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IS-MPMI Directory

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IS-MPMI REPORTER DEADLINE
Deadline for submitting items
for the next issue is August 15,
2009.

Submission of materials as electronic files, on disk or as e-mail attachments, will speed processing. For information on submitting electronic images, contact Joel Berg at jberg@scisoc.org.

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Rendez-vous à Québec! XIV International Congress
on Molecular Plant-Microbe Interactions Preview

This July 19–23, attendees from 46 countries around the world will gather to present their research, network, and discuss the future of molecular genetics and molecular biology at the XIV International Congress on Molecular Plant-Microbe Interactions in Québec, Canada. IS-MPMI, together with the Local Organizing Committee in Québec, have put together an outstanding program, featuring an impressive list of confirmed speakers, as well as networking opportunities and unique cultural events. As the chair of the Local Organizing Committee, I could not be more proud of the program we've put together and I'm excited to invite you to join us in Québec.

More than 800 abstracts have been submitted for the XIV Congress, ensuring that there will be something for everyone in the scientific and technical program. The program will also include eight plenary sessions and 18 concurrent sessions on topics related to pathogenic interactions, symbiotic interactions, common host mechanisms, signaling and molecular dialogues, dynamics of plant responses to microbes, plant-microbe interactions and technology transfer, RNA silencing, and plant immunity.

This year's program also includes a special guest lecture on Sunday, July 19. **Luis Sequeira** will present the opening lecture, "Facing the Issues in Agricultural Biotechnology." Sequeira, University of Wisconsin-Madison, will address the importance of our society's role in developing translational research in plant diseases and how IS-MPMI can become a leader in agricultural biotechnology.

The program also includes ample time to take part in networking and cultural activities. In these lean economic times, face-to-face meetings with other scientists in the field are more critical than ever and will help ensure that your career is on the right path. There is no better opportunity to meet top-level speakers and researchers within specific areas of interest than at the XIV Congress. Networking opportunities at the meeting include the Welcome Reception on Sunday, a two-hour boat cruise on the St. Lawrence River on Wednesday afternoon, and the Final Banquet and Show on Thursday evening. These activities provide an important venue for making personal and professional connections that will last throughout your career.

Finally, Québec offers beautiful July weather, historic charm, and exciting cultural and outdoor activities for you and your guests at the meeting. Visit the congress website at www.mpmi2009.ulaval.ca for complete information on the program, accommodations, travel, registration, and more. We're looking forward to seeing everyone in Québec!

Hani Antoun
Chair, Local Organizing Committee



Photo courtesy of J.F. Bergerson, Enviro Foto

A Message from the President



Federico Sanchez

The XIV IS-MPMI Congress to be held in Québec City is only a few weeks ahead and **Hani Antoun**, the congress organizer, and the Local Organizing Committee are ready to receive participants from more than 30 countries, making it possible once again to gather together members of our society. Multiple benefits are also ahead of us, including hearing first-hand science on the cutting edge, establishing new collaborations, and last but not least, making new friends and meeting dear colleagues in the diverse field of plant-microbe interactions. All of this will happen in one of the most emblematic and charming venue cities of the Northern Hemisphere.

A sign of robustness and driving vitality is evident when a scientific community is able to attract and congregate a substantial number of graduate students and young scientists to do stimulating work in their fields of competence, assuring both its prevalence and evolution. In this regard, it is rewarding and extremely pleasing to know that, at the XIV edition of the congress, almost 50% of the attendees will be students and post-docs working in different fields of plant-microbe interactions. In light of the increasing interest of young scientists in the different aspects of plant-microbe interactions, it is vital that authorities support research in new developments, create opportunities by opening positions and scholarships for young scientists, and commend the renewed interest, creativity, and professionalism of those professors and mentors who teach and lead with enthusiasm and scientific rigor, while unraveling the frontiers of knowledge. One example of such a remarkable person, is the outstanding professor, mentor, and scientist, **Jeff Dangl**.

This year, IS-MPMI is proud to honor Dangl, one of its most distinguished members, for his pioneering and passionate commitment to unraveling the puzzling mysteries of signaling pathways that plants use in defense of pathogens with this year's IS-MPMI Award. In 2009, he also received the important Stephen Hales Prize from the American Society of Plant Biologists. (See adjacent article for more on Dangl and this notable award.)

My term as president of IS-MPMI is coming to an end at the XIV Congress, and it has been an honor and a great pleasure to serve our society. I would like to express my deepest gratitude for the help and continuous support received from the staff members of IS-MPMI and my colleagues from the Board of Directors. I wish most success to the new president of IS-MPMI, **Felice Cervone**.

Federico Sanchez
President

2009 IS-MPMI Award Winner Announced

Jeff Dangl, a plant-genome scientist and the John N. Couch Professor in the Department of Biology at the University of North Carolina (UNC), Chapel Hill, is this year's recipient of the IS-MPMI Award. The award, only the second to be given, honors outstanding innovative research. The first award went to **Thomas Boller**, Institute of Botany, University of Basel, Switzerland, in 2007.

This year's winner joined UNC in 1995, after receiving a B.S. degree in biology, a B.A. degree in modern literature (1981), and M.S. and Ph.D. degrees (1986) from Stanford University. He is an elected member of the U.S. National Academy of Sciences, a fellow in the American Association for the Advancement of Science, and associate director of the Carolina Center for Genome Science. He served on the *MPMI* Editorial Board for 11 years. This self-proclaimed "hard-core geneticist" focuses on host-pathogen interactions, mainly looking at how plants recognize and fend off pathogens; overall, how to make plants more disease resistant.

His lab mostly uses *Arabidopsis thaliana* (thale cress)—the first plant genome to be sequenced—to study disease resistance. The lab's research centers on 1) understanding the structure and function of plant NB-LRR disease resistance proteins, especially RPM1; 2) the molecular control of hypersensitive programmed cell death that accompanies disease resistance responses; and 3) understanding the molecular mechanism by which pathogenic bacteria cause disease in plants using diverse suites of type III effector proteins.

Dangl has made significant contributions to the understanding of plant defense from pathogens and the molecular basis of the innate immune response in plants. For these and other important contributions to the field of molecular-plant interactions and following an internal vote process by the IS-MPMI Board of Directors, Dangl was elected as the recipient of the 2009 award.

Read more about the award and the previous awardee's work at www.ismpminet.org/members/award.asp.

XIV International Congress on Molecular Plant-Microbe Interactions

PRELIMINARY PROGRAM

Speaker listings are preliminary and subject to change.



XIV
International Congress
On Molecular
Plant-Microbe Interactions
July 19-23, 2009, Quebec City, Canada

Sunday, July 19

- 12:00 – 15:00 Board of Directors Meeting
12:00 – 20:00 Registration Open
15:00 – 19:00 Exhibitor Set-up
15:00 – 19:00 Poster Set-up (authors place posters in Exhibit Hall)
18:00 – 20:00 **Opening Welcome and Address by Luis Sequeira and Award Presentation**
20:00 – 22:30 **Welcome Reception**

Monday, July 20

- 7:00 – 9:00 Cash Breakfast Concessions Open
7:00 – 16:30 Registration
7:00 – 23:00 Poster Viewing
8:00 – 10:00 **Plenary I**—PATHOGENIC INTERACTIONS, P. de Wit, N. Talbot, R. Kahmann, and P. Dodds
10:00 – 10:30 Coffee Break
10:30 – 12:30 **Plenary II**—SYMBIOTIC INTERACTIONS, S. Long, D. Bogusz, M. Kahn, and U. Mathesius
12:30 – 14:00 Lunch Break with Exhibits/Poster Viewing—*cash concessions available*
14:00 – 16:00 **Concurrent 1**—Secondary Metabolites
14:00 – 16:00 **Concurrent 2**—Symbiotic Interactions
14:00 – 16:00 **Concurrent 3**—ROS/NO Signaling and PCD
16:00 – 16:30 Coffee Break
16:30 – 18:30 **Concurrent 4**—Quorum Sensing
16:30 – 18:30 **Concurrent 5**—Molecular Dialogues and Signaling I
16:30 – 18:30 **Concurrent 6**—Plant Cell Wall Modifications
18:30 – 20:00 Dinner Break
20:00 – 23:00 Exhibits Open and Poster Viewing
20:00 – 21:30 Odd-Numbered Poster Authors Present

Tuesday, July 21

- 7:00 – 9:00 Cash Breakfast Concessions Open
7:00 – 16:30 Registration
7:00 – 23:00 Poster Viewing
8:00 – 10:00 **Plenary III**—COMMON HOST MECHANISMS, J. Jones, S.-Y. He, M. Harrison, and K. Szczygłowski
10:00 – 10:30 Coffee Break
10:30 – 12:30 **Plenary IV**—SIGNALING AND MOLECULAR DIALOGUES, B. Staskawicz, R. Innes, U. Bonas, and F. Cervone
12:30 – 14:00 Lunch Break with Exhibits/Poster Viewing—*cash concessions available*
14:00 – 16:00 **Concurrent 7**—Molecular Dialogues and Signaling II
14:00 – 16:00 **Concurrent 8**—Genomics of Diazotrophic Plant-Associated Bacteria
14:00 – 16:00 **Concurrent 9**—Theory and Practice of Biocontrol
16:00 – 16:30 Coffee Break
16:30 – 18:30 **Concurrent 10**—Interactions with Invertebrates
16:30 – 18:30 **Concurrent 11**—Gene Silencing
16:30 – 18:30 **Concurrent 12**—Practical Application Bioinoculants
18:30 – 20:00 Dinner Break
20:00 – 23:00 Exhibits Open and Poster Viewing
20:00 – 21:30 Even-Numbered Poster Authors Present

Wednesday, July 22

- 7:00 – 9:00 Cash Breakfast Concessions Open
7:00 – 14:00 Poster Viewing
7:00 – 16:00 Registration
8:00 – 10:00 **Plenary V**—DYNAMICS OF PLANT RESPONSES TO MICROBES, E. Kombrink, B. Mauch-Mani, R. Wise, and S. Kamoun

Preliminary Program continued on page 4

Preliminary Program *continued from page 3*

- 10:00 – 10:30 Coffee Break
10:30 – 12:30 **Plenary VI**—PLANT-MICROBE INTERACTIONS AND TECHNOLOGY TRANSFER, R. Belanger, L. Matteo, D. Gonsalves, and J. Loper
12:30 – 14:00 Lunch Break—*cash concessions available*
Exhibits Open
14:00 – 15:30 Poster Take-down
14:00 – 20:00 Exhibitor Move-out
15:30 – 17:30 Boat Cruise
17:30 Dinner Break and Free Evening

Thursday, July 23

- 7:00 – 9:00 Cash Breakfast Concessions Open
7:00 – 16:30 Registration
8:00 – 10:00 **Concurrent 13**—Effector Proteins and the Endocytic Pathway
8:00 – 10:00 **Concurrent 14**—Plant-Virus Interactions
8:00 – 10:00 **Concurrent 15**—Microbe-Induced Plant Organogenesis
10:00 – 10:30 Coffee Break
10:30 – 12:30 **Concurrent 16**—Tree-Microbe Interactions
10:30 – 12:30 **Concurrent 17**—Model Organisms and the Post-Genomic Area
10:30 – 12:30 **Concurrent 18**—Emerging Technologies
12:30 – 14:00 Lunch Break—*cash concessions available*
14:00 – 16:00 **Plenary VII**—RNA SILENCING, O. Voinnet, H. Jin, M. Crespi, and P. Moffett
16:00 – 16:30 Coffee Break
16:30 – 18:30 **Plenary VIII**—PLANT IMMUNITY AND CLOSING CEREMONY, P. Schulze-Lefert, J. Parker, X. Dong, and S. Dinesh-Kumar
19:30 – 24:00 **Final Banquet and Show**

Concurrent sessions and invited speakers who have confirmed their participation to date:

P. Birch, P. Bonfante, W. J. Broughton, J. Burgyan, X. Chen, E. L. Davis, J. Ellis, M. T. Esquerré-Tugayé, D. Faure, E. Giraud, M. Grant, R. Hammerschmidt, A. Herrera-Estrella, A. Hirsch, M. Hofte, S. A. Hogenhout, E. Kondorosi, K. Korth, M. Leggett, X. Li, C. Lo, M. Parniske, A. Pühler, G. Reuter, S. Robatzek, A. Seguin, K. Shimamoto, S. Smith, G. Stacey, J. Stenlid, J. Walton, and D. Wendehenne.

Xanthomonas Genomics Conference



Xanthomonas Genomics Conference 2009 will assemble scientists from around the world who are engaged in research to better understand how this important group of bacteria has adapted to exploit an extraordinary diversity of plant hosts and host tissues; develop needed improvements in disease control and prevention; and understand and better harness the capacity of members of this genus to produce useful compounds. The goals of the conference are to share and discuss recent discoveries; identify important challenges and opportunities; discuss new technologies and approaches; and determine key resources for continued, community-driven advances. The conference will be held at the beautiful and bucolic Pingree Park Campus of Colorado State University. Registration is \$200; \$100 for students. Reasonable housing packages are available on a first-come basis. For more information, please see www.plantpath.iastate.edu/xgc2009 or contact **Adam Bogdanove**, chair Iowa State University (ajbog@iastate.edu); **Jan Leach**, Colorado State University (jan.leach@colostate.edu); or **Frank White**, Kansas State University (fwhite@ksu.edu).

Visit IS-MPMI Central at the 2009 IS-MPMI Congress!

IS-MPMI Central is your place to learn about everything IS-MPMI has to offer. Speak with IS-MPMI Headquarters staff to learn about your many member benefits.

Stay on top of leading research with *MPMI* Online. Learn about how *MPMI* Online can save you time when researching and sign up for a saved search alert to receive a FREE 2GB *MPMI* flash drive (while supplies last). Staff will be on hand to help you create or update your online profile. Take advantage of 10–30% discounts when you purchase books published or recommended by IS-MPMI.

IS-MPMI connects you with scientists and organizations around the world. You will have the opportunity to share your thoughts on the society and the congress and to meet some of the IS-MPMI board members. Look for jobs or post a position opening in your institution on the Job Board.

Find out everything IS-MPMI can do for you and your research! Be sure to visit IS-MPMI Central in the entry hall of the Québec City Convention Center.

St. Louis Hosts 9th International Plant Molecular Biology Congress

Elite scientists from around the world will gather in St. Louis, MO, U.S.A., for one of the most prestigious plant research events—the 9th International Plant Molecular Biology (IPMB) Congress. Scheduled for October 25–30, 2009, the congress is making its debut in St. Louis, returning to the United States after being hosted abroad for more than 20 years.

Internationally renowned scientists will be hosted here by **Perry Gustafson**, professor of plant sciences at the University of Missouri in Columbia; Monsanto's Chief Technology Officer **Robert T. Fraley**; and **Peter Raven**, president of the Missouri Botanical Garden.

The congress will feature more than 50 forums, abstract poster displays and presentations, evening workshops, and other events to chronicle the advances in plant molecular biology. Topics will include agricultural productivity, phytotherapy for human health, generating biofuels, and the future of biodiversity, further reinforcing the need to find solutions for today's global challenges.

"The Congress is considered one of the most important events to share worldwide progress in discovering the potential that plants hold to help solve the issues facing the planet," says Gustafson. "Our guest speakers are discovering solutions for fuel, food, and medicine and will offer a fascinating look into our sustainable future."

The IPMB Congress has already secured commitments from more than 300 scientists who will present during the course of the event. Registration for the 2009 IPMB Congress is now open online. Please visit www.ipmb2009.org for more information.

Receive a FREE MPMI 2GB Flash Drive



Receive a **FREE MPMI** 2GB flash drive when you set up a saved search in *MPMI* Online during the IS-MPMI Congress in Québec City. You can easily customize online content from *MPMI* to match your interests. Learn more during the IS-MPMI Congress, where staff will be available to help you create your profile in *MPMI* Online. Set up a saved search and receive a FREE *MPMI* 2GB flash drive (while supplies last)! Visit IS-MPMI Central in the entry hall of the Québec City Convention Center.

Funding Opportunity: The National Science Foundation and the Gates Foundation Announce BREAD Program



BILL & MELINDA
GATES foundation

The National Science Foundation (NSF) has announced a nearly \$50 million partnership with the Bill & Melinda Gates Foundation to support innovative solutions to critical agricultural challenges in developing countries. Each organization will provide \$24 million over five years to support a competitive awards program for science research projects addressing drought, pests, disease, and other serious problems facing small farmers and their families who rely on their crops for their food and income.

The award program—Basic Research to Enable Agricultural Development (BREAD)—will support a competitive award program for science research projects developing innovative approaches and technologies to boost agricultural productivity in developing countries.

The NSF will manage the program using its peer-review process in consultation with the Gates Foundation. NSF funds will be used to support research carried out in eligible U.S. academic institutions, such as nonprofit research organizations, including museums, research laboratories, and professional societies. The Gates Foundation's funds will be used to support eligible international partners via subawards from the U.S. awardees.

BREAD program solicitations will be nonprescriptive, inviting a broad scope of applications. A solicitation for funding proposals under the BREAD program will be available in early June on the BREAD website, accessible through www.nsf.gov/funding/pgm_summ.jsp?pims_id=5338&org=BIO.

This funding program is part of the Gates Foundation's Agricultural Development Initiative. The foundation is working to strengthen the entire agricultural value chain—from seeds and soil to farm management and market access—so that progress against hunger and poverty is sustainable over the long term.

Meet IS-MPMI Members

IS-MPMI's diverse membership spans the globe and includes professionals who have been in their field for decades, as well as those who are just starting out. To help members learn more about their colleagues, the *IS-MPMI Reporter* includes profiles of randomly chosen members at different career stages.



Caroline Gutjahr

Student Member

Caroline Gutjahr

University of Lausanne
Lausanne, Switzerland

I studied biology at the University of Freiburg (Germany) and the University of Aberdeen (Scotland) with a main focus on plant physiology and minors in microbiology, tropical ecology,

and philosophy. My diploma (equivalent to an M.Sc.) finished with a thesis in the lab of **Peter Nick** at the University of Freiburg (now Karlsruhe). I studied the role of phytohormone gradients and the cytoskeleton in gravitropism of rice coleoptiles and found that the classically well-known auxin gradient, first proposed by Cholodny and Went, is opposed by a gradient of jasmonic acid during gravitropism. During undergraduate courses in tropical ecology at Aberdeen, I had been introduced to the importance of rhizobial and mycorrhizal symbioses for plant nutrition, especially on poor soils. A research expedition to Sri Lanka shortly after strengthened the impression these courses left on me. I saw how farmers recovered soils that were badly degraded because of intensive tea cultivation for small-scale, self-subsistence agriculture. They induced an artificial succession by planting legume trees, which are hosts to rhizobia and arbuscular mycorrhizal fungi. Enriching the soil with nutrients in this way, they prepared the ground for more demanding crops which were planted a few years later. Although I am fascinated by the potential of plant root symbioses to help establish sustainable agricultural practices, I am mostly interested in basic research and I started to ask myself questions, such as "How can a plant distinguish between pathogenic and symbiotic microbes and react accordingly?" and "What are the signals and mechanisms that lead to reprogramming of host cells allowing changes in shape (as during formation of an arbuscule) or formation of a new organ (like a nodule) to occur?"

I decided to work on the arbuscular mycorrhizal symbiosis and joined the well-established lab of **Paola Bonfante** at the University of Turin (Italy) to study physiological changes occurring in roots of the model legume *Lotus japonicus* in response to presymbiotic signals exuded by arbuscular mycorrhizal fungi. After having spent one year in Italy, I started a Ph.D. thesis (supported by fellowships from the German National Academic Foundation and the Roche Foundation, Switzerland) in the relatively young lab of **Uta Paszkowski** at the University of Geneva (Switzerland) and moved with her to the University of

Lausanne (Switzerland) one year later. The initial phase of my Ph.D. degree turned out to be quite exciting, as we not only set up a new lab but also established rice as a new model plant for mycorrhiza research. Using qPCR and in situ hybridization, I first developed a set of robust marker genes in rice that could be used as readouts for arbuscular mycorrhiza-specific signaling when characterizing rice mutants or specific physiological scenarios. I found that 18 rice genes, which had emerged from a microarray study previously performed in Paszkowski's lab, were exclusively expressed in mycorrhizal rice roots and fell into several classes of spatio-temporal expression patterns during arbuscular mycorrhizal colonization. Two of these marker genes encode putative small apoplastic proteins containing an LysM domain. This domain is known to bind chitin oligomers and is present in legume Nod-factor receptors, as well as in Arabidopsis and rice receptors for chitin elicitors released by pathogenic fungi. Also, arbuscular mycorrhizal fungi are predicted to release chitin oligomers of some sort, for example, during hyphal growth. Being very intrigued by the possible function of these LysM domain-containing proteins, we created transgenic rice perturbed in expression of these genes via RNAi-mediated silencing and over-expression and are currently analyzing the transgenic lines. As a second step, I determined how far signaling components known to be required for the mycorrhizal symbiosis in dicotyledonous legumes are conserved in the phylogenetically distant monocotyledonous rice. Taking advantage of the extensive reverse genetics collections of rice insertion mutants, I could demonstrate (in parallel with the labs of **Haruko Imaizumi-Anraku**, Japan, and **Hongyan Zhu**, United States) that four components (CASTOR, POLLUX, CCAMK, and CYCLOPS) of the legume common SYM pathway are conserved in rice. In mutants of all four signaling components, arbuscular mycorrhizal colonization stops at the rhizodermis and the cortex is never colonized. Nevertheless, the mutants still express some of the above-mentioned marker genes. The gene expression pattern differs between the mutants depending on their position within the common SYM pathway. We concluded that mycorrhizal-specific signaling must be a complex signaling network with signaling cues that should be independent and such that are deviating from the main common SYM-signaling pathway. Using transcriptomics as readout for underlying signaling events, I am currently investigating more deeply into some recently discovered peculiarities of common SYM signaling in rice.

I joined IS-MPMI after the congress in Mérida, Mexico, where the temples of *Uxmal* remained "in the dark and silence" but the scientific part of my mind did not. I perceive that IS-MPMI is a great platform for information exchange in our rapidly evolving field.



Akiko Sugio

Post-Doctoral/Early Career Member

Akiko Sugio

John Innes Centre
Norwich, United Kingdom

During my studies, plant-microbe interactions were not my focus. I was interested in the application of science in industries, and I took a job in Novo Nordisk Japan, currently

known as Novozymes Japan, just after I obtained a master's degree in biotechnology at the Tokyo Institute of Technology, Japan. The company specialized in producing industrial enzymes, and my project was the cloning, expression, and characterization of the genes that encode enzymes. I enjoyed the new environment and my work, but gradually I was attracted to the world of plant-microbe interactions because I cloned a few genes from plant-pathogenic fungi and wondered about their functions in nature.

In 2001, I started a Ph.D. degree in plant pathology at Kansas State University, U.S.A., as a Tillman scholar. The reasons I went to the United States were to improve my English skills and to get financial support, which we didn't normally get in Japan at that time. I joined **Frank White's** lab and studied the interaction of rice and *Xanthomonas oryzae* pv. *oryzae*. In the beginning, I focused on the bacteria side and studied *hrpF* in *X. oryzae* pv. *oryzae*. Although HrpF was known as a translocator of type III secretion system (TTSS), we found that HrpF was not absolutely necessary for *X. oryzae* pv. *oryzae* to secrete avirulence proteins and trigger hypersensitive response.

Next, I studied rice responses to *X. oryzae* pv. *oryzae* ingress. I examined several rice genes that are induced by *X. oryzae* pv. *oryzae* infection, and with **Bing Yang** (he became an assistant professor at Iowa State University) in the lab, we discovered that the activities of specific *X. oryzae* pv. *oryzae* virulence factors were correlated with the inductions of corresponding rice genes. We have also shown that the induction of those rice genes made rice more susceptible to some *X. oryzae* pv. *oryzae* strains. The *X. oryzae* pv. *oryzae* virulence factors have a common structure that looks like a transcription activator and are secreted in TTSS-dependent manner. Therefore, we hypothesized that those effectors activate transcription of rice genes to create a suitable environment for bacteria growth. We don't know which kind of condition is actually "suitable" for the bacteria yet. It might be achieved by suppression of host defense or higher availability of nutrients.

In 2004, I received a Sarachek pre-doctoral honors fellowship which financially helped me a lot and

encouraged me to explore science in a different country. After I obtained my Ph.D. degree in 2005, I was interested in studying plant systems by using pathogen infection as a tool. I left Kansas and started my first post-doc in **Andy Maule's** lab at the John Innes Centre, United Kingdom, as a Marie Curie Incoming International Fellow.

It has been known that virus infection leads to the induction of plant cytosolic Hsp70. It was presumed that the response was triggered by accumulation of virus proteins (some misfolded), but no one had formally examined this idea. We named this response as cytosolic protein response (CPR) and investigated it as a parallel response to unfolded protein response (UPR), which is a response to the accumulation of misfolded protein in ER. By comparing plant responses to different treatments that create misfolded proteins, we succeeded in dissecting CPR from larger heat shock response and identified a part of the pathway that triggers the induction of Hsp70. In this two-year post-doc period, I experienced the power of *Arabidopsis* and *Nicotiana benthamiana* in plant research.

After training in the Maule lab, I decided to return to study plant-microbe interactions closely and am currently working as a post-doc in **Saskia Hogenhout's** lab in the John Innes Centre. We are studying plant-pathogenic bacteria, phytoplasma strain aster yellows witches'-broom (AY-WB), which can infect the model plant *Arabidopsis*. Phytoplasmas are transmitted by leafhoppers and cause severe damage on multiple host plants. The symptoms are diverse and often show disturbance of normal plant development. For example, phytoplasma-infected *Arabidopsis* produce a cluster of young leaves and green-colored flowers. Interestingly, phytoplasma-infected plants are more attractive to insect vectors and increase the fecundity of the insects. We hypothesized that phytoplasmas secrete multiple effector proteins that target plant development and/or defense system against insects and the bacteria. To prove this hypothesis, we are currently studying the function of AY-WB-secreted proteins and identified a few proteins that change the morphology of *Arabidopsis*.

As I was educated in a university that specialized in engineering, I tend to be attracted to research that can be useful in future industry. Molecular plant-microbe interactions is a perfect research field because many studies ask fundamental biological questions but they are not too far from the applications. I became a member of IS-MPMI and attended the IS-MPMI Congress held in Sorrento, Italy, in 2006. I appreciated the financial assistance provided by IS-MPMI. I enjoyed the talks and poster presentations, and in addition to that, I really enjoyed meeting with old and new friends at the congress. I am looking forward to seeing them this year in Québec City.

Meet IS-MPMI Members continued on page 8



Robert Dudler

Distinguished Member

Robert Dudler

Institute of Plant Biology
University of Zurich
Zurich, Switzerland

Being fascinated by nature and bird watching and strongly influenced as a high school student by books of the behavioral scientist **Konrad**

Lorenz, I studied zoology at the University of Zurich, obtaining a (master's) diploma in zoology in 1977. Soon I became captivated by molecular biology and decided to concentrate on this exciting and expanding field for my further education. I stayed at the Institute of Zoology working on tRNA genes of *Drosophila melanogaster* and obtained a Ph.D. degree in 1981. From 1982 to 1984, I joined the Laboratory of Molecular Biology (LMB) of the Medical Research Council (MRC) in Cambridge, United Kingdom, as a post-doctoral fellow. The method to transform *D. melanogaster* had just been published, and working under the supervision of **Andrew Travers** and **Hugh Pelham**, I took the opportunity to perform one of the first in vivo promoter analyses in transgenic flies. My stay at the MRC had a permanent impact on my scientific thinking and independence that still is invaluable for me today. After my post-doctoral stay, I took a job as a lecturer at the Institute of Plant Biology, University of Zurich, with **Roger Parish**. Although at a plant research institute, he ran a cancer project and had just published a paper describing a monoclonal antibody that inhibited the invasiveness of metastatic mouse cancer cells. Free to choose and fascinated by the project, I decided to clone the gene encoding the mouse epitope recognized by the antibody. After many difficulties we succeeded, surprisingly discovering that the epitope belonged to a periplasmic binding protein homolog of a high-affinity transport system (an ABC transporter) in *Mycoplasma hyorhynnis*, discovering such transporters in Gram-positive bacteria. This led to two publications in *EMBO* journal in 1988 that are still cited, since a connection between this mycoplasma species and certain cancer forms has materialized also clinically. Being at a plant biology institute, where I was promoted to adjunct professor in 2001, I finally chose to change my research to acquired resistance in cereals, at the beginning mainly focusing on the interaction of powdery mildew with wheat, but also on the interaction of rice blast with rice. We began by cloning transcripts accumulating at the onset of acquired resistance. It became quickly obvious that, to test whether the respective genes were involved in resistance, a quick and reliable assay system for candidate genes was needed. We thus established a transient assay system based on the biolistic transformation of cereal epidermal cells. This widely used system works well and allows scoring of the phenotypic effect on powdery mildew-cereal interactions of transient over-expression as well as suppression by RNA interference of genes of interest. From this period, two diverging research projects evolved. One was triggered by the cloning of a pathogen-

induced glutathione-S-transferase (GST) gene from wheat in the early 1990s. At that time, the first multiple drug resistance (MDR) protein was cloned in mammals. Being familiar with ABC transporters from the cancer cell work and hypothesizing that GSTs would conjugate drugs and perhaps herbicides that then would be transported out of the cell or into the vacuole likely by an MDR homologue, we cloned the first plant ABC transporter (which we named PGP1) from *Arabidopsis*. Although the hypothesis was later shown to be correct, we of course cloned the wrong one. Plants with altered *PGP1* expression exhibited a developmental phenotype and PGP1 was later shown to be involved in auxin transport in a collaboration with **Enrico Martinoia's** laboratory at our institute.

Another project had its origins in results we could not reproduce while studying acquired resistance of rice. As in wheat, we had cloned a number of rice transcripts that accumulated at the onset of acquired resistance (AR) of rice to rice blast, although only locally and not systemically as we thought they should. As AR was triggered by infiltration of a particular strain of the nonhost pathogen *Pseudomonas syringae* pv. *syringae*, we tried another *P. syringae* pv. *syringae* strain. We still did not see systemic accumulation of our transcripts of interest, but one of them (named *Pir7b*) no longer accumulated at all with the new strain. The *Pir7b* product exhibited strong sequence similarity to a class of hydroxynitrile lyases of certain cyanogenic plants (rice of course is not cyanogenic) but had esterase activity with a very restricted set of model substrates. I thought that we could elucidate the biological function of *Pir7b* if we identified the difference between the two *P. syringae* pv. *syringae* strains responsible for *Pir7b* transcript accumulation. Using *Pir7b* activation as an assay, we showed that the relevant difference was the secretion of a cyclic peptide derivative we named syringolin A (SylA). One of the intriguing properties of SylA was that, upon spray application, it triggered hypersensitive cell death of wheat (and *Arabidopsis*) cells colonized by powdery mildew, thus converting a compatible interaction into an incompatible one. Transcriptome analyses of SylA-sprayed and untreated plants revealed very complex SylA-induced changes whose interpretation eventually led to the hypothesis that SylA may target the proteasome. Recently, we showed together with our collaborators that SylA is a virulence factor synthesized by many strains of *P. syringae* pv. *syringae* (but not by other pathogens as far as tested) that irreversibly inhibit the eukaryotic proteasome by a novel mechanism. Cloning of the SylA synthetase gene cluster allowed to build a SylA biosynthesis model, which in turn allowed the cloning of the gene cluster responsible for the biosynthesis of glidobactin A (GlbA), a structurally similar antifungal and antitumor agent isolated from an unknown species belonging to the *Burkholderiales* 20 years ago. We showed that GlbA is also a proteasome inhibitor acting by the same mechanism as SylA, and thus we dubbed this class of inhibitors as syrbactins. Homologous gene clusters potentially responsible for the synthesis of syrbactin-like compounds were identified in an intriguing set of human and insect pathogens (*Burkholderia pseudomallei*, *B. oklahomensis*, *Photobacterium*

asymbiotica, *P. luminescens*). Currently, we are engaged in a number of collaborations to elucidate the biology of this new class of proteasome inhibitors. Why is SylA a virulence factor? Are the putative syrbactin-like compounds of the mentioned pathogens also virulence factors? And, because proteasome inhibitors are a promising class of anticancer therapeutics, is there a medical potential of syrbactins? These and other questions will keep us busy for some time.

In 1990, I attended the first IS-MPMI Congress in Interlaken which, being new to the field, was tremendously instructive

for me. Since I have entered the field of plant-pathogen interactions, IS-MPMI and its journal have been important for me. This is also reflected by the fact that part of the research output of my laboratory was published in *MPMI*.

Since 2000, I am fortunate to be involved in an interesting and rewarding connection with Zhejiang Agricultural University in Hangzhou, China, as a visiting professor. This interaction will be intensified in the coming years as I was elected Pao Yu-Kong Chair Professor at Zhejiang University for 2009–2011.

Welcome New Members

Our community is growing! From January 1 to April 30, 2009, nearly 400 new members have joined IS-MPMI!
Please join us in welcoming them to our society!

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Maria Rosa Marano	Antonio Molina	Anne S. Prabhu	Pinnamaneni Srinivasa Rao	Ye Wang
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Recently published research in *Molecular Plant-Microbe Interactions*

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March 2009, Volume 22, Number 3

Overexpression of *Brassica napus* *MPK4* Enhances Resistance to *Sclerotinia sclerotiorum* in Oilseed Rape.

Tomato Transcriptional Responses to a Foliar and a Vascular Fungal Pathogen Are Distinct.

TOO MUCH LOVE, a Root Regulator Associated with the Long-Distance Control of Nodulation in *Lotus japonicus*.

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Identification of a New Biocontrol Gene in *Trichoderma atroviride*: The Role of an ABC Transporter Membrane Pump in the Interaction with Different Plant-Pathogenic Fungi.

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Symbiosis-Related Plant Genes Modulate Molecular Responses in an Arbuscular Mycorrhizal Fungus During Early Root Interactions.

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Changes in Disease Resistance Phenotypes Associated With Plant Physiological Age Are Not Caused by Variation in *R* Gene Transcript Abundance.

April 2009, Volume 22, Number 4

CURRENT REVIEW—Population Genetics of Fungal and Oomycete Effectors Involved in Gene-for-Gene Interactions.

Unusual Long-Distance Movement Strategies of *Potato mop-top virus* RNAs in *Nicotiana benthamiana*.

Regulation of Tomato Prf by Pto-like Protein Kinases.

Conidiophore *Stalk-less1* Encodes a Putative Zinc-Finger Protein Involved in the Early Stage of Conidiation and Mycelial Infection in *Magnaporthe oryzae*.

The *Magnaporthe oryzae* Avirulence Gene *AvrPiz-t* Encodes a Predicted Secreted Protein That Triggers the Immunity in Rice Mediated by the Blast Resistance Gene *Piz-t*.

Induction of Distinct Defense-Associated Protein Patterns in *Aphanomyces euteiches* (Oomycota)–Elicited and –Inoculated *Medicago truncatula* Cell-Suspension Cultures: A Proteome and Phosphoproteome Approach.

Higher Copy Numbers of the Potato *RB* Transgene Correspond to Enhanced Transcript and Late Blight Resistance Levels.

Correlation Between Transcript Abundance of the *RB* Gene and the Level of the *RB*-Mediated Late Blight Resistance in Potato.

Insights into the Defense-Related Events Occurring in Plant Cells Following Perception of Surfactin-Type Lipopeptide from *Bacillus subtilis*.

The *Arabidopsis* Patatin-Like Protein 2 (PLP2) Plays an Essential Role in Cell Death Execution and Differentially Affects Biosynthesis of Oxylipins and Resistance to Pathogens.

May 2009, Volume 22, Number 5

CURRENT REVIEW—Primary Metabolism and Plant Defense—Fuel for the Fire.

Analysis of Temperature Modulation of Plant Defense Against Biotrophic Microbes.

Impaired Colonization and Infection of Tomato Roots by the Δ *frp1* Mutant of *Fusarium oxysporum* Correlates with Reduced CWDE Gene Expression.

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Different Regulation and Roles of Lactonases AiiB and AttM in *Agrobacterium tumefaciens* C58.

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A Gall-Inducing Caterpillar Species Increases Essential Fatty Acid Content of Its Host Plant Without Concomitant Increases in Phytohormone Levels.

Characterization of Two Unusual Features of Resistance to *Soilborne cereal mosaic virus* in Hexaploid Wheat: Leakiness and Gradual Elimination of Viral Coat Protein from Infected Root Tissues.

Sinorhizobium fredii HH103 *cgs* Mutants Are Unable to Nodulate Determinate- and Indeterminate Nodule-Forming Legumes and Overproduce an Altered EPS.

Rpi-vnt1.1, a *Tm-2* Homolog from *Solanum venturii*, Confers Resistance to Potato Late Blight.

Mapping and Cloning of Late Blight Resistance Genes from *Solanum venturii* Using an Interspecific Candidate Gene Approach.

June 2009, Volume 22, Number 6

Nitric Oxide as a Partner of Reactive Oxygen Species Participates in Disease Resistance to Necrotrophic Pathogen *Botrytis cinerea* in *Nicotiana benthamiana*.

Exploiting Knowledge of *R/Avr* Genes to Rapidly Clone a New LZ-NBS-LRR Family of Late Blight Resistance Genes from Potato Linkage Group IV.

The Role of the *Cucumber mosaic virus* 2b Protein in Viral Movement and Symptom Induction.

The *Xanthomonas campestris* pv. *vesicatoria* Type III Effector Protein XopJ Inhibits Protein Secretion: Evidence for Interference with Cell Wall-Associated Defense Responses.

A Host-Selective Toxin of *Pyrenophora tritici-repentis*, Ptr ToxA, Induces Photosystem Changes and Reactive Oxygen Species Accumulation in Sensitive Wheat.

The N-terminal Region of the *Plantago asiatica* mosaic virus Coat Protein Is Required for Cell-to-Cell Movement but Is Dispensable for Virion Assembly.

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A Key Enzyme of the Leloir Pathway Is Involved in Pathogenicity of *Leptosphaeria maculans* Toward Oilseed Rape.

Interaction with a Host Ubiquitin-Conjugating Enzyme Is Required for the Pathogenicity of a Geminiviral DNA β Satellite.

Adhesion and Fitness in the Bean Phyllosphere and Transmission to Seed of *Xanthomonas fuscans* subsp. *fuscans*.

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Only we can determine that. In the almost 20 years since the society was established, IS-MPMI has grown and seen many significant advancements in the science. Now IS-MPMI has reached a new milestone—1,000 members. Together we are creating a strong community of leaders in our unique science; a community posed to make great achievements for molecular plant-microbe interactions.

IS-MPMI wants to extend a special welcome to all new members to the society. This is the true home for molecular plant-microbe interactions. Your contributions are requested to help advance our science and IS-MPMI. Be sure to become involved and stay connected with this global village of scientists.

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By helping each other, our potential is unlimited.**

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June 28-July 2

Ascochyta 2009: 2nd International Ascochyta Workshop
Pullman, WA, U.S.A.

<http://capps.wsu.edu/conferences/ascochyta>

July 5-10

XXI International Conference on Virus and Other Graft Transmissible Diseases of Fruit Crops

Neustadt an der Weinstrasse, Germany
www.phytomedizin.org/index.php?id=193

July 8-10

Plant ROS 2009

Helsinki, Finland
<http://pog2009.org>

July 13-15

Xanthomonas Genomics Conference 2009

Fort Collins, CO, U.S.A.
www.plantpath.iastate.edu/xgc2009/welcome

July 19-23

XIV International Congress on Molecular Plant-Microbe Interactions

Québec City, Québec, Canada
www.ismpminet.org/meetings

August 1-5

APS Annual Meeting

Portland, OR, U.S.A.
<http://meeting.apsnet.org>

September 13-16

22nd New Phytologist Symposium: Effectors in Plant-Microbe Interactions

Paris, France
www.newphytologist.org/effectors/default.htm

October 25-30

9th International Plant Molecular Biology Congress

St. Louis, MO, U.S.A.
www.ipmb2009.org

Include your meeting in IS-MPMI's printed and online event calendar. Submit online at
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As of May 12, 2009

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Associate in Research, Molecular Plant Pathology/Nematology

An associate in research position is open to conduct research on molecular interactions between parasitic nematodes and their host plants. Position responsibilities and duties include maintaining nematode cultures in a greenhouse; preparing nematodes for plant inoculations; designing and executing experiments; managing data; and facilitating lab operations. In addition to greenhouse work, laboratory techniques will involve molecular techniques (e.g., PCR; real-time PCR; DNA, RNA, and protein purification; DNA cloning; gel electrophoresis; sequencing; western, southern, and northern blotting;

and in situ hybridizations) and plant tissue culture. Applicant must be willing to learn additional techniques and problem solve as needed for the advancement of the laboratory. Qualifications include an M.S. degree in plant pathology, plant science, molecular genetics, nematology, or a related field; a strong demonstrated background in molecular techniques; interest in molecular plant-microbe interactions; and the ability to work in a team, including interdisciplinary collaborators. Submit application, including cover letter, statement of research interests (addressing all qualifications listed above), CV, and transcripts electronically and three letters of reference (direct from source) to Axel Elling. **Closing Date:** Screening begins October 1, 2009. (Position open until filled.) **Contact:** Axel Elling, Department of Plant Pathology, Washington State University, P.O. Box 646430, Pullman, WA, 99164-6430 U.S.A. **E-mail:** elling@wsu.edu; **Web:** <http://plantpath.wsu.edu/people/faculty/elling.htm>.

Ph.D. Student, Molecular Plant Pathology/Nematology

A position for a Ph.D. student is open to conduct research on molecular interactions between the Columbia root-knot nematode (*Meloidogyne chitwoodi*) and potato at Washington State University (WSU). Position responsibilities and duties include design and execution of experiments; data analysis; preparation of manuscripts; and presentation of results at meetings. Research projects may include molecular techniques (e.g., PCR; real-time PCR; DNA, RNA, and protein purification; DNA cloning; gel electrophoresis; sequencing; and western, southern, and northern blotting); microarray experiments; microscopy; and plant tissue culture and greenhouse work. Qualifications include an M.S. degree in plant pathology, plant science, molecular genetics, nematology, or a related field; a strong demonstrated background in molecular techniques and plant-molecular biology essential; interest in molecular plant-microbe interactions; the ability to work in a team, including interdisciplinary collaborators; excellent oral/written communication skills; the ability to work independently; excellent record-keeping abilities; and the ability to manage multiple experimental approaches simultaneously. Prior experience in nematology is an advantage, but not required. The position is available spring or fall semester 2010. For more information, please visit <http://plantpath.wsu.edu/people/faculty/elling.htm>. Applicants should e-mail Axel Elling

directly to indicate interest in this position. A research assistantship (i.e., health insurance, tuition waiver, and other benefits) will be provided. **Contact:** Axel Elling. **E-mail:** elling@wsu.edu; **Web:** <http://plantpath.wsu.edu/students/prostudents.htm>.

Post-Doctoral Position in Plant-Microbe Interactions

A post-doctoral research associate position is available in the laboratory of Paola Veronese, North Carolina State University, Center for Integrated Fungal Research, Raleigh, NC, U.S.A. Qualifications include a Ph.D. degree in molecular biology/genetics or a related field. Responsibilities include conducting research on the molecular genetics of plant interaction with phytopathogenic fungi of the genus *Verticillium*, causal agents of vascular diseases. The incumbent will be assigned projects aimed at the identification of the genetic and molecular basis of host defense and fungal pathogenicity mechanisms. In particular, he/she will study the interaction of model plants, such as *Arabidopsis thaliana* and tomato with *V. dahliae* and *V. longisporum*, by applying functional genomics approaches.

The candidate is required to have practical expertise in PCR-based technologies, including molecular cloning and gene expression analysis. Other specific molecular techniques include construct assembly, plant and fungal transformation, nucleic acid analysis, and protein expression. Familiarity with commonly used bio-informatics tools, a strong ability to work as a team member, and excellent oral and written communication skills are required. The position has secured funding for a minimum of two years and the continuity will depend on successful performance. The start date is July 1, 2009. Applicants should e-mail their CV, a brief description of past research and career goals, and the names and contact information of three references to Paola Veronese. **Salary:** Commensurate with experience. **Closing Date:** June 30, 2009, or until suitable candidate is found. **Contact:** Paola Veronese, Center for Integrated Fungal Research, Department of Plant Pathology, North Carolina State University, Campus Box 7251, Raleigh, NC, 27695-7251 U.S.A. **E-mail:** paola_veronese@ncsu.edu; **Phone:** +1.919.513.8219.

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