The interplay between *Agrobacterium* and the host plant – From the perspective of the host

Rosalia DEEKEN

Julius-von-Sachs Institute of biosciences, Botany 1, Department of Molecular Plant Physiology and Biophysics, University of Wuerzburg, Germany

deeken@botanik.uni-wuerzburg.de

Agrobacterium species deliver native as well as constructed T-DNAs into the genome of various plant species. Plant cells harbouring a native T-DNA undergo substantial reprogramming and give rise to crown gall disease. Constructed T-DNAs are used as a biological tool to manipulate the plant genome for studying gene functions and improving plant performance. In both cases, the interacting partners, the Agrobacterium and the plant cell, need to communicate and respond to signals emanating from each other. Our focus is the response of the host plant to Agrobacterium in Agrobacterium-mediated tumorigenesis. We analyse the changes in the methylome, transcriptome and the metabolome as well as solute content to assess the morphological and physiological changes that occur during Agrobacterium infection and crown gall development.

At this meeting, I will present our findings that pathogen defence signalling is largely dampened in the first hours after infection before the T-DNA is integrated. I will also show that after T-DNA integration and during crown gall development, the host plant activates biotic and abiotic stress responses, thereby controlling the degree of transformation and crown gall growth.