


TOPIC: Transcriptomic Fingerprint Technology in Food Safety

Hanspeter NAEGELI

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BIOGRAPHY

Toxicology is the study of the adverse effects of chemicals and biological substances on living organisms. The Naegeli group is active in different areas ranging from molecular toxicology (DNA repair, endocrine disruptors), to food toxicology (detection of phytoestrogens, mycotoxins and pesticides), and to clinical toxicology. They are using advanced methods of molecular biology combined with live-cell-imaging and DNA microarrays to study typical reaction patterns of human cells in response to toxic insults.

Education and academic positions

- 1985 Graduation from School of Veterinary Medicine, Zürich
- 1988-1989 Participant at the “Postgraduate Course in Experimental Medicine and Biology” at the University of Zürich
- 1989 Doctoral degree, Institute of Pharmacology and Biochemistry, University of Zürich
- 1990 Postdoctoral fellow, Department of Pathology, Stanford University Medical School, Stanford CA, USA
- 1991-1992 Postdoctoral fellow, Department of Pathology, The University of Texas Southwestern Medical Center at Dallas, Dallas TX, USA
- 1993 “Oberassistent”, Institute of Veterinary Pharmacology and Toxicology, University of Zürich
- 1995 Participant at the Postgraduate Training “Course on Veterinary Toxicology”, Wuppertal, Germany
- 1996 Promotion to “Abteilungsleiter”, Head Toxicology Division
- 1998 Degree of “Privatdozent” (Habilitation) for Pharmacology and Toxicology
- since 2000 Delegate of the Vetsuisse-Faculty at the Swiss Academy of Medical Sciences
- 2002 Promotion to Professor *ad personam* at the University of Zürich
- 2006 Nomination to Professor of Toxicology at the University of Zürich (“Lehrstuhl Toxikologie”)

ABSTRACT

Transcriptomic fingerprinting technology in food safety

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The goal of transcriptomic fingerprinting is to improve food safety by using cultured human cells as versatile biological detectors (“cytosensors”) of toxic contaminants. This new strategy is prompted by the finding that living cells respond to toxic chemicals by changing the pattern of genes that are converted into messenger RNA transcripts. Each individual RNA transcript carries the information for the synthesis of a particular protein product. The term “transcriptome” refers to the entire spectrum of such messenger RNA intermediates in a given biological system. Accordingly, “transcriptomics” stands for large-scale analytical methods that can be used to monitor complex RNA profiles consisting of thousands of transcripts. The general scheme of contaminant detection by transcriptomic fingerprinting is as follows. Cultured human cells are exposed to extracts prepared from food samples (meat, milk, cereals, etc.). Following an incubation time of 3-24 hours, messenger RNA transcripts are isolated, labelled and detected on high-throughput DNA microchips. Different contaminants generate characteristic changes in the transcriptome pattern, thus giving rise to diagnostic RNA fingerprints that can be used to recognise and quantify hazardous constituents. In the context of the BioCop project, we have adapted a miniaturised DNA microarray platform to determine transcriptome fingerprints induced by estrogenic endocrine disruptors as well as type A trichothecenes. With exception of a portable reader, this transcriptomic platform requires no specialised equipment and, hence, can be easily disseminated. A key advantage is that this novel test exploits health-relevant parameters in a toxicologically significant target system. With the widespread use of rapid screening tests, which are not related to any toxicological endpoint, effect-driven in vitro bioassays will become of increasing importance to support risk assessment and monitor the success of risk management actions.

Keywords: Mycotoxin, Trichothecene, Phytoestrogen, Endocrine Disruptor

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