

## **HD-ZIP II TRANSCRIPTION FACTOR GENES CONTROL ADAXIAL-ABAXIAL PATTERNING IN *ARABIDOPSIS* LEAF MORPHOGENESIS**

TURCHI L.\* , CARABELLI M.\* , SASSI M.\* , POSSENTI M.\*\* , RUZZA V.\* , MELATTI C.\* , MORELLI G.\*\* , RUBERTI I.\*

\*) Institute of Molecular Biology and Pathology, National Research Council (Italy)

\*\*\*) National Research Institute for Food and Nutrition (Italy)

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The *Arabidopsis* genome encodes for 10 Homeodomain-Leucine Zipper II (HD-Zip II) proteins. It has been previously shown that *ATHB2*, *HAT1*, *HAT2* (HD-Zip II  $\gamma$  subfamily), *HAT3* and *ATHB4* ( $\delta$  subfamily) are induced by changes in the Red/Far Red ratio of the light environment. However, these genes are also tightly regulated during plant development with both distinct and overlapping patterns (Ciarbelli et al., *Plant Mol. Biol.* 2008, 68: 465-78). In order to understand the role of the light-regulated HD-Zip II genes in plant development, we have analyzed single and multiple mutants within  $\gamma$  and  $\delta$  subfamilies. Here we present the phenotype of mutants in the *HAT3* and *ATHB4* genes. Young seedlings show a gradual loss of cotyledon and leaf expansion, up to completely radialized organs. The pattern of vascular development is also profoundly altered, in a manner that is tightly linked to lamina expansion. Fully radialized leaves lack procambial cells whereas trumpet shaped leaves show hyperproliferation of phloem with respect to xylem, a feature that is found in the vasculature of abaxialized leaves. *In situ* and GUS/GFP reporter analyses of the  $\delta$  subfamily genes show that they are expressed in the adaxial side of cotyledons and leaves. Taken together, these data demonstrate that *HAT3* and *ATHB4* are required to specify adaxial identity in leaf morphogenesis. We are currently analyzing the molecular and genetic relationships between the d HD-Zip II genes and members of the HD-Zip III family genes, such as *PHB*, *PHV* and *REV*, key determinants of adaxial leaf identity.