Yearbook 2007/08

MSc/PhD Molecular Biology Program
at the University of Göttingen

International Max Planck Research School

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Letter from the University

The international Master's / PhD Programs Molecular Biology and Neurosciences were established by the Georg August University Göttingen, together with the Max Planck Society for the Advancement of Science, in the year 2000 to attract excellent students from all over the world and provide them with an

outstanding, research-oriented graduate program. Both programs are taught in English by internationally renowned scientists and offer a high level of services and individual support.

Several hundred students from all over the world apply for the 20 study places available in each of the programs every year. Both programs have introduced and combined elements of international recruitment, competitive admission procedures, advanced curricula, research training, social integration programs, extracurricular support and evaluation procedures into successful working structures. They have achieved excellent recommendations in several external evaluations and have been awarded the 2004 prize for excellent support services for foreign students by the German Federal Foreign Office. For the newly established Georg August University School of Science (GAUSS) and two other graduate schools in Göttingen, the Molecular Biology and Neuroscience Programs are considered exemplary and serve as best practice models.

In October 2006, the two programs were awarded the label "Top 10 International Master's Degree Courses made in Germany" by "Stifterverband für die Deutsche Wissenschaft" and the German Academic Exchange Service (DAAD) in a national contest, in which 121 Master's programs of 77 universities participated. The Göttingen Molecular Biology and Neuroscience programs were the only Master's programs in the natural sciences and medicine which received this award. Both programs are members of the Göttingen Graduate School for Neurosciences and Molecular Biosciences (GGNB), which was successful in the recent Excellence Competition by the German Federal and State Government to promote science and research at German universities.

Five Göttingen University faculties, three Göttingen Max Planck Institutes as well as the German Primate Center participate in the programs. International guest lecturers are also involved. The Max Planck Society contributes through its newly established International Max Planck Research Schools. Both programs keep close contacts with the relevant industries to further enhance the chances of the graduates for a successful professional career.

I would very much like to thank all scientific bodies and institutions for their committed support in establishing these international programs and, last but not least, the German Academic Exchange Service (DAAD), the Lower Saxony Ministry of Science and Culture and the various generous donors.

The Georg August University of Göttingen is proud of its long-standing international experience the two attractive and innovative programs have already become an integral part of. The university will continue to support these programs within the setting of Göttingen's lively urban, cultural and social life, in itself a prerequisite for creative teaching and research.

Prof. Dr. Kurt von Figura
(President of the Georg August University Göttingen)

Letter from the Max Planck Society

The mission of the Max Planck Society is to conduct basic research in science and humanities at the highest level. More than 80 Max Planck Institutes are located on scientific campuses across Germany, most of them close to universities.

Scientific ties between Max Planck Institutes and universities are traditionally strong. In 1998, during the 50th year celebration of the Max Planck Society in Göttingen, the Max Planck Society - together with the Hochschulrektorenkonferenz - launched the International Max Planck Research Schools as a new joint program to further intensify cooperation.

The goals of the International Max Planck Research Schools are

- to attract excellent students from all around the world to intensive Ph.D. training programs in Germany, preparing them for careers in science,
- to integrate Max Planck scientists in top-level scientific training of junior scientists,
- to intensify the ties to the Universities owing to the participation of internationally renowned Max Planck scientists in joint teaching activities, and
- to strengthen international relationships by providing individual support to each student and by exposing foreign students to German culture and the German language.

By now, 49 International Max Planck Research Schools have been established involving 63 Max Planck Institutes, 47 German universities with participating 73 faculties and more than 15 universities abroad. More than 1900 (mostly PhD-) students from 87 countries are presently enrolled. Approximately 800 PhD students have graduated to date from an International Max Planck Research School.

Since resuming operation in the year 2000, the Göttingen International Max-Planck Research Schools in Molecular Biology and Neurosciences have met with extraordinary success. Every year, the programs receive hundreds of applications, with the quality of the students consistently being very high. Most students graduated so far have moved on to postdoctoral positions, many at prestigious international institutions. In the past years, the Göttingen Schools received unanimous acclaim during external evaluations and won national awards. For instance they are the only Life Science Programs within Germany that were selected for the “Top Ten International Master’s Degree Courses”. The Schools have also re-shaped the local scientific community, strengthening the ties between the participating institutions, and initiated new scientific collaborations that augment the international reputation of Göttingen as a center for scientific excellence. Furthermore, the Schools served as role models and founding members of the Göttingen Graduate School for Neurosciences and Molecular Biosciences, thus were being instrumental for the success of the University in the German Excellence Initiative. We hope that in the years to come the students of the International Max-Planck Research Schools will be successful in their professional careers. We also hope that they will remember their training period in Göttingen as an exciting and stimulating phase in their lives.

Peter Gruss
President
Max Planck Society

Reinhard Jahn
Dean of the IMPRS
Molecular Biology

Overview

This yearbook is intended to provide information on the International MSc/PhD Molecular Biology Program in Göttingen, Germany, which was established in 2000. In addition to general information on the program, the yearbook introduces the current year's students, the faculty members, the program committee and the coordination team.

The program is member of the recently founded Göttingen Graduate School for Neurosciences and Molecular Biosciences (GGNB), which is funded by the Excellence Initiative of the German Federal and State Governments. It is offered by the Göttingen Center for Molecular Biosciences (GZMB), a newly established scientific center of excellence at the University of Göttingen, the Max Planck Institute for Biophysical Chemistry, the Max Planck Institute for Experimental Medicine, and the German Primate Center. Further to their active participation in the Molecular Biology Program and the research activities of the GZMB, the abovementioned partners closely cooperate in several research alliances, collaborative research centers and interdisciplinary doctoral programs. An example for cooperation with research institutes abroad are joint activities and student exchange with the Feinberg Graduate School at the Weizmann Institute of Science in Rehovot, Israel.

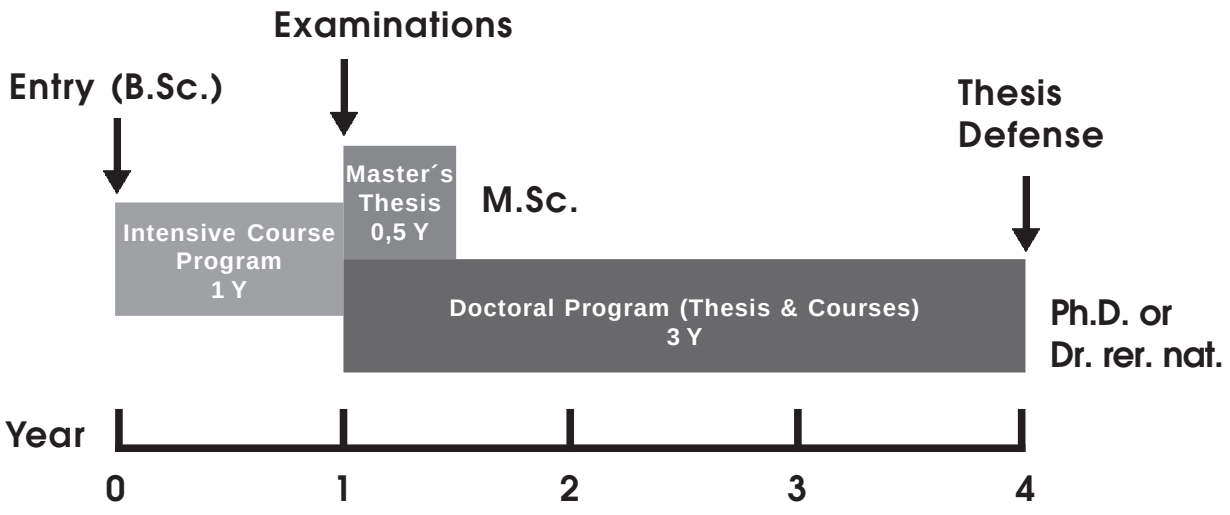
The intensive, research-oriented curriculum of the International MSc/PhD Molecular Biology Program qualifies students for professional work in the fields of molecular and cellular biosciences. The program is open to students from Germany and from abroad, who hold a Bachelor's degree (or equivalent) in the biosciences, chemistry, medicine, or related fields. Scholarships are available. All courses are held in English. The academic year starts in October and is preceded by three week orientation program. Applications may be submitted until January 15 of the year of enrollment. To ensure a high standard of individual training, the number of participants is limited to 20 students per year.

All students initially participate in one year of intensive course work. This first segment of the program comprises lectures, tutorials, seminars, methods courses, and individually supervised research projects (laboratory rotations). The traditional German structure of academic semesters is not followed. The condensed schedule allows students to accumulate 90 credits (ECTS) within one year, which would normally require three semesters.

Subsequently, two separate segments are offered:

- **PhD Program:** Good to excellent results after the first year qualify for direct admission to a three-year doctoral project in one of the participating research groups. The Master's thesis requirement is waived in this case. After successful defense of a doctoral thesis, the degree *Doctor of Philosophy* (Ph.D.) or the equivalent title *Doctor rerum naturalium* (Dr. rer. nat.) is conferred.

- **MSc Program:** Alternatively, students may conclude the program with a Master's thesis, based on six months of experimental scientific research. The degree Master of Science (MSc) is awarded upon successful completion of the Master's thesis.



Funding of the Program

The Molecular Biology Program thanks the following institutions and funding initiatives, who contributed to the success of the Molecular Biology Program:

DAAD

German Academic Exchange Service (DAAD),
Bonn, Germany, http://www.daad.de

International Degree Programs -
Auslandsorientierte Studiengänge (AS)

IPP made in Germany

International Postgraduate Programs –
Internationale Promotionsprogramme (IPP)

Max Planck Society for the Advancement of Science,
Munich, Germany, http://www.mpg.de

International Max Planck Research Schools

Niedersächsisches Ministerium für Wissenschaft und Kultur

Ministry of Lower Saxony for Science and Culture,
Hannover, Germany, http://www.mwk.niedersachsen.de/home/

Innovationsoffensive

Doctoral Programs - Promotionsprogramme

Stifterverband für die Deutsche Wissenschaft

Stifterverband für die Deutsche Wissenschaft,
Essen, Germany, http://www.stifterverband.org

Donors

The Molecular Biology Program thanks the following companies for their donations, which were used to financially support students during the first year of studies:

Bayer AG, Leverkusen, Germany

Carl Zeiss Lichtmikroskopie, Göttingen, Germany

Degussa AG, Düsseldorf, Germany

DeveloGen AG, Göttingen, Germany

Heka Elektronik GmbH, Lambrecht / Pfalz, Germany

Hellma GmbH & Co. KG, Müllheim / Baden, Germany

KWS Saat AG, Einbeck, Germany

Leica Microsystems GmbH, Bensheim, Germany

Luigs & Neumann, Ratingen, Germany

Olympus Deutschland GmbH, Hamburg, Germany

Roche Diagnostics GmbH, Penzberg, Germany

Sartorius AG, Göttingen, Germany

Solvay Pharmaceuticals, Hannover, Germany

Springer Verlag, Heidelberg, Germany

Vossius & Partner, München, Germany

Intensive Course Program (First Year)

Throughout the first year, current topics in molecular biology are covered by

- lectures
- tutorials
- methods courses
- laboratory rotations
- seminars

Lectures and Tutorials

A comprehensive lecture series is offered in a sequence of 8-11 week units. The following topics are taught on an advanced level throughout the first year (36 weeks, 4 hours per week):

A. Biochemistry and Structural Biology

- The Prokaryotic and Eukaryotic Cell
- Enzyme Mechanisms and Regulation
- Energy Metabolism, Lipid Metabolism
- Metabolic Networks
- Protein Modification and Degradation
- Protein Structure and Folding, NMR, Crystallography

B. Molecular Genetics

- DNA and Chromatin Structure
- DNA Replication and Repair
- Transcription
- RNA-processing and Translation
- Signal Transduction
- Genomics, Bioinformatics

C. Functional Organization of the Cell / Neurobiology / Immunology

- Nucleocytoplasmic Transport
- Protein Sorting and Processing
- Vesicular Transport, Organelle Biogenesis
- Cytoskeleton
- Cell Adhesion
- Nervous Systems, Sensory Systems
- Immunology
- Infectious Diseases, Principles of Pathogenicity
- Cell Cycle, Apoptosis, Cancer

D. Model Systems of Molecular Biology/Biotechnology

- Prokaryotes
- Biotechnology
- Fungi
- *Arabidopsis*
- *Drosophila*, *C. elegans*
- *Xenopus*, Zebrafish
- Chicken, Mouse
- Human Genetics

Each lecture is accompanied by a tutorial session, where students meet with a tutor in small groups. Tutorials involve exercises, review of lecture material, and discussion of related topics.

Methods Courses

During the first months of the Molecular Biology Program, students participate in a series of methods courses to introduce them to principles and practical aspects of basic scientific techniques and the handling of model organisms. The methods comprise 18 two-day experiments in small groups.

A. Proteins

- Protein preparation and characterization by gel electrophoresis and Western blot
- Chromatographic protein separation
- NMR spectroscopy
- Structural analysis of proteins and protein structure validation
- Proteomics
- Microarrays
- Analysis of protein-protein and nucleic acid-protein interaction

B. Nucleic Acids

- Purification and electrophoresis of nucleic acids
- Polymerase chain reaction I
- cDNA-synthesis, cloning
- DNA sequence analysis and bioinformatics
- Chemical and enzymatic analysis of RNA structure
- Spectroscopic characterization of nucleic acids

C. Cell Biology and Genetics

- Light microscopy
- Electron microscopy
- Biochemical cell fractionation
- Cell culture
- Expression analysis

Laboratory Rotations

Starting in January, every student conducts three independent research projects (laboratory rotations) in the participating departments. Each project is individually supervised. These involve seven weeks of experimental work, followed by one week for data analysis and presentation. For each project, a report must be completed in the format of a scientific publication. The laboratory rotations must cover three different subjects.

Seminars

Seminars start in March. The class meets weekly for two hours to discuss two student presentations. The presentations are research reports based on work from the laboratory rotations.

Examinations

After the first year of intensive training, all students take one written and two oral Master's examinations. The Master's examinations explore the students' theoretical background in topics covered by lectures and tutorials. Each oral examination investigates the qualification in two of the following disciplines:

- biochemistry
- structural biology
- genetics
- microbiology
- cell biology
- immunology
- developmental biology

PhD Program

Students who have passed the Master's examinations with good or excellent results qualify for direct admission to a three-year doctoral project in one of the participating research groups without being required to complete a Master's thesis first.

The PhD program emphasizes independent research on the part of the students. Doctoral students select three faculty members as their doctoral committee which closely monitors progress and advises students in their research project. Laboratory work is accompanied by seminars and lecture series, a wide variety of advanced methods courses, training in scientific writing and oral presentation skills, courses in intercultural communication, bioethics and research ethics, elective courses, and participation in international conferences or workshops.

Doctoral students of the program organize the international PhD student symposium "Horizons in Molecular Biology" every year with great success, outstanding speakers and, by now, more than 300 participants from all over the world. The meeting was designed by the students to promote scientific exchange between young researchers from different disciplines.

At the end of the PhD training program, a doctoral thesis is submitted either in the traditional format, or as a collection of scientific publications in internationally recognized journals along with a general introduction and a discussion of the results. The degree PhD or, alternatively, Dr. rer. nat. will be awarded after the successful defense of the doctoral thesis.

Master's Program

After the first year of intensive training, students may conclude the program with a six-month thesis project, leading to a Master of Science degree. The thesis project involves experimental work under the supervision of faculty member of the Molecular Biology Program. Students have the opportunity to conduct their Master's thesis project at a research institution abroad.

Orientation, Language Courses, Social Activities

A three-week orientation prior to the program provides assistance and advice for managing day-to-day life in Germany, including arrangements for bank account, health insurance, residence permit, housing, and enrolment. Students have the opportunity to meet faculty members and visit laboratories of the participating institutions. In addition, the orientation program informs students about computing and library facilities, the city and university of Göttingen, sports facilities, and cultural events.

Prior to the start of lectures and courses, basic knowledge in mathematics, chemistry and physics is refreshed in a one-week crash course, the so-called "Week Zero".

An intensive basic language course in German is offered in cooperation with *Lektorat Deutsch als Fremdsprache* to facilitate the first weeks in Göttingen. Additional language courses and social activities accompany the program.

Application, Selection and Admission 2007

Applicants must hold a Bachelor's degree or equivalent in biology, biochemistry, chemistry, medicine, or related fields. Applicants who are not native speakers of English should demonstrate adequate competence of the English language by acceptable results of an internationally recognized test.

In the year 2007, the Molecular Biology program received 412 applications from 56 countries.

Continent	Applications	Admissions
Europe (total)	**82**	**15**
Germany	26	7
other West Europe	10	0
East Europe	46	8
America (total)	**18**	**0**
North America	1	0
Central/South America	17	0
Africa (total)	**32**	**0**
North Africa	12	0
Central/South Africa	20	0
Asia (total)	**280**	**5**
Near East	45	3
Central Asia/ Far East	235	2

Students 2007/2008

Name		Home Country
Shweta	Aggarwal	India
Alwaleed	Alkhaja	Bahrain
Adi	Bar-Shalom	Israel
Janina	Berghoff	Germany
Neva	Caliskan	Turkey
Annette	Denker	Germany
Aliaksandr	Dzementsei	Belarus
Liudmila	Filonava	Belarus
Mandy	Hannemann	Germany
Sebastian	Hogl	Germany
Chandini	Kadian	India
Frederik	Köpper	Germany
Andrius	Kraskauskas	Lithuania
Nadya	Mitova	Bulgaria
Sina	Mozaffari Jovin	Iran
Miroslav	Nikolov	Bulgaria
Doris	Petroi	Romania
Volkan	Sakin	Turkey
Broder	Schmidt	Germany
Cornelius	Schneider	Germany

Shweta Aggarwal

First Name
Shweta

Last Name
Aggarwal

Date of Birth
10 March 1984

Country
India

EDUCATION

College / University
2005 - 2007: School of Biotechnology, Jawaharlal Nehru University, New Delhi, India
2002 - 2005: Daulat Ram College, University of Delhi, New Delhi, India

Highest Degree
M.Sc.

Major Subjects
Biochemistry, Biotechnology

Lab Experience
Recombinant DNA technology (DNA isolation, PCR, cloning, transformation, constructing gene knock outs in bacteria), DNA/protein gel electrophoresis, Western blotting

Projects / Research
2006 - 2007 Bioprocess and genetic strategies to generate quiescent cells for high level recombinant protein production

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society
2007 Qualified CSIR- Junior Research Fellowship in the field of life sciences
2007 Qualified GATE (Graduate Aptitude Test in Engineering) in the field of life sciences with a percentile of 99.66

SCIENTIFIC INTERESTS AND GOALS

The arena of cellular signaling and the regulation of gene expression fascinate me. It is interesting to know that simple small molecules participate in diverse functions ranging from cell survival to cell death and thus integrate the external signals with the internal signal transduction pathway that ultimately controls the gene expression profile. Also intriguing is the cross talk between different signaling pathways. Through this program I would like to gain experience in theoretical and experimental application to develop a better understanding of the field of biology.

Alwaleed Alkhaja

First Name
Alwaleed

Last Name
Alkhaja

Date of Birth
3 February 1982

Country
Bahrain

EDUCATION

College / University
2003 - 2007: University of Leeds, United Kingdom

Highest Degree
B.Sc. (with industrial placement)

Major Subjects
Biochemistry and Molecular Biology

Lab Experience
Various techniques in biochemistry and molecular biology

Projects / Research
09/2006 - 12/2007 The characterization of an ACE2-like enzyme in HEK cells, School of Biochemistry, University of Leeds, United Kingdom
08/2005 - 08/2006 Retinoic acid resistance in embryonic carcinoma cells, Department of Pharmacology and Toxicology, Dartmouth College, U.S.A.

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society
2004 - 2007 University of Leeds: International Student Scholarship

SCIENTIFIC INTERESTS AND GOALS

Currently, I am interested in the application of molecular biology in medical research and therapeutics. I am interested in epigenetics and post-translational modifications and how they relate to developmental aberrations and disease. With that said, I am excited about expanding my horizons by learning new techniques and discovering new interests and passions.

Adi Bar-Shalom

First Name
Adi

Last Name
Bar-Shalom

Date of Birth
2 August 1983

Country
Israel

EDUCATION

College / University
2004 - 2007 Ben-Gurion University of the Negev, Israel

Highest Degree
B.Sc.

Major Subjects
Biochemistry, Molecular Biology

Lab Experience
Southern-blot, DNA purification and agarose gel electrophoresis, basic techniques in microbiology and molecular biology, several bioinformatic tools

Projects / Research
10/2006 - 06/2007: A glance into a scorpion's venom gland: Identification of 73 novel toxins-coding cDNA from the venom of the scorpion *Buthus occitanus israelis*

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

I find many cellular and molecular processes fascinating, such as signal transduction, the molecular basis of diseases, proliferation and apoptosis and RNA functions. I feel quite astonished by the depths of the evolving biosciences, and undecided about where to focus my further interests. Many of nature's riddles are still unsolved and so much is still waiting to be unveiled. Through the Molecular Biology program I wish to deepen and broaden my theoretical knowledge and practical skills in the diverse fields of molecular biology and to explore areas of biosciences which are new to me such as developmental and stem cell biology. During the program I hope to gain an insight into different promising research areas, while meeting colleagues from different scientific and cultural backgrounds.

Janina Berghoff

First Name
Janina

Last Name
Berghoff

Date of Birth
7 December 1983

Country
Germany

EDUCATION

College / University
2003 - 2007 Hochschule Mannheim, University of Applied Sciences, Germany

Highest Degree
B.Sc.

Major Subjects
Biotechnology

Projects / Research
Chaim Sheba Medical Center, Institute of Hematology, Laboratory for Molecular Cytogenetics, Tel Hashomer, Israel, with Prof. Nagler: “Definition of a DNA Index DI of Plasma Cells (PCs) by the Use of Image Processing on 3D Images of FISH Preparations in Cases of Multiple Myeloma”
Hochschule Mannheim, University of Applied Sciences, Institute of Molecular Biology and Cell Culture Technology with Prof. Dr. M. Hafner: Evaluation of a human, putative autonomously replicating sequence

Publications
Koren-Michowitz M, Hardan I, Berghoff J, Yshoev G, Amariglio N, Rechavi G, Nagler A, Trakhtenbrot L (2007) Chromosome 13q deletion and IgH abnormalities may be both masked by near-tetraploidy in a high proportion of multiple myeloma patients: A combined morphology and I-FISH analysis. Cancer Letters, 255(2): 307-314

Scholarships
2003 - present: Studienstiftung des Deutschen Volkes (German National Academic Foundation)

SCIENTIFIC INTERESTS AND GOALS

Although I do not have any experience in this particular field, I am interested in the use of stem cells in medical treatment of various diseases and injuries. However, I find molecular biology in general fascinating. Through this program I hope to obtain insights into different fields as well as into techniques used in medical research.

Neva Caliskan

First Name
Neva

Last Name
Caliskan

Date of Birth
12 August 1983

Country
Turkey

EDUCATION

College / University
2002 - 2007 Middle East Technical University, Ankara, Turkey

Highest Degree
B.Sc.

Major Subjects
Molecular Biology and Genetics

Lab Experience
Various techniques in molecular biology, cell culture and biochemistry

Projects / Research
09/2006 - 09/2007: Investigating the mechanism of anti-tumorigenic effects of 15-Lox-1 in colon cancer, Middle East Technical University, Ankara, Turkey
06/2005 - 09/2005: Identification of novel growth regulating genes in *Drosophila melanogaster*, EMBL, Developmental Biology Research Unit, Heidelberg, Germany

Publications
Banerjee S, Bozkurt O, Caliskan Neva G, Donertas Derya, Erdog A, Severcan F (2007) Effects of bioactive lipids on the cellular characteristics of colorectal cancer cells HCT-116. Bioactive Lipids in Cancer, Inflammation and related diseases. Montreal, Canada. September 16-19, 2007

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society
2006 - 2007 Scholarship from the Scientific and Technological Research Council of Turkey

SCIENTIFIC INTERESTS AND GOALS

My current interests include mechanisms of signal transduction in the cell, epigenetic regulations of chromatin structure and post transcriptional regulations of gene expressions. Through this program, I desire to have a deeper knowledge on molecular and cellular biology which will enable me to decide on the subject which I will further study on.

Annette Denker

First Name
Annette

Last Name
Denker

Date of Birth
29 January 1985

Country
Germany

EDUCATION

College / University
2004 - 2007 Ruprecht-Karls-University Heidelberg, Germany

Highest Degree
B.Sc.

Major Subjects
Molecular Biotechnology

Projects / Research
Bayer AG, Wuppertal, Department of Microbiology (HealthCare-Pharma-Biotechnology) with Dr. H. Wehlmann: Fermentation (Saccharomyces cerevisiae), drug production
German Cancer Research Center (DKFZ), Heidelberg, Department of Molecular Genetics with PD Dr. H. Herrmann-Lerdon: Laminopathies
Michigan State University, Lansing, MI, USA, Department of Biochemistry & Molecular Biology with Dr. W. Wedemeyer: Toxin secretion system of cholera bacteria
Ruprecht-Karls-University Heidelberg, Institute for Pharmacy and Molecular Biotechnology with Prof. Dr. S. Wölfl: Apoptosis in yeast and mammalian cells

Scholarships
2007 - 2008 International Max Planck Research School support
Since 2004 Studienstiftung des deutschen Volkes
2004 Kölner Gymnasial- und Stiftungsfonds

SCIENTIFIC INTERESTS AND GOALS

Although being not absolutely certain about my possible future research interests, I am currently especially fascinated by the process of apoptosis and the ways in which a cancer cell can escape this regulatory mechanism. I also find the change of cell fate (from a healthy to a tumor cell or from a stem cell to a differentiated cell) very interesting and further research on these topics will hopefully also be of great benefit in medical aspects, which, together with pure curiosity, is my main motivation for doing research.

Aliaksandr Dzementsei

First Name
Aliaksandr

Last Name
Dzementsei

Date of Birth
23 May 1985

Country
Belarus

EDUCATION

College / University
2002 - 2007 Belarusian State University, Minsk, Belarus
Highest Degree
Diploma
Major Subjects
Biotechnology, Microbiology
Lab Experience
Techniques in microbiology, molecular biology and molecular genetics
Projects / Research
2003 - 2006 The study of phenomenon caused by the mutation in gene pelW of bacterium *Erwinia carotovora* subsp. *atroseptica* that results in unusual production of pectinolytic enzymes.
2006 - 2007 The study of the minor phytopathagenicity and virulence factors of *Erwinia carotovora* subsp. *artroseptica*
Scholarships/Awards
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

I find molecular biology very interesting in all of its aspects. Although my research experience so far is mainly in microbiology, I am also very interested in molecular biology of nucleic acids and developmental biology. My main goal is to get advanced knowledge and research experience in a wide range of fields in modern molecular biology, to become a good scientist, and maybe one day to receive a Nobel Prize (why not?).

Liudmila Filonava

First Name
Liudmila

Last Name
Filonava

Date of Birth
14 February 1982

Country
Belarus

EDUCATION

College / University
2001- 2006 Belarusian State University, Minsk, Belarus
Highest Degree
Diploma
Major Subjects
Molecular Biology
Lab Experience
Techniques in microbiology, molecular biology and molecular genetics
Projects / Research
2006 - 2007: RAPD Analysis of the Flax Genome
2004 - 2006: The Obtaining of *Pantoea agglomerans* Strains with the Increased Carotenoid Pigments Production
Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

Based in my experience in microbiology, I find bacteria to be a fascinating object to study so that I might continue my research in that area. I am glad that I will have an opportunity to further broaden my theoretical and practical knowledge in other fields through my participation in this program.

Mandy Hannemann

First Name
Mandy

Last Name
Hannemann

Date of Birth
12 November 1978

Country
Germany

EDUCATION

College / University
09/2003 - 02/2007 Lausitz University of Applied Sciences, Senftenberg, Germany

Highest Degree
B.Sc.

Major Subjects
Biotechnology

Lab Experience
Techniques in molecular biology, microbiology, biochemistry and cell biology

Projects / Research
08/2006 – 01/2007: Molecular and structural characterization of a putative virulence chaperone in *Salmonella enterica* serovar Typhimurium, McMaster University, Canada
08/2005 – 01/2006: Ring Exported Protein 4 (REX4) from the human malaria parasite *Plasmodium falciparum*. Queensland Institute of Medical Research, Brisbane, Australia

Scholarships
10/2007 - 09/2008: International Max Planck Research School support
08/2006 - 12/2006: Scholarship from the German Academic Exchange Service (DAAD)
09/2005 - 02/2006: Scholarship from the Internationale Weiterbildung und Entwicklung gGmbH (InWEnt)

SCIENTIFIC INTERESTS AND GOALS:

I have a great interest in molecular genetics and microbiology. I am fascinated by pathogens and their ability to adapt to changing environments by gene regulation. I am looking forward to study how they can cause diseases.

Sebastian Hogl

First Name
Sebastian

Last Name
Hogl

Date of Birth
27 January 1985

Country
Germany

EDUCATION

College / University
2004 - 2007 Ruprecht-Karls-Universität, Heidelberg, Germany

Highest Degree
B.Sc.

Major Subjects
Drug Research, Bioinformatics, Biophysical Chemistry

Lab Experience
Experience in yeast (gene knockouts, protein expression, fluorescence microscopy), protein biochemistry, molecular biology and cell culture

Projects / Research
02/2006 - 08/2007 Analysis of spindle asymmetry in yeast, Biochemistry Center, University of Heidelberg, Germany

Scholarships
2007 - 2008 International Max Planck Research School support
Since 2005 Scholarship of the Foundation of German Business (Stiftung der Deutschen Wirtschaft)

SCIENTIFIC INTERESTS AND GOALS

I am especially interested in molecular cell biology as well as in molecular (tumor) pharmacology. Further I am interested in proteomics and signal transduction.

Chandini Kadian

First Name
Chandini

Last Name
Kadian

Date of Birth
3 September 1985

Country
India

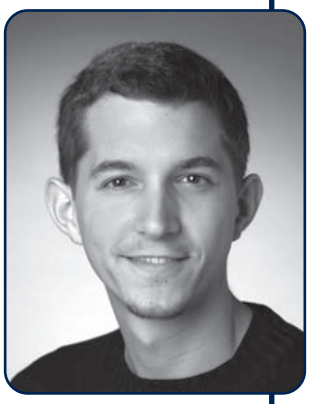
EDUCATION

College / University
2003 - 2006 Sri Venkateswara College, Delhi University
Highest Degree
B.Sc.
Major Subjects
Biochemistry
Lab Experience
Basic techniques in biochemistry
Projects / Research
Down regulation of thin filament proteins in actin mutants in indirect flight muscles of *Drosophila melanogaster*. IISc, Bangalore under Dr. Upendra Nongthomba (2005)
PCR amplification of the gene RV1211 from *Mycobacterium tuberculosis* and comparing the PCR product with the insert incised from recombinant pQE 32 – CAMLP: Dept of Biochemistry, Sri Venkateswara College under Dr. Hemlatha Reddy (2005)
Standardization of spectrophotometric NBT assay for detection of phagocytic activity of isolated immune cell suspensions. Dept of Biochemistry, Sri Venkateswara College under Dr. Nandita Narayanasamy (2004)
Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society
2003 - 2006 Dhirubhai Ambani Scholarship

SCIENTIFIC INTERESTS AND GOALS

Currently, DNA structure, chromatin modification and regulation of gene expression interest me. I would also like to learn more in the areas of protein structural analysis and RNA processing. Participating in this program gives me an excellent opportunity to further my knowledge in my areas of interest and explore many new exciting areas of research, something which I am eagerly looking forward to.

Frederik Köpper



First Name
Frederik

Last Name
Köpper

Date of Birth
20 July 1983

Country
Germany

EDUCATION

College / University
2003 - 2006 University of Bielefeld, Germany
Highest Degree
B.Sc.
Major Subjects
Molecular Biology and Genetics
Lab Experience
Basic methods in molecular biology, e.g. real time and RT-PCR, radiochemical assays, cell culture, biolistics
Projects / Research
3/2006 - 8/2006 Stable genetic transformation of the green alga *Gonium pectorale*. Bachelor's thesis, University of Bielefeld, Germany
4/2007 - 6/2007 Studies of the structure and kinetics of Xylosyltransferase I + II. Institut für Laboratoriums- und Transfusionsmedizin, Herz- und Diabeteszentrum Bad Oeynhausen, Germany
Scholarships
2007 - 2008 International Max Planck Research School support

SCIENTIFIC INTERESTS AND GOALS

Finding all fields of molecular biology highly fascinating, I have not yet decided where to put my main scientific focus, but I am particularly interested in gene regulation, cell differentiation, human genetics, biomedicine and its applications.

Andrius Krasauskas

First Name
Andrius

Last Name
Krasauskas

Date of Birth
15 May 1985

Country
Lithuania

EDUCATION

College / University
Jacobs University Bremen (former International University Bremen), Germany

Highest Degree
B.Sc.

Major Subjects
Biochemistry and Cell Biology

Lab Experience
Biochemistry, biophysical chemistry, chemistry, molecular and cell biology techniques

Projects / Research
02/2007 - 05/2007 Investigation of fis promoter *in vivo*; purification of H-NS. Jacobs University Bremen, Germany
09/2006 - 01/2007 Creating plasmid encoding Dnmt1-GFP fusion protein. Jacobs University Bremen, Germany
06/2006 - 09/2006 Investigation of the binding of Munc18 with mutant and wild-type syntaxin 1a; studies of SNARE complex assembly. Department of Neurobiology, MPI for Biophysical Chemistry, Göttingen, Germany

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

Structural biology: from the 3D view of a biological molecule to the 3D view of the cell.

Nadya Mitova

First Name
Nadya

Last Name
Mitova

Date of Birth
14 August 1985

Country
Bulgaria

EDUCATION

College / University
Jacobs University Bremen (former International University Bremen), Germany

Highest Degree
B.Sc.

Major Subjects
Biochemistry and Cell Biology

Lab Experience
Techniques in biochemistry, molecular biology, microbiology and embryology

Projects / Research
06/2007 - 09/2007 Characterization of a GT1/GT2-Nanog reporter mouse line during embryonal development, Max Planck Institute of Immunobiology, Freiburg, Germany
09/2006 - 04/2007 Guided research - Cloning, purification and biochemical characterization of the PHD domains of Dnmt3 family of mammalian DNA methyltransferases, Jacobs University Bremen, Bremen, Germany
06/2006 - 08/2006 Construction of in-frame deletion mutants in the translocation domain of the virulent effector YopE, in *Yersinia pseudotuberculosis*, Umeå University, Umeå, Sweden

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

My main interests are in the areas of developmental biology, stem cell research and epigenetics. My goal is to do research on the molecular networks and mechanisms that govern cellular behaviors.

Sina Mozaffari Jovin

First Name
Sina

Last Name
Mozaffari Jovin

Date of Birth
3 April 1983

Country
Iran

EDUCATION

College / University
2001 - 2005 University of Shiraz, Shiraz, Iran
2005 - 2007 National Institute of Genetic Engineering and Biotechnology

Highest Degree
M.Sc.

Major Subjects
Cell and Molecular Biology

Lab Experience
Techniques in molecular biology, biochemistry and microbial biotechnology

Projects / Research
2006 High-yield extracellular expression (in shake-flask and bioreactor), and efficient purification of *E.coli* asparaginase II, an enzyme used in leukemia treatment.
2007 Reduction of glutaminase activity in an engineered *E.coli* asparaginase II with protein engineering techniques.

Publication:
Mozaffari Jovin S, et al. (2007) Study of hypervitaminosis A and histological effects on mice testis. Iranian Journal of Basic Medical Sciences, 10(1): 33
Mozaffari Jovin S, et al. (2007) Pilot-scale production of asparaginase for reduction of acrylamide in fried foods. Oral presentation Tarbiat Modares University, Tehran, Iran
Mozaffari Jovin S, et al. (2007) The effects of hypervitaminosis A on alkaline phosphatase activity in mice testis. Oral presentation at the 18th Iranian Congress of Physiology & Pharmacology, Mashad, Iran

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society
2005 University of Shiraz award for the first rank in B.Sc. of Biological Science

SCIENTIFIC INTERESTS AND GOALS

Investigation of the molecular aspects of diseases to develop new therapies. Besides, I am keen to investigate the rules of protein structure-function relationships in order to design and engineer proteins with new functions or desirable features.

Miroslav Nikolov

First Name
Miroslav

Last Name
Nikolov

Date of Birth
18 June 1984

Country
Bulgaria

EDUCATION

College / University
2003 - 2007 Sofia University “St.Klement Ohridski”

Highest Degree
B.Sc.

Major Subjects
Molecular Biology, Biochemistry

Lab Experience
Basic techniques in molecular biology, phage display, plant tissue cultures

Projects / Research
“Investigation of the structure and composition of the cell wall of NaCl-stressed callus cultures from *Dactylis glomerata* L.” and another related project
“Effects of synthetic low- and high-molecular weight zwitterions on the ligand-binding ability of C1q and its globular fragments.” All at Faculty of Biology, Sofia University

Publications
Nikolov M, Tsacheva I, Petrova S, Tchorbadjieva M, Ivanov I, Kamenova I, Kamenska E, Georgiev G (2006) Effects of synthetic low- and high-molecular weight zwitterions on the IgG-binding ability of C1q and its globular fragments. V National Conference in Chemistry for students and PhD-students, Sofia University
Zagorchev L, Tsanov K, Nikolov M, Odjakova M (2006) Protein markers for *in vitro* regeneration of somatic embryoids from *D. glomerata* L., IBSC, Riga, Latvia

Scholarships
2007-2008 Stipend of the Excellence Foundation of the Max Planck Society
2004 - 07/2007 Scholarship for academic excellence, Sofia University
2003 - 07/2007 Scholarship for high achievements on IBO, “Evrika” Foundation

SCIENTIFIC INTERESTS AND GOALS

Currently, I am interested in epigenetic control of gene expression and silencing, some aspects of immunology and especially instrumental methods applied in (prote)omics.

Doris Petroi

First Name
Doris

Last Name
Petroi

Date of Birth
3 June 1985

Country
Romania

EDUCATION

College / University
2004 - 2007 Jacobs University Bremen, Germany

Highest Degree
B.Sc.

Major Subjects
Biochemistry and Cell Biology

Lab Experience
Techniques in biochemistry, molecular biology, cell biology, and chemistry

Projects / Research
09/2006 - 12/2006 Dnmt3a and Dnmt3b expression in Bac-to-Bac system and further molecular analysis
06/2006 - 08/2006 Finding, amplifying and overexpressing alcohol dehydrogenase genes from *Clostridium perfringens*
03/2005 - 04/2006 Metabolic compensation of temperature acclimation in *Saccharomyces cereviciae*

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

My main research interests are molecular medicine and biotechnology. I am very keen to gain more insight into the cellular and molecular processes that underlie the development of diseases, and I would like to apply new molecular biology techniques to the medical field.

Volkan Sakin

First Name
Volkan

Last Name
Sakin

Date of Birth
1 July 1984

Country
Turkey

EDUCATION

College / University
2002 - 2007 Middle East Technical University, Ankara, Turkey

Highest Degree
B.Sc.

Major Subjects
Molecular Biology and Genetics

Lab Experience
Cloning/ Subcloning, assay for telomerase activity, genotyping of transgenic mice, paper and thin layer chromatography, electroporation, isolation of mouse embyo fibroblasts and some other basic molecular biology techniques

Projects / Research
07/2006 - 09/2006 Cloning of TIMP3 and OSF2 into pcDNA3-Neomycin to investigate the in vivo effect of expression of these genes on oncogenic phenotype
10/2006 - 01/2007 Investigation of hTERT mRNA levels and telomerase activity in drug resistant MCF-7 cells

Publications
Sakin V, Eskiocak U, Demirel Kars M, Darcansoy Iseri O, Gündüz U (2007) Investigation of hTERT mRNA levels and telomerase activity in drug resistant MCF-7 cells (e-poster at II. Congress of Molecular Medicine, March 2007, Istanbul, Turkey)

Scholarships
2007 - 2008 Stipend of the Excellence Foundation of the Max Planck Society

SCIENTIFIC INTERESTS AND GOALS

Many branches of biological sciences are fascinating to me such as posttranslational modifications of proteins of the SUMO family, stem cell biology, molecular oncology, and epigenetics. The Master's year will be exciting to explore which is best for me. With this program. I hope that I will be well suited to ask my own questions and direct my own research into their answers.

Broder Schmidt

First Name
Broder

Last Name
Schmidt

Date of Birth
30 April 1985

Country
Germany

EDUCATION

College / University
2004 - 2007 Ruprecht-Karls-Universität Heidelberg, Heidelberg, Germany
Highest Degree
B.Sc.
Major Subjects
Cell and Molecular Biology
Lab Experience
Various techniques in biochemistry, cell and molecular biology
Projects / Research
04/2007 - 06/2007 Analysis of the cell cycle in GET-gene mutant yeast (diploma thesis). Center for Molecular Biology Heidelberg (ZMBH), Heidelberg, Germany
07/2007 - 09/2007 Cloning, purification and crystallization of the G-domain of the chloroplast translocon subunit Toc159. Center for Biochemistry of the University of Heidelberg (BZH), Heidelberg, Germany
Scholarships
2007 - 2008 International Max Planck Research School support

SCIENTIFIC INTERESTS AND GOALS

I am generally interested in trying to understand the essential processes of life in molecular detail. Among all the different cellular events, protein trafficking and insertion fascinate me most at the moment. However, as cellular life is much more complex and intriguing, I find it difficult to decide on a single specific topic. By providing the opportunity to deepen my knowledge in the major research areas of molecular biology both in theory and practice, this program promises to prepare me thoroughly to address the many interesting unsolved questions remaining in nature.

Cornelius Schneider

First Name
Cornelius

Last Name
Schneider

Date of Birth
Germany

Country
Russian Federation

EDUCATION

College / University
2004 - 2007 Alma Mater Studiorum Bologna
Highest Degree
B.Sc.
Major Subjects
Molecular Biology
Lab Experience
Various techniques in molecular biology and microbiology
Projects / Research
2006 Two month summer project on improving the detection and identification of viral contamination in shellfish.
2007 Bachelor's thesis: gene transfer of alk gene of Rhodococcus aeterovorans BCP1 in *Eschericha coli* DH5a and Pseudomonas putida KF707
Scholarships
2007 - 2008 International Max Planck Research School support

SCIENTIFIC INTERESTS AND GOALS

One of my major interests is the study of metabolic pathways that allow different organisms to degrade or synthesize molecules of specific interest. I am also quite interested in structural biochemistry such as, for example, the relation between structure and function of biomolecules.

Susana	Andrade	Structural Biology	U Göttingen
Donna J.	Arndt-Jovin	Molecular Biology	MPI bpc
Mathias	Bähr	Neurology	U Göttingen
Botho	Bowien	Microbiology	U Göttingen
Gerhard H.	Braus	Molecular Microbiology	U Göttingen
Bertram	Brenig	Molecular Biology of Livestock	U Göttingen
Nils	Brose	Molecular Neurobiology	MPI em
Matthias	Dobbelstein	Molecular Oncology	U Göttingen
Detlef	Doenecke	Biochemistry	U Göttingen
Stefan	Eimer	Molecular Neurogenetics	ENI
Wolfgang	Engel	Human Genetics	U Göttingen
Dirk	Fasshauer	Neurobiology	MPI bpc
Ivo	Feußner	Plant Biochemistry	U Göttingen
Ralf	Ficner	Molecular Structural Biology	UGöttingen
Wolfgang	Fischle	Chromatin Biochemistry	MPI bpc
Christiane	Gatz	General and Developmental Physiology of the Plant	U Göttingen
Dirk	Görlich	Cellular Logistics	MPI bpc
Christian	Griesinger	NMR-based Structural Biology	MPI bpc
Uwe	Groß	Bacteriology	U Göttingen
Heidi	Hahn	Human Genetics	U Göttingen
Herbert	Jäckle	Molecular Developmental Biology	MPI bpc
Reinhard	Jahn	Neurobiology	MPI bpc
Thomas	Jovin	Molecular Biology	MPI bpc
Michael	Kessel	Molecular Biology	MPI bpc
Dieter	Klopfenstein	Biochemistry	U Göttingen
Wilfried	Kramer	Molecular Genetics	U Göttingen
Kerstin	Krieglstein	Neuroanatomy	U Göttingen
Wolfgang	Liebl	Microbiology	U Göttingen
Reinhard	Lührmann	Cellular Biochemistry	MPI bpc
Ahmed	Mansouri	Molecular Developmental Genetics	MPI bpc
Frauke	Melchior	Biochemistry	U Göttingen
Burkhard	Morgenstern	Bioinformatics	U Göttingen
Klaus-Armin	Nave	Neurogenetics	MPI em
Erwin	Neher	Membrane Biophysics	MPI bpc
Tomas	Pieler	Developmental Biochemistry	U Göttingen
Stefanie	Pöggeler	Genetics of Eukaryotic Organisms	U Göttingen
Silvio	Rizzoli	STED Microscopy of Synaptic Function	ENI
Reinhard	Schuh	Molecular Organogenesis	MPI bpc
George Michael	Sheldrick	Structural Chemistry	U Göttingen
Holger	Stark	3D Electron Cryomicroscopy	MPI bpc
Jörg	Stülke	General Microbiology	U Göttingen
Michael	Thumm	Molecular Cell Biology	U Göttingen
Henning	Urlaub	Bioanalytical Mass Spectrometry	MPI bpc
Markus	Wahl	X-Ray Crystallography	MPI bpc
Lutz	Walter	Primate Genetics	DPZ
Jürgen	Wienands	Immunology	U Göttingen
Ernst	Wimmer	Developmental Biology	U Göttingen
Andreas	Wodarz	Stem Cell Biology	U Göttingen

U Göttingen = Georg August University, MPI bpc = Max Planck Institute for Biophysical Chemistry, MPI em = Max Planck Institute for Experimental Medicine, DPZ = German Primate Center

Susana Andrade

Group Leader within the Emmy Noether Program

- Biochemistry Diploma, Faculty of Sciences, University of Lisboa, Portugal, 1996
- PhD, Faculty of Sciences and Technology, University Nova of Lisboa, 2001
- Postdoc, California Institute of Technology, Pasadena, CA, USA, 2002
- Marie Curie Postdoc Fellow, Dept. of Molecular Structural Biology, Göttingen, 2003 - 2005
- Independent Emmy Noether Group Leader since 2006

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fax: + 49-551-39 14082
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Further Information

http://www.img.bio.uni-goettingen.de/ms-www/index.html

Major Research Interests

Our group focuses on understanding the mechanisms that regulate the function of membrane proteins at a molecular level. Most of the proteins we investigate belong to the ubiquitous Ammonium Transport family (Amt). From prokaryotes to plants, these proteins supply nitrogen to the cell, in its bio-available form NH_4^+/NH_3, which is required for the synthesis of molecules like amino acids or nucleic acids. Their homologous in mammals are the Rhesus proteins, from which the most prominent member is the Rhesus protein from erythrocytes that determines our blood type. Also present in kidney and liver tissues, Amt proteins in such organisms are fundamentally involved in acid/basic homeostasis. The precise mechanism of transport remains, however, unclear.

We determined the high-resolution structure of an archaeal Amt protein, in various ammonium-containing soak solutions, by X-ray crystallography. To clarify the transport mechanism of this class of proteins, we take a multidisciplinary approach using in combination with such structural studies, molecular biology, biochemistry, modeling and *in vitro* transport assays of Amt proteins reconstituted into lipid bilayers (proteoliposomes).

Selected Recent Publications

Andrade SLA, Einsle O (2007) The Amt/Mep/Rh Family of Ammonium Transport Proteins. Molec Memb Biol 24(5-6): 357-365

Einsle O, Andrade SLA, Dobbek H, Meyer J, Rees DC (2007) Assignment of Individual Metal Redox States in a Metalloprotein by Crystallography Refinement at Multiple X-ray Wavelengths. J Am Chem Soc 129: 2210-2211

Andrade SLA, Dickmanns A, Ficner R, Einsle O (2005) Crystal Structure of the Archaeal Ammonium Transporter Amt-1 from *Archaeoglobus fulgidus*. Proc Natl Acad Sci USA 102: 14994-14999

Andrade SLA, Dickmanns A, Ficner R, Einsle O (2005) Expression, Purification and Crystallization of the Ammonium Transporter Amt-1 from *Archaeoglobus fulgidus*. Acta Cryst F61: 861-863

Andrade SLA, Cruz F, Drennan CL, Ramakrishnan V, Rees DC, Ferry JG, Einsle O (2005) Structures of the Iron-Sulfur Flavoproteins from *Methanosarcina thermophila* and *Archaeoglobus fulgidus*. J Bacteriology 187(11): 3848-3854

Andrade SLA, Brondino CD, Kamenskaya EO, Levashov AV, Moura JJG (2003) Kinetic behavior of *Desulfovibrio gigas* Aldehyde Oxidoreductase encapsulated in Reverse Micelles. Biochem Biophys Res Comm 308: 73-78

Einsle O, Tezcan FA, Andrade SLA, Schmid B, Yoshida M, Howard JB, Rees CD (2002) The Nitrogenase MoFe Protein at 1.16 Å Resolution: A Central Ligand in the FeMo Cofactor. Science 297: 1696-1700

Donna J. Arndt-Jovin

Professor, Group Leader at the Max Planck Institute for Biophysical Chemistry

- A.B., Chemistry, Hiram College, 1963
- Ph.D., Biochemistry, Yale University, 1969
- Fellow of the Jane Coffin Childs Memorial Fund for Medical Research, Department of Biochemistry, Stanford University School of Medicine, 1969 - 1971
- Research Scientist, Max Planck Institute for Biophysical Chemistry, 1971 - 1993
- Senior Research Scientist, Max Planck Institute for Biophysical Chemistry, 1993 - present

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Biophysical Chemistry
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Further Information

http://www.mpibpc.gwdg.de/abteilungen/060/people/donna/index.html

Major Research Interests

Chromatin structure and function *in vivo*,

(a) the study of nuclear architecture using immunochemistry, *in situ* hybridization, and *in vivo* 3-D image microscopy

(b) the role of epigenetic regulation in gene expression and development of Dipteran embryos with focus on polycomb group proteins

Signal transduction processes: cell surface antigen-receptor proximities and mobilities focused on the erb B receptor family in living tissue culture cells.

DNA structure and function. Biological roles of unusual helical DNA structures.

Development of new fluorescence imaging modalities for rapid, *in vivo* cell and organism imaging.

Selected Recent Publications

Ficz G, Heintzmann R, Arndt-Jovin DJ (2005) Polycomb group protein complexes exchange rapidly in living *Drosophila*. Development 132: 3963-3976

Hanley QS, Lidke KA, Heintzmann R, Arndt-Jovin DJ, Jovin TM (2005) Fluorescence lifetime imaging in an optically sectioned Programmable Array Microscope (PAM). Cytometry 67A: 112-118

Lidke DS, Lidke KA, Rieger B, Jovin TM, Arndt-Jovin DJ (2005) Reaching out for signals: filopodia sense EGF and respond by directed retrograde transport of activated receptors. J Cell Biol 170: 619-626

Post JN, Lidke KA, Rieger B, Arndt-Jovin DJ (2005) One- and two-photon photoactivation of a paGFP-fusion protein, a phototoxicity study in live Drosophila embryos. FEBS Lett 579: 325-330

Lidke DS, Nagy P, Heintzmann R, Arndt-Jovin DJ, Post JN, Grecco H, Jares-Erijman EA, Jovin TM (2004) Quantum dot ligands provide new insights into erbB/HER receptor-mediated signal transduction. Nat Biotechnol 22: 198-203

Shchyolkina A, Kaluzhny DN, Borisova OF, Hawkins ME, Jernigan RL, Jovin TM, Arndt-Jovin DJ, Zhurkin VB (2004) Formation of an intramolecular triple-stranded DNA structure monitored by fluorescence of 2-aminopurine or 6-methylisoxanthopterin. Nucl Acids Res 32: 432-440

Mathias Bähr

Professor of Neurology

- 1985 MD, University of Tübingen Medical School, Training in Neurology at University Hospitals in Tübingen and Düsseldorf
- DFG and Max Planck Fellow at the Max Planck Institute for Developmental Biology Tübingen and at the Department of Anatomy and Cell Biology, Washington University St.Louis
- Schilling-Foundation Professor for Clinical and Experimental Neurology, University of Tübingen
- Director at the Department of Neurology, University of Göttingen since 2001

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Neurological Medicine
Neurology
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Further Information

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Major Research Interests

We are interested to understand 2 basic questions in cellular and molecular neurobiology:
1. Which factors support survival of adult CNS neurons?
2. What kills these cells under pathological conditions?

Up to now, only little is known about the mechanisms that support survival of a postmitotic cell like a human neuron for eventually more than 100 years under physiological conditions. However, by examining the molecular regulation of cell survival and cell death during development and in the lesioned adult CNS, one may get some clues to answer this question.

In our group, several *in vitro* and *in vivo* model systems are used which allow examination of neuronal de- and regeneration. Our basic model is the rodent retino-tectal projection. Here, we can study development, de- and regeneration of the respective projection neurons, the retinal ganglion cells (RGCs) in single cell cultures, explants or *in vivo*. Transection or crush-axotomy of the optic nerve induces retrograde death more than 80% of RGCs within two weeks. This secondary cell loss is mainly apoptotic and involves specific changes in gene expression pattern of transcription factors (e.g. c-jun or ATF-2), pro- and anti-apoptotic genes (e.g. bcl-2 or bax) and growth-associated genes (like GAP-43). Thus, long term survival and initiation of regeneration programmes of RGCs critically depends on inhibition of apoptotic cell death. To that end, we have used a variety of techniques to interfere with the cell death cascades that follow lesions of the optic nerve in adult rats. Inhibition of neuronal apoptosis can be afforded by pharmacological administration of trophic factors or by gene therapy approaches using adeno- or adeno-associated virus vectors that can deliver neurotrophic or anti-apoptotic factors directly into neurons or into surrounding glial cells. These, and other new strategies like using peptide-transduction-domains to deliver anti-apoptotic proteins across the blood-brain-barrier are now used to develop new experimental therapy strategies in animal models of human neurological disorders like stroke, trauma, multiple sclerosis or neurodegenerative diseases (e.g. Alzheimer´s or Parkinson´s disease).

Selected Recent Publications

Meyer R, Weissert R, de Graaf K, Diem R, Bähr M (2001) Acute neuronal apoptosis in a rat model of multiple sclerosis. J Neurosci 21: 6214-6220

Kilic E, Dietz GPH, Herrmann DM, Bähr M (2002) Intravenous TAT-Bcl-XL is protective when delivered before and after middle cerebral artery occlusion in mice. Ann Neurol 52(5): 617-22

Diem R, Hobom M, Maier K, Weissert R, Storch MK, Meyer R, Bähr M (2003) Methyprednisolone increases neuronal apoptosis during autoimmune CNS inflammation by inhibition of an endogenous neuroprotective pathway. J Neurosci 23(18): 6993-7000

Dietz GPH, Bähr M (2004) Delivery of Bioactive Molecules into the Cell: The Trojan Horse Approach. Mol Cell Neurosci 27(2): 85-131

Diem R, Sättler MB, Merkler D, Demmer I, Maier K, Stadelmann C, Ehrenreich H, Bähr M (2005) Combined therapy with methylprednisolone and erythropoietin in a model of multiple sclerosis. Brain 128: 375-85

Lingor P, Koeberle P, Kügler S, Bähr M (2005) Downregulation of apoptosis mediators by RNA interference inhibits axotomy-induced retinal ganglion cell death *in vivo*. Brain 128: 550-558

Botho Bowien

Professor of Microbiology

- Dr. rer. nat., Georg-August-Universität Göttingen, 1970
- Postdoc, Case Western Reserve University, Cleveland, Ohio, USA, 1973 - 1975
- Habilitation (Microbiology), Georg-August-Universität Göttingen, 1978
- Professor of Microbiology, Georg-August-Universität Göttingen, 1983

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Grisebachstr. 8

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fax: +49-551-39 9842
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Major Research Interests

Carbon dioxide (CO_2) is an essential gas for all organisms. Assimilation of CO_2 by autotrophs such as the photosynthetic higher plants, algae and cyanobacteria constitutes the primary biosynthetic activity in the biosphere. In addition to these organisms there is a great diversity of photo- and/or chemoautotrophic bacteria and archaea. Such organisms are often facultative autotrophs, i.e. they are able to grow either autotrophically or heterotrophically. The mutual shift between autorophy and heterotrophy requires a sophisticated regulation on the metabolic as well as genetic level.

Ralstonia eutropha H16 is an aerobic, facultatively chemoautotrophic bacterium that assimilates CO_2, like the majority of autotrophs, via the Calvin-Benson-Bassham (CBB) carbon reduction cycle. A main interest of our laboratory concerns the transcriptional control of the *cbb* operons encoding most of the CBB enzymes in *R. eutropha*. The regulatory components of the *cbb* system, their response to metabolic signals and the interlocking of the *cbb* control with larger regulatory networks are the prime research subjects. The recently elucidated genome sequence of the organism provides an excellent basis to study these questions.

Functional genomics of *R. eutropha* H16 with the goal of assessing and developing the metabolic potential of the organism for future biotechnological applications -particularly under autotrophic growth conditions- is another field of research. The genetics and control of sugar and sugar acid utilization in *R. eutropha* H16 are also being investigated.

Selected Recent Publications

Pohlmann A, Fricke WF, Reinecke F, Kusian B, Liesegang H, Cramm R, Eitinger T, Ewering C, Pötter M, Schwartz E, Strittmatter A, Voß I, Gottschalk G, Steinbüchel A, Friedrich B, Bowien B (2006) Genome sequence of the bioplastic-producing "Knallgas" bacterium *Ralstonia eutropha* H16. Nature Biotechnol 24: 1257-1262

Pötter M, Müller H, Reinecke F, Wieczorek R, Fricke F, Bowien B, Friedrich B, Steinbüschel A (2004) The complex structure of polyhydroxybutyrate (PHB) granules: four orthologous and paralogous phasins occur in *Ralstonia eutropha*. Microbiology 150: 2301-2311

Kusian B, Sültemeyer D, Bowien B (2002) Carbonic anhydrase is essential for growth of *Ralstonia eutropha* at ambient CO_2 concentrations. J Bacteriol 184: 5018-5026

Bowien B, Kusian B (2002) Genetics and control of CO_2 assimilation in the chemoautotroph *Ralstonia eutropha*. Arch Microbiol 178: 85-93

Gerhard H. Braus

Professor of Microbiology and Genetics

- Diploma (Biology), Albert-Ludwig University, Freiburg i. Br. (Germany), 1983
- Dr.sc.nat., Swiss Federal Institute of Technology (ETH), Zürich (Switzerland), 1987
- Habilitation (Microbiology), Swiss Federal Institute of Technology (ETH), Zürich (Switzerland), 1991
- Associate Professor of Biochemistry, Friedrich Alexander University, Erlangen (Germany), 1993 - 1996
- Since 1996 Professor of Microbiology (since 2001 Professor of Microbiology and Genetics) in Göttingen

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Further Information

http://www.gwdg.de/~molmibio

Major Research Interests

The major focus of our laboratory is on molecular microbiology of eukaryotic microorganisms (yeasts and filamentous fungi): Control of developmental programs, pathogenicity (humans and plants) and interplay with metabolism.

(i) We are interested in the molecular control of adhesion as initial step in infection and biofilm formation.

(ii) Amino acid starvation activates a complex fungal genetic network (transcription factor Gcn4p/CpcA) which also controls adhesion in yeast, virulence in *Aspergillus fumigatus* and fruitbody formation in *A. nidulans*. We study the translational control and the protein stability regulation which is involved.

(iii) Nedd8 is a ubiquitin-like protein which is involved in the control of protein turnover. We study the Nedd8-system including the COP99 signalosome using fungi as model systems.

(iv) We study alpha-synuclein from Parkinson patients expressed in fungi to explore the potential to learn more about neuronal diseases (Collaborative research within the DFG Research Center of Molecular Physiology of the Brain, CMPB)

Selected Recent Publications

Busch S, Schwier EU, Nahlik K, Bayram Ö, Draht OW, Helmstaedt K, Krappmann S, Valerius O, Lipscomb WN, Braus GH (2007) An eight-subunit COP9 signalosome with an intact JAMM motif is required for fungal fruit body formation. Proc Natl Acad Sci USA 104: 8125-8130.

Valerius O, Kleinschmidt M, , Rachfall N, Schulze F, Marin SL, Hoppert M, Bömeke K, Fischer C, Braus GH (2007) The *S. cerevisiae* homolog of mammalian RACK1, CPC2/ASC1, is required for FLO11 dependent adhesive growth and dimorphism. Mol Cell Proteomics, in press.

Bömeke K, Pries R, Korte V, Scholz E, Herzog B, Schulze F, Braus GH (2006) Yeast Gcn4p stabilization is innitiated by the dissociation of the nuclear Pho85p/Pc15p complex. Mol Biol Cell 17: 2952-2962

Helmstaedt K, Strittmatter A, Lipscomb WN, Braus GH (2005) Evolution of 3-deoxy-D-arabino-heptulosonate-7-phosphate synthase-encoding genes in the yeast *Saccharomyces cerevisiae*. Proc Natl Acad Sci USA 102: 9784-9789

Galagan JE, …, Braus GH (18th of 50 authors), …Birrer B (2005) Sequencing of *Aspergillus nidulans* and comparative analysis with *A. fumigatus* and *A. oryzae*. Nature: 428: 1105-1115

Bertram Brenig

Full Professor of Molecular Biology of Livestock

- Director of the Institute of Veterinary Medicine
- Dr. med. vet., University of Munich, Munich 1987

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Major Research Interests

The main interest of the laboratory is in the structural and functional analysis of mammalian genes and genomes. We are investigating the cause of different economical important genetic defects in livestock and other domesticated animals. So far our main focus was on porcine genes and their function, e.g. we are currently analysing the molecular origin of porcine hernia inguinalis and scrotalis. Using a whole genome scan we have identified several chromosomal regions that are linked to this disorder. Fine mapping, positional cloning and candidate gene analysis are used for further elucidation. However, in recent years we have also started to look at genes in other species, e.g. cattle, dog, sheep, and buffalo. Especially, in cattle, we are interested in the molecular analysis of bovine spongiform encephalopathy (BSE). The differences in oral uptake of prion protein between cattle and pig is studied *in vivo* and *in vitro*. We have identified humoral microvesical encapsulated nucleic acids that are altered during a prion infection and can be used as diagnostic tool. Humoral nucleic acids are also studied in several other diseases, e.g. liver carcinoma in dog and pancreatic neoplasias in cat.

Selected Recent Publications

Beck J, Bornemann-Kolatzki K, Knorr C, Täubert H, Brenig B (2006) Molecular characterization and exclusion of porcine GUSB as a candidate gene for congenital hernia inguinalis/scrotalis. BMC Vet Res 2: 14

Chen K, Knorr C, Bornemann-Kolatzki K, Huang L, Rohrer GA, Brenig B (2006) Characterization of the PGK2 associated microsatellite S0719 on SSC7 suitable for parentage and QTL diagnosis. Anim Biotechnol 17: 43-49

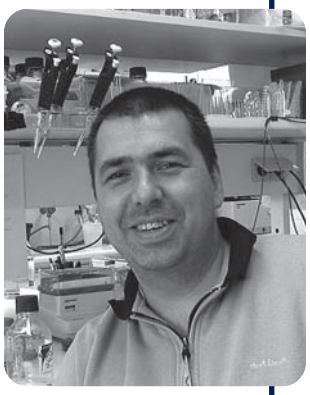
Drögemüller C, Giese A, Martins-Wess F, Wiedemann S, Andersson L, Brenig B, Fries R, Leeb T (2006) The mutation causing the black-and-tan pigmentation phenotype of Mangalitza pigs maps to the porcine ASIP locus but does not affect its coding sequence. Mamm Genome 17: 58-66

Schütz E, Scharfenstein M, Brenig B (2006) Genotyping of ovine prion protein gene (PRNP) variants by PCR with melting curve analysis. Clin Chem 52: 1426-1429

Knorr C, Beuermann C, Beck J, Brenig B (2007) Characterization of the porcine multicopy ribosomal protein SA/37-kDa laminin receptor gene family. Gene 395: 135-143

Thomzig A, Schulz-Schaeffer W, Wrede A, Wemheuer W, Brenig B, Kratzel C, Lemmer K, Beekes M (2007) Accumulation of pathological prion protein PrPSc in the skin of animals with experimental and natural scrapie. PLoS Pathog 3: e66

Nils Brose



Professor, Director at the Max Planck Institute for Experimental Medicine

- Dr. rer. nat. (Ph.D.) 1990, Ludwig Maximilians University Munich
- Appointed as Director at the Max Planck Institute for Experimental Medicine 2001

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Major Research Interests

Research in the Department of Molecular Neurobiology focuses on the molecular mechanisms of synapse formation and function in the vertebrate central nervous system. Typically, synapses are formed between cellular processes of a sending and a receiving nerve cell. They are the central information processing units in the vertebrate brain where some 10^{12} nerve cells are connected by 10^{15} synapses to form an elaborate and highly structured neuronal network that is the basis for all forms of behaviour. Signal transmission at synapses is mediated by the regulated release of signal molecules (neurotransmitters) which then diffuse to the receiving nerve cell and change its physiological state. In the Department of Molecular Neurobiology, we combine biochemical, morphological, mouse genetic, behavioural, and physiological methods to elucidate the molecular basis of synapse formation and transmitter release processes. Our synaptogenesis research concentrates on synaptic cell adhesion proteins and their role in synapse formation. Studies on the molecular mechanisms of neurotransmitter release focus on components of the presynaptic active zone and their regulatory function in synaptic vesicle fusion.

Selected Recent Publications

Varoqueaux F, Aramuni G, Rawson RL, Mohrmann R, Missler M, Gottmann K, Zhang W, Südhof TC, Brose N (2006) Neuroligins determine synapse maturation and function. Neuron 51: 741-754

Reim K, Wegmeyer H, Brandstätter JH, Xue M, Rosenmund C, Dresbach T, Hofmann K, Brose N (2005) Structurally and functionally unique Complexins at retinal ribbon synapses. J Cell Biol 169: 669-680

Junge H, Rhee J-S, Jahn O, Varoqueaux F, Spiess J, Waxham MN, Rosenmund C, Brose N (2004) Calmodulin and Munc13 form a Ca^{2+}-sensor/effector complex that controls short-term synaptic plasticity. Cell 118: 389-401

Roßner S, Fuchsbrunner K, Lange-Dohna C, Hartlage-Rübsamen M, Bigl V, Betz A, Reim K, Brose N (2004) Munc13-1-mediated vesicle priming contributes to secretory APP processing. J Biol Chem 279: 27841-27844

Rhee J-S, Betz A, Pyott S, Reim K, Varoqueaux F, Augustin I, Hesse D, Südhof TC, Takahashi M, Rosenmund C, Brose N (2002) b Phorbol ester- and diacylglycerol-induced augmentation of transmitter release is mediated by Munc13s and not by PKCs. Cell 108: 121-133

Professor of Molecular Oncology

- Dr. med., University of Munich, 1993
- Postdoctoral fellow, Princeton University, USA, 1993 - 1996
- Group leader, University of Marburg, 1997 - 2004
- Professor of Molecular Oncology, University of Southern Denmark, Odense, since 2004
- Head of the Department of Molecular Oncology, Georg-August-Universität Göttingen, since 2005

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Major Research Interests

We are focussing our research on the tumor suppressor p53, trying to elucidate its mechanisms of action, its regulation and its suitability as a target for cancer therapy. p53 operates as a transcription factor and prevents uncontrolled cell proliferation. This activity is regulated through a sophisticated regulatory network that responds to DNA damage. Despite our knowledge concerning the molecular biology of p53, an integrated concept of its regulation, and its translation into rational diagnostics and therapy, are still in their infancy. The tumor suppressor gene TP53 is mutated or deleted in approximately 50% of malignant tumors. However, this does not mean that p53 is active in the remaining cases. It appears that in the vast majority of the remaining 50% of tumors, p53 is inactivated through malfunction of its modulators, such as Mdm2, p14ARF, deltaNp73, and others. We are therefore pursuing the unique opportunity to re-establish p53's "dormant" tumor-suppressive activity by targeting its modulators as a potential avenue to therapy.

Selected Recent Publications

Kranz D, Dobbelstein M (2006) Non-genotoxic p53 activation protects cells against S phase specific chemotherapy. Cancer Research, in press

Schürmann M, Dobbelstein M (2006) Adenovirus-induced ERK phosphorylation during the later phase of infection enhances viral protein levels and virus progeny. Cancer Research 66: 1282-1288

Roth J, Lenz-Stöppler C, Contente A, Löhr K, Koch P, Dobbelstein M (2003) Reactivation of mutant p53 by a one-hybrid adaptor protein. Cancer Research 63, 3904-3908

Contente A, Dittmer A, Koch MC, Roth J, Dobbelstein M (2002) A polymorphic microsatellite that mediates induction of PIG3 by p53. Nature Genetics 30: 315-320

Roth J*, Dobbelstein M*, Freedman D, Shenk T, Levine AJ (1998) Nucleo-cytoplasmic shuttling of the hdm2-oncoprotein regulates the levels of the p53 protein via a pathway used by the human immunodeficiency virus rev protein. EMBO Journal 17: 554-564 (*equal contributors)

Dobbelstein M*, Roth J*, Kimberly WT, Levine AJ, Shenk T (1997) Nuclear export of the adenoviral oncoproteins E1B-55 kD and E4-34 kD. EMBO Journal 16: 4276-4284 (*equal contributors)

Detlef Doenecke

Professor of Biochemistry

- MD, 1967, University Saarland Medical School
- Postdoc at the Universities of San Francisco (UCSF) and Marburg
- Professor of Biochemistry, 1987, University of Göttingen
- Head of the Department of Molecular Biology at the Institute of Biochemistry and Molecular Cell Biology

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Major Research Interests

The main interest of the laboratory is in the structure, function and regulation of synthesis of nuclear proteins including chromosomal proteins and other protein factors involved in the control of transcription. DNA replication during the S-phase of the cell cycle requires the coordinate synthesis of histones in stoichiometric amounts for the assembly of chromatin on replicated DNA. The major human histone gene cluster has been mapped to chromosome 6p21.1-6p22.2, and more than 50 histone genes were identified and sequenced within that gene cluster. Several S-phase independent histone genes map as solitary genes to other chromsomes. Current work in this project area deals with the function and of individual H1 histone subtype genes. A second major project deals with the factors mediating the transport of histone-related transcriptional regulators from the cytoplasm to the cell nucleus. This work concentrates on the differential role of nuclear import receptors and specific protein-protein interactions during the nuclear transport of these proteins. The third topic of research deals with the structural transitions of chromatin during programmed cell death and with the regulation of factors involved in apoptotic chromatin cleavage and histone modification.

Selected Recent Publications

Bäuerle M, Doenecke D, Albig W (2002) The requirement of H1 histones for a heterodimeric nuclear import receptor. J Biol Chem 277: 32480-32489

Schliephake T, Meinl A, Kratzmeier M, Doenecke D, Albig W (2004) The telomeric region is excluded from nucleosomal fragmentation during apoptosis, but the bulk nuclear chromatin is randomly degraded. Cell Death Differ 11: 693-703

Kahle J, Baake M, Doenecke D, Albig W (2005) Subunits of the heterotrimeric transcription factor NF-Y are imported into the nucleus by distinct pathways involving importin β and importin 13. Mol Cell Biol 25: 5339-5354

Goebel W, Obermeyer N, Bleicher N, Kratzmeier M, Eibl HJ, Doenecke D, Albig W (2007) Apoptotic DNA fragmentation is not related to the phosphorylation state of histone H1. Biol Chem 388: 197-206

Stoldt S, Wenzel D, Schulze E, Doenecke D, Happel N (2007) G1 phase dependent nucleolar accumulation of human histone H1x. Biol Cell 99: 541-552

Stefan Eimer

Group Leader Molecular Neurogenetics/ Neurodegeneration

- Ph.D. 2003 at the Gene Center of the Ludwig-Maximilian University (LMU) in Munich
- 2003 Postdoc at the Ecole Normale Superieure in Paris, France
- since Oct 2005 independent group leader of the Center for Molecular Physiology of the Brain (CMPB) at the European Neuroscience Institute (ENI) in Goettingen

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Major Research Interests

Neuotransmitter gated ion channels are involved in a large subset of neuronal events ranging from fast synaptic transmission to the modulation of neuronal circuits that lead to memory formation and cognition. En route to the cell surface these multimeric receptors have to undergo multiple assembly, quality control, and sorting steps to eventually reach the synapse.

Our group aims to understand the mechanisms and rules that control the trafficking and sorting of ligand gated ion channels within the secretory apparatus. In particular, we are focusing on the nicotinic acetylcholine receptor family of ligand gated ion channels, which have been implicated in numerous neurological and neurodegenerative diseases.

To find new molecules involved in these processes, we take advantage of the nematode *Caenorhabditis elegans* as a main model system, and use a combination of genetic, cell biological, and biochemical approaches as well as electro-physiology and electron-microscopy. As our main model system were are studying cholinergic neurotransmission at the neuro-muscular junction (NMJ) of *C. elegans*. Through genetic screens we have identified novel evolutionary conserved integral membrane proteins that regulate nAChR sorting at the Golgi-Endosomal interface. Further studies have implicated these molecules in the regulation and activation of small GTPases at Golgi complex. Based on these findings we have also started to study systematically how these GTPases are required for structure and function of the Golgi apparatus and how their activity affects the trafficking and neurotransmission at the NMJ of *C. elegans*.

Selected Recent Publications

Marza E, Long T, Saiardi A, Sumakovic M, Eimer S, Hall DH, Lesa GM (2007) Polyunsaturated fatty acids influence synaptojanin localization to regulate synaptic vesicle recycling. Mol Biol Cell, in press

Eimer S, Gottschalk A, Richmond JE, Hengartner M, Schafer W, Bessereau J-L (2007) Regulation of nicotinic receptor trafficking by the transmembrane Golgi protein UNC-50. EMBO J 26: 4313-23

Yamasaki A, Eimer S, Okochi M, Smialowska A, Kaether C, Baumeister R, Haass C, Steiner H (2006) The GxGD motif of presenilin contributes to catalytic function and substrate identification of gamma-secretase. J Neurosci 26: 3821-8

Gally C, Eimer S, Richmond JE, Bessereau J-L (2004) A transmembrane protein required for acetylcholine receptor clustering in *C. elegans*. Nature 431: 578-582

Eimer S, Lakowski B, Donhauser R, Baumeister R (2002) Loss of spr-5 bypasses the requirement for the presenilin sel-12 by stage-specific derepression of hop-1. EMBO J 21: 5787-5796

Wolfgang Engel

Professor of Human Genetics

- Dr. med., Universität Freiburg, 1967
- Physician, Hospital Schorndorf, 1966 - 1968
- Postdoc, Institute of Human Genetics and Anthropology, Universität Freiburg, 1968 - 1977
- Habilitation (Human Genetics), Universität Freiburg, 1974
- Professor of Human Genetics and Director of the Institute, Universität Göttingen, 1977

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Major Research Interests

Our research is focussed on the molecular analysis of normal human variability and genetic disturbances of development and differentiation.
Isolated genes are being analysed in detail with respect to their functional properties by animal models (transgenic and knock-out-mice). For suitable genetic diseases therapeutic strategies (substitution; gene therapy) are being developed and initial evaluation of such strategies is done in the mouse. - We are working on the genotype – phenotype correlations in neurological and cardiovascular diseases (e. g. Spastic paraplegia, Rett syndrome, mental retardation by subtelomeric microdeletions, molybdenum cofactor deficiency; cardiomypathies, Noonan syndrome) and several genetically determined malformation syndromes (e.g. Townes-Brocks syndrome, Okihiro syndrome, Morbus Osler).We are also engaged in the molecular and cellular basis of initiation events of cancer, specifically in prostate cancer, medulloblastoma and rhabdomyosarcoma. - One main interest in our institute is the analysis of structure, expression and function of genes involved in differentiation of male gametes. The knowledge of the function of those genes can help us to clarify the genetic causes of male infertility. We have isolated spermatogonial stem cells (SSCs) from adult mouse testis and demonstrated that these cells are as pluripotent as embryonic stem cells (ESCs). Our main interest is now to isolate and proliferate SSCs from adult human testis. These cells would be of great interest for regenerative medicine.

Selected Recent Publications

Nayerniaa K, Lee JH, Drusenheimer N, Nolte J, Wulf G, Dressel R, Gromoll J, Engel W (2006) Derivation of male germ cells from bone marrow stem cells. Laboratory Investigation 86: 654-663

Nayernia K, Nolte J, Michelmann HW, Lee JH, Rathsack K, Drusenheimer N, Dev A, Wulf G, Ehrmann IE, Elliott DJ, Okpanyi V, Zechner, Haaf T, MeinhardtA, Engel W (2006) *In vitro*-differentiated embryonic stem cells give rise to male gametes that can generate offspring mice. Developmental Cell 11: 125-132

Lee JH, Engel W, Nayernia K (2006) Stem cell protein Piwil2 modulates expression of murine spermatogonial stem cell expressed genes. Molecular Reproduction and Development 73: 173-179

Guan K, Nayernia K, Maier LS, Wagner S, Dressel R, Lee JH, Nolte J, Wolf, F, Li M, Engel W, Hasenfuß G (2006) Pluripotency of spermatogonial stem cells from adult mouse testis. Nature 440, 1199-1203

Lee HJ, Göring W, Ochs M, Mühlfeld C, Steding G, Paprotta I, Engel W, Adham IM (2004) Sox 15 is required for skeletal muscle regeneration. Molecular and Cellular Biology 19: 8428-8436

Nayernia K, Li, M, Jaroszynski L, Khusainow R, Wulf G, Schwandt I, Korbiowska M, Michelmann HW, Meinhardt A, Engel W (2004) Stem cells based therapeutical approach of male infertility by teratocarcinoma derived germ cells. Human Molecular Genetics 13: 1451-1460

Dirk Fasshauer

Independent Research Group Leader - Structural Biochemistry

- 1994 Doctoral degree (Dr. rer. nat.) University of Göttingen
- 1995-97 Postdoctoral fellow, Yale University
- since 1997 Postdoctoral fellow, Dept. for Neurobiology, Max Planck Institute for
- Biophysical Chemistry, Göttingen
- since 2002 Group leader within the Dept. for Neurobiology, Max Planck Institute for Biophysical Chemistry, Göttingen
- since 2006 Independent Research Group Leader, Structural Biochemistry, Max Planck Institute for Biophysical Chemistry, Göttingen

Major Research Interests

The mechanism by which eukaryotic cells transport material between intracellular organelles is of fundamental importance in cell biology. Transport is mediated by vesicles that bud from a donor organelle and afterwards fuse with a target organelle. Currently, it is becoming clear that the underlying molecular machineries involved in the principal aspects of vesicular trafficking are highly conserved among all eukaryotes. Key players during the final step in vesicle trafficking, the fusion of a vesicle with its acceptor membrane, are the so-called SNARE proteins. SNARE proteins are thought to assemble into a tight complex between the fusing membranes, pulling them together (the 'zipper' model). To come to a better understanding of the molecular events during vesicular fusion, we focus on a detailed structural, kinetic, thermodynamic, and phylogenetic characterization of the underlying protein-protein interactions. In particular, we want to investigate how SNARE assembly takes place, how this process is controlled and catalyzed by other factors. Next to standard biochemical techniques, we employ spectroscopic (Circular Dichroism and Fluorescence Spectroscopy) and calorimetric (Isothermal Titration Calorimetry) methods.

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Selected Recent Publications

Kloepper TH, Kienle CN, Fasshauer D (2007) An elaborate classification of SNARE proteins sheds light on the conservation of the eukaryotic endomembrane system. Mol Biol Cell 18: 3463-71

Pobbati A, Stein A, Fasshauer D (2006) N- to C-terminal SNARE complex assembly promotes rapid membrane fusion. Science 313: 673-6

Soerensen JB, Wiederhold K, Müller EM, Milosevic I, Nagy G, de Groot BL, Grubmüller H, Fasshauer D (2006) Sequential N- to C-terminal 'zipping-up' of the SNARE complex drives priming and fusion of secretory vesicles. EMBO J 25: 955-66

Pobbati A, Razeto A, Böddener M, Becker S, Fasshauer D (2004) A structural basis for the inhibitory role of tomosyn in exocytosis. J Biol Chem 279: 47192-200

Fasshauer D, Antonin W, Subramaniam V, Jahn R (2002) SNARE assembly and disassembly exhibit a pronounced hysteresis. Nat Struct Biol 9: 144-151

Ivo Feußner

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- Diploma (Chemistry), Philipps-University, Marburg (Germany), 1990
- Dr. rer. nat., Philipps-University, Marburg (Germany), 1993
- Leader of an independent research group at the Institute for Plant Biochemistry (IPB), Halle/Saale (Germany), 1997 - 1999
- Habilitation (Biochemistry), Martin-Luther-University, Halle/Saale (Germany), 2000
- Leader of an independent research group at Institute for Plant Genetics and Crop Plant Research (IPK), Gatersleben (Germany), 2000 - 2002
- Since 2002 Professor for Biochemistry, Georg-August-University, Göttingen (Germany)
- Award: Habilitation-Prize of the Ernst Schering Research Foundation (2001)

Major Research Interests

Plant Metabolic Pathways: Our laboratory is currently studying the primary metabolism of plants with main focus on the metabolism of lipids. For this purpose, different approaches ranging from analytical
chemistry to biochemistry and molecular biology were used.
Plant Lipid Metabolism: We are interested in physiological functions of specific lipoxygenases, i.e. their involvement in the degradation of storage lipids during germination and in the destruction of organellar membranes during stress. Another research topic is the analysis of their catalytic mechanism. In addition, lipid peroxidation reactions were analysed in general by metabolomic approaches and by studying the biosynthesis of aldehydes (fruit aromas), hydroxy fatty acids and divinyl ether fatty acids (plant defence). Moreover, enzymes which introduce new functionalities (i.e. conjugated double bonds) in the fatty acid backbone were isolated and characterized in order to obtain new seed oils for biotechnological and medical purposes. In relation to that we are manipulating the primary metabolism and organelle development of seeds in order to increase the oil content of seeds.
Metabolic transport processes: Another research topic is the analysis of the mechanism and regulation of transport processes across the peroxisomal membrane. The biochemistry of phosphoinositides and the transfer of enzymes facilitate the metabolic pathways for lcPUFAs from donor organisms into plants.

Selected Recent Publications

Hoffmann M, Hornung E, Busch S, Kassner N, Ternes P, Braus GH, Feussner I (2007) A Small Membrane-peripheral Region Close to the Active Center Determines Regioselectivity of Membrane-bound Fatty Acid Desaturases from *Aspergillus nidulans*. J Biol Chem 282: 26666-26674

Stumpe M, Göbel C, Demchenko K, Hoffmann M, Klösgen RB, Pawlowski K, Feussner I (2006) Identification of an allene oxide synthase (CYP74C) that leads to formation of α-ketols from 9-hydroperoxides of linoleic and linolenic acid in below ground organs of potato. Plant J 47: 883-896

Ochsenbein C, Przybyla D, Danon A, Landgraf F, Göbel C, Imboden A, Feussner I, Apel K (2006) The role of EDS1 (Enhanced Disease Susceptibility) during singlet oxygen-mediated stress responses of Arabidopsis. Plant J 47: 445-456

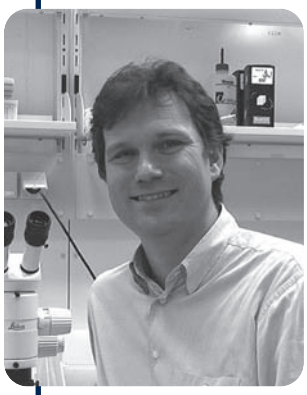
Liavonchanka A, Hornung E, Feussner I, Rudolph MG (2006) Structure and mechanism of the *Propionibacterium acnes* polyunsaturated fatty acid isomerase. Proc Natl Acad Sci USA 103: 2576-2581

Senger T, Wichard T, Kunze S, Göbel C, Lerchl J, Pohnert G, Feussner I (2005) A multifunctional lipoxygenase with fatty acid hydroperoxide cleaving activity from the moss *Physcomitrella patens*. J Biol Chem 280: 7588-7596

Ralf Ficner

Professor of Structural Biology

- Dr. rer. nat. (1992) and Postdoc (1993), Max Planck Institute for Biochemistry, Martinsried
- Postdoctoral fellow, EMBL Heidelberg, 1994 - 1996
- Junior Group Leader, University of Marburg, 1997 - 2000
- Appointed 2001 as Head of the Department of Molecular Structural Biology at the University of Göttingen



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Major Research Interests

In order to understand the relationship between the three-dimensional structure and the cellular function of biological macromolecules we determine the structures of proteins and protein-RNA complexes by means of X-ray crystallography. Our current projects concern proteins involved in the splicing and modification of RNA and, as well, proteins required for the nucleocytoplasmic transport, and enzymes of the polysialic acid metabolism.

Selected Recent Publications

Strasser A, Dickmanns A, Lührmann R, Ficner R (2005) Structural basis for m3G-cap-mediated nuclear import of spliceosomal UsnRNPs by snurportin1. EMBO J 24: 2235-43

Dierks T, Dickmanns A, Preusser-Kunze A, Schmidt B, Mariappan M, von Figura K, Ficner R, Rudolph MG (2005) Molecular basis for multiple sulfatase deficiency and catalytic mechanism for formylglycine generation of the human formylglycine generating enzyme. Cell 121 541-552

Stummeyer K, Dickmanns A, Mühlenhoff M, Gerardy-Schahn R, Ficner R (2005) Crystal structure of endosialidase NF - the polysialic acid degrading tailspike of bacteriophage K1F. Nature Struct Mol Biol 12: 90-96

Rudolph MG, Kraus I, Dickmanns A, Eickmann M, Garten W, Ficner R (2003) Crystal structure of the Borna Disease Virus nucleoprotein. Structure 11, 1219-1226

Vidovic I, Nottrott S, Hartmuth K, Lührmann R, Ficner R (2000) Crystal structure of the spliceosomal 15.5kD protein bound to a U4 snRNA fragment. Mol Cell 6: 1331-1342

Wolfgang Fischle

Group Leader at the MPI for Biophysical Chemistry

- Dr. rer. nat. (PhD), University of Tübingen, Germany, 2001
- Graduate Research Fellow, The J. David Gladstone Institute (UCSF), San Francisco, CA, USA, 1997 - 2001
- Postdoctoral Fellow, The Rockefeller University, New York, NY, USA, 2001 - 2005
- Damon Runyon Cancer Research Fellow, 2002 - 2005
- Independent Group Leader, Max Planck Institute for Biophysical Chemistry, Göttingen, Germany, since 2005

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Major Research Interests

Chromatin is the physiological template of genetic information controlling the capacity of a cell's genome to store, release, and inherit biological information. The fundamental unit of chromatin is the nucleosome: a stretch of DNA wrapped around a core of histone proteins (H2A, H2B, H3 and H4). Post-translational modifications of histones have emerged as key for regulating chromatin structure and are thought to crucially control chromatin dynamics and genome activity. Whereas more and more histone modification marks are being identified that alone or in combination could mediate distinct biological conditions of a cell and while correlative studies have begun to establish unambiguous links between several states of chromatin, various histone modifications, and diverse biological processes, our knowledge of how certain histone modifications exert their biological effects on a molecular/biochemical level is still very li-mited.

Due to their long-term stability, histone lysine methyl-marks are of particular interest to us, since they might be involved in establishing and maintaining durable and inheritable gene expression profiles (so called 'epi-genetic' regulation). Current projects include the study of Polycomb, HP1, and MBT proteins that bind to and act as effectors of distinct histone lysine methyl-marks. We are especially interested in the interplay of these factors and their cognate histone marks in regulating chromatin organization and dynamics. Furthermore, we are trying to identify and characterize novel binding proteins of various other histone modifications.

The long-term goal of our research is to gain mechanistic insight(s) into the signaling mechanisms and biological role of single but also combinations of histone modification marks and to understand how certain states of chromatin regulate the functionality of a cells genome. To this end, we aim to reconstitute chromatin-signaling pathways in recombinant and cell free systems and study their epi-genetic regulatory circuits in various biological model systems (i.e. *Xenopus laevis*, mice, tissue culture).

Selected Recent Publications

Fischle W, Tseng BS, Dormann H, Ueberheide BM, Garcia BA, Shabanowitz J, Hunt DF, Funabiki H, Allis CD (2005) Regulation of HP1-chromatin binding by histone H3 methylation and phosphorylation. Nature 438: 1116-22

Yamada T, Fischle W, Allis CD, Grewal SIS (2005) The nucleation and maintenance of heterochromatin by a histone deacetylase in fission yeast. Molecular Cell 20: 1-13

Fischle W, Wang Y, Allis CD (2003) Binary switches and modification cassettes in histone biology and beyond. Nature 425: 475-479

Fischle W, Wang Y, Jacobs SA, Kim Y, Allis CD, Khorasanizadeh S (2003) Molecular basis for the discrimination of repressive methyl-lysine marks in histone H3 by Polycomb and HP1 chromodomains. Genes & Development 17: 687-698

Fischle W, Dequiedt F, Hendzel M, Guenther MG, Lazar MA, Voelter W, Verdin E (2002) Enzymatic activity associated with class II HDACs is dependent on a multiprotein complex containing HDAC3 and SMRT/N-CoR. Molecular Cell 9: 45-57

Chen Lf, Fischle W, Verdin E, Greene WC (2001) Duration of nuclear NF-kappaB action regulated by reversible acetylation. Science 293: 1653-1657

Christiane Gatz

Professor of Plant Molecular Biology

- Dr. rer.nat. (1985) at the Institute for Biochemistry, Technical University Darmstadt
- Postdoctoral fellow at the University of Wisconsin, Madison, USA (1985 - 1987)
- Habilitation in Molecular Genetics at the Freie Universität Berlin in 1992
- Professor at the University of Bielefeld (1993 – 1995)
- Alfried Krupp von Bohlen und Halbach-Award for young university professors (1994)
- Professor at the University of Göttingen since 1996

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Major Research Interests

Our laboratory is interested in signal transduction pathways that regulate plant defense. Our research is focused on two main topics: 1) The molecular genetic basis of defense responses against bacteria and fungi 2) The molecular mechanism of volatile emission upon herbivore feeding.

Defense responses against bacteria and fungi.
Plants have developed multiple layers of defense responses against pathogens. In general, infection of *Arabidopsis* plants with biotrophic pathogens (pathogens that exploit resources of living cells) leads to the activation of salicylic acid (SA)-mediated defense responses, whereas infection with necrotrophic pathogens (pathogens that kill cells to obtain access to nutrients) elicits (jasmonic acid/ethylene) JA/ET-dependent responses (Glazebrook, 2005). Transcription factors that have been identified as essential regulators for both responses are proteins of the TGA family. These proteins reside in the cell in an inactive state before pathogen infection. We are interested in the SA- and JA/ET-mediated mechanisms that activate the function of TGA factors. More recent results indicate that redox signaling plays a major role in these processes. Moreover, we are interested in the cross-talk between both pathways.

Volatile emission upon herbivore feeding
When insects feed on plants, plants emit volatiles to attract the enemies of the insect. We are interested in the transcriptional regulation of biosynthetic genes leading to the synthesis of volatiles. Again, the JA/ET pathway plays a major role in this response.

Selected Recent Publications

Ndamukong I, Al Abdallat A, Thurow C, Fode B, Zander M, Weigel R, Gatz C (2007) SA-inducible *Arabidopsis* glutaredoxin interacts with TGA factors and suppresses JA-responsive PDF1.2 transcription. Plant J 50: 128-139

Butterbrodt T, Thurow C, Gatz C (2006) Chromatin immunoprecipitation analysis of the tobacco PR-1a- and the truncated CaMV 35S promoter reveals differences in salicylic acid-dependent TGA factor binding and histone acetylation. Plant Mol Biol 61: 665-674

Weigel RR, Pfitzner UM, Gatz C (2005) Interaction of NIMIN1 with NPR1 modulates PR gene expression in *Arabidopsis*. Plant Cell 17: 1279-1291

Thurow C, Schiermeyer A, Krawczyk S, Butterbrodt T, Nickolov K, Gatz C (2005) Tobacco bZIP transcription factor TGA2.2 and related factor TGA2.1 have distinct roles in plant defense responses and plant development. Plant J 44: 100-113

Dirk Görlich

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Professor, Director at the Max Planck Institute for Biophysical Chemistry

- 1989 Diploma (Biochemistry), Martin-Luther-Universität in Halle
- 1990 - 1993 Graduate studies (Laboratory of T.A. Rapoport, Berlin)
- 1993 Dr. rer. nat. (Biochemistry) Humboldt-Universität Berlin
- 1993 Postdoc (Laboratory of T.A. Rapoport, Berlin)
- 1993 - 1995 Postdoc (Laboratory of R.A. Laskey, Cambridge, England)
- 1996 - 2007 Research group leader at the ZMBH Heidelberg
- 2001 - 2007 Professor for Molecular Biolology (Universität Heidelberg)
- 2001 - 2006 Deputy Director of the ZMBH
- 2005 - Director, Dept. Cellular Logistics, MPI for Biophysical Chemistry, Göttingen

Major Research Interests

- Nuclear transport
- Importins and Exportins
- RanGTPase-system
- Nuclear pore complexes (NPCs), NPC-assembly, Mechanism of NPC-passage
- Hydrogels
- Integral membrane proteins, Translation
- Systems biology
- Spermiogenesis

Selected Recent Publications

Frey S, Görlich D (2007) A saturated FG-repeat hydrogel can reproduce the permeability-properties of nuclear pore complexes. Cell 130: 512-523

Bohnsack MT, Stüven T, Kuhn C, Cordes VC, Görlich D (2006) A selective block of nuclear actin export stabilizes the giant nuclei of *Xenopus* oocytes. Nat Cell Biol 8: 257-263

Frey S, Richter RP, Görlich D (2006) FG-rich repeats of nuclear pore proteins form a three-dimensional meshwork with hydrogel-like properties. Science 314: 815-817

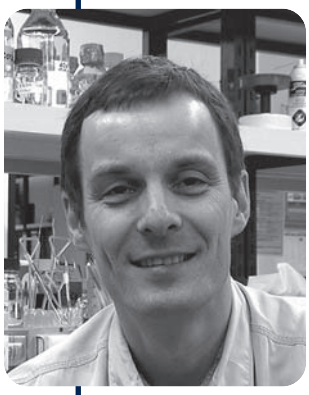
Mingot JM, Bohnsack MT, Jäkle U, Görlich D (2004) Exportin 7 defines a novel general nuclear export pathway. EMBO J 23: 3227-3236

Görlich D, Seewald MJ, Ribbeck K (2003) Characterization of Ran-driven cargo transport and the RanGTPase system by kinetic measurements and computer simulation. EMBO J 22: 1088-1100

Jäkel S, Mingot JM, Schwarzmaier P, Hartmann E, Görlich D (2002) Importins fulfil a dual function as nuclear import receptors and cytoplasmic chaperones for exposed basic domains. EMBO J 21: 377-386

Ribbeck K, Görlich D (2001) Kinetic analysis of translocation through nuclear pore complexes. EMBO J 20: 1320-1330

Christian Griesinger



Professor, Director at the Max Planck Institute for Biophysical Chemistry, Göttingen

- Dr. phil. nat. University of Frankfurt (1986, Prof. Dr. H. Kessler)
- Postdoctoral Fellow at Lab. for Physical Chemistry, ETH Zürich (1986 - 1989, Prof. Dr. R. R. Ernst)
- Full Professor for Organic Chemistry at the University of Frankfurt (1990 - 2000)
- Appointed as Director at the Max Planck Institute for Biophysical Chemistry (1999)

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030

Major Research Interests

Our group focuses on the structure and dynamics of biomolecules and their complexes. We use to this end nuclear magnetic resonance (NMR) spectroscopy as well as X-ray crystallography. We apply solution state and solid state NMR spectroscopy to biomolecules and their complexes in their physiological environment, be it water for cytosolic proteins or lipids for membrane proteins. The methods developments are pursuing the following goals:
Relaxation compensated methods to increase the molecular weight of biomolecules that are amenable for NMR investigations. This is tackled by new NMR pulse techniques, novel schemes for labelling the biomolecules with isotopes and the usage of optimized expression schemes. New NMR derived parameters that allow to define biomolecular structure and dynamics better are derived and applied e.g. to DNA binding proteins, spliceosomal RNA, a bacterial sensor and proteins involved in signal transduction and apoptosis and for the investigation of enzyme mechanisms.
Development of NMR methods to assign and determine the structure of isotopically labeled membrane-proteins and peptides with solid state NMR spectroscopy on oriented samples or using magic angle sample spinning on several systems including a 150 kD membrane-Protein and the complex of a peptide and a G-protein coupled receptor.

Selected Recent Publications

Reif B, Hennig M, Griesinger C (1997) Direct measurement of angles between bond vectors in high resolution NMR. Science 276: 1230-1233

Marino JP, Schwalbe H, Griesinger C (1999) J-coupling restraints for structural refinements of RNA. Acc Chem Res 32: 614-632

Bartoschek S, Johannson M, Geierstanger BH, Okun JG, Lancaster CRD, Humpfer E, Yu L, Yu CA, Griesinger C, Brandt U (2001) Three molecules of ubiquinone bind specifically to mitochondrial cytochrome bc1 complex. J Biol Chem 276: 35231-35234

Peti W, Meiler J, Brüschweiler R, Griesinger C (2002) Model free analysis of protein backbone motion from residual dipolar couplings. J Am Chem Soc 124: 5822-5833

Carlomagno T, Blommers MJJ, Meiler J, Jahnke W, Schupp T, Petersen F, Schinzer D, Altmann K-H, Griesinger C (2003) The high-resolution solution structure of epothilone a bound to tubulin: An understanding of the structure-activity relationships for a powerful class of antitumor agents. Angew Chem 115: 2615-2619, Angew Chem Int Ed 42: 2511-2515

Pappalardo L, Janausch IG, Vijayan V, Zientz E, Junker J, Peti W, Zweckstetter M, Unden G, Griesinger C (2003) The NMR structure of the sensory domain of the membrancous two-component fumarate sensor (histidine protein kinase) DcuS of *Escherichia coli*". J Biol Chem 278: 39185 - 39188

Uwe Groß

Professor of Medical Microbiology

- M.D., University of Hamburg 1987
- Postdoctoral fellow, UC Los Angeles, California, 1987 - 1989
- Professor of Medical Parasitology, University of Würzburg 1998/1999
- Appointed 1999 as head of the Department of Medical Microbiology, University of Göttingen

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Major Research Interests

The protozoan parasite *Toxoplasma gondii* usually causes asymptomatic infections in immunocompetent adults leading to lifelong persistence especially in the brain and in muscle tissue. Life-threatening reactivation of such infection might occur in immunocompromised individuals (i. e. patients suffering from AIDS). This parasite serves as a model organism for studying evasion mechanisms of intracellular pathogens.
We are interested in the cross-talk between the parasite and its host cell on a molecular level. We could demonstrate that the parasite (i) modulates the host cell capacity for MHC-restricted antigen presentation and (ii) inhibits apoptosis of the infected cell. Both mechanisms allow intracellular persistence. Vice versa, the host´s immune response determines the fate of the parasite by direct interference with differentiation processes of *Toxoplasma gondii*. The precise molecular events for these strategies of intense interplay between both partners are currently under our investigation.
Recently, we also started to investigate host-pathogen interactions of *Campylobacter jejuni*. This pathogen is the most prominent bacterial species that causes diarrhoea followed eventually by the development of neurological complications. Currently, we are focusing on how the pathogen is inducing host-cell apoptosis, thereby promoting disease of epithelial-layered tissues, such as the intestine.
In addition, we are appointed the National Reference Center for Systemic Mycoses. In this respect, we are inverstigating fungal factors and mechanisms that are involved in pathogenesis of mycoses; i.e. cell wall structure and differentiation processes.

Selected Recent Publications

Holpert M, Groß U, Bohne W (2006) Disruption of the bradyzoite-specific P-type (H+)-ATPase /PMA1/ in *Toxoplasma gondii* leads to decreased bradyzoite differentiation after stress stimuli but does not interfere with mature tissue cyst formation. Mol Biochem Parasitol 146:129-33

Fasshauer V, Groß U, Bohne W (2005) The parasitophorous vacuole membrane of *Encepalitozoon cuniculi* lacks host cell membrane proteins immediately after invasion. Eukaryot Cell 4: 221-224

Lüder CGK, Groß U (2005) Apoptosis and its modulation during infection with Toxoplasma gondtii: molecular mechanisms and role in pathogenesis. Curr Topics Microbiol Immunol 289: 219-238

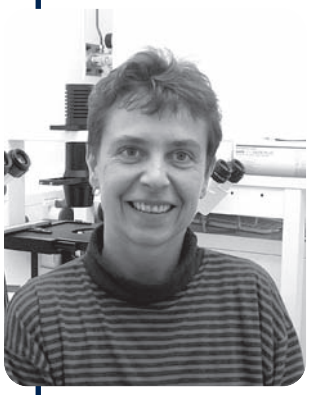
Weig M, Jäntsch L, Groß U, de Koster CG, Klis FM, de Groot PWJ (2004) Systematic identification in silico of covalently bound cell wall proteins and analysis of protein-polysaccharide linkages of the human pathogen *Candida glabrata*. Microbiology 150: 3129-3144

Lüder CGK, Lang C, Giraldo-Velasquez M, Algner M, Gerdes J, Groß U (2003) *Toxoplasma gondii* inhibits MHC class II expression in neural antigen-presenting cells by down-regulating the class II transactivator CIITA. J Neuroimmunol 134:12-24

Heidi Hahn

Professor of Molecular Developmental Genetics

- Dr. med., University of Würzburg, 1992
- Postdoctoral Fellow, National Institutes of Health, Bethesda, Maryland, USA (1993 - 1998)
- Junior Group Leader (BioFuture), Technical University of Munich (1999 - 2000)
- Professor of Molecular Developmental Genetics, University of Göttingen since 2001



Major Research Interests

Hedgehog (Hh) signaling molecules play a key role in the patterning of numerous tissues during development. Hh signaling is initiated by binding of Hh to its receptor Patched (Ptch). This binding suspends the inhibitory action of Ptch on its signaling partner Smo. Smo is activated and the signaling pathway is turned on. The pathway can also be activated by mutational inactivation of Ptch or by activating mutations in either Hh or Smo and we were able to show that this pathological activation results in developmental defects and in tumor formation in humans and mice.

The goal of our group is to characterize the role of the Hh signaling pathway in the diseased state by identification of its cellular targets and by characterization of its interaction with other signaling pathways. This is achieved by the application of modern genetic techniques (e.g. microarray analysis) to human and murine tumors and cell lines with mutations in one or more components of the pathway. This approach should help to develop molecular diagnostics for Hh-related malignancies as well as to identify targets for therapeutic interventions.

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Selected Recent Publications

Chang-Claude J, Dunning A, Schnitzbauer U, Galmbacher P, Tee L, Wjst M, Chalmers J, Zemzoum I, Harbeck N, Pharoah PDP, Hahn H (2003) The Patched Polymorphism Pro1315Leu (C3944T) may modulate the association between use of oral contraceptives and breast cancer risk. Int J Cancer 103(6): 779-83

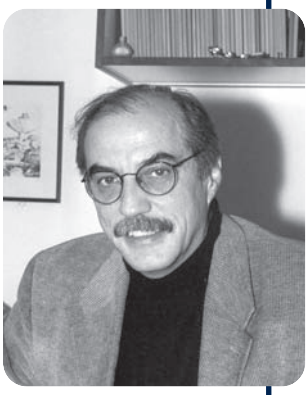
Kappler R, Bauer R, Calzada-Wack J, Rosemann M, Hemmerlein B, Hahn H (2004) Profiling the molecular difference between Patched1- and p53-dependent rhabdomyosarcoma. Oncogene 23(54): 8785-95

Koleva M, Kappler R, Vogler M, Herwig A, Fulda S, Hahn H (2005) Pleiotropic effects of sonic hedgehog on muscle satellite cells. Cell Mol Life Sci 62(16): 1863-1870

Pazzaglia S, Tanori M, Mancuso M, Rebessi S, Leonardi S, Di Majo V, Covelli V, Atkinson MJ, Hahn H, Saran S (2006) Linking DNA damage to medulloblastoma tumorigenesis in Patched heterozygous mice. Oncogene 25(8): 1165-73

Uhmann A, Dittmann K, Nitzki F, Dressel R, Koleva M, Frommhold A, Zibat A, Binder C, Adham I, Nitsche M, Heller T, Armstrong V, Schulz-Schaeffer W, Wienands J, Hahn H (2007) The Hedgehog receptor Patched controls lymphoid lineage commitment. Blood 110(6): 1814-23

Herbert Jäckle



Professor, Director at the Max Planck Institute for Biophysical Chemistry

- Faculty member at the EMBL, Heidelberg (1980 - 1982)
- Head of the group (associate professor), Max Planck Institute for Developmental Biology, Tübingen (1982 - 1988)
- Professor and Chairman, Dept. of Genetics and Microbiology, Univ. of Munich (1988 - 1991)

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Major Research Interests

How is the embryo generated from a single cell, the egg? We address this question by using the *Drosophila* embryo as an experimental system, applying the combined tools of classical embryology, genetics, molecular biology and biochemistry. We have focussed our efforts to isolate and characterize the factors underlying early pattern formation along the anterior-posterior axis of the embryo. We sought to unravel their mode of action und the molecular mechanism in which they function.

Many of the factors required to establish the basic body plan are also necesssary for organ formation, a process which involves local inductive interactions between groups of cells and/or epithelial cell layers. We have started to identify the genetic components and regulatory circuitries involved in organogenesis as well as in neural conductivity and function. We also use the fly to identify the components of novel biochemical pathways and cellular key components that control and maintain homeostasis and energy balance, and we initiated a gene discovery program to systematically characterize the function of genes on the *Drosophila* X-chromosome.

Selected Recent Publications

Peter A, Schöttler P, Werner M, Beinert N, Dowe G, Burkert P, Mourkioti F, Dentzer L, He Y, Deak P, Benos PV, Gatt MK, Murphy L, Harris D, Barrell B, Ferraz C, Vidal S, Brun C, Demaille J, Cadieu E, Dreano S, Gloux S, Lelaure V, Mottier S, Galibert F, Borkova D, Miñana B, Kafatos FC, Bolshakov S, Sidén-Kiamos I, Papagiannakis G, Spanos L, Louis C, Madueño E, de Pablos B, Modolell J, Bucheton A, Callister D, Campbell L, Henderson NS, McMillan PJ, Salles C, Tait E, Valenti P, Saunders RDC, Billaud A, Pachter L, Klapper R, Janning W, Glover DM, Ashburner M, Bellen HJ, Jäckle H, Schäfer U (2002) Mapping and identification of essential gene functions on the X chromosome of *Drosophila*. EMBO Reports 3: 34-38

Carrera P, Moshkin YM, Grönke S, Silljé HHW, Nigg EA, Jäckle H, Karch F (2003) Tousled-like kinase functions with the chromatin assembly pathway regulating nuclear divisions. Genes Dev 17: 2578-2590

Steigemann P, Molitor A, Fellert S, Jäckle H, Vorbrüggen G (2004) Heparan sulfate proteoglycan Syndecan promotes axonal and myotube guidance by Slit/Robo signaling. Curr Biol 14: 225-230

Grönke S, Mildner A, Fellert S, Tennagels N, Petry S, Müller G, Jäckle H, Kühnlein RP (2005) Brummer lipase is an evolutionary conserved fat storage regulator in *Drosophila*. Cell Metabolism 1: 323-330

Reinhard Jahn

Professor, Director at the Max Planck Institute for Biophysical Chemistry

- Dr. rer. nat. 1981, University of Göttingen
- Assistant Professor, The Rockefeller University, New York (USA) 1985
- Junior Group leader, Max Planck Institute for Psychiatry, Martinsried, 1986
- Associate Professor of Pharmacology and Cell Biology, Yale University, and Investigator, Howard Hughes Medical Institute, New Haven (USA) 1991
- Professor of Pharmacology and Cell Biology, Yale University, New Haven, 1995
- Director, Max Planck Institute for Biophysical Chemistry, Göttingen, 1997

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Major Research Interests

Our group is interested in the mechanisms of membrane fusion, with the main emphasis on regulated exocytosis in neurons. Since recent years it is known that intracellular membrane fusion events are mediated by a set of conserved membrane proteins, termed SNAREs. For fusion to occur, complementary sets of SNAREs need to be present on both of the fusing membranes. The neuronal SNAREs are among the best characterized. They are the targets of the toxins responsible for botulism and tetanus. To understand how these proteins make membranes fuse, we studied their properties in detail using biochemical and biophysical approaches. We found that they assemble into a tight complex which ties the membrane closely together and thus probably initiates bilayer mixing.

In our current approaches, we study membrane fusion at the level of isolated proteins as well as in semi-intact and intact cells. Thus, we are investigating conformational changes of the SNARE proteins before and during fusion. Furthermore, we use reconstitution of membrane fusion in cell-free assays and in proteoliposomes. Other projects of the group include the study of neurotransmitter uptake by synaptic vesicles and the function of Rab-GTPases in neuronal exocytosis.

Selected Recent Publications

Jahn R, Lang T, Südhof TC (2003) Membrane fusion. Cell 112: 519-533

Schuette CG, Hatsuzawa K, Margittai M, Stein A, Riedel D, Küster P, König M, Seidel CAM, Jahn R (2004) Determinants of liposome fusion mediated by synaptic SNARE proteins. Proc Natl Acad Sci 101: 2858-2863

Graf C, Riedel D, Schmitt HD, Jahn R (2005) Identification of functionally interacting SNAREs using complementary substitutions in the conserved '0' layer. Mol Biol Cell 16: 2263-2274

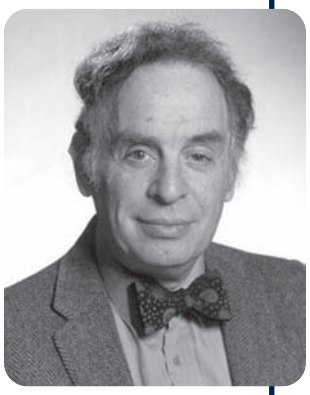
Willig KI, Rizzoli SO, Westphal V, Jahn R, Hell S (2006) STED-microscopy reveals that the synaptic vesicle protein synaptotagmin remains clustered after exocytosis. Nature 440: 935-939

Jahn R, Scheller RH (2006) SNAREs – engines for membrane fusion. Nature Reviews Mol Cell Biol 7: 631-643

Takamori S, Holt M, Stenius K, Lemke EA, Grønborg M, Riedel D, Urlaub H, Schenck S, Brügger B, Ringler P, Müller SA, Rammner B, Gräter F, Hub JS, De Groot BL, Mieskes G, Moriyama Y, Klingauf J, Grubmüller H, Heuser J, Wieland F, Jahn R (2006) Molecular anatomy of a trafficking organelle. Cell 127: 831-846

Zwilling D, Cypionka A, Pohl W, Fasshauer D, Walla PJ, Wahl MC, Jahn R (2007) Early endosomal SNAREs form a structurally conserved SNARE complex and fuse liposomes with multiple topologies. EMBO J 26: 9-18

Thomas M. Jovin



Chairman, Department of Molecular Biology and Director at the Max Planck Institute for Biophysical Chemistry

- B.S. California Institute of Technology, Pasadena, CA 1960
- M.D. Johns Hopkins Medical School, Baltimore, MD 1964
- Scientific Member, Max Planck Society 1969

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Current Research Interests

Structural studies of nucleic acids; complexes with proteins and ligands
Exotic helical structures: parallel-stranded DNA; triple helices; Z-DNA. Protein–DNA interactions: p53, α-synuclein, snRNPs.

Signal transduction of eukaryotic cells
Receptor tyrosine kinase activation, transport, and internalization; downstream signaling (MAPK cascade); and mechanism of antibody-based tumor therapy. Further development of Fluorescence Resonance Energy Transfer (FRET) as a probe of protein-protein interactions in the cellular application of quantitative microscopy. Quantum dot ligands and functional expression probes for proteins and nucleic acids in the microscopy of live cells.

Optical and scanning-probe microscopy
Development and application of novel microscopes for cellular and molecular studies: temparature-controlled atomic force (AFM), fluorescence lifetime (FLIM), fluorescence correlation (FCM), programmable optical sectioning (PAM), and single molecule dynamics.

Structure and function of α-synuclein (protein involved in Parkinson's disease)
Biochemical, biophysical, spectroscopic, and cell biological studies: intrinsic structure, ligand binding, and mechanism of aggregation

Selected Recent Publications

Cojocaru V, Nottrott S, Klement R, Jovin TM (2005) The snRNP 15.5K protein folds its cognate K-Turn RNA. A combined theoretical and biochemical study. RNA 11: 197-209

Hanley QS, Lidke KA, Heintzmann R, Arndt-Jovin DJ, Jovin TM (2005) Fluorescence lifetime imaging in an optically sectioned Programmable Array Microscope (PAM). Cytometry 67A: 112-118

Jares-Erijman EA, Jovin TM (2006) Imaging molecular interactions in living cells by FRET microscopy. Curr Opin Chem Biol 10: 1-8

Lidke DS, Lidke KA, Rieger B, Jovin TM, Arndt-Jovin DJ (2005) Reaching out for signals: filopodia sense EGF and respond by directed retrograde transport of activated receptors. J Cell Biol 170: 619-626

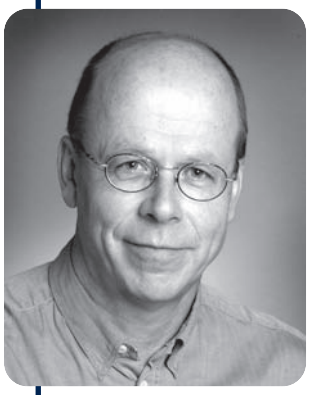
Nagy P, Friedländer E, Tanner M, Kapanen AI, Carraway KL, Isola J, Jovin TM (2005) Decreased accessibility and lack of activation of erbB2 in JIMT-1, a Herceptin-resistant, MUC-4-expressing breast cancer cell line. Cancer Res 65: 473-482

Pelah A, Ludueña SJ, Jares-Erijman EA, Szleifer I, Pietrasanta LI, Jovin TM (2006) Nanoscale memory provided by thermoreversible stochastically structured polymer aggregates on mica. Langmuir 22: doi 10.1021/la053431

Michael Kessel

Professor of Molecular Biology

- Until 1981 Biochemical Institute, Kiel University
- 1981 - 1983 National Cancer Institute, NIH, Bethesda,USA
- 1983 - 1986 Center for Molecular Biology (ZMBH), Heidelberg University
- Since 1987 Max Planck Institute for Biophysical Chemistry, Göttingen



Major Research Interests

The group studies patterning processes in chick and mouse embryos. We apply biochemical, genetic and embryological techniques, including expression analysis, transplantation in embryo culture, in vivo gene transfer by electroporation, and gene knock-out technology.

We identified the Geminin protein as a mediator between cell cycle progression and the control of axial specification. Geminin regulates homeodomain proteins of the Hox family both on a transcriptional and a chromatin level. We are currently studying a conditional mouse knock-out model.

We further analyze the homeobox gene Hesx1 and its role during the development of the pituitary. Hesx1 protein interacts with Mad2l2, a regulator of the APC/C complex, and a subunit of translesion DNA polymerase zeta. We study the involvement of Mad2l2-Hesx1 in the progression of the cell cycle in conditional knock-out mice.

Our goal is an understanding of the coordination between proliferation and pattern formation.

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Selected Recent Publications

Luo L, Kessel M (2004) Geminin coordinates cell cycle and developmental control. Cell Cycle 3: 711-714

Luo L, Yang X, Takihara Y, Knoetgen H, Kessel M (2004) The cell-cycle regulator geminin inhibits Hox function through direct and polycomb-mediated interactions. Nature 427: 749-53

Spieler D, Baumer N, Stebler J, Koprunner M, Reichman-Fried M, Teichmann U, Raz E, Kessel M, Wittler L (2004) Involvement of Pax6 and Otx2 in the forebrain-specific regulation of the vertebrate homeobox gene ANF/Hesx1. Developmental Biology 269: 567-79

Wittler L, Kessel M (2004) The acquisition of neural fate in the chick. Mechanisms of Development 121: 1031-42

Pitulescu M, Kessel M, Luo L (2005) The regulation of embryonic patterning and DNA replication by Geminin. Cellular and. Molecular. Life Science 62: 1425-1433

Luo L, Uerlings Y, Happel N, Asli NS, Knoetgen H Kessel M (2007) Regulation of geminin functions by cell cycle dependent nuclear-cytoplasmic shuttling. Molecular and Cellular Biology 27: 4737-4744

Dieter Klopfenstein

Junior Group Leader at the Centre for Molecular Physiology of the Brain, University of Göttingen

- Dr. phil. nat. (Ph.D.) University of Basel, 1999
- Postdoctoral fellow at the University of California San Francisco, 1999 - 2003
- Since 2003 head of an independent Junior Research Group

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Major Research Interests

The long-range transport of membrane organelles in neurons depends primarily upon microtubules and motor proteins that move unidirectionally along these tracks. One type of microtubule-based motor proteins powering membrane transport is the kinesin superfamily. We are interested in how these motors achieve specificity in cargo binding, elicit membrane transport, and the regulation of transport activity. One example of a kinesin motor is UNC-104/KIF1A, which specifically transports presynaptic vesicle to the synaptic terminal and binds with its tail domain directly to membrane lipids in vitro. This unique cargo-interaction mechanism help us to understand how lipids and their membrane environment contribute to cargo transport, how motor-lipid interaction could be regulating transport, and how accessory proteins contribute to membrane motility. Using fluorescently tagged motor and vesicle markers we investigate these questions in the nervous system of the nematode *C. elegans* serves us as a model system for microscopic tools (confocal, TIRF, FRET FLIM) and biochemical transport assays *in vitro*.

Selected Recent Publications

Klopfenstein DR, Vale RD (2004) The Lipid Binding Pleckstrin Homology Domain in UNC-104 Kinesin is Necessary for Synaptic Vesicle Transport in *Caenorhabditis elegans*. Mol Biol Cell 15(8): 3729-39

Al-Bassam J, Cui Y, Klopfenstein D, Carragher BO, Vale RD, Milligan RA (2003) Distinct conformations of the kinesin Unc104 neck regulate a monomer to dimer motor transition. J Cell Biol 163(4): 743-53

Tomishige M, Klopfenstein DR, Vale RD (2002) Dimerization triggers fast, processive movement of single Unc104/KIF1A kinesin motor along microtubules. Science 297(5590): 2263-2267

Klopfenstein DR, Tomishige M, Stuurman N, Vale RD (2002) Role of phosphatidylinositol(4,5)bisphosphate organization in membrane transport by the Unc104 kinesin motor. Cell 109(3): 347-58

Klopfenstein DR, Vale RD, Rogers SL (2000) Motor protein receptors: Moonlighting on other jobs. Cell 103(4): 537-40

Wilfried Kramer

Privatdozent Molecular Biology and Genetics

- Diploma (Biology), University of Cologne, Germany, 1982
- Dr. rer. nat., University of Cologne, Germany, 1986
- Postdoctoral Fellow, University of California, Berkeley, USA, 1986 - 1989
- Habilitation in Molecular Biology and Genetics, University of Göttingen, Germany, 2000
- At the Dept. of Molecular Genetics since 1989

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Major Research Interests

Besides being fast and highly accurate, the most important demand on replication of DNA is that is has to be completed. While this may sound trivial on first glance, many obstacles like protein-DNA complexes and damaged nucleotides on the template strand can prevent replication fork progression. It is estimated that at least one fork arrest occurs per replication round in *E. coli*. Therefore, all organisms analysed so far in detail possess several pathways to reactivate stalled replication forks. We discovered that the baker´s yeast Mph1 protein defines a hitherto unknown pathway for replication restart, which is apparently also operating in higher eukaryotes including humans. One question we are interested in is the exact mechanism, by which this pathway works. We are also interested in positioning this pathway within the complex cellular network of replication reinitiation mechanisms, where two principle possibilities for fork reactivation can be found: one being quite safe, but acting on the expense of replicational fidelity, whereas the other is error-free, but bears the inherent danger of genomic rearrangements. Therefore, we are also interested in the regulatory mechanisms that guide the choice of the cell for one or the other possibility as well as the conditions that are sensed by the regulatory proteins.

Selected Recent Publications

Rudolph C, Schürer KA, Kramer W (2005) Facing stalled replication forks: The intricacies of doing the right thing. In: Genome Dynamics and Stability: Facets of Genome Integrity. Lankenau DH (Ed). Springer Verlag Heidelberg. in press (Review)

Prakash R, Krejci L, van Komen S, Schürer KA, Kramer W, Sung P (2005) *Saccharomyces cerevisiae* MPH1 gene, required for homologous recombination-mediated mutation avoidance, encodes a 3' to 5' DNA helicase. J Biol Chem 280: 7854-7860

Schürer KA, Rudolph C, Ulrich HD, Kramer W (2004) Yeast MPH1 gene functions in an error-free DNA damage bypass pathway that requires genes from homologous recombination, but not from postreplicative repair. Genetics 166: 1673-1686

Laging M, Lindner E, Fritz H-J, Kramer W (2003) Repair of hydrolytic DNA deamination damage in thermophilic bacteria: Cloning and characterization of a Vsr endonuclease homolog from *Bacillus stearothermophilus*. Nucl Acids Res 31: 1913-1920

Meyer C, Scheller J, Kramer W (2001) Transcription of mutS- and mutL-homologous genes during meiosis in S*accharomyces cerevisiae* and identification of a regulatory cis-element for meiotic induction of MSH2. Mol Gen Genomics 265: 826-836

Kerstin Krieglstein

Professor of Anatomy/Neuroanatomy

- Dr. rer. nat., University of Gießen, Germany, 1990
- Postdoctoral fellow, University of California, Irvine, 1990 - 1992
- Professor of Anatomy, University of Saarland, 1999 - 2001
- Appointed 2001 as head of the Department of Anatomy/Neuroanatomy, University of Göttingen

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Major Research Interests

The nervous system is a complex network of billions of neurons building appropriate connections and transmitting the information required. Although the nervous system has a lifelong synaptic plasticity, it is essentially built just once with very little regenerative capacity, meaning that neurons have to survive and function for lifetime. Loss of neurons will eventually lead to functional impairments such as those found in Alzheimer's, Parkinson's or ALS patients.

We are interested in the understanding of the regulation of neuronal survival and death. Recent advancements in the field have provided clear evidence that neuronal survival is caused by synergistic actions of neurotrophic factors along with other cytokines most prominently from the TGF-ß superfamily. Synergisms of TGF-ß in combination with neurotrophic factors, like GDNF or NGF, will be studied to establish their role in nervous system development and their therapeutic potential in brain repair. Specifically, we shall investigate such synergisms by utilising mouse mutants to understand the developmental role and by emplying genomic screens to identify new target genes for the establishment of new therapeutic strategies for human neurodegenerative disorders. Furthermore, as growth factors function not only in the decision of neuron survival or death, we shall explore their morphogenetic and differentiation capacities employing the powerful potential of embryonic (ES) and CNS stem cells.

Selected Recent Publications

Krieglstein K, Henheik P, Farkas L, Jaszai J, Galter D, Krohn K, Unsicker K (1998) GDNF requires TGF-ß for establishing its neurotrophic activity. J Neurosci 18: 9822-9834

Schober A, Hertel R, Arumäe U, Farkas L, Jaszai J, Krieglstein K, Saarma M, Unsicker K (1999) GDNF rescues target-deprived spinal cord neurons but requires TGF-ß as co-factor *in vivo*. J Neurosci 19: 2008-2015

Krieglstein K, Richter S, Farkas L, Schuster N, Dünker N, Oppenheim R W, Unsicker K (2000) Reduction of endogenous transforming growth factor beta prevents ontogenetic neuron death. Nature Neuroscience 3: 1085-1091

Peterziel H, Unsicker K, Krieglstein K (2002) TGFbeta induces GDNF responsiveness in neurons by recruitment of GFRalpha1 to the plasma membrane, J Cell Biol 159: 157-167

Farkas L, Dünker N, Roussa E, Unsicker K, Krieglstein K (2003) Transforming growth factor-beta(s) are essential for the development of midbrain dopaminergic neurons *in vitro* and *in vivo*. J Neurosci 23: 5178-5186

v Bohlen und Halbach O, Schober A, Krieglstein K (2004) Genes, proteins, and neurotoxins involved in Parkinson's disease. Prog Neurobiol 73: 151-177

Wolfgang Liebl

Professor of Microbiology

- 1984 Diploma (Biology), Technische Universität München
- 1986 Ph.D. (Dr. rer. nat.), Technische Universität München
- 1986 - 1988 Postdoctoral Fellow, Massachusetts Institute of Technology, Cambridge, MA, USA
- 1997 Habilitation (Microbiology), Technische Universität München
- 1997 - 2003 Associate Professor of Microbiology, Georg-August-Universität, Göttingen
- Since 2003 Full Professor (Applied Microbiology), Georg-August-Universität, Göttingen

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Major Research Interests

One of the main interests of our group is the analysis of polysaccharide and oligosaccharide breakdown and utilization by microorganisms adapted to extreme habitats. In particular, we are interested in cellulose, xylan and starch degrading enzyme systems from hyperthermophiles, i. e. organisms that grow optimally at 80°C or higher. These organisms represent very deep branches within the prokaryotic lineages of the phylogenetic tree of organisms. We are interested in the biochemical properties, the molecular structure and catalytic mechanism, the function(s) of non-catalytic domains, and the cellular localization of unusual glycosyl hydrolases and transferases from *Thermotoga maritima*, the model organism of hyperthermophilic bacteria. Other projects in the field of extremophilic microorganisms deal with the enzymology and molecular biology of thermoalkaliphiles and thermoacidophiles. We have completed the genome sequence of the extreme thermoacidophilic archaeon *Picrophilus torridus* with the objective to better understand the evolutionary, metabolic and molecular mechanisms that allow this organism to thrive at up to 65°C at pH values close to pH 0.

Another group of bacteria studied in the laboratory are the Gram-positive bacteria with a high G+C content. We employ molecular biological techniques to study and modify physiological traits of amino acid-producing corynebacteria and micrococci.

Also, the group is interested in the characterization of genome (metagenome) structures of various microbial habitats (PD Dr. R. Daniel). DNA libraries are constructed from microbial consortia and biofilms in order to explore the genetic diversity of the different environments. Also, classical activity-based screens are used for the isolation of novel enzymes useful for biotechnology.

Selected Recent Publications

Angelov A, Fütterer O, Valerius O, Braus GH, Liebl W (2005) Properties of the recombinant glucose/galactose dehydrogenase from the extreme thermoacidophile, *Picrophilus torridus*. FEBS J (Eur J Biochem) 272:1054-1062

Daniel R (2005) The metagenomics of soil. Nature Rev Microbiol 3: 470-478

Fütterer O, Angelov A, Liesegang H, Gottschalk G, Schleper C, Schepers B, Dock C, Antranikian G, Liebl W (2004) Genome sequence of *Picrophilus torridus* and its implications for life around pH 0. Proc Natl Acad Sci USA 101: 9091-9096

Liebl W (2004) Genomics taken to the extreme. Nature Biotechnology 22: 524-525

Tzvetkov M, Klopprogge C, Zelder O, Liebl W (2003) Genetic dissection of trehalose biosynthesis in *Corynebacterium glutamicum*: inactivation of trehalose production leads to impaired growth and an altered cell wall lipid composition. Microbiology 149: 1659-1673

Lodge JA, Maier T, Liebl W, Hoffmann V, Sträter N (2003) Crystal structure of *Thermotoga maritima* α-glucosidase AglA defines a new clan of NAD^+-dependent glycosidases. J Biol Chem 278: 19151-19158

Reinhard Lührmann

Professor, Director at the Max Planck Institute for Biophysical Chemistry

- Dr. rer. nat (Ph. D.), University of Münster (1975)
- Research group leader, Max Planck Institute for Molecular Genetics, Berlin (1981 - 1988)
- Professor of Biochemistry and Molecular Biology at the University of Marburg (1988 - 1999)
- Director, Dept. of Cellular Biochemistry, Max Planck Institute for Biophysical Chemistry, Göttingen (since 1999)

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Major Research Interests

Most metazoan pre-mRNAs contain multiple introns and exons. In order to generate mature mRNA, the introns must be excised from the pre-mRNA, a process termed pre-mRNA splicing. In many cases, alternative splicing generates different mRNAs from a single pre-mRNA by the regulated removal of different sections of the RNA, a process which greatly expands the complexity of the repertoire of proteins that can be expressed from relatively small genomes. Splicing is catalysed by a large macromolecular machine, termed the spliceosome which consists of the small nuclear RNAs (U1, U2, U4, U5 and U6) and more than 150 proteins, 50 of which are associated with the snRNAs to form snRNPs.

In our laboratory, intense efforts are focussed on understanding how the spliceosome recognizes and binds the intron ends and discriminates them from exons. This is an especially confounding problem in metazoans because, in contrast to lower eucaryotes such as yeast, pre-mRNA introns are often extremely long (104-105 nucleotides), while exons are generally small (less than 300 nucleotides). Another major goal of our research is the elucidation of the mechanisms by which the spliceosome assembles into a catalytically active machine and catalyses intron excision. None of the building blocks of the spliceosome contains an active site. Instead, the catalytically active domain must be assembled anew on to each intron, a highly dynamic process which entails dramatic structural rearrangements of the RNP structure of the spliceosome, and which is orchestrated by the successive action of more than 10 enzymes such as RNA helicases and GTPases, as well as by posttranslational phosphorylation of a multitude of spliceosomal proteins. Our studies involve a large number of experimental approaches, including biochemical purification of entire spliceosomes or large protein ensembles, and characterization of their proteins by mass spectrometry; RNA biology methods such as enzymatic engineering of RNA molecules, RNA structure probing and RNA interference methods; production of recombinant proteins and antibodies; procedures for the investigation of protein-protein and protein-RNA interactions in vitro and *in vivo*; and biophysical methods such as fluorescence spectroscopy. Finally, we are investigating the 3D structure of purified spliceosomes or major building blocks thereof using electron microscopic approaches and X ray crystallography. Our studies on the regulatory mechanisms of constitutive and alternative pre-mRNA splicing involve mainly mammalian systems. As the basic mechanisms of splicing catalysis appear to be evolutionarily highly conserved, we are also taking advantage of molecular genetic approaches in bakers yeast to elucidate the structure and function of the catalytic core domain of the spliceosome.

Selected Recent Publications

Dönmez G, Hartmuth K, Kastner B, Will C L, Lührmann R (2007) The 5'End of U2 snRNAs is in close proximity to U1 and functional sites of the pre-mRNA in early spliceosomal complexes. Mol Cell 25: 399-411

Liu S, Li P, Dybkov O, Nottrott S, Hartmuth K, Lührmann R, Carlomagno T, Wahl MC (2007) Binding of the Human Prp31 Nop Domain to a Composite RNA-Protein Platform in U4 snRNP. Science 316: 115-120

Deckert J, Hartmuth K, Böhringer D, Behzadnia N, Will CL, Kastner B, Stark H, Urlaub H, Lührmann R (2006) Protein composition and electron microscopy structure of affinity-purified human spliceosomal B complexes isolated under physiological conditions. Mol Cell Biol 26: 5528-5543

Will CL, Lührmann R (2005) Spliceosome structure and function. RNA World III (CSH Laboratory Press) (R F Gesteland, T R Cech, J F Atkins eds.): 369-400

Böhringer D, Makarov EM, Sander B, Makarova OV, Kastner B, Lührmann R, Stark H (2004) Three-dimensional structure of a pre-catalytic human spliceosomal complex B. Nature Struct. Mol Biol 11: 463-468

Makarov EM, Makarova OV, Urlaub H, Gentzel M, Will CL, Wilm M, Lührmann R (2002) Small nuclear ribonucleoprotein remodeling during catalytic activation of the spliceosome. Science 298: 2205-2208

Watkins NJ, Segault V, Carpentier B, Nottrott S, Fabrizio P, Bachi A, Wilm M, Rosbash M, Branlant C, Lührmann R (2000) A common core RNP structure shared between the small nuclear box C/D RNPs and the spliceosomal U4 snRNP. Cell 103: 457- 466

Ahmed Mansouri

Molecular Developmental Genetics

- Diploma (Chemistry), Technical University, Braunschweig (Germany) 1975
- Dr. rer. nat. Chemical Technology Institute, Technical University, Braunschweig (Germany), 1978
- Postdoc at the Institute of Human Genetics in Göttingen (1982 - 1986)
- Postdoc at the Miescher Institute in Tübingen (MPI) and at the Max Planck
- Institute of Immunbiology in Freiburg (Germany) (1986 - 1989)
- Since 1989 Dept of Molecular Cell Biology at the MPI for Biophysical Chemistry in Göttingen
- Habilitation (Molecular Developmental Genetics), University of Göttingen, Germany, 1999
- Since 2005: Dr. Helmut Storz Stiftungsprofessur for “dopaminerge Stammzelltherapie”, Dept. of Clinical Neurophysiology at the University of Göttingen

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Major Research Interests

Molecular mechanisms of mammalian development and stem cell biology

In order to understand the molecular mechanisms governing mammalian development, we are using the mouse as a model system. We are focusing on the role of transcription factors in development. Using embryonic stem cells mouse loss-of-function mutants were generated. Specifically, we have shown that Pax and homeobox-containing genes are required for early decisions in organogenesis and cell differentiation. In addition, we are currently taking advantage of the *in vitro* differentiation potential of embryonic stem cells to search for molecules that are involved in dopaminergic neuron induction, differentiation, and/or survival.

Selected Recent Publications

Schindehütte J, Fukumitsu H, Collombat P, Griesel G, Brink C, Baier PC, Capecchi MR, Mansouri A (2005) *In vivo* and *in vitro* tissue-specific expression of GFP using the Cre-lox system in mouse embryonic stem cells. Stem Cells 23: 10-15

Collombat P, Hecksher-Soerensen J, Broccoli V, Krull J, Ponte I, Mundiger T, Smith J, Gruss P, Serup P, Mansouri A (2005) The simultaneous loss of Arx and Pax4 genes promotes a somatostatin-producing cell fate specification at the expense of the alpha- and beta-cell lineages in the mouse endocrine pancreas. Development 132 (13): 2969-80

Thinyane K, Baier PC, Schindehütte J, Mansouri A, Paulus W, Trenkwalder C, Flügge G, Fuchs E (2005) Fate of pre-differentiated mouse embryonic stem cells transplanted in unilaterally 6-hydroxydopamine lesioned rats: histological characterization of the grafted cells. Brain Res 1045 (1-2): 80-87

Relaix F, Rocancourt D, Mansouri A, Buckingham M (2005) A Pax3/Pax7-dependent population of skeletal muscle progenitor cells. Nature 435: 948-953

Baier PC, Schindehütte J, Thinyane K, Flugge G, Fuchs E, Mansouri A, Paulus W, Gruss P, Trenkwalder C (2004) Behavioral changes in unilaterally 6-hydroxy-dopamine lesioned rats after transplantation of differentiated mouse embryonic stem cells without morphological integration. Stem Cells 22(3): 396-404

Frauke Melchior

Professor of Biochemistry

- 1990 Dr. rer. nat., University of Marburg, Germany
- 1990 - 1992 Postdoctoral fellow at the Max Planck Institute for Biophysical Chemistry, Göttingen
- 1992 - 1998 Postdoctoral fellow at the Scripps Research Institute, La Jolla, CA, USA
- 1998 - 2004 Independent group leader at the Max Planck Institute of Biochemistry, Martinsried
- 1998 BioFUTURE young investigator award Since 2004 Professor of Biochemistry, Georg-August University Göttingen

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Major Research Interests

Research in our group centers around posttranslational modification with small ubiquitin-related proteins of the SUMO family. SUMO proteins are ubiquitously expressed in eukaryotic cells, and are essential for life. They are reversibly coupled to a large number of cellular targets, and thereby modulate protein / protein or protein / DNA interactions, alter intracellular localization, or protect from ubiquitin mediated degradation. Higher organisms often express several distinct SUMO proteins (e.g., three in humans). Those are conjugated to different targets under normal growth conditions, or conjugated preferentially upon stress. Most of the known targets for sumoylation can be associated with a few specific pathways: signal transduction, transcription, chromatin remodelling, DNA repair, mitosis, viral infection, and nucleocytoplasmic trafficking. Projects in the lab aim to understand basic mechanisms, regulation, and function of SUMOylation in mammalian cells. This involves, e.g., characterization of SUMO enzymes, analysis of SUMO conjugation under stress conditions, and the identification and characterization of novel SUMO targets. Special emphasis is also given to the interplay between SUMOylation and nucleocytoplasmic trafficking.

Selected Recent Publications

Bossis G, Melchior F (2006) Regulation of SUMOylation by reversible oxidation of SUMO conjugating enzymes. Mol Cell 21: 349-357

Pichler A, Knipscher P, Saitoh H, Sixma T, Melchior F (2004) SUMO E3 ligase is neither Hect nor Ring type. Nat Struct Mol Biol 11: 984-991

Swaminathan S, Kiendl F, Körner R, Lupetti R, Hengst L, Melchior F (2004) RanGAP1*SUMO-1 is phosphorylated at the onset of mitosis and remains associated with RanBP2 upon NPC disassembly. J Cell Biol 164:965-971

Melchior F, Schergaut M, Pichler A (2003) SUMO: ligases, isopeptidases and nuclear pores. Trends Biochem Sci 28: 612-618

Pichler A, Gast A, Seeler JS, Dejean A, Melchior F (2002) The nucleoporin RanBP2 is a SUMO1 E3 Ligase. Cell 108: 109-120

Professor of Bioinformatics

- 1993 Diploma (Mathematics), LMU München
- 1996 PhD (Dr. Math.), Universität Bielefeld
- 1997 - 1998 Visiting Scientist, North Carolina State University, Raleigh, NC, USA
- 1998 - 2000 RPR/Aventis, Dagenham, Essex, UK
- 2000 - 2001 MIPS, MPI fuer Biochemie, Martinsried and GSF, Neuherberg
- 2001 - 2002 Group leader and faculty member at International Graduate
- School in Bioinformatics and Genome Research, Univertität Bielefeld
- Since 2002 Professor of Bioinformatics, Universität Göttingen

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Major Research Interests

The focus of our work is on algorithm development for nucleic acid and protein sequence analysis. We are particularly interested in multiple sequence alignment and gene prediction; the software programs DIALIGN and AUGUSTUS are developed and maintained by our department.

In recent years, alignment of large genomic sequences became a powerful tool for genome analysis and annotation. Cross-species alignment of genomic sequences has been used for gene prediction, to detect regulatory sites or to identify signature sequences for pathogen microorganisms. These novel alignment approaches are also used to improve our gene-finding software tools.

Other areas of research include: metabolomics and mass spectroscopy data analysis, phylogeny reconstruction, RNA structure analysis, motif discovery and remote homology detection using machine-learning methods, genome annotation for prokaryotes, recombinations in viral genomes and grid computing.

Selected Recent Publications

Chen et al (2007) Nature Biotechnology 25: 1007-1014 http://www.nature.com/nbt/journal/v25/n9/abs/nbt1325.html

Stanke M, Tzvetkova A, Morgenstern B (2006) AUGUSTUS+ at EGASP: using EST, protein and genomic alignments for improved gene prediction in the human genome. Genome Biology 7: S11

Schultz A-K, Zhang M, Leitner T, Kuiken C, Korber B, Morgenstern B, Stanke M (2006) A jumping profile Hidden Markov Model and applications to recombination sites in HIV and HCV genomes. BMC Bioinformatics 7: 265

Stanke M, Keller O, Gunduz I, Hayes A, Waack S, Morgenstern B (2006) AUGUSTUS: ab initio prediction of alternative transcripts. Nucleic Acids Res 34: W435 - W439

Subramanian AR, Weyer-Menkhoff J, Kaufmann M, Morgenstern B (2005) DIALIGN-T: An improved algorithm for segment-based multiple sequence alignment. BMC Bioinformatics 6: 66

Brudno M, Chapman M, Göttgens B, Batzoglou S, Morgenstern B (2003) Fast and sensitive multiple alignment of large genomic sequences. BMC Bioinformatics 4: 66

Klaus-Armin Nave

Professor of Molecular Biology, Director at the Max Planck Institute of Experimental Medicine

- PhD 1987, University of California, San Diego, Postdoc, The Salk Institute, La Jolla, California
- 1991 Junior Group Leader, ZMBH,University of Heidelberg
- 1998 Professor of Molecular Biology (C4), ZMBH
- 2000 Director, Department of Neurogenetics Max Planck Institute for Experimental Medicine, Göttingen, and Professor of Biology, University of Heidelberg

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Major Research Interests

We are interested in the mechanisms of neuron-glia interactions in the higher nervous system, and in the genes that are required for normal glial cell function. Here, transgenic and mutant mice have become important to study developmental processes as well as genetic diseases. For example, oligodendrocytes are glial cells highly specialized for enwrapping CNS axons with multiple layers of membranes, known to provide electrical insulation for rapid impulse propagation. We found that oligodendrocytes are also essential for maintaining the long-term integrity of myelinated axons, independent of the myelin function itself. The mechanisms by which oligodendrocytes support long-term axonal survival are still under investigation. The importance of glial cells as the “first line of neuroprotection”, however, is illustrated by several myelin-associated diseases in which axonal neurodegeneration contribute to progressive disability. These range in humans from peripheral neuropathies (CMT1) to spastic paraplegia (SPG2), and presumably multiple sclerosis (MS) and certain forms of psychiatric disorders. We are developing transgenic animal models for some of these diseases, in order to dissect the underlying disease mechanisms and, in the case of CMT1A, have used these models to design novel therapeutic strategies.
The glial “decision” to myelinate an axonal segment is partly controlled by the axon itself, but the signaling mechanism is not understood. We have found that axonal neuregulin-1 (NRG1) is the major determinant of myelination in the peripheral nervous system. We are now investigating NRG1 dysregulation also in CNS myelination, using quantifiable behavioural functions in mice. By combining genetics with enviromental risk factors for schizophrenia (in collaboration with H. Ehrenreich) we will explore the hypothesis that NRG1, a known human schizophrenia susceptibility gene, points to an important role of myelinating glia in some psychiatric disorders.

Future Projects and Goals
Mechanisms of neuron-glia signalling; function of myelin proteins and lipids; transcriptional profiling of single cells *in vivo*; novel mouse models of neuropsychiatric disorders.

Selected Recent Publications

Dhaunchak A, Nave K-A (2007) A common mechanism of proteolipid protein misfolding leading to cysteine-mediated ER retention in oligodendrocytes and Pelizaeus-Merzbacher disease. Proc Natl Acad Sci USA (in press)

Kassmann CM, Lappe-Siefke C, Baes M, Brügger B, Mildner A, Werner HB, Natt O, Michaelis Th, Prinz M, Frahm J, Nave K-A (2007) Axonal loss and neuroinflammation caused by peroxisome-deficient oligodendrocytes. Nature Genetics 8: 969-976

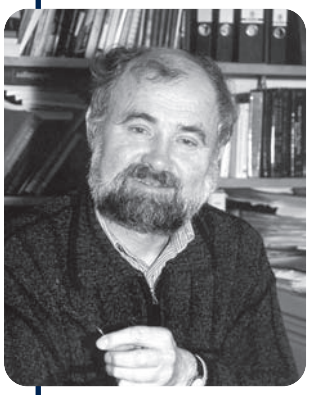
Kramer-Albers EM, Gehrig-Burger K, Thiele C, Trotter J, Nave K-A (2006) Perturbed interactions of mutant proteolipid protein/DM20 with cholesterol and lipid rafts in oligodendroglia: implications for dysmyelination in spastic paraplegia. J Neurosci. 26: 11743-11752

Saher G, Brügger B, Lappe-Siefke C, Möbius W, Tozawa R, Wehr M, Wieland F, Ishibashi S, Nave K-A (2005) Cholesterol is essential and rate-limiting for myelin membrane growth. Nature Neurosci 8: 468-475

Michailov GV, Sereda MW , Brinkmann BG, Fischer TM, Haug B, Birchmeier C, Role L, Lai C, Schwab MH, Nave K-A (2004) Axonal neuregulin-1 regulates myelin sheath thickness. Science 304: 700-703

Sereda MW, Meyer zur Hörste G, Suter U, Uzma N, Nave K-A (2003) Therapeutic administration of anti-progesterone in a PMP22-transgenic model of Charcot-Marie-Tooth disease (CMT1A). Nature Medicine 9: 1533-1537

Erwin Neher



Professor, Director at the Max Planck Institute for Biophysical Chemistry

- M.Sc. (Physics), University of Wisconsin, (1967)
- Ph.D. (Physics), Institute of Technology, Munich (1970)
- Research associate at the Max Planck Institute for Biophysical Chemistry in Göttingen, Germany (1972 - 1975 and 1976 - 1982) and as a guest in the
- laboratory of Dr. Ch.F. Stevens at Yale University, Dept. of Physiology, New Haven, Conn. (1975 - 1976)
- Fairchild Scholar, California Institute of Technology; Pasadena, USA (1989)
- Director of the Membrane Biophysics Department at the Max Planck Institute for Biophysical Chemistry, Göttingen, Germany, since 1983

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Major Research Interests

Molecular Mechanisms of Exocytosis, Neurotransmitter Release, and Short Term Synaptic Plasticity

In order to understand how the brain handles its information flow and adjusts synaptic connections on the second and subsecond timescale, one has to understand all aspects of synaptic transmission ranging from availability of vesicles for exocytosis, presynaptic electrophysiology, Ca^{++} signalling, the process of exocytosis, and postsynaptic neurotransmitter action. Our work concentrates on presynaptic aspects. We study the basic mechanisms of exocytosis, using adrenal chromaffin cells as a model system and the patch-clamp method. This work, in which intracellular Ca^{++} is manipulated (caged Ca^{++}) and measured on the single cell level aims at understanding the role of specific synaptic proteins in the maturation and exocytosis of secretory vesicles. We use neuronal cell cultures and brain slices for studying mechanisms of short term plasticity, such as depression and paired pulse facilitation. The Calyx of Held, a specialized synapse in the auditory pathway, offers unique possibilities for simultaneous pre- and postsynaptic voltage clamping. This allows a quantitative analysis of the relationship between $[Ca^{++}]$ and transmitter release.

Selected Recent Publications

Klingauf J, Neher E (1997) Modeling buffered Ca^{2+} diffusion near the membrane: Implications for secretion in neuroendocrine cells. Biophys J 72: 674-690

Neher E (1998) Vesicle pools and Ca^{2+} microdomains: new tools for understanding their roles in neurotransmitter release. Neuron 20: 389-399

Schneggenburger R, Neher E (2000) Intracellular calcium dependence of transmitter release rates at a fast central synapse. Nature 406: 889-893

Rettig J, Neher E (2002) Emerging roles of presynaptic proteins in Ca^{++}-triggered exocytosis. Science 298: 781-785

Sakaba T, Neher E (2003) Direct modulation of synaptic vesicle priming by $GABA_B$ receptor activation at a glutamatergic synapse. Nature 424: 775-778

Soerensen J, Nagy G, Varoqueaux F, Nehring RB, Brose N, Wilson MC, Neher E (2003). Differential control of the releasable vesicle pools by SNAP-25 splice variants and SNAP-23. Cell 114: 75-86

Sakaba T, Stein A, Jahn R, Neher E (2005) Distinct kinetic changes in neurotransmitter release after SNARE protein cleavage. Science 309: 491-494

Tomas Pieler

Professor of Biochemistry

- Dr. rer. nat. Biochemistry, Freie Universität Berlin, 1984
- Guest Investigator, Rockefeller University, New York (1985/86)
- Heisenberg fellow, Freie Universität Berlin and Rockefeller University, New York (1986/87)
- Junior group leader, Max-Planck-Institut für Molekulare Genetik, Berlin (1988 - 92)
- Professor of Biochemistry, Georg-August-Universität Göttingen (since 1992)
- Head of the Department of Developmental Biochemistry, Georg-August-Universität Göttingen

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Major Research Interests

The differentiation of complex organisms has its origin in the asymmetric distribution of regulatory proteins or of the corresponding mRNAs in the egg, as well as in a complex system of cell/cell communication events via extracellular signalling molecules during early stages of embryogenesis. The genes that encode for these different activities form functional networks which provide the basis for the genetic programming of embryonic development. Our primary research interest is in the identification of such regulatory genes and networks in vertebrates, as well as in the definition of their regulation and function on the molecular level. For this purpose, we use *Xenopus laevis*, a frog from South Africa, as a model system. As a traditional object in experimental embryology and in comparison with other experimental systems such as the mouse, use of *Xenopus* offers a number of practical advantages. Oocytes and embryos are easy to collect in large numbers, they are easy to manipulate by relatively simple techniques, also because embryonic development proceeds in the petridish, and, more recently, it has even become possible to generate hundreds of transgenic frogs within a single experimental day. The research topics that we are focussing on are:

Transport and function of vegetally localized maternal mRNAs
Organogenesis: formation of pancreas and liver in vertebrate embryos
Early neural development: primary neurogenesis

Selected Recent Publications

Chen Y*, Pan FC*, Brandes N, Afelik S, Sölter M, Pieler T (2004) Retinoic Acid signaling is essential for pancreas development and promotes endocrine at the expense of exocrine cell differentiation in *Xenopus*. Dev Biol 271: 144-160 *equal contribution

Claußen M, Horvay K, Pieler T (2004) Evidence for overlapping but not identical protein mechineries to operate in vegetal localisation along early and late pathways in *Xenopus* oocytes. Development 131: 4263-4273

Loop S, Katzer M, Pieler T (2005) mPer1 mediated nuclear export of Cry ½ is an important element in establishing the circadian rhythm. EMBO Reports 4:341-347

Afelik S, Chen Y, Pieler T (2006) Combined ectopic expression of Pdx1 and Ptfa/p48 results in the stable conversion of posterior endoderm into endo- and exocrine pancreatic tissue. Genes and Dev 20:1441-1446

Sölter M, Locker M, Boy S, Taelman V, Bellefroid E, Perron M, Pieler T (2006) Characterization and function of the bHLH-O protein XHes2: Insight into the mechanisms controlling retinal cell fate decision. Development (in press)

Stefanie Pöggeler

Professor of Genetics of Eukaryotic Microorganisms

- 1993 Dr. rer. nat., Ruhr-Universität Bochum
- 1993 - 1995 Research associate, Laboratory of Prof. U. Kück, Ruhr-Universität Bochum
- 1995 - 2001 Postdoctoral research fellow and group leader, Ruhr-Universität Bochum
- 1997 Visiting Scientist, Institut de Génétique et Microbiologie, Laboratory of Dr. D. Zickler, Université Paris-Sud, Orsay, France
- 2000 Habilitation (Botany), Ruhr-Universität Bochum
- 2001 - 2003 Associate Professor of Botany (stand-in), University of Münster
- 2003-2006 University lecturer (Hochschuldozentin) and group leader, Ruhr-Universität Bochum
- since 2006 Associate Professor of Genetics of Eukaryotic Microorganisms, Georg-August-Universität Göttingen

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Major Research Interests

Fruiting-body development in filamentous ascomycetes

Fruiting-body development in filamentous ascomycetes is a complex cellular differentiation process that requires special environmental conditions and is controlled by many developmentally regulated genes. We are interested in the genes regulating this development process. We use the homothallic (self-fertile) ascomycete *Sordaria macrospora* as a model organism. Since *S. macrospora* is able to complete the sexual cycle without a mating partner, recessive mutations affecting fruiting body development are directly visible. Numerous mutants which are blocked at various stages of fruiting-body development have been generated and molecular genetic procedures have been applied to isolate genes involved in fruiting-body development.

Fungal inteins

An intein is a self-catalytic protein-intervening sequence that catalyses its precise excision from a host protein and the ligation of its flanking sequences, termed N- and C-exteins, to produce the mature spliced product. Protein splicing is a posttranslational event that releases an internal intein sequence from a protein precursor. Projects in the lab aim to analyse the splicing activity of inteins detected in the *prp8* gene of fungi. The PRP8 protein is one of the largest and most highly conserved nuclear proteins occupying a central position in the catalytic core of the spliceosome. Because of their compactness and high splicing activity inside foreign proteins, fungal PRP8 inteins may be used for the development of new intein-mediated protein-engineering applications.

Selected Recent Publications

Nolting N, Pöggeler S (2006) A STE12 homologue of the homothallic ascomycete *Sordaria macrospora* interacts with the MADS box protein MCM1 and is required for ascosporogenesis. Mol Microbiol 62: 853-868

Mayrhofer S, Weber JM, Pöggeler S (2006) Pheromones and pheromone receptors are required for proper sexual development in the homothallic ascomycete *Sordaria macrospora*. Genetics 172: 1521-1533

Elleuche, S, Nolting N, Pöggeler S (2006) Protein splicing of PRP8 mini-inteins from species of the genus *Penicillium*. Appl Microbiol Biotechnol 72: 959-967

Pöggeler S, Kück U (2004) A WD40 repeat protein regulates fungal cell differentiation and can functionally be replaced by the mammalian homologue striatin. Eukaryotic Cell 3: 232-240

Van Heemst D, James F, Pöggeler S, Bertaux-Lecellier V, Zickler D (1999) Spo76p is a conserved chromosome morphogenesis protein that links the mitotic and meiotic programs. Cell 98: 261-271

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Group Leader STED Microscopy of Synaptic Function

- 2000 - 2004 Research assistant with William Betz at the Dep. of Physiology and Biophysics, University of Colorado Health Sciences Center (USA)
- 08/2004 PhD degree (Physiology) awarded by the University of Colorado
- 2004 - 2007 Post doctoral fellow with Reinhard Jahn at the Neurobiology Department of the Max Planck Institute for Biophysical Chemistry in Göttingen (Germany)
- since 2007 Group Leader (STED Microscopy) at the European Neuroscience Institute Göttingen (ENI-G)

Major Research Interests

Conventional fluorescence microscopy is limited by the diffraction of light: fluorescent objects that are close together cannot be discerned. Stimulated emission depletion (STED) is a recent advancement in optical physics that breaks the diffraction barrier, allowing microscopes to obtain much clearer images.

The diffraction barrier has been particularly problematic for imaging synaptic vesicles, which are among the smallest known organelles (30-50 nm in diameter). They are located in small areas in the synapses (about 1 micron in diameter). The group takes advantage of the increased imaging resolution provided by STED to investigate synaptic vesicle function, with an emphasis on synaptic vesicle recycling. Since STED microscopy also allows imaging of protein domains, the group aims at studying the patterning of protein domains in the synapse, in order to understand its molecular architecture.

Selected Recent Publications

Bethani I, Lang T, Geuman U, Sieber JJ, Jahn R, Rizzoli SO (2007) The specificity of SNARE pairing in biological membranes is mediated by both proof-reading and spatial segregation. EMBO J, in press

Rizzoli SO, Bethani I, Zwilling D, Wenzel D, Siddiqui TJ, Brandhorst D, Jahn R (2006) Evidence for early endosome-like fusion of recently endocytosed synaptic vesicles. Traffic 7(9): 1163-76

Gaffield MA, Rizzoli SO, Betz WJ (2006) Mobility of synaptic vesicles in different pools in resting and stimulated frog motor nerve terminals. Neuron 51(3): 317-25

Willig KI, Rizzoli SO, Westphal V, Jahn R, Hell SW (2006) STED microscopy reveals that synaptotagmin remains clustered after synaptic vesicle exocytosis. Nature 440(7086): 935-9

Rizzoli SO, Betz WJ (2005) Synaptic vesicle pools. Nat Rev Neurosci 6(1): 57-69. Review

Reinhard Schuh

Research Group Leader at the MPI for Biophysical Chemistry

- Dr. rer. nat., University of Tübingen, Germany, 1986
- Postdoctoral Fellow at the Max Planck Institute for Developmental Biology, Tübingen, Germany, 1986 - 1988
- Postdoctoral Fellow at the University of Munich, Germany, 1989 - 1991
- Group leader in the Department of Molecular Developmental Biology at the Max Planck Institute for Biophysical Chemistry, Göttingen, Germany, 1992 - 2004
- Habilitation in Cellular and Molecular Biology, Technical University of Braunschweig, Germany, 2001
- Leader of the Research Group Molecular Organogenesis at the Max Planck Institute for Biophysical Chemistry, since 2005

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Major Research Interests

Branched tubular networks are a fundamental structural design of many organs including lung, vascular system and kidney. Critical for organ function, i.e. the transport of fluids or gases, is the proper size and diameter of the tubular branches as well as an elaborated network formation. How do these networks develop? How do the branches grow out, detect their fusion partners and interconnect? How are tube size and diameter controlled? How can the system respond to different physiological needs? How do epidermal sheets control the paracellular passage of solutes?

We investigate the development of the *Drosophila* tracheal (respiratory) system since it provides an ideal model to address such questions, because of its simple stereotypic architecture, accessible genetics and molecular tools.

Selected Recent Publications

Krause C, Wolf C, Hemphälä J, Samakovlis C, Schuh R (2006) Distinct functions of the leucine-rich repeat transmembrane proteins Capricious and Tartan in the *Drosophila* tracheal morphogenesis. Dev Biol 296: 253-264

Adryan B, Schuh R (2004) Gene Ontology-based clustering of gene expression data. Bioinformatics 20: 2851-2852

Behr M, Riedel D, Schuh R (2003) The claudin-like Megatrachea is essential in septate junctions for the epithelial barrier function in *Drosophila*. Dev Cell 5: 611-620

Wolf C, Gerlach N, Schuh R (2002) *Drosophila* tracheal system formation involves FGF-dependent cell extensions contacting bridge-cells. EMBO Reports 3: 563-568

Wolf C, Schuh R (2000) Single mesodermal cells guide outgrowth of ectodermal tubular structures in *Drosophila*. Genes Dev 14: 2140-2145

George M. Sheldrick

Professor of Structural Chemistry and part-time programming technician at the University of Göttingen

- PhD (1966) University of Cambridge with E.A.V. Ebsworth; thesis entitled "NMR Studies of Inorganic Hydrides"
- 1966 - 1978: University Lecturer and Fellow of Jesus College, Cambridge
- Since 1978 Professor at the University of Göttingen
- Author of more than 750 scientific papers and of a computer program called SHELX (http://shelx.uni-ac.gwdg.de/SHELX/)

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Major Research Interests

Interested in methods of solving and refining crystal structures (both small molecules and proteins) and in structural chemistry.
Holy Grail: the Crystallographic Phase Problem. If only there was an easy way of measuring the phases of X-ray reflections as well as their intensities, crystal structures could be determined directly. At resolutions of better than about 2.5A, there are more measured intensities than atomic coordinates, so the problem is overdetermined and there should be a solution. Recently we were able to increse the size of structures that can be solved from the intensity data alone by 'ab initio direct methods' from about 200 to 1000 unique atoms, given data to 'atomic resolution', but most interesting macromolecular structures are still out of the reach of such methods. Indirctly however the same techniques are proving very useful for the solution of large macromolecular structures when a little starting phase information is available, e.g. by incorporating heavy atoms into the crystal.

Selected Recent Publications

Bunkóczi G, Vértesy L, Sheldrick GM (2005) The antiviral antibiotic feglymycin: First direct-methods solution of a 1000+ equal-atom structure. Angew Chem Int Ed 44: 1340-1342

Lehmann C, Bunkóczi G, Vértesy L, Sheldrick GM (2002) Structures of glycopeptide antibiotics with peptides that model Bacterial Cell-Wall Precursors. J Mol Biol 318: 723-732

Sheldrick GM (2002) Macromolecular phasing with SHELXE. Z Kristallogr 217: 644-650

Debreczeni JÈ, Bunkóczi G, Ma Q, Blaser H, Sheldrick GM (2003) In-house measurement of the sulfur anomalous signal and its use for phasing. Acta Crystallogr D59: 688-696

Debreczeni JÈ, Girmann B, Zeeck A, Krätzner R, Sheldrick GM (2003) Structure of viscotoxin A3: dislulphide location from weak SAD data. Acta Crystallogr D59: 2125-2132

Holger Stark

Group Leader 3D-Cryo Electron Microscopy

- 1996 Dr. rer. nat. (Biochemistry) Free University of Berlin
- 1997 - 1998 Postdoc (Laboratory of Marin van Heel, Imperial College, London)
- 1998 - 1999 Junior group leader, University of Marburg
- 2000 - 2004 Junior group leader, Max-Planck-Institute for Biophysical Chemistry
- 2005 - BioFuture group leader, Max-Planck-Institute for Biophysical Chemistry

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Major Research Interests

- 3D Structure determination of large macromolecular complexes
- Splicing and Translation
- Electron Microscopy
- Image processing

Selected Recent Publications

Stark H, Rodnina MV, Wieden HJ, van Heel M, Wintermeyer W (2000) Large-scale movement of elongation factor G and extensive conformational change of the ribosome during translocation. Cell 3: 301-9

Stark H, Dube P, Lührmann R, Kastner B (2001) The 3-D arrangement of RNA and proteins in the spliceosomal U1 snRNP. Nature 409: 539-542

Golas MM, Sander B, Will CL, Luhrmann R, Stark H, (2003) Molecular architecture of the multiprotein splicing factor SF3b. Science 300(5621): 980-4

Dube P, Herzog F, Gieffers C, Sander B, Riedel D, Muller SA, Engel A, Peters JM Stark H (2005) Localization of the Coactivator Cdh1 and the Cullin Subunit Apc2 in a Cryo-Electron Microscopy Model of Vertebrate APC/C. Mol Cell 20: 867-879

Sander B, Golas MM, Makarov EM, Brahms H, Kastner B, Luhrmann R, Stark H (2006) Organization of core spliceosomal components U5 snRNA loop I and U4/U6 Di-snRNP within U4/U6.U5 Tri-snRNP as revealed by electron cryomicroscopy. Mol Cell 24: 267

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Professor of Microbiology

- 1990 Diploma (Biology), Ernst-Moritz-Arndt-Universität Greifswald
- 1994 Dissertation (Dr. rer. nat.), Ernst-Moritz-Arndt-Universität Greifswald
- 1994 - 1996 Postdoctoral Fellow at the Institut Pasteur, Paris
- 1996 - 2003 Group leader at the Chair of Microbiology, University Erlangen-Nürnberg
- 2000 Habilitation (Microbiology), University Erlangen-Nürnberg
- Since 2003 Professor of General Microbiology, Head of the Department of General Microbiology at the Institute of Microbiology and Genetics, University of Göttingen

Major Research Interests

Our group studies the regulation of metabolism in the pathogenic bacterium *Mycoplasma pneumoniae* and the model organism *Bacillus subtilis*. We are following global ("post-genomic") and gene-specific approaches. In *Mycoplasma pneumoniae*, we study the regulation of gene expression in this pathogenic bacterium and its relation to pathogenicity. This is highly interesting because this bacterium is an important cause of pneumonia. Moreover, *M. pneumoniae* is one of the organisms with the smallest genetic equipment that is capable of independent life. Understanding *M. pneumoniae* means understanding life! So far, we have studied one of the few regulatory proteins of *M. pneumoniae* and determined its crystal structure. Interestingly, the mode of action of this protein is opposed to that of homologous proteins from other bacteria: a hint to the parasitic lifestyle of *M. pneumoniae*! We are now starting to study the metabolic responses of *M. pneumoniae* to the infection process. If we understand what happens upon infection, we may subsequently try to interrupt this chain of events. Metabolism in *Bacillus subtilis* is studied by transcriptomics, protein arrays, and metabolome and fluxome analyses. Our specific interests are focussed on two key pathways: glycolysis and glutamate biosynthesis, the decisive link between carbon and nitrogen metabolism. We discovered recently that genes for glutamate biosynthesis in *B. subtilis* are only expressed if rich carbon sources are available and we identified a regulatory protein-protein interaction that governs this sugar induction. The regulation of glycolysis is studied at the level of a controlled protein-RNA interaction. Regulation through RNA has become widely recognized in the past few years. In the framework of a national priority program, we will analyze the adaptation of RNA-based regulatory processes in organisms that live at very low or very high temperatures.

Selected Recent Publications

Halbedel S, Busse J, Schmidl S, Stülke J (2006) Regulatory protein phosphorylation in *Mycoplasma pneumoniae*: A PP2C-type phosphatase serves to dephosphorylate HPr(Ser-P). J Biol Chem 281:26253-26259

Schilling O, Herzberg C, Hertrich T, Vörsmann H, Jessen D, Hübner S, Titgemeyer F, Stülke J (2006) Keeping signals straight in transcription regulation: specificity determinants for the interaction of a family of conserved bacterial RNA-protein couples. Nucl Acids Res 34: 6102-6115

Stülke J (2007) Regulation of virulence in *Bacillus anthracis*: the phosphotransferase system transmits the signals. Mol Microbiol 63: 626-628

Halbedel S, Eilers H, Jonas B, Busse J, Hecker M, Engelmann S, Stülke J (2007) Transcription in *Mycoplasma pneumoniae*: Analysis of the promoters of the ackA and ldh genes. J Mol Biol 371: 596-607

Commichau FM, Herzberg C, Tripal P, Valerius O, Stülke J (2007) A regulatory protein-protein interaction governs glutamate biosynthesis in *Bacillus subtilis*: The glutamate dehydrogenase RocG moonlights in controlling the transcription factor GltC. Mol Microbiol 65: 642-654.

Herzberg C, Flórez Weidinger LA, Dörrbecker B, Hübner S, Stülke J, Commichau FM (2007) SPINE: A method for the rapid detection and analysis of protein-protein interactions *in vivo*. Proteomics, in press.

Michael Thumm

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- Center of Biochemistry and Molecular Cell Biology, University of Göttingen
- 1987 Dr. rer. nat., University of Stuttgart
- 1997 Habilitation (Biochemistry), University of Stuttgart

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Major Research Interests

We are studying the molecular mechanism of autophagy in the yeast *Saccharomyces cerevisiae*. Autophagy is a starvation induced transport pahway, which delivers cytosolic material for degradation to the lysosome (vacuole). It is highly conserved in all eukaryots from yeast to human and helps the cells to survive periods of nutrient limitation. Autophagy further plays an important role in ageing, the development of breast cancer and cardiomyopathy and it was linked to neurodegenerative diseases like Alzheimer´s, Huntington´s and Parkinson´s disease. Autophagy is mechanistically unique, since its transport intermediates, the autophagosomes, are surrounded by two individual membranes. It starts at the newly-discovered preautophagosomal structure, where autophagosomes are formed. Autophagosomes unspecifically enclose parts of the cytoplasm including organelles like mitochondria, peroxisomes and parts of the ER. When the autophagosomes reach the vacuole, their outer membrane-layer fuses with the vacuolar membrane and a still membrane-enclosed autophagic body is released into the vacuolar lumen. In the vacuole autophagic bodies are lysed and broken down together with their cytosolic content.
The intravacuolar breakdown of autophagic bodies requires the selective lysis of their limiting membrane. The intracellular lysis of a membrane is a very interesting feature of eukaryotic cells and implies a high risk for cellular integrity. In a genetic screen, we identified Aut5 as an essential component of this lysis process. We found that Aut5 is an integral membrane protein and that its lipase active site motive is essential for lysis of autophagic bodies.

Selected Recent Publications

Thumm M (2002) Hitchhikers guide to the vacuole-mechanisms of cargo sequestration in the Cvt and autophagic pathways. Mol Cell 10: 1257-1258

Epple UD, Eskelinen E-L, Thumm M (2003) Intravacuolar membrane lysis in *Saccharomyces cerevisae*: Does vacuolar targeting of Cvt17/Aut5p affect its function? J Biol Chem 278: 7810-7821

Regelmann J, Schüle T, Josupeit FS, Horak J, Rose M, Entian K-D, Thumm M, Wolf DH (2003) Catabolite degradation of fructose-1,6-bisphosphatase in the yeast *Saccharomyces cerevisiae*: A genome-wide screen identifies eight novel GID genes and indicates the existence of two degradation pathways. Mol Biol Cell 14: 1652-63

Meiling-Wesse K, Barth H, Voss C, Eskelinen EL, Epple UD, Thumm M (2004) Atg21 is required for effective recruitment of Atg8 to the preautophagosomal structure during the Cvt pathway. J Biol Chem 279: 37741-37750

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Group Leader - Bioanalytical Mass Spectrometry Group

- since 2005: Independent research group “Bioanalytical Mass Spectrometry Group” at the Max Planck Institute for Biophysical Chemistry, Göttingen
- 2004, 2005, 2006: Organizer of the 1st, 2nd, and 3rd BMBF Summer School “Proteomic Basics”
- since 2001: Establishment and management of the mass spectrometry in the Department of Cellular Biochemistry at the Max Planck Institute for Biophysical Chemistry, Göttingen
- 2000-2001: Guest researcher at the EMBL, Heidelberg, Protein Analytical Group of Dr. Matthias Wilm
- 2000: Senior scientist in the Department of Cellular Biochemistry at the Max Planck Institute for Biophysical Chemistry, Göttingen
- 1997-2000: Post-Doc in the group of Prof. Dr. Reinhard Lührmann at the Institut für Molekularbiologie und Tumorforschung (IMT) of the Philipps-Universität Marburg
- 1996: Dr. rer. nat. at Faculty of Chemistry, Frei Universität Berlin
- 1993 – 1996: Doctoral thesis project in the group of Prof. Dr. Brigitte Wittmann-Liebold at the Max-Delbrück-Centre of Molecular Medicine, Berlin

Major Research Interests

Modern mass-spectrometric methods are key technologies in the life sciences to elucidate changes at the protein level. Nonetheless, the detection of post-translational modification, reliable MS-quantification procedures, MS-based detection of protein–protein and protein–nucleic acid interactions and, importantly, the identification of proteins that escape detection under standard conditions (e.g., protein isoforms and membrane proteins) are still far from being routine matters.

Our own projects are centered around the establishing of methods for the mass-spectrometric analysis of post-translational modifications and protein-nucleic acid contact sites in ribonucleoprotein (RNPs) particles, such as the spliceosome (collaboration with Reinhard Lührmann at the Max Planck Institute for Biophysical Chemistry (http://www.mpibpc.gwdg.de/english/research/dep/luehrmann/index.html). For that purpose we are developing novel analytical techniques including mass-spectrometric methods (MALDI- and ESI-MS) and chromatographic enrichment strategies.

In collaboration with the Neurobiology Department of Reinhard Jahn at the Max Planck Institute for Biophysical Chemistry (http://www.mpibpc.mpg.de/groups/jahn/), we are developing methods suitable for the reliable MS-based identification of membrane proteins. We use different gel-based purification strategies and liquid-chromatographic approaches to identify novel membrane proteins, for exmple from synaptic vesicles.

Selected Recent Publications

Merz C, Urlaub H, Will CL., Lührmann R (2007) Protein composition of human mRNPs spliced *in vitro* and differential requirements for mRNP protein recruitment. RNA 13: 116-128

Deckert J, Hartmuth K, Boehringer D, Behzadnia N, Will CL, Kastner B, Stark H, Urlaub H, Lührmann R (2006) Protein composition and electron microscopy structure of affinity-purified human spliceosomal B complexes isolated under physiological conditions. Mol Cell Biol 26: 5528-5543

Holt M, Varoqueaux F, Wiederhold K, Takamori S, Urlaub H, Fasshauer D, Jahn R (2006) Identification of SNAP-47, a novel Qbc-SNARE with ubiquitous expression. J Biol Chem 281: 17076-17083

Kuhn-Holsken E, Lenz C, Sander B, Lührmann R, Urlaub H (2005) Complete MALDI-ToF MS analysis of cross-linked peptide-RNA oligonucleotides derived from nonlabeled UV-irradiated ribonucleoprotein particles. RNA 11: 1915-1930

Markus Wahl

Ph.D. - Group Leader at the Max Planck Institute for Biophysical Chemistry

- 1996 Ph.D., The Ohio State University, Columbus, OH, USA
- 1997 - 2002 Post-Doc, Max-Planck-Institute for Biochemistry, Martinsried, Germany
- 2002 Habilitation and Privatdozentur, Technical University München, Faculty of Chemistry, München, Germany
- Since 2002 group leader, Max Planck Institute for Biophysical Chemistry

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Major Research Interests

Gene expression pathways encompass a number of RNA metabolic steps such as transcription, splicing, editing, post-transcriptional nucleotide modification, nuclear export, RNA surveillance, cytoplasmic localization, translation and RNA degradation, each of which is orchestrated by a multi-component cellular machine. In a traditional view, RNAs are shuttled along simple linear arrays of such steps that all represent discrete events. However, in recent years it became evident that the various machineries are extensively interconnected and that the route of a given RNA through this network of processes is determined by its particular set of decorating proteins, RNAs or RNA-protein complexes. The general interest of our group lies in exploring the molecular mechanisms underlying the modes of action and functional coupling of RNA-protein machineries involved in these gene expression networks. In particular, we are interested in the regulation of termination and antitermination of transcription by RNA polymerase, in the inner workings of the spliceosome and its constituents and in selected aspects of translation. We employ X-ray crystallography to elucidate the structures of proteins, RNAs and macromolecular complexes, which are part of these molecular machines. Structure-guided hypotheses are then tested by biochemical, molecular biological and genetics assays.

Selected Recent Publications

Liu S, Li P, Dybkov O, Nottrott S, Hartmuth K, Lührmann R, Carlomagno T, Wahl MC (2007) Binding of the human Prp31 Nop domain to a composite RNA-protein platform in U4 snRNP. Science 316: 115-20

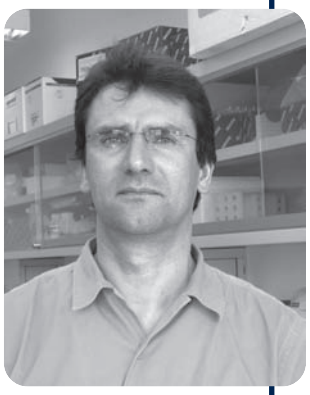
Pena V, Liu S, Bujnicki JM, Lührmann R, Wahl MC (2007) Structure of a multipartite protein-protein interaction domain in splicing factor Prp8 and its link to Retinitis pigmentosa. Mol Cell 25: 615-24

Spadaccini R, Reidt U, Dybkov O, Will C, Frank R, Stier G, Corsini L, Wahl MC, Lührmann R, Sattler M (2006) Biochemical and NMR analyses of an SF3b155-p14-U2AF-RNA interaction network involved in branch point definition during pre-mRNA splicing. RNA 12: 410-25

Diaconu, M, Kothe, U, Schlünzen, F, Fischer, N, Harms, JM, Tonevitsky, A, Stark, H, Rodnina, MV, Wahl, MC (2005) Structural basis for the function of the ribosomal L7/12 stalk in factor binding and GTPase activation. Cell 121: 991-1004

Bonin I, Mühlberger R, Bourenkov GP, Huber R, Bacher A, Richter G, Wahl MC (2004) Structural basis for the interaction of *Escherichia coli* NusA with protein N of phage Lambda. Proc Natl Acad Sci USA 101: 13762-13767

Lutz Walter



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Head of Department of Primate Genetics at the German Primate Center

- Dr. rer. nat. (PhD), University of Göttingen, 1994
- Postdoctoral fellow and group leader at the Division of Immunogenetics, University of Göttingen, 1994 - 2004
- Head of Department of Primate Genetics, German Primate Center, Göttingen, since 2004
- Habilitation (Immunology and Immunogenetics), Medical Faculty of the University of Göttingen, 2005

Major Research Interests

The main interests of the laboratory are immunology, the genetic control of immune responses, functional genomics, molecular evolution, and population genetics. The research is focussed on the functional, evolutionary, and genomic analysis of genes of the major histocompatibility complex (MHC) as well as the natural killer cell receptor and leukocyte receptor complexes (NKC, LRC). The analyses are carried out in various organisms that are used as models of human diseases such as certain nonhuman primates (rhesus macaque, common marmoset) and rodents (mouse, rat). Functional studies aim at elucidating the role of certain genes of the MHC, NKC, and LRC in innate and adaptive immunity.

In a further research focus, the molecular evolution and population genetics of various nonhuman primate taxa are analyzed on the basis of molecular data. These studies aim at clarifying the phylogenetic relationship of various primates (molecular phylogeny) and to determine their geographic distribution (phylogeography), particularly of highly endangered primate species (conservation biology).

Selected Recent Publications

Averdam A, Kuhl H, Sontag M, Becker T, Hughes AL, Reinhardt R, Walter L (2007): Genomics and diversity of the common marmoset monkey natural killer complex (NKC). J Immunol 178: 7151-7161

Walter L (2007): Pas de deux: natural killer receptors and MHC class I ligands in primates. Curr Genomics 8: 51-57

Rölleke U, Flügge G, Plehm S, Schlumbohm C, Armstrong VW, Dressel R, Uchanska-Ziegler B, Ziegler A, Fuchs E, Czeh B, Walter L (2006) Differential expression of major histocompatibility complex class I molecules in the brain of a New World monkey, the common marmoset (*Callithrix jacchus*). J Neuroimmunol 176: 39-50

Kelley J, Walter L, Trowsdale J (2005) Comparative genomics of natural killer cell receptor gene clusters. PLoS Genetics 1: e27

Roos C, Dressel R, Schmidt B, Günther W, Walter L (2005) The rat expresses two complement factor C4 proteins, but only one isotype is expressed in the liver. J Immunol 174: 970-975

Hurt P, Walter L, Sudbrak R, Klages S, Müller I, Shiina T, Inoko H, Lehrach H, Günther E, Reinhardt R, Himmelbauer H (2004) The genomic sequence and comparative analysis of the rat major histocompatibility complex. Genome Res 14: 631-639

Jürgen Wienands

Professor of Cellular and Molecular Immunology

- 1982-89 Study of Biology at the University of Cologne; graduated at the Institute of Genetics, Dept. of Immunology
- 1989-92 Ph.D. poject at the Max Planck Institute for Immunobiology, Freiburg, Germany
- 1992-94 Postdoctoral fellow at the Dept. of Preclinical Research at Sandoz Pharma Ltd., Basel, Switzerland
- 1994-96 Postdoctoral fellow at the Max Planck Institute for Immunobiology, Freiburg, Germany
- 1996-2001 Group leader at the University of Freiburg, Institute of Biology III
- 2001 “Habilitation” and Venia Legendi in “Molecular Immunology and Biochemistry”
- 2001-2004 Full Professor for “Biochemistry and Molecular Immunology” at the University of Bielefeld
- since August 2004 Full Professor for “Molecular and Cellular Immunology” at the University of Göttingen

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Major Research Interests

The signature structure of B lymphocytes is their clonotypic antigen receptor (BCR). Our major research focuses on the elucidation of intracellular BCR signaling pathways that regulate the development and activation of B cells in health and disease. We have identified enzymatically inert adaptor proteins such as SLP-65 (for: SH2 domain-containing leukocyte adaptor of 65 kDa), which nucleate the formation of multi-molecular protein complexes to integrate and amplify BCR signals. A key function of these signaling modules is to orchestrate the mobilization of the second messenger Ca^{2+}. Interference with expression and/or function of one the signaling components can cause severe immunodeficiencies in mouse and man. Moreover, viruses such as the Epstein-Barr virus (EBV) abuse BCR effector proteins to reorganize signaling cascades for their own benefit. Biochemical and genetic methods, which are applied to study these events *in vitro* and *in vivo*, include protein purification by affinity chromatography and immunoprecipitation, analysis of protein interactions, flow cytometry, targeted gene disruption in cell culture and embryonic stem cells followed by reconstitution experiments using electroporation techniques or retroviral gene transfer.

Selected Recent Publications

Grabbe A, Wienands J (2006) Human SLP-65 isoforms contribute differently to activation and apoptosis of B lymphocytes. Blood 108: 3761-3768

Connert S, Wienand S, Thiel C, Krikunova M, Glyvuk N, Tsytsyura Y, Hilfiker-Kleiner D, Bartsch JW, Klingauf J, Wienands J (2006) SH3P7/mAbp1 deficiency leads to tissue and behavioral abnormalities and impaired vesicle transport. EMBO J 25(8): 1611-22

Stork B, Engelke M, Frey J, Horesjsí V, Hamm-Baarke A, Schraven B, Kurosaki T, Wienands J (2004) Grb2 and the non-T cell activation linker NTAL constitute a Ca^{2+}-regulating signal circuit in B lymphocytes. Immunity 21: 681-691

Wakabayashi C, Adachi T, Wienands J, Tsubata T (2002) A distinct signaling pathway used by the IgG-containing B-cell antigen receptor. Science 298: 2392-2395

Engels N, Merchant M, Pappu R, Chan AC, Longnecker R, Wienands J (2001) Epstein-Barr virus LMP2A employs the SLP-65 signaling module. J Exp Med 194: 255-264

Wienands J (2000) The B cell antigen receptor: Formation of signaling complexes and the function of adaptor proteins. Current Topics Microbiol Immunol 245: 53-76

Ernst Wimmer

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Professor of Developmental Biology

- 1991 Diplom (Biology), Ludwig Maximilians University, Munich (Germany)
- 1995 Dr. rer. nat., Max-Planck-Institute for Biophysical Chemistry, Göttingen (Germany) and Howard Hughes Medical Institute, Baylor College of Medicine, Houston (USA)
- 1995 - 1998 Postdoctoral Fellow and Associate, Howard Hughes Medical Institute, The Rockefeller University, New York (USA)
- 1998 - 2003 Assistant Professor and Robert Bosch Foundation 'Junior Professor' Department of Genetics, University of Bayreuth, Bayreuth (Germany)
- Since 2003 Professor of Developmental Biology at the Johann Friedrich Blumenbach Institute of Zoology and Anthropology, Georg August University, Göttingen (Germany)

Major Research Interests

A key question in developmental biology is how diverse animal body plans are specified. Early developmental decisions determine the coordinates of the embryo and activate the genetic circuitry that sequentially subdivides and regionalizes the animal body. For insects, only in *Drosophila* the early developmental events are known in molecular detail. However, insects with varied life histories must compensate different reproductive strategies by adjusting the regulatory networks within the developmental program. Therefore, phylogenetic differences between diverse species must be manifested in the genetic circuitries regulating embryogenesis.

To identify the plasticity in early developmental processes, we study their conservation and divergence in different arthropod species. Developmental regulatory genes code for signal transduction molecules and transcription factors. But of equal importance to the coding part of these genes are their cis-regulatory sequences, which serve as integration points for originally distinct signals. By insect transgenesis and functional genomics approaches, we analyze genetic interactions within the regulatory network of early embryogenesis in diverse insect species. This will help us to understand how animal evolution is based on changes in gene regulation governing early pattern formation.

Furthermore we apply our knowledge about developmental processes to insect pest management. Current control efforts rely mostly on insecticides, but the costs for developing new chemical products to overcome the problem of insecticide resistance are escalating. Genetic control based on the sterile-insect technique (SIT) uses the release of sterile males to cause infertile matings which reduce pest population levels. Due to the species specificity, SIT is considered an ecologically safe procedure. However, conventional sterilization by ionizing radiation also decreases the competitiveness of sterilized males. To overcome this problem, we design transgenic approaches to selectively produce vigorous and potent sterile males by generating conditional male sterility in combination with conditional female lethality.

Selected Recent Publications

Wimmer EA, Carleton A, Harjes P, Turner T, Desplan C (2000) Bicoid-independent formation of thoracic segments in *Drosophila*. Science 287: 2476-2479

Ito J, Ghosh A., Moreira LA, Wimmer EA, Jacobs-Lorena M (2002) Transgenic anopheline mosquitoes impaired in transmission of a malaria parasite. Nature 417: 452-455

Horn C, Wimmer EA (2003) A transgene-based embryo-specific lethality system for insect pest management. Nature Biotechnology 21: 64-70

Häcker U, Nystedt S, Padash-Barmchi M, Horn C, Wimmer EA (2003) piggyBac-based insertional mutagenesis in the presence of stably integrated P elements in *Drosophila*. PNAS 100: 7720-7725

Wimmer EA (2003) Applications of Iinsect transgenesis. Nature Reviews Genetics 4: 225-232

Andreas Wodarz

Professor of Stem Cell Biology

- Diploma Biology, University of Cologne, 1990
- Dr. rer. nat. Developmental Biology, University of Cologne, 1993
- Postdoc, Howard Hughes Medical Institute, Stanford University, 1994 - 1997
- Junior Group Leader, Heinrich Heine University Düsseldorf, 1997 - 2004
- Habilitation in Genetics, Heinrich Heine University Düsseldorf, 2001
- Appointed as Head of the Department of Stem Cell Biology at the University of Göttingen, 2004

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Major Research Interests

At the center of my research interests is the question of how neural stem cells divide asymmetrically to produce another stem cell and a progenitor cell that will differentiate and give rise to neurons and glia cells. One important aspect of asymmetric cell division is the establishment of an intrinsic polarity which is the prerequisite for the asymmetric localization of proteins and mRNAs that serve as cell fate determinants. Our model system for the asymmetric division of stem cells is the embryonic neuroblast of Drosophila. Here we study the function of genes that control cell polarity, asymmetric localization of cell fate determinants and orientation of the mitotic spindle. The knowledge obtained in the Drosophila system has stimulated intense research on the participation of the orthologous genes and proteins in the asymmetric division of vertebrate stem cells.

Selected Recent Publications

Wodarz A, Stewart DB, Nelson WJ, Nusse R (2006) Wingless signaling modulates cadherin-mediated cell adhesion in *Drosophila* imaginal disc cells. J Cell Sci 119: 2425-2434

Wodarz A (2005) Molecular control of cell polarity and asymmetric cell division in *Drosophila* neuroblasts. Curr Opin Cell Biol 17: 475-481

von Stein W, Ramrath A, Grimm A, Müller-Borg M, Wodarz A (2005) Direct association of Bazooka/PAR-3 with the lipid phosphatase PTEN reveals a link between the PAR/aPKC complex and phosphoinositide signaling. Development 132: 1675-1686

Wodarz A, Ramrath A, Grimm A, Knust E (2000) *Drosophila* atypical protein kinase C associates with Bazooka and controls polarity of epithelia and neuroblasts. J Cell Biol 150: 1361-1374

Wodarz A, Ramrath A, Kuchinke U, Knust E (1999) Bazooka provides an apical cue for Inscuteable localization in *Drosophila* neuroblasts. Nature 402: 544-547

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