

Welcome

MBB 604: Advances in crop biotechnology (3+0)

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Transgenic approaches for improvement of plant abiotic stress tolerance

Environmental stress physiology

- **Drought stress**
- **Salt stress**
- **High temperature**
- **Cold stress**

Introduction

- A variety of abiotic stresses causing crop loss of about >50%
- Due to the declining water availability, by 2025, 30% crop production will be at risk
- World Bank projects that the climate change will reduce crop yield by 20% or more by the year 2050
- So for that reason transgenic plant have to be developed

APPROACHES

1. Engineering of genes for **Osmolyte biosynthesis**
2. Engineering of genes encoding enzymes for **scavenging active oxygen**
3. Engineering of genes encoding **LEA Proteins**
4. Engineering of genes encoding heterologous enzymes with different **temperatures**
5. Genetic engineering of **molecular chaperones**
6. Engineering of genes encoding **transcription factors**
7. Genetic engineering of **cell membranes**

1. Engineering of Genes for Osmolyte biosynthesis

- Tolerance to abiotic stresses has mainly been achieved at cellular levels of osmotically-active solutes
 - Proline, glycine betaine
 - Sugars such as sucrose, trehalose and fructans
 - Sugar alcohols like sorbitol, mannitol, ononitol, pinitol and polyols
- These osmolytes are uniformly neutral with respect to cellular functions

Accumulation of these molecules helps plants to –

- Retain water within cells and protects cellular compartments from injury caused by dehydration
- Maintains turgor pressure during water stress
- Stabilize the structure and function of certain macromolecules
- Signalling functions or induction of adaptive pathways
- Scavenge reactive oxygen species

Proline

- Proline accumulates in many organisms in response to drought and salinity
- Proline is encoded by a nuclear gene ***Pyrroline-5-carboxylate synthetase (P5CS)***
- Proline may serve as a **hydroxyl radical scavenger** reducing the acidity of the cell
- It may also function as an **osmolyte** and **molecular chaperone**
- Protect protein integrity and enhance the activities of different enzymes

proline...

- Glutamate can be achieved by using the *P5CS* gene from Moth bean (*Vigna aconitifolia*)
- Antisense proline dehydrogenase (*PDH*) in transgenic *Arabidopsis*
- However, *P5CS*, limiting enzyme in proline biosynthesis, feedback inhibition by proline
- Removed this feedback inhibition by site-directed mutagenesis and the resulting gene ***P5CSF129A***

Transgenic plants for Proline

Gene	Molecular function	Source	Transformed plant	Performance of transgenic plant	Additional note
P5CS	<i>D1-pyrroline-5-carboxylate synthase</i>	<i>V. aconitifolia</i>	<i>O. sativa</i>	Drought & salt tolerance	Increased biomass (higher fresh shoot & root weight)
P5CS F129 A	<i>Mutated P5CS</i>	<i>V. Aconitifolia</i>	<i>N. tabacum</i>	Oxidative & osmotic stress tolerance	Removal of feedback inhibition resulted in 2-fold high proline than the plants expressing P5CS
PDH	<i>Proline dehydrogenase</i>	<i>A. thaliana</i>	<i>G. max</i>	Drought & heat tolerance	Rapid increase in proline resulted in least water loss under drought stress

Transgenic plants for Proline

Gene	Molecular function	Source	Transformed plant	Performance of transgenic plant	Additional note
<i>P5CSF129A</i>	<i>Mutated P5CS</i>	<i>V. aconitifolia</i>	<i>O. sativa</i>	Salt stress tolerance	Transgenic plants with high proline level showed higher biomass & growth performance under salinity stress
<i>P5CSF129A</i>	<i>Mutated P5CS</i>	<i>V. aconitifolia</i>	<i>P. vulgaris</i>	Drought tolerance	Elevated free proline resulted in better adaptation to water stress
<i>δ-OAT</i>	<i>Ornithine-diaminotransferase</i>	<i>A. thaliana</i>	<i>N. Plumbaginifolia</i>	Osmotic stress tolerance	Transgenic plant showed higher germination and increased biomass

Glycine Betaine

- GB accumulates in the **chloroplasts** and **plastids** and increases the tolerance of plants to various abiotic stresses (drought, salinity and freezing)
- The physiological role of GB in **alleviating osmotic stress**
- It can also **protects proteins** and **enzyme** activities under water deficits and **stabilize membranes** during freezing

- It can help stabilize the protein tertiary structure and prevent or reverse disruption of the tertiary structure
- Accumulation of GB is limited due to **choline** supply
- Transgenic potato plants expressing a bacterial choline oxidase (*betA*) gene leads to high levels of GB under drought stress

Transgenic plants for Glycine betaine

Gene	Molecular function	Source	crop	Function	Additional note
BADH	<i>Betaine aldehyde dehydrogenase</i>	<i>A. Hortens is</i>	<i>O. sativa</i>	Improved growth & salinity tolerance	10% transgenic plants set seed in 0.5% NaCl solution in greenhouse
BADH	<i>Betaine aldehyde dehydrogenase</i>	<i>H. Vulgare</i>	<i>N. tabacum</i>	Salt tolerance	Tolerance was judged by high biomass & photosynthetic
CHO	<i>Choline dehydrogenase</i>	<i>E. coli</i>	<i>O. sativa</i>	Drought & salt tolerance	Higher yield under stresses
betA	<i>Choline dehydrogenase</i>	<i>E. coli</i>	<i>Z. mays</i>	Drought tolerance at seedling stage	Grain yield was significantly higher compared with wild-type

Transgenic plants for Glycine Betaine

Gene	Molecular function	Source	Crop transformed	Performance of transgenic plant	Additional note
<i>betA</i>	<i>Choline dehydrogenase</i>	<i>E. coli</i>	<i>G. hirsutum</i>	Drought tolerance	GB accumulation was positively correlated with drought stress
<i>codA</i>	<i>Choline oxidase</i>	<i>E. coli</i>	<i>S. tuberosum</i>	Oxidative, Drought & salt Stress tolerance	Stress-inducible GB production in non accumulators resulted in tolerance
<i>CMO</i>	<i>Choline monooxygenase</i>	<i>A. hortensis</i>	<i>G. hirsutum</i>	Salt tolerance	Seed cotton yield was significantly higher in transgenic plants

Sugars and Sugar Alcohols

- Accumulation of sugar-related compounds, response to osmotic stress
- These compounds **stabilize** the **membranes** and **proteins** during dehydration
- Sugars can replace the water molecules and stabilize the proteins or membranes in a similar of water molecules
- They can form a glass phase in the dry state of high viscosity have capable of slowing down chemical reactions lead to long-term stability in a living system

Trehalose

- Trehalose, a rare **non-reducing sugar**
- Trehalose protects the biological molecules in response to different stress conditions
- It does not accumulate to high enough levels in most plants, probably because of the presence of trehalase activity

trehalose...

- Trehalose synthesized in two steps from **glucose-6-phosphate** and **uridine diphosphoglucose**, *via* trehalose-6-phosphate
- The first step is catalyzed by **trehalose phosphate synthase (TPS)**, and the second by **trehalose-6-phosphatase (TPP)**

Transgenic plants for Trehalose

Gene	Molecular function	Source	Transformed plant	Function	Additional note
<i>TPS1</i>	Trehalose-6-phosphate synthase	<i>S. cerevisiae</i>	<i>N. tabacum</i>	Enhanced drought tolerance	A few transgenic plants showed phenotypic alteration along with stress tolerance
<i>otsA & otsB</i>	<i>TPS & TPP</i> fusion gene	<i>E. coli</i>	<i>O. sativa</i>	Drought, salt & cold tolerance	Sustained growth, less photo-oxidative damage, enhanced photosynthesis under various stresses
<i>TPP1</i>	Trehalose-6-phosphate phosphatase	<i>O. sativa</i>	<i>O. sativa</i>	Enhanced salt & cold stress Tolerance	Analysis of transgenic plants suggested a possible role of TPP in transcriptional regulation pathways

Transgenic plants for Trehalose

Gene	Molecular function	Source	Transformed plant	Function	Additional note
TPS1	<i>Trehalose-6-phosphate synthase</i>	<i>S. cerevisiae</i>	<i>L. esculentum</i>	Drought, salt & oxidative stress tolerance	Sustained yield under all stress
TP	<i>Trehalose phosphorylase</i>	<i>P. sajor-caju</i>	<i>S. officinarum</i>	Drought tolerance	Transgenic plants did not withered even after 10 days of water deficit & also exhibited no pleiotropic effects
TPS1+ TPS2	<i>Bifunctional gene</i>	<i>S. cerevisiae</i>	<i>M. sativa</i>	Multiple abiotic stress tolerance	Improved growth with significant increase in biomass

Fructans and other Carbohydrates

- Fructan or polyfructose molecules serve as the main **storage carbohydrate** in many plant species
- However, fructans are to be involved in abiotic stress tolerance by virtue of their presence in vacuoles
- Fructans are also thought to protect plants against drought and cold stress evidence of this was derived from transgenic plants

- It act as regulators or signal molecules, thus influencing plant metabolism, as scavengers of ROS
- It promote the process of root branching, thus increasing the root surface and subsequent water uptake

Fructans and other Carbohydrates

Gene	Molecular function	Source	Transformed plant	Function
sacB	Levan sucrose, a fructosyl transferase	<i>B. subtilis</i>	<i>N. tabacum</i> , <i>B. vulgaris</i>	PEG-induced drought tolerance
lmt1	Myo-inositol-o methyl transferase	<i>E. coli</i>	<i>N. tabacum</i>	Drought & salt tolerance
mtID	Mannitol-1-phosphate dehydrogenase	<i>E. coli</i>	<i>S. melongena</i>	Salt, drought & chilling tolerance
lpk2 β	Inositol polyphosphate 6-/3-kinase	<i>A. thaliana</i>	<i>N. tabacum</i>	Enhanced drought, salinity & freezing tolerance

2. Engineering of genes encoding enzymes for scavenging active oxygen

- Plants suffering from various stresses, which leads to overproduces active oxygen species in cells
- To minimize the damaging effects of active oxygen, plants produces antioxidant to detoxifying harmful oxygen
- Genes encoding enzymes with antioxidant capacity, such as ascorbate peroxidase, superoxide dismutase and glutathione reductase

Example:

- Transgenic alfalfa overproducing **superoxide dismutase (SOD)** showed reduced injury from water deficit and freezing stresses in field conditions
- Engineering chloroplastic **superoxide dismutase (SOD)** in tobacco led to an increase of chilling tolerance of photosynthesis
- Tobacco plants that overproduced cytosolic **ascorbate peroxidase** showed increased tolerance to oxidative stress
- Over expression of **glutathione S-transferase** or **glutathione peroxidase** provided some protection against cold and salt stress in tobacco

3. Engineering of genes encoding LEA Proteins

- A genes encoding LEA (**Leaf Embryo Abundance**) proteins is activated under osmotic stress
- LEA proteins play a role in desiccation tolerance during seed development and in response to dehydration, salinity and cold stress
 - **Maintenance of protein or membrane structure**
 - **Sequestration of ions,**
 - **Binding of water**
 - **Operating as molecular chaperones**

Example:

- *HVA1*, a group 3 LEA protein from barley
- *LE25*, a group 4 LEA protein from tomato
- Rice plants transformed with barley LEA gene *HVA1* possessed increased tolerance to water deficit and salinity
- Improved salinity and freezing resistance has been observed in yeast transformed with *LE25*

4. Engineering of genes encoding heterologous enzymes with different temperature optima

- Homologous proteins and their specific enzymatic activities from different plant species differ in terms of their temperature
 - E.g: When a gene of **NADH-hydroxypyruvate reductase** from cucumber (higher optimum growth temperature) was transformed into tobacco (lower optimum growth temperature), the optimum temperature of transgenic tobacco was increased

5. Genetic Engineering of molecular Chaperones

‘Chaperone’ : assist and maintain correct folding and trafficking of cell proteins, are crucial for plants to survive in abiotic stresses

- E.g: An antisense inhibition of the *HSP70* family prevented Arabidopsis plants from acquiring thermotolerance
- Heat shock proteins *HSP101* from soybean, corn, wheat and tobacco complemented a thermo tolerance defect in yeast caused by deletion of yeast *HSP104*

6. Engineering of genes encoding Transcription Factors

- Responses of plants to abiotic stresses are multigenic and a single gene is not likely to induce the whole cascade
- Recently, transcription factors for abiotic stress-induced genes have been identified, cloned and used in transgenic experiments
- Simultaneous expression of downstream stress-inducible genes have been achieved with parallel increase in stress tolerance

Example:

- Over expression of the transcriptional activator ***CBF1*** ('C-repeat binding factor') induced the expression of four **COR** ('cold-regulated') genes and increased freezing tolerance of *Arabidopsis* plant
- By over expressing a single stress-inducible transcription factor (***DREB1A***), lead to plant tolerance to freezing, salinity and dehydration

7. Genetic Engineering of Cell Membranes

- Membranes are critical sites of injury by chilling, freezing, heat and osmotic stresses
- Increased levels of unsaturated membrane lipids leads to higher membrane fluidity which improves plant-chilling tolerance and photosynthetic parameters

Example:

- Tobacco transformed with acyl-ACP:glycerol-3-phosphate acyltransferase (*GPAT*) from chilling-tolerant species of *Arabidopsis* acquired chilling tolerance

Thank you !!!