

**In the Condition of Salt Stress, Differential Expression of Akvaporin Genes in
Mesembryanthemum crystallinum L. Plant**

Elfira AGAYEVA¹, Nargiz BAYRAMOVA²

Anatomy, physiology and zoology /GanjaStateUniversity, Azerbaijan

Anatomy, physiology and zoology / GanjaStateUniversity, Azerbaijan

shanur@rambler.ru

Aim of the study: The purpose of our study to determinate differential expression of genes in crystal flower plant and it consists of learning role of this process to adaptation time to chloride salinity of plants. To explore Natrium chlorid impact of basic physiological parameters which characterize the water status of plants. Assess the role of akvaporin in the adaptation of plants to high concentrations of natrium chloride. To explore the differential expression of six genes of akvaporin in the bodies of *M. crystallinum* plants.

Materials and methods: Crystal flower (*M. crystallinum* L.) was used as an object of study. Plants were grown around the 23-25°C in the afternoon and 18-20°C in the night in the fitotron cell. *M. crystallinum* 4-10 weakly plants have been used in the experiment.

The practise is calculated from the beginning. For this purpose, 200 and 400 mM concentration was used. The length of each experiment, 3, 9, 24, 72, 168 hours. 3-4 leaves from the bottom layer was fixed at every point of the experiment. Determination of transpiration intensity of leaves was taken to generally accepted gravitational method Ivanova. For determine free proline as mainly Bates et al., (1973) method the ninhydrin reactive was used. The RNA separation of crystal flower leaves and roots have been carried out by RNeasy Mini Kit. Polymerase chain reaction was carried out. For getting microsomal membranes, appropriate authorities of plant 300 mM sucrose, 100 mM Tris - HCl (pH 8.0), 10 mM EDTA, 5mM potassium metabisulfite, 5mM dithioeritol, 1mM phenylmethylsulfonyl fluoride and were homogenized in 0.6% polyvinylpyrrolidone environment. It was homogenized at 1000D filtration. 10 min centrifuge with it. Determination of protein Bradford (1976) was conducted in accordance with the procedure. Electrophoretic splitting of proteins Laemmli (1970) basis was taken 12.5% PPA buffer mini-Protein cell 3Cell.

Results: The expression of six genes of akvaporin was studied by us: MIPA, MIPB, MIPC, MIPH, MIPF, MIPK. The analysis of amino acid sequences showed that, 4 of them plasma membrane (MIP A, B, C, H), and 2 - tonoplast (MIP F, K) has been localized. The study of *Mesembryanthemum crystallinum* plant, throughout the day in response to the intensity and length of the effect NaCl in various organs McPIP1; 1; McPIP2; 1; McPIP2; 3; McTIP1 2 and McTIP2 2 allows to determine expression of akvaporin genes. 4 gene was Plasma membrane aquaporins McPIP1 ruled out. Its constitutive expression is not depend on the body's location, what time of the day, on the nature of the impact of stressor. In the stress conditions of changing the main physiological parameters of water status of plants. Associated with drought the water deficit irrigation of leaf intensity transpiration, SNH, is accompanied by a decrease in osmotic potential. Akvaporin genes of plasma membrane decreasing of transcellular water transmission in succulent bodies. At the same time, it is reflected less in the intensity distribution of water intracellular, so the evidence the stressor tonoplast akvaporin genes would show the weak impact to the expression.

Keywords: differential expression, akvaporin, expression of genes, salt shock