

## Plant transformation in Vienna

# Intruder Alert

*Agrobacterium* transforms plants, that's a well-known fact. But how exactly does the transforming T-DNA get into the plant's nucleus? Heribert Hirt and his team now know: *Agrobacterium* abuses the host's own defence response.

A plant-focused laboratory barely copes without it; every plant geneticist admires its services and those learning about this tool for the first time generally are fairly impressed. *Agrobacterium tumefaciens*, also referred to as a natural trans-kingdom genetic engineer, has made the transformation of thousands and thousands of plants possible. In its free time, the soil-borne bacterium lives as a plant pathogen that causes tumorous growth and demotes its host to a kind of bed and breakfast. In the lab it functions as a handy gene ferry, introducing scientifically relevant pieces of DNA into the plant genome. In both cases *Agrobacterium* fools the plant cell the ancient Greeks' way – an exciting and important strategy of the *Agrobacterium*-mediated transformation which was recently published by Heribert Hirt's group from the University of Vienna, Austria, and the URGV Plant Genomics Laboratory, Evry, France (*Science*, 318, 2007: 453-6).

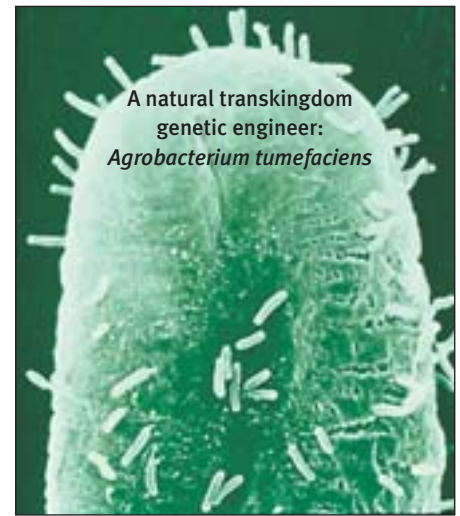
"Botany used to be a thorn in my side. All that dissection and drawing seemed awfully boring to me," laughs Hirt at his former attitude towards plants. Hirt was born in Persia and grew up in Germany. He first studied biochemistry at the University of Cape Town in South Africa, followed by a PhD. His interest was in the medical topics of neurobiology and human genetics; later he switched to yeast as a model sys-

tem for cell cycle control. "All I knew about plants was that they have RNA, DNA and proteins. Still, we decided to look for parallels in their regulation of the cell cycle compared to yeast and other eukaryotes", he says, remembering the early stages of his plant research.

## Not boring at all

In the early nineties, Hirt and his team identified several homologues of regulatory components of the eukaryotic cell cycle in alfalfa (*Medicago sativa*). They found cdc2 protein kinases, phosphoprotein phosphatases, cyclins and mitogen-activated protein kinases (MAPKs). Closer investigation revealed that it was the latter that built a bridge to the regulation of stress response and tolerance in plants. MAPK-pathways are known to transfer information from sensors to responses in all eukaryotes. Now they have been shown to be involved in development, cell proliferation, and hormone physiology, as well as in biotic and abiotic stress signalling. "We looked for MAPKs as regulators of the cell cycle and we found many of them, but they were all involved in the plant stress response," Hirt explains. "This is how we finally got on that track."

Thus were green and silent plants revealed to be less boring than they appeared. As sessile organisms, plants cannot move away from adverse environmental condi-



tions. Instead, they have to show a good sense of detection of and adaptation to perturbations. "With *Arabidopsis* we have a fascinating, complex organism and a perfect genetic model system. Yeast is a great system too but it is not that impressive in its appearance," Hirt laughs. How do plants react to too much salt and heat, to too little water and light, to the attacks of pests and pathogens? How do they communicate with the latter? Have they experienced a co-evolution with their enemies? If so, what has been the outcome?

These are the questions Hirt and his group ask every day – and which finally lead them to *Agrobacterium*. Hirt: "During our search for targets of the *Arabidopsis thaliana* MAP kinase MPK3 we repeatedly isolated the VirE2 interacting protein (VIP1). This protein was only known to be involved in *Agrobacterium*-mediated transformation; its function in plant cells had not been resolved yet."

## Abuse of a transcription factor

However, let's remember the course of a typical transformation first: *Agrobacterium* introduces a partial copy of its tumour-inducing Ti plasmid into the host cell. Bacterial virulence (Vir) proteins, together with several plant proteins, stabilize this so-called transfer DNA (T-DNA) and guide it to the plant cell nucleus. The T-DNA is inserted into the plant genome where it can fulfil *Agrobacterium*'s ultimate goal of hijacking the plant's metabolic system. The result is the synthesis of T-DNA encoded opines and plant-type hormones. While the former constitute an ideal carbon and nitrogen source for the bacteria, the latter lead to a tumour-like proliferation of the plant cells – altogether creating a spacious and cosy habitat for *Agrobacterium*.

"The process of transporting the T-DNA from the plasma membrane to the nucleus is all but trivial", Hirt stresses. "It is comparable to an asylum-seeker who tries to take on the hurdles of bureaucracy." The protein VIP1 had already been known to facilitate T-DNA transport by acting as an adaptor between T-DNA and the plant's nuclear import



