Abstract

Potassium is essential macroelement and large amount of potassium is taken up by plants,

because it's the major osmotic of plant cell. Due to various potassium availability in different

kinds of soil, plants have evolved transport system that can maintain sufficient K⁺ uptake

between several orders of magnitude of potassium concentration. In Arabidopsis thaliana

mechanism of K⁺ acquisition is well understood and it's known that AtHAK5 is mainly

involved in K⁺ uptake in its very low concentrations. AtHAK5 belongs to KT/HAK/KUP family

that consist 13 members in A. thaliana. There's known function for some members for example

in auxin transport in root tip (TRH1) or in K⁺ efflux in stomatal closure (KUP6).

In this thesis possible involvement of yet unstudied transporters KUP5, KUP7, KUP9 and

KUP10 in K⁺ acquisition and homeostasis in A. thaliana was investigated. In vitro cultivation

showed that kup9 mutant has very short lateral roots in K⁺ deficiency. On the other hand kup5

mutant showed significantly higher dry weight in K⁺ deficiency than Col-0. Kup9 phenotype

was successfully replicated in subsequent cultivations and anatomy of lateral root apical

meristems was investigated. Root tips of kup9 were differently organized in K⁺ deficiency and

they showed signs of early termination of meristems. To localize the expression of KUP9 plants

of A. thaliana were transformed with pKUP9::GUS construct, although it wasn't possible to

evaluate the results yet.

When the plants were exposed to draught all lines showed decrease in relative water content

of shoot moreover kup9 had lower dry weight and fresh weight. These results may suggest that

all transporters have function in K⁺ homeostasis, although these results has to be checked

deeper.

When cultivated with ¹³⁴Cs radioisotope *kup7* mutants showed reduced Cs⁺ accumulation in

roots, this may suggest that KUP7 is involved in K⁺ uptake. In addition *hak5* mutant was studied

and it also accumulated less Cs⁺ in roots. If this result could be replicated it will be the final

evidence of relevant involvement of HAK5 in cesium uptake from environment.

Key words: Cesium, KT/HAK/KUP, lateral roots, potassium, uptake