic acids to the root apoplast and rhizosphere. Such aspects are beginning to be understood and will be the main focus of this review.

Strategies for detoxification of Al

Crop species and cultivars exhibit wide genetic variability, both within and between species, in their response to Al resistance of plants, suggesting that Al-tolerant species or cultivars possess mechanisms for detoxifying Al. There have been two strategies for the detoxification of Al by plants cells^{5,6}. One is the exclusion of Al from the root tips and the other is tolerance to Al that absorbs the plant cells⁶. The main difference between these two strategies is the site of Al detoxification: apoplasm (external) and symplasm (internal). Thus organic acids play an important role in the external and internal neutralization of Al.

Two patterns of secretion of organic acids have been proposed in terms of time required in plants (Figure 1)³: The release of organic acids is quick in pattern I, whereas it is delayed for several hours in pattern II after the addition of Al in nutrient solution. In the first pattern, there is no discernable delay between the moment of Al addition and the onset of organic acid efflux. Activation of an anion channel located on the plasma membrane by Al is a possible mechanism responsible for quick secretion⁷⁻⁹. This quick response suggests that the necessary metabolic machinery is constitutively expressed in the root cells and that organic anion efflux is simple triggered by Al^{3+} and induction of a novel protein is not required¹⁰. In pattern I, as observed in wheat^{9,10}, buckwheat^{8,11} and barley¹², the secretion of organic acids is rapidly activated (15–30 min) after exposure of the plants to Al solution and the rate of release remains constant with time. In Al-tolerant

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genotype of wheat, Al-stimulated secretion of malate from both intact roots and excised root apices was observed within 20 min exposure to Al¹³. Similarly, Osawa and Matsumoto¹⁰ showed that malate efflux started 5 min after the addition of Al in wheat. The efflux rate in this pattern is the same at any time after the exposure to Al. In the second pattern, organic acid efflux is delayed for several hours after exposure to Al¹⁴⁻¹⁶ and protein induction is required¹⁷. These induced proteins could be involved in organic acid metabolism or in the transport of organic acid anions. Such secretion pattern was observed in *Cassia tora*¹⁸, rye¹⁵, triticale¹⁹ and rice bean¹⁶. For example, in *C. tora* maximal efflux of citrate occurs after 4 h of exposure to Al¹⁸; rice bean roots released citrate to alleviate Al toxicity and the efflux was delayed by at least 3 h (ref. 16), and in rye citrate and malate efflux increased steadily during a 10-h period¹⁵. In Al-resistant cultivars of maize, a considerable lag phase before the maximal citrate efflux was observed^{19,20}. Al-induced secretion of malate and citrate was found to significantly increase after 6 and 12 h respectively, in triticale line²¹. At sufficient concentrations, these organic acids can form complexes with Al ions, prevent the Al ions from binding to the fixed negative sites of the cell wall and plasma membrane, and confer Al tolerance to plants to maintain the normal functions of the cell wall and plasma membrane¹⁹.

External exclusion for Al tolerance

Secretion of organic acids from roots under Al stress has been identified as the most important mechanism avoiding Al toxicity²⁰ as result of the formation of Al chelates with organic acids^{22,23}. Among organic acids, citrate, oxalate, malate and acetate have been identified as important Al-chelating ligands secreted by the roots^{21,24,25}.

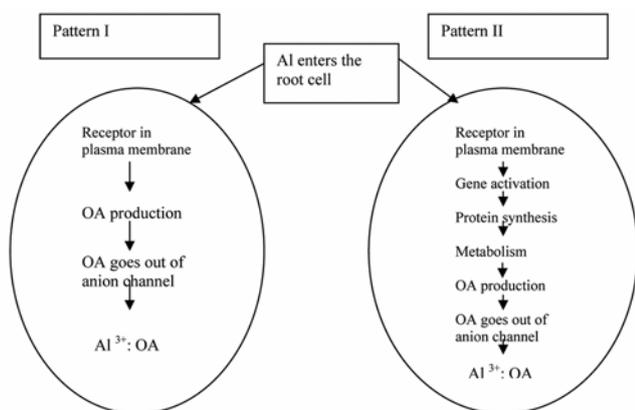


Figure 1. Two patterns of aluminium-induced secretion of organic acid (OA) anions in roots. Pattern I: There is no discernible delay between the addition of Al and the onset of secretion. Al activates the anion channel present in the plasma membrane resulting in specific release of organic acid anions to roots. Pattern II: Organic acid secretion is delayed for several hours after Al exposure. Induction of genes for release and/or biosynthesis of organic acid anions may be involved.

These organic acids have different capabilities for detoxifying Al in the soils. Citrate has the maximum ability followed by malate and oxalate to alleviate Al toxicity. Secretion of organic acids tends to be localized to the root apex and is associated with the level of Al resistance and the presence of genes conferring Al resistance^{19,26}. The organic acid exudation from different crops is summarized in Table 1.

Citrate

Citrate is produced in the mitochondria through the tricarboxylic acid (TCA) or Krebs cycle, and citrate carrier, an intrinsic protein of the inner mitochondrial membrane, plays a vital role in exporting citrate out of the mitochondria (Figure 2). Citrate is one of several organic acids exuded by plants, and correlation between organic acid exudation and Al tolerance was initially detected by Miyasaka *et al.*²⁷ in snapbean and found to play an important role in many other plants^{21,28}. The Al-tolerant

Table 1. Organic acid exudation from different plant roots

Organic acid produced	Crop plants
Citrate	Barley ¹² , carrot ⁶⁴ , citrus ⁶¹ , <i>Hydrangea</i> ⁴⁵ , maize ¹⁹ , pea ⁶⁶ , oat ⁶⁷ , pineapple ⁶⁸ , rape ⁶⁷ , radish ⁶⁷ , rice ¹⁷ , rice bean ¹⁶ , rye ¹⁵ , snapbean ²⁷ , soybean ⁶⁹ , tobacco ⁵⁶ , eucalyptus ⁶⁵ , triticale ²¹ .
Malate	<i>Arabidopsis</i> ⁴⁹ , eucalyptus ⁶⁵ , oat ⁶⁷ , pineapple ⁶⁸ , rape ⁶⁷ , radish ⁶⁷ , rye ¹⁵ , sunflower ⁷⁰ , triticale ²¹
Oxalate	Buckwheat ²⁹ , spinach ⁷¹

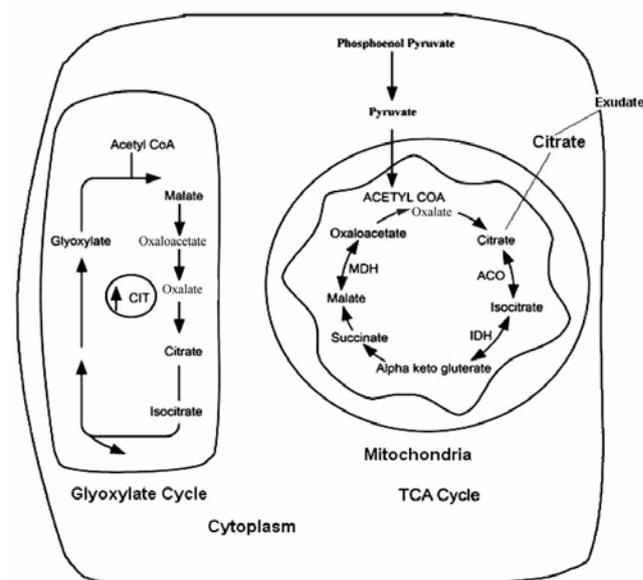


Figure 2. Metabolic map of tricarboxylic acid and glyoxylate cycles.

snapbean cultivar secreted 10-fold more citrate than the Al-sensitive cultivar. Similar observations of 3.5–7-fold citrate release were made from the roots of Al-tolerant than Al-sensitive maize with Al exposure¹⁹. Ma *et al.*¹⁸ observed that Al exposure resulted in 2.5–3.0 times more citrate secretion from roots of the Al-tolerant shrub *Cassia tora* than the Al-sensitive *Cassia occidentalis*. Ma *et al.*¹⁷ observed that citrate was detected in the presence of Al in secretion from *japonica* rice cultivar Koshihikari and *indica* rice cultivar Kasalath, while no citrate was detected in the absence of Al. The amount of citrate secreted was low ranging from 0.74 to 1.33 $\mu\text{mol g}^{-1}$ root day⁻¹ for Koshihikari and from 0.68 to 1.08 $\mu\text{mol g}^{-1}$ root day⁻¹ for Kasalath at various Al concentrations.

Oxalate

Organic acid is dominant in buckwheat leaves. Secretion of oxalate from the roots indicates that external Al-detoxification mechanisms contribute to the high Al resistance seen in buckwheat. One of the mechanisms responsible for this high resistance in buckwheat is proposed to be rapid and specific secretion of oxalate by the roots^{8,29}. The secretion of oxalate can prevent Al³⁺ from entering the roots. Oxalate was secreted in the region 0–10 mm from the root tip of buckwheat²⁹. It was observed that in buckwheat, the secretion of oxalate occurred within 30 min after the exposure to Al. Zhang *et al.*⁸ found that buckwheat had higher resistance to Al compared with an Al-tolerant cultivar of wheat, cv. Atlas 66.

Malate

Removal of aluminium from the roots results in a rapid decline in malate secretion to non-Al level, indicating responsive Al and malate-secreting mechanisms. The differences in the degree of tolerance to aluminium depend on transport of malate out of the apical root cells via Al-activated malate-permeable channel³⁰. Three possibilities have been proposed on the activation of the anion channel⁷: (i) Al interacts directly with the channel protein to trigger opening; (ii) Al interacts with a specific receptor on the membrane surface or with the membrane itself to initiate secondary messenger cascade, which then activates the channel; (iii) Al enters the cytoplasm and activates the channel directly or indirectly via secondary messengers.

The role of malate in Al³⁺ tolerance was first reported by Kitagawa *et al.*³¹ in wheat. Later, Christiansen-Weniger *et al.*³² found that an Al-tolerant wheat cultivar excreted more malate from its roots than a sensitive cultivar. However, the most convincing findings came from near-isogenic wheat lines⁹. It was found that Al³⁺ stimulated up to 10-fold greater efflux of malate from the roots of the Al-tolerant line than from the roots of the Al-

sensitive line. Malate was mostly exuded from the terminal 3 mm of the root, which is the part of the root most susceptible to Al toxicity². Salazar *et al.*³³ observed that Al-tolerant genotypes secreted about 10-fold higher malate and about 3–5 fold higher succinate than Al-sensitive seedlings over 24 h exposure to 50 μM Al. Malate efflux has been confirmed for other wheat cultivars differing in Al tolerance^{34–36}. Li *et al.*¹⁵ found that alteration in the metabolism of organic acid was involved in the Al-induced secretion of organic acids in rye but only activation of an anion channel seems to be responsible for the rapid secretion of malate in wheat.

Internal detoxification of Al by organic acids

In internal tolerance mechanisms, absorbed Al is detoxified by the organic acids in the cytosol. About 100 plant species accumulate Al in their stem and leaves without showing symptoms of Al toxicity³⁷. The formation of a non-toxic Al complex with organic acids or other chelators, and sequestering these complexes in the vacuoles play an important role in internal detoxification of Al in Al-accumulating plants³⁸. Foy³⁹ defines Al accumulator plants as those with more than 1000 mg kg⁻¹ of Al in the leaves. Al tolerance is associated with Al accumulation in plant shoot as seen in *Arnica montana*, *Deschampsia flexuosa* L.⁴⁰, *Melastoma malabathricum*⁴¹ and *Camellia sinensis*⁴². These species are well-known Al accumulators and collect a large amount of Al in the leaves. This suggests that Al accumulator plants detoxified the internal Al³⁺ by forming Al organic complexes. *M. malabathricum* L. is a woody plant that accumulates more than 10,000 mg kg⁻¹ Al, in its leaves in the form of monomeric Al and Al-oxalate complexes⁴¹. In tea (*Camellia sinensis* L.) plants, Al is taken up and stored in the central vacuole as complexes with organic acids⁴². High Al tolerance in buckwheat has been associated with internal⁴³ and external⁸ detoxification mechanisms by the formation of oxalate complexes non-phytotoxic with Al. Buckwheat accumulates (1500 mg kg⁻¹) high levels of Al in the leaves without showing any symptoms of toxicity¹¹ and most of this Al is complexed with oxalate²⁹. Most of the Al in both roots and leaves of buckwheat was complexed with oxalate in a 1:3 Al-oxalate complex²⁹. Later, it was observed that the Al being transported to the shoot in the xylem sap was complexed with citrate and not oxalate⁴³. These results suggest that Al undergoes a ligand exchange from oxalate to citrate when it is transported into the xylem, and is exchanged back with oxalate when in the leaves. Leaf compartmental analysis showed that 80% of Al in buckwheat leaves was stored in vacuoles as a 1:3 Al-oxalate complex⁴⁴. *Hydrangea macrophylla* can also accumulate more than 3000 mg kg⁻¹ Al dry weight in its leaves⁴⁵ and Al is complex with citrate at 1:1 ratio. These findings indicate that Al accumulating

species detoxify the internal Al by forming aluminium organic complexes.

Exogenous application of organic acids to reduce Al toxicity

Organic acids play an important role in both internal and external aluminium detoxification. Meriga *et al.*⁴⁶ observed that Al treatment reduced the root and shoot lengths of the control seedlings (without citrate) in rice-tolerant cultivar by 26% and 21% respectively, and those of sensitive cultivars by 51% and 23% respectively. However, corresponding seedlings grown in citrate-supplemented solution exhibited better root and shoot growth particularly at a citrate concentration of above 100 μM . At 200 μM citrate concentration, the root and shoot lengths of tolerant and sensitive cultivars improved by 60% over their respective controls. Li *et al.*⁴⁷ reported that exogenous organic acids such as citrate at 1:1 ratio with Al can completely detoxify Al, but the ratio of organic acid:Al that resulted in the same degree of Al detoxification for oxalate and malate was 2:1 and 8:1 respectively. These findings suggest that citrate is a better chelator of aluminium toxicity than oxalate and malate.

Molecular-assisted biotechnology to enhance secretion of organic acids

Several genes encoding organic acid production and chelation of aluminium in the rhizosphere are available. These discoveries will open up new avenues in the understanding of physiological and molecular mechanisms. The most commonly documented mechanisms of Al tolerance are the Al chelating ligands such as citrate and malate from the root tips and subsequent formation of non-toxic Al complexes in the apoplast or rhizosphere. The four cloned Al tolerance genes all encode organic anion transporters involved in this tolerance mechanism. The first two plant Al tolerance genes isolated were those encoding an Al-activated malate transporter in wheat and *Arabidopsis*^{48,49} and subsequently two Al-activated citrate transporters belonging to the multidrug and toxic compound extrusion (MATE) family of membrane transporters were encoded by Al tolerance gene in barley and sorghum^{50,51}. In wheat, the *TaALMT1* gene was located on the 4DL chromosome arm and was completely sequenced⁵². In barley, the expression of *ALMT1* significantly increased the flow of Al-activated malate, and Al tolerance⁵³. Ma *et al.*²¹ found that release of organic acid is linked to the genes on the short arm of chromosome 3R in triticale. Ma *et al.*⁵⁴ observed that tightly linked marker Bmag 353 explained 51.3% of phenotypic variance for citrate secretion in barley. Further, Furukawa *et al.*⁵⁰ identified a gene (*HvAACT1*) responsible for the Al-activated citrate secretion by fine-mapping combined with microarray analysis in barley.

Genetic engineering technology for the release and synthesis of organic acids

Identification and isolation of genes that specify Al tolerance are prerequisites for direct engineering of crop plants to increase crop production in acidic soils. The production of transgenic plants with an increased capacity to produce and/or excrete organic acids that chelate and detoxify Al in the rhizosphere, is an appealing strategy to produce Al-tolerant plants. Genetically engineered plants that overexpress genes involved in the biosynthesis and transport of organic acids and Al toxicity events at the cell level have been produced. Plant breeders can take advantage of genetic engineering, by which useful genes are made available in any species. Researchers have manipulated the biosynthesis capacity of cells which produce and accumulate higher amounts of organic acids. As a result, this will change the root exudation profile and Al resistance of a genotype. Table 2 summarizes the attempts to obtain transgenic plants with higher Al resistance by enhancing organic acid exudation.

Overproduction of citrate appears to be an important strategy to produce Al-tolerant transgenic plants. The production of citrate by the condensation of acetyl CoA and oxaloacetate is the first step in the TCA cycle and is catalysed by the enzyme citrate synthase (Figure 2). The most well-known example of successful achievement in this direction is the work of de la Fuente *et al.*⁵⁵. They introduced a *Pseudomonas auruginosa CSb* gene into tobacco and papaya. As a result, the transgenic plants showed enhanced citrate efflux and greater Al tolerance than non-transformant lines. The transformed lines of tobacco expressing *CSb* had up to 10-fold greater internal citrate in their root tissues, whereas in papaya citrate level in the roots was only 2–3 fold. Increased production of citrate was shown to result in Al tolerance in both the species. However, an attempt to repeat the work using the same transgenic lines as those of de la Fuente *et al.*⁵⁵ and tobacco transgenics expressing the *P. aeruginosa CS* gene to 100-fold greater levels, has shown neither increased citrate concentration in the roots nor increased citrate efflux. Thus no improvement of Al tolerance was achieved⁵⁶.

Tolerance to Al was also conferred on other crops, viz. rice⁵⁷ and carrot cells⁵⁸ by overexpression of the *CSb* gene, and on alfalfa by overexpression of the malate dehydrogenase gene⁵⁹. These promising findings strengthen the possibility of engineering plants with superior resistance to aluminium toxic acid soils.

The strategy of overexpressing enzymes involved in organic acid metabolism has proven to be effective to enhance organic acid exudation and to increase Al resistance with other transgenic plants. The overexpression of a mitochondrial *CSb* gene from *Arabidopsis* into carrot cells resulted in higher *CSb* activity and higher secretion of citrate compared with wild-type cells⁵⁸. Koyama *et al.*⁶⁰

Table 2. Aluminium tolerance in transgenic plants expressing genes involved in organic acid synthesis and release

Gene	Gene product	Source plant	Target	Aluminium-resistance
<i>CSb</i>	Citrate synthase	<i>Pseudomonas aeruginosa</i>	Tobacco, papaya ⁵⁵	Increases
<i>CSb</i>	Citrate synthase	<i>P. aeruginosa</i>	Tobacco, Alfalfa ⁵⁶	Does not change
<i>neMDH</i>	Malate dehydrogenase	Alfalfa	Alfalfa ⁵⁹	Increases
<i>PEPC</i>	Phosphoenolpyruvate carboxylase	Alfalfa	Alfalfa ⁵⁹	Does not change
<i>At-mtCS</i>	Citrate synthase	<i>Arabidopsis</i>	Oilseeds ⁷²	Does not change
<i>ALMT1</i>	Malate channel	Wheat	Barley ⁵³	Increases

introduced a mitochondrial *CSb* gene isolated from *Daucus carota* into *Arabidopsis thaliana* causing the same effect in transgenic plants. The transformants showed up to three fold increase in citrate synthase activity and 1.6-fold increase in citrate secretion compared with controls.

Deng *et al.*⁶¹ observed that exposure to Al triggered the exudation of citrate from the yuzu root (*Citrus junos* Sieb. ex Tanaka). Al also elicited an increase in citrate content and increased the expression level of mitochondrial citrate synthase (*CjCS*) gene and enzyme activity in yuzu. The *CjCS* gene was cloned from yuzu and overexpressed in *Nicotiana benthamiana* through *Agrobacterium tumefaciens*. The transgenic *Nicotiana* plants showed increased levels of citrate in roots compared to wild-type plants. The exudation of citrate from the roots of transgenic plants significantly increased when exposed to Al. Increased expression level of the *CjCS* gene and enhanced enzyme activity were observed in transgenic plants compared with wild-type plants. The results with transgenic plants suggest that overexpression of mitochondrial citrate synthase can be a useful tool to achieve Al tolerance.

In cultured tobacco, the functions of the BnALMT1 and BnALMT2 (*Brassica napus* Al-activated malate transporter) proteins were studied by heterologous expression. Such transfection system showed an enhanced capacity for malate efflux but not citrate efflux when exposed to Al. Transgenic tobacco cells grow significantly better than control cells. This indicated that expression of BnALMT1 and BnALMT2 increased the resistance of these plant cells to Al stress⁶². Thus use of citrate synthase gene may prove to be an effective strategy for the production of Al-tolerant crop species without undesirable effects on other agronomic traits.

Other approaches aim to increase organic acid exudation by overexpressing phosphoenolpyruvate carboxylase (PEPC) and malate dehydrogenase (MDH)⁶³. Overexpression of plant genes for such enzymes (PEPC, MDH) enhanced organic acid synthesis and secretion and greater Al tolerance in alfalfa⁵⁹. Selected transgenic plants with a 1.6-fold increase in MDH specific activity showed a 4.2-fold increase in citrate, oxalate, malate, succinate and acetate in root tissues compared to the control untransformed line. A plant line containing the *PEPC* transgene

with a two fold increase in PEPC activity had increased amounts of malate compared to the control. Plants expressing *MDH* or *PEPC* transgene showed enhanced root elongation compared with the control untransformed line in solution culture assay⁵⁹. In alfalfa, overexpression of MDH resulted in enhanced organic acid synthesis and secretion and greater Al resistance⁵⁹. Begum *et al.*⁵⁷ reported that PEPC transgenic rice was more tolerant to Al than the wild type, because root tips of transgenic rice accumulated less Al than those of the wild type. Al-induced oxalate exudation from the roots occurred at increased rates in the transgenic line. Overexpression of C4-PEPC drastically increased PEPC activity in the leaves of transgenic rice, and resulted in enhanced Al tolerance in transgenic rice causing higher organic acid concentration in the leaves and roots.

Conclusion

There is evidence that some plant species secrete organic acids to protect their root tips from Al³⁺ toxicity in acidic soil. Thus organic acids such citrate, oxalate and malate play an important role as a response to aluminium stress. Al-tolerant species secrete more organic acids than sensitive plants. However, research on organic acids has also resulted in many questions that remain unanswered. For example, why do species release different organic acids as a response to Al stress? How much organic acid secretion is sufficient to detoxify Al? How does Al exposure activate or induce the secretion of specific organic acids out of root cells? Are there any other defence mechanisms for Al resistance? We are just releasing internal detoxification of Al with organic acids and its sequestration as Al-organic complexes in the vacuole in tolerating Al toxicity. It is also likely that other additional resistance mechanisms exist.

The cloning of genes whose products increase the synthesis of organic acids and their transport across the plasma membrane of roots are needed to enhance conventional breeding approaches for improving Al tolerance. Several genes are expressed in Al-tolerant genotypes of wheat, soybean and *Arabidopsis*. These genes are good candidates to be studied in other crops to find mechanisms of Al tolerance in plants. Citrate synthase gene has

been identified, and there are many yet to be discovered before targeted genetic modifications can be effectively designed. The molecular aspects underlying Al resistance mechanisms via Al-induced expression of some other enzymes catalysing organic acid synthesis and metabolisms need to be studied.

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