GLOBAL KNOWLEDGE CENTER ON CROP BIOTECHNOLOGY

Biotechnology with Salinity for Coping in **Problem Soils**



Salinity and Agriculture

and salinity. Land degradation due to increased it has also resulted in large-scale water lodging increases in tood production of the last 40 years, and has been partly responsible for the large agriculture to semi-and and and areas of land, irrigation has made it possible to extend which accumulate in the soil over time. While water of poor quality leaves behind sait solutes practices, since the evaporation of irrigated a problem since the beginning of cultivation High salinity in agricultural fields has been

water retention properties, and can eventually render fields unsuitable for agriculture. presence of salt also has a very negative effect on the soil structure, affecting porosity and of other stresses, diseases and pollutants; and can be lethal to the plant. The excessive (1). Most crops are very sensitive to salt, which severely attects yield; increases the severity into account and areas or deserts, which comprise a quarter of the total land of the planet salinity presently affects about 20 percent of world's area under irrigation, without taking

soils are all needed. cultivation, and the use of mixed cropping systems to mitigate the accumulation of salt in Changes in farming practices, such as the selection of suitable species and varieties for essential to reverse the degradation of the environment and to ensure sustained productivity. A more rational and sustainable use of natural resources- land and water- is therefore

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APPLICATIONS

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Salt flat in Utah. United States





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and salinity is therefore an important strategy to this end (2). development of crop varieties with increased tolerance to abiotic stresses such as drought water scarcity and high salt concentrations represent important constraints to yield. The of tood production in both irrigated and rain-ted lands, including those areas where Meeting current and future tood demands necessitates nevertheless a short term increase

Salt Stress



to the production of toxic reactive oxygen species. ot cell membranes, inhibit photosynthesis, and lead structure and permeability

> intormation on drought tolerance, 3). Generally, varieties developed responses to drought and to salinity (see pocket K 30 for more to plants, and hence there is a substantial overlap between plant Salt stress effectively decreases the availability of water in the soil

protein synthesis, affect the enzyme tunction, inhibit tatal. Salt ions impair cells is toxic, and potentially accumulation of salt ions in problem to plants: excess to affecting the water balance of the plant, salt poses another will also be more resilient to salt stress (4; 5). However, in addition to be more tolerant to drought and that use water more efficiently,

Development of Salt-Tolerant Crops by Conventional Breeding

The existence of plants that thrive in soils with high level of salts (termed halophytes), and the occurrence of variation between crop cultivars in salt sensitivity, indicate that salt tolerance is to a large extent under genetic control. Halophytes represent only about 2 percent of plant species, however, they can be found among half of the terrestrial plant families and are very variable and diverse. Although the development of tolerance to salt is believed to have occurred independently several times during the evolution of land plants, halophytes seem to have evolved the same basic method for dealing with salinity: storing harmful salt ions in the cell vacuole and accumulating organic solutes (which act as osmoprotectants) in the cell cytoplasm (*6*).



Exposure to high salinity is killing Yecoro Rojo (right), a wheat cultivar that has moderate salt tolerance. Plants resulting from hybrid crosses with W4910 (left) show much greater tolerance of high salinity. Photo by Richard Wang, USDA.



Conventional breeding requires the identification of genetic variability to salinity among different varieties or cultivars of a crop, or in sexually compatible species, and breeding this tolerance into lines with suitable agronomic characteristics. Conventional breeding programs for salinity tolerance include the development of rice, wheat and Indian mustard varieties tolerant to salt and to alkali soils by the Central Soil Salinity Research Institute in Karnal, India (7) and efforts to incorporate salt tolerance to wheat from wild related species (8). A number of

Plants growing in a salt marsh, Canada

genomic tools, such as molecular markers and gene profiling methods, can greatly improve the efficiency of breeding programs, and should be fully exploited for conventional breeding initiatives.

Engineering Salt-Tolerant Crops by Genetic Modification



Although not a crop plant, *Arabidopsis* has played a vital role in the elucidation of the basic processes underlying stress tolerance, and the knowledge obtained has been transferred to a certain degree to important food plants (4). Several features make *Arabidopsis* an ideal model organism: a small fully-sequenced genome, a small size, and a short life cycle. In addition, a wealth of genomic resources is available for Arabidopsis, and insights gained can be used to improve our understanding of the same processes in crop plants, which are less amenable to genetic studies. Many of the genes known to be involved in stress tolerance have been isolated initially in *Arabidopsis*.

Tissue-cultured *Arabidopsis thaliana*. Photo by Keith Weller, USDA.

Mutant analysis- the screening for mutations that affect the plant's response to stress- has been a crucial tool in the discovery of genes acting in the network. Screens designed include those

aimed to identify mutations with increased or decreased sensitivity to drought, salinity and cold stresses. Also important has been the use of DNA microarray technology, which allows monitoring changes in gene expression in response to stress, and to identify genes that are either induced or repressed by the treatment (*5*).

The development of salt-tolerant crops by genetic engineering have focused on the following strategies: increasing the plant's ability to limit the uptake of salt ions from the soil; increasing the active extrusion rate of salt ions; and improving the compartmentalization of salt ions in the cell vacuole where they do not affect cellular functions. Genes encoding osmoprotectants have also been the targets of genetic modification experiments, but although their over-expression in some cases improves salt tolerance, in general they also affect plant growth in the absence of stress with negative effects in yield, a highly undesirable trait for farmers (2, 9).



Salt intake is controlled by low and high affinity ion transporters: trans-membrane proteins that move ions across the cell membrane, which are also required for

the intake of potassium ions (K+). The efflux of ions from the plant depends on the activity of the *SOS1* gene (for *Salt Overly Sensitive1*), initially characterized in Arabidopsis but recently identified in rice, and shown to be functionally conserved between dicots and monocots (*10*). Vacuolar membrane transporters, including the one encoded by *AtNHX1* gene of *Arabidopsis*, play a role in the sequestration of ions into the vacuole. NHX1 proteins are also conserved across species, and have been isolated from several crops. Over-expression of *NHX1* genes in *Arabidopsis*, rice, canola and tomato have been reported to increase the tolerance to salt stress (*3*).

Tomato plants

Conclusion

Salt tolerance is a very complex trait, both at the physiological and at the genetic levels, and is also very influenced by other environmental factors acting on the plant at the same time. In addition, the genetic control to salt stress differs in different stages of the plant's life cycle: tolerance at the adult stage does not necessarily correlate with tolerance at the seedling and juvenile stages, or to the ability to germinate in the presence of salts (*2*, *9*). Rice, for example, is much more affected in grain filling than in vegetative growth by the presence of salt in the soil. To complicate matters further, it is very difficult to design field trials to test the agronomic performance of improved salt-tolerant varieties, as the salt concentration in soils is very variable, and is complicated by the presence of additional pollutants and inland water intrusion.





Plant genomes need to be very plastic, a feature required to cope with a variable environment that requires a constant adjustment of the plant's metabolism. It is therefore essential to test newly developed stresstolerant varieties to multiple stresses in laboratory conditions, and the importance of carrying out extensive field studies in a large range of conditions that assess tolerance as absolute yield increases cannot be over-emphasized (*11*).



Wheat