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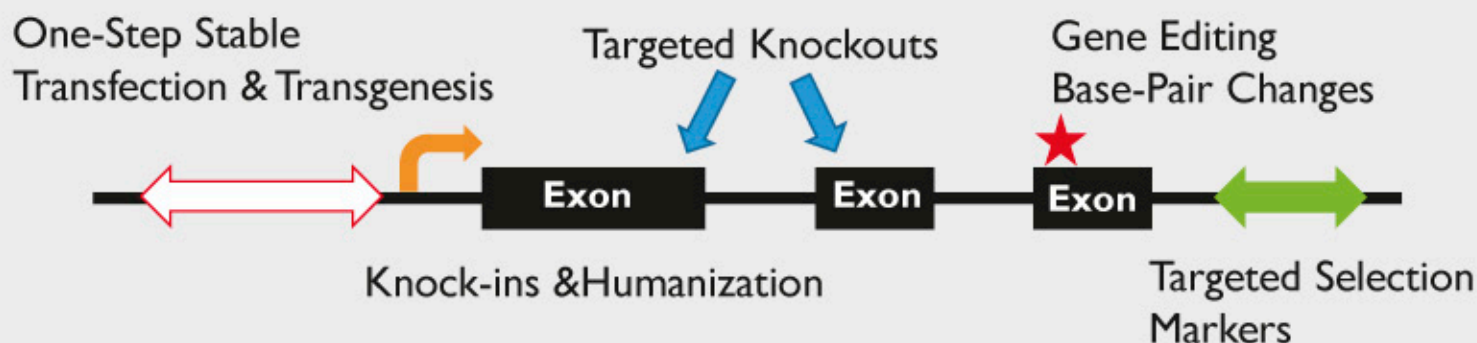
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immunogenomics

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Mary Ellen Conley
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Mark Davis
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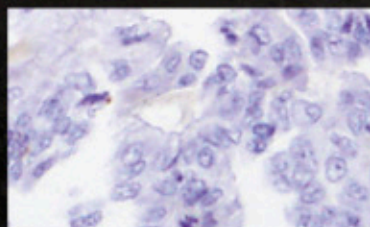


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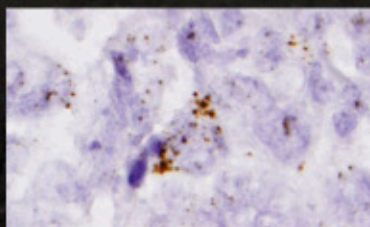


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SCIENTIFIC CONFERENCES 2014-2015

AACR
American Association
for Cancer Research

Presenting the most significant research on cancer etiology, prevention, diagnosis, and treatment

**Marsha Rivkin Center for Ovarian
Cancer Research-AACR 10th Biennial
Ovarian Cancer Research Symposium**

*Co-Chairpersons: Kathleen Cho, Sandra Orsulic,
Mary L. "Nora" Disis, and Saul E. Rivkin*
September 8-9, 2014
Seattle, WA

Targeting PI3K-mTOR Networks in Cancer

*Co-Chairpersons: Lewis C. Cantley, Jose Baselga,
Joan S. Brugge, Brendan D. Manning,
and Malte Peters*
September 14-17, 2014
Philadelphia, PA

Hematologic Malignancies:

Translating Discoveries to Novel Therapies

Chairperson: Kenneth C. Anderson
*Co-Chairpersons: Scott Armstrong
and Riccardo Dalla-Favera*
September 20-23, 2014
Philadelphia, PA

Advances in Melanoma: From Biology to Therapy

*Co-Chairpersons: Suzanne L. Topalian,
Keith T. Flaherty, and Levi A. Garraway,*
September 20-23, 2014
Philadelphia, PA

**13th Annual International Conference
on Frontiers in Cancer Prevention Research**

Program Committee Chairperson:
Phillip A. Dennis
September 28-October 1, 2014
New Orleans, LA

**Seventh AACR Conference on the Science of Can-
cer Health Disparities in Racial/Ethnic Minorities
and Medically Underserved**

*Co-Chairpersons: Ethan Dmitrovsky, Rick A. Kit-
tles, Electra D. Paskett, and Victoria L. Seewaldt*
November 9-12, 2014
San Antonio, TX

**EORTC-NCI-AACR International Symposium on
Molecular Targets and Cancer Therapeutics**

Scientific Committee Co-Chairpersons:
*Jean-Charles Soria, Lee J. Helman,
and Jeffrey A. Engelman*
November 18-21, 2014
Barcelona, Spain

**Tumor Immunology and Immunotherapy:
A New Chapter**

*Co-Chairpersons: Robert H. Vonderheide,
Nina Bhardwaj, Stanley Riddell,
and Cynthia L. Sears*
December 1-4, 2014
Orlando, FL

San Antonio Breast Cancer Symposium

*Co-Directors: Carlos L. Arteaga,
Ismail Jatoi, and C. Kent Osborne*
December 9-13, 2014 • San Antonio, TX

Myc: From Biology to Therapy

*Co-Chairpersons: James E. Bradner,
Martin Eilers, Dean W. Felsher,
and Carla Grandori*
January 7-10, 2015 • La Jolla, CA

Translation of the Cancer Genome

February 7-9, 2015
*Co-Chairpersons: William Hahn,
Lynda Chin, and William Sellers*

Computational and Systems Biology of Cancer

February 9-11, 2015
*Co-Chairpersons: Andrea Califano,
Brenda Andrews, and Peter Jackson*
The Fairmont, San Francisco, CA

**AACR-Society of Nuclear Medicine and
Molecular Imaging Joint Conference: Molecular
Imaging in Cancer Biology and Therapy**

*Co-Chairpersons: Carolyn J. Anderson,
Christopher H. Contag, and David Piwnica-Worms*
February 11-14, 2015 • San Diego, CA



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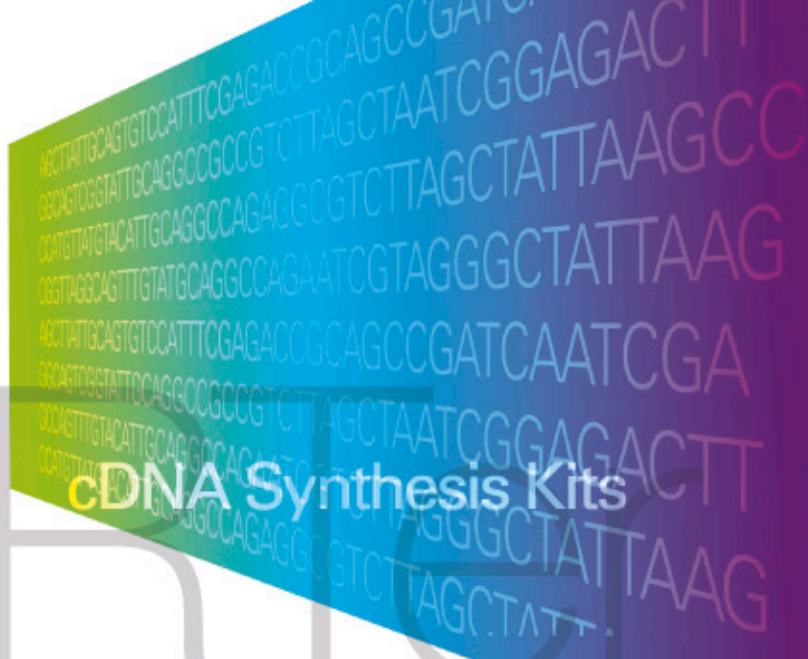


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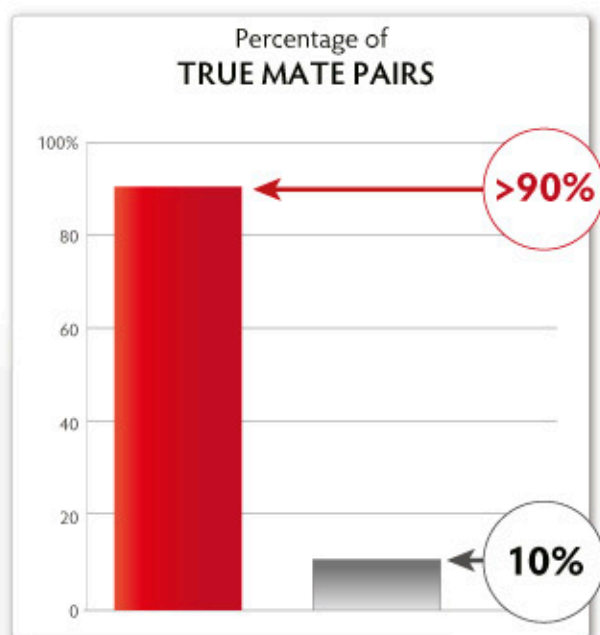
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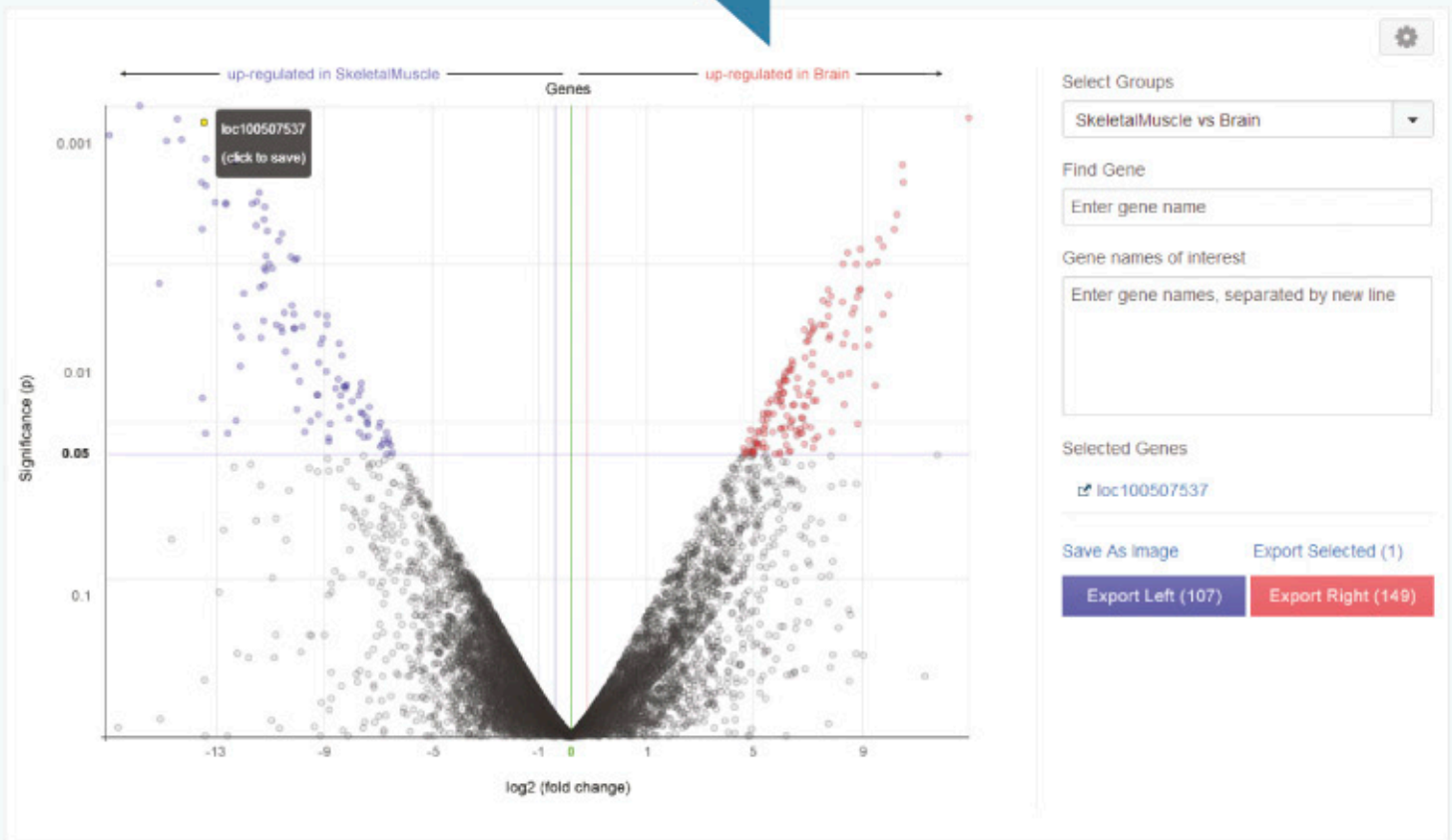
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Web-based analysis

- Real-time integration with public databases; most up-to-date annotations
- Share analysis with collaborators, without duplicating files

Integrated analysis environment

- Preserves link from the final figure to the raw data, including all computational methods used

Advanced user interface

- Multiple views of the same data enable efficient transcript filtering and hypothesis generation
- Export views for presentations
- Search for genes using multiple identifiers

A man with a full grey beard and glasses, wearing a white lab coat over a yellow shirt, is looking directly at the camera in a laboratory setting. In the background, there are shelves with various lab equipment and a window.

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Postdoc Positions in Bioinformatics and Computational Biology

The Center for Systems Biology Dresden calls for applications for several open postdoc positions in **Bioinformatics and Computational Biology** within the prestigious

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We seek outstanding candidates with a degree in Bioinformatics, Computational Biology or Computer Science and a strong interest in working in a multi-disciplinary environment on **Comparative and Evolutionary Genomics, Biological Sequence Analysis, Bioimage Informatics or Network Biology**.

The Center for Systems Biology Dresden focuses on developing and using computational methods to address key questions in biology and to transform data into knowledge. The center is a joint activity by the Max Planck Institute of Molecular Cell Biology and Genetics (MPI-CBG) and the Max Planck Institute of the Physics of Complex Systems (MPI-PKS). The center is a highly interactive and collaborative workplace with an international atmosphere where English is the working language, and has a strong commitment to interdisciplinary training and career development. The center provides access to cutting-edge computer infrastructure, genome sequencing technology and wet lab facilities. The MPI-CBG has been awarded one of the "Best Places To Work for Postdocs" in 2011.

ELBE Fellowships are awarded on a competitive basis to outstanding young researchers with a doctoral degree. To foster collaboration, fellows are usually affiliated with two hosting groups working in different disciplines.

Dresden was awarded "City of Science" in 2006 and is one of the leading scientific centers in Europe with 3 Max Planck and 13 other research institutes, and the Dresden University of Technology as one of eleven Universities of Excellence in Germany. Dresden has half a million inhabitants and is considered one of the most beautiful cities of Germany, located 2 hours from Berlin and Prague.

The Max Planck Society is an equal opportunity employer: handicapped individuals are strongly encouraged to apply. The Center for Systems Biology, the MPI-CBG and the MPI-PKS aim to increase the number of women in scientific positions. Female candidates are therefore particularly welcome.

Deadline for applications is September 5, 2014.

For details about the application procedure, please visit our website <http://mpg-sysbio.de/index.php?id=jobs>



CBG
Max Planck Institute
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MAX-PLANCK-GESELLSCHAFT

THE FACULTY OF BIOLOGY AND MEDICINE OF THE
UNIVERSITY OF LAUSANNE, SWITZERLAND, INVITES
APPLICATIONS FOR THE POSITION OF:

TENURE-TRACK ASSISTANT PROFESSOR TO A LEVEL OF ASSOCIATE PROFESSOR IN THE DOMAIN OF ONCOGENETICS / ONCOGENOMICS

The appointed person shall be responsible for:

- Lead fundamental, translational and applied research in the field of oncogenetics /oncogenomics. The research should be of high level and subsidised by independent organisms.
- Participate actively to the education in the field of oncogenomics.

The candidates should:

- Have a PhD and/or an MD title.
- For a MD, have a FMH (Swiss Medical Association) title in oncology and/or in genetics, or equivalent.
- Have a laboratory experience in the field of oncology and genetics.
- Have adequate teaching capacities for performing pre-graduate, post-graduate and continuous education.
- Be able to lead an independent research programme competitively supported by external peer-reviewed funding.
- Demonstrate ability to carry out interdisciplinary collaborations.
- Have strong communication skills and a good knowledge of French or be able to acquire it quickly.

The job description is available on the Internet site
<http://www.unil.ch/fbm/page64812.html>.

Further information may be obtained from Prof. Michielin
(Olivier.Michielin@unil.ch), chairman of the search committee.

The applications, in English, will include the curriculum vitae, the list of publications with a copy of the five most significant ones, a brief statement of the research programme and teaching experience and a copy of diplomas. They should be sent by September 1st, 2014 to www.unil.ch/iafbm/application.



Seeking to promote an equitable representation of women and men among its staff, the University encourages applications from women.

Keystone Symposia 2015 Genomics Meetings

Precision Genome Engineering and Synthetic Biology

January 11–16, 2015 | Big Sky Resort | Big Sky, Montana | USA

Scientific Organizers: Philip D. Gregory, Jennifer A. Doudna and Ron Weiss

Building on published genome sequences from a range of organisms, new enzymatic tools for the purposeful, precise modification of these genomes in vivo are enabling researchers to interrogate the complexity of genetic variation and determine the cause or consequence of mutations as never before. This meeting brings together scientists developing tools for precision genome engineering, engineering cells with novel pathways and properties, and creating whole synthetic genomes with potential for wholesale reengineering.

Session Topics:

- Editing and Control at Whole-Genome Scale: The Challenge
- Design Principles and Practice of Basic Parts to Large-Scale Circuits
- Beyond On-and-Off: Analog Circuits, Non-Transcriptional Regulation
- Found in Translation: Disease Modeling in Cells and Animals
- The Genome-Edited Cell as a Window onto Biological Mechanism
- Health 2025: On the Path to the Clinic
- Protein, Enzyme and Metabolic Engineering: "Tools and Fuels"
- Earth 2025: The Edited Life Forms for a Better World

www.keystonesymposia.org/15A1

Deadlines: Discounted Abstract/Scholarship – Sep 22, 2014; Abstract – Oct 14, 2014; Discounted Registration – Nov 11, 2014

Genomic Instability and DNA Repair

joint with the meeting on *DNA Replication and Recombination*

March 1–6, 2015 | Whistler Conference Center | Whistler, British Columbia | Canada

Scientific Organizers: Daniel Durocher, Jiri Lukas and Agata Smogorzewska

The meeting highlights the latest trends including how genomic technologies have revolutionized our view of genome integrity and led to the discovery of new types of genome rearrangements whose mechanistic underpinnings remain largely unknown. The unusually broad scope of the conference will bring together scientists operating at the leading edge of multiple disciplines ranging from basic cell biology to pre-clinical oncology.

Session Topics:

- Animal Models and Syndromes
- Telomeres, Aging and Stem Cells
- Recombination Repair
- Mitotic Progression and DNA Damage
- Interface between Chromatin and Genome Maintenance
- Genome Integrity in the Immune System
- DNA Repair Pathway Decisions
- Transcription and Genome Integrity

www.keystonesymposia.org/15X4

Deadlines: Discounted Abstract/Scholarship – Nov 5, 2014; Abstract – Dec 4, 2014; Discounted Registration – Jan 7, 2015

Epigenomics

joint with the meeting on *DNA Methylation*

March 29–April 3, 2015 | Keystone Resort | Keystone, Colorado | USA

Scientific Organizers: Bing Ren and Daniel Zilberman

Empowered by recent technological advances in DNA sequencing technologies, epigenetic regulatory mechanisms have been investigated in a growing number of organisms at increasingly high resolution and scope. The results are an explosion in new genomic technologies, large-scale datasets and computational resources. This meeting is designed to capture the recent progresses in the field while bridging technological gaps, such as dataset availability, method adoption and data interpretation.

Session Topics:

- Genome-Wide DNA Methylation
- Nucleosome Dynamics
- Functional Chromatin Domains
- Chromatin Regulation of Development
- Higher-Order Chromatin
- Small RNA
- Long Noncoding RNA

Deadlines: Discounted Abstract/Scholarship – Dec 3, 2015; Abstract – Jan 8, 2015; Discounted Registration – Jan 29, 2015



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