

Program
qPCR Satellite Symposium

at the BioCity Leipzig, Deutscher Platz 5, 04103 Leipzig, Germany

Symposium Homepage: <http://leipzig05.gene-quantification.info/>

organized by Michael W. Pfaffl and powered by

BioScience Events

An international Symposium open to everybody who is interested in qPCR, from 10-11th March 2005.
 The main focus of the qPCR symposium is related to the key problems in quantitative real-time PCR (qPCR):

Optimization, Normalization & Standardization

Leading European academic researchers and **international industrial contributors from the R&D** will participate in the symposium and present **23 innovative and interesting talks**. The symposium will be an arena for fruitful discussions between researchers of different backgrounds and is divided in **6 sessions** offering an overview of the present knowledge and future developments in qPCR technology and its new applications:

Session 1: Pre-analytical steps	Moderator: M. W. Pfaffl
Session 2: Single cell PCR	Moderator: M. Kubista
Session 3: Probes Technologies	Moderator: S. Bustin
Session 4: Quantification strategies & Optimization	Moderator: S. Bustin
Session 5: Applications	Moderator: N. Zoric
Session 6: Post PCR & Bioinformatics	Moderator: M. W. Pfaffl

An **Industrial Exhibition** covering 11 booth from 18 companies and a small **Poster Session** will be held in the Foyer of the BioCity Leipzig. Plan of the industrial exhibition: <http://www.bioscience-events.com/leipzig/exhibition-zoom.gif>

qPCR Symposium Sponsors:



Agilent Technologies



BioTechniques

Hotels close to the BioCity Leipzig

In the hotels around the BioCity Leipzig we have special discounts. Please do your own reservations with the PDF registration forms. Booking with this form and applying the quoted special rates will only be possible by **18th February 2005**. The prices apply per night and room and include breakfast and statutory VAT.

Thursday 10th March 2005

8:00 – 9:00 Arrival & Registration

9:00 – 9:30 **Welcome by the Organizers & Opening of the Satellite-Symposium**
Michael W. Pfaffl & Katharina Spanel-Borowski**9:30 – 12:45 Session 1: Pre-analytical steps Moderator: M. W. Pfaffl**9:30 – 10:15 **Neven Zoric**
Coordinator of the TATAA Biocenter, Göteborg, Sweden
What to do and not to do in reverse transcription !10:15 – 10:45 **Thomas Kaiser**
Corbett Research R&D, Sydney, Australia
DNA Preparation on the X-tractor-Gene and pre-PCR setup on the CAS 1200.

10:45 – 11:15 Coffee break

11:15 – 11:45 **Debra Ann Nickson**
European Senior Product Manager, Genomics, Invitrogen Europe, Paisley, Scotland
Improving cDNA yields using SuperScript III.11:45 – 12:15 **Marc Valer**
Agilent Technologies, Waldbronn, Germany
Chip based analysis of nucleic acids combined with RNA Integrity Number (RIN) algorithm enables fast, independent measurement of RNA quality.12:15 – 12:45 **MarcValer**
Agilent Technologies, Waldbronn, Germany
Practical demonstration of the Bioanalyzer 2100.**12:45 – 13:45 Lunch and Poster Session: ABI, Artus Biotech, Exiqon, Premier Biosoft, Qiagen****13:45 – 15:30 Session 2: Single cell PCR Moderator: M. Kubista**13:45 – 14:30 **Stephen A. Bustin**
Institute of Cell & Mol. Science, Queen Mary's School of Medicine & Dentistry, University of London
Laser capture micro dissection for biologically meaningful quantification of mRNA.14:30 – 15:00 **Anders Stahlberg**
Department of Chemistry and Biosciences, Chalmers University of Technology, Sweden
Gene Expression Profiling - From Tissue to Single cell.15:00 – 15:30 **Tanja Nolan**
Sigma-Aldrich Chemie GmbH, Munich, Germany
Accurate gene expression profiling from limiting material - facing the issues of normalization, efficiency and mRNA to protein correlation.

15:30 – 16:15 Coffee break

16:15 – 18:15 Session 3: Probes Technologies Moderator: S. Bustin16:15 – 16:45 **Olfert Landt**
TIB MOLBIOL Syntheselabor GmbH, Berlin, Germany
Clamped-Probe-Assay: Detection and quantification of minor/minimal variants.16:45 – 17:15 **Ralf Mauritz**
Roche Applied Science, Diagnostics, Penzberg, Germany
ProbeLibrary: One Transcriptome - One Kit.17:15 – 17:45 **Andres Jarrin**
Eppendorf, Germany
Improved Real Time PCR Application using Novel Eppendorf Amplification Technologies.17:45 – 18:15 **Uta Möllers**
Application Specialist, Applied Biosystems
TaqMan Assays -not only Primers and Probes.20:00 – 23:30 Banquet in the **Auerbachs Keller** in the city center of Leipzig
www.auerbachs-keller-leipzig.de

Friday 11th March 2005**9:00 – 12:15 Session 4: Quantification strategies & Optimization Moderator: S. Bustin**

9:00 – 9:45 **Michael W. Pfaffl**
Center of Life Science, Weihenstephan, Technical University of Munich, Germany
Quantification strategies in real-time RT-PCR.

9:45 – 10:15 **Mary Span**
EUROGENTEC, Liege science park, Seraing, Belgium
Pitfalls in qPCR.

10:15 – 10:45 **Thomas Kaiser**
Corbett Research R&D, Sydney, Australia
qPCR normalization and optimization in the Rotor-Gene 3000.

10:45 – 11:15 **Coffee break**

11:15 – 11:45 **Jim Huggett**
Centre for Infectious Diseases & International Health, University College London, UK
Real time RT-PCR normalisation: methods and considerations.

11:45 – 12:15 **Reinhold Müller**
Stratagene R&D, La Jolla, USA
Considerations for standardizing qPCR assays.

12:15 – 13:15 Lunch and Poster Session: ABI, Artus Biotech, Exiqon, Premier Biosoft, Qiagen**13:15 – 14:45 Session 5: Applications Moderator: N. Zoric**

13:15 – 13:45 **Tomasz Czechowski**
Molecular Plant Nutrition Group, MPI for Molecular Plantphysiology Golm, Germany
Real time RT-PCR profiling of over 1,400 Arabidopsis transcription factors: unprecedented sensitivity reveals novel tissue-specific genes.

13:45 – 14:15 **Hilary Sreere**
Bio-Rad Laboratories Europe, Munich, Germany
The Ups and Downs of Gene Expression: Characterization of New Tools Used for Gene Expression and Knockdown.

14:15 – 14:45 **Linda Vigilant**
Max Planck Institute for Evolutionary Anthropology, Leipzig, Germany
Applications of qPCR to genetic analyses of wild primates.

14:45 – 15:15 **Coffee break**

15:15 – 17:15 Session 6: Post PCR & Bioinformatics Moderator: M. W. Pfaffl

15:15 – 16:00 **Mikael Kubista**
Department of Chemistry and Biosciences, MultiD Analyses AB, Göteborg, Sweden
Gene expression analysis.

16:00 – 16:30 **Jo Vandesompele**
Center for Medical Genetics Ghent, Ghent University Hospital
Gene expression profiling: Accurate normalization and automated data-analysis.

16:30 – 17:00 **Yevhen Vainshtein**
EMBL Heidelberg, Meyerhofstrasse, Heidelberg, Germany
REST-384: an improved way of simple and precise qRT-PCR data evaluation.

17:00 – 17:15 **Closing of the Symposium**
Michael W. Pfaffl & Katharina Spänzel-Borowski

Plan of the Bio City Leipzig lecture hall and industrial exhibition:

