CHARACTERIZATION OF A NITRATE HIGH-AFFINITY TRANSPORT SYSTEM IN BRACHYPODIUM DISTACHYON

Laure David¹, Mathilde Grégoire¹, Patrick Berquin¹, Elodie Phommabouth¹, Anne Krapp¹, Françoise Daniel-Vedele¹, Thomas Girin¹ and <u>Sylvie Ferrario-Méry¹</u>

¹Institut Jean-Pierre Bourgin, INRA, AgroParisTech, CNRS, Université Paris-Saclay, RD10, 78026 Versailles Cedex, France

sylvie.ferrario-mery@inrae.fr

Nitrogen is an essential element for growth and plant development. Improving the nitrogen use efficiency of crops is a priority for the future in order to limit the fertilizer use and environmental impacts. An efficient nitrate uptake system contributes to improve the nitrogen use efficiency of crops in conditions of low N availability. The High-Affinity nitrate Transport System (HATS) in plants is active in low range of external nitrate and is mediated by a two-component system (the high affinity transporters NRT2 associated to a partner protein NRT3).

In Brachypodium, the model plant for C3 cereals, seven BdNRT2 and two BdNRT3 have been identified (Girin et al. 2014). Expression of the BdNRT2 and BdNRT3 genes in response to nitrate availability fits with the characteristics of the HATS components already described in other species (David, Girin et al. 2019). Likewise, co-expression of BdNRT2A and BdNRT3.2 is required for an effective nitrate transport in the heterologous expression system Xenopus oocytes. Functional interaction between BdNRT2A-GFP and BdNRT3.2-RFP has been observed at the plasma membrane in Arabidopsis protoplasts transiently expressing both constructs, while a cytosolic localization was observed when BdNRT2A-GFP or BdNRT3.2-RFP were transiently expressed separately, suggesting а monocotyledonous/dicotyledonous specificity for the functional interaction. Site directed mutagenesis on a Ser residue of the C-terminal extension of BdNRT2A specifically conserved in monocotyledonous plants and mimicking a constitutive phosphorylation status of this Ser was used to study the species specificity of the NRT2/NRT3 interaction. Finally, ¹⁵NO3 influx measurements with Brachypodium mutants, defective in *BdNRT2A* expression, confirmed that BdNRT2A is a major contributor of the HATS in Brachypodium.

Girin T, David LC, Chardin C, Sibout R, Krapp A, Ferrario-Méry S, Daniel-Vedele F. 2014 Brachypodium: a promising hub between model species and cereals. J Exp Bot. 65(19):5683-96.

David LC, Girin T, Fleurisson E, Phommabouth E, Mahfoudhi A, Citerne S, Berquin P, Daniel-Vedele F, Krapp A, Ferrario-Méry S. 2019 Developmental and physiological responses of Brachypodium distachyon to fluctuating nitrogen availability. Sci Rep. 9(1):3824.