



# Human Genetics & Genomics

September 6 - 10, 2010

Suzhou, China

Arranged by:

**Paul Liu**, National Human Genome Research Institute, USA

**Maynard Olson**, University of Washington, USA

**Svante Paabo**, Max Planck Inst. for Evolutionary Anthropology, Germany

**Huanming Yang**, Beijing Genomics Institute, China

We are pleased to announce the inaugural Cold Spring Harbor Asia conference on Human Genetics & Genomics which will be held at the Suzhou Dushu Lake Conference Center in Suzhou, China. The conference will begin at 7.30pm on the evening of Monday September 6th, and will conclude after lunch on Friday, September 10th, 2010.

This is the first genomics-themed meeting organized by CSH Asia, a salute to the rapid advancement of Asian countries within the fields of genetics and genomics. This conference will survey a wide range of topics including **genomic structural variations and polymorphisms, population genetics, complex genetic traits, new technology and challengers...** Like the Human Genome Project, which was a large-scale international collaboration, this meeting will foster new international cooperation at the forefront of genomics research within the most dynamic region of the world.

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### Abstract submission:

Many talks will be selected from the openly submitted abstracts on the basis of scientific merit and relevance. Abstracts should contain only new and unpublished material and must be submitted electronically by the abstract deadline. Selection of material for oral and poster presentation will be made by the organizers.

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### Stipends:

We are eager to have as many young people as possible attend since they are likely to benefit most from this meeting. We have applied for funds from government and industry to partially support graduate students and postdocs. For more details, please visit <http://www.csh-asia.org/stipends.html>.

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**FOR THE MOST UPDATED INFO, PLEASE VISIT [www.csh-asia.org](http://www.csh-asia.org)**

### Keynote Speakers

**Sydney Brenner**,

Salk Institute for Biological Studies, USA

**Edison Liu**,

Genome Institute of Singapore, Singapore

**Mathias Uhlen**,

Royal Institute of Technology, Sweden

### Invited Speakers

**Lars Bolund**,

University of Aarhus, Denmark

**Yali Cong**, Peking University Health Science Center, China

**Laurent Duret**,

Université Claude Bernard-Lyon 1, France

**Laura Elnitski**, NHGRI, USA

**Li Jin**, Fudan University, China

**Henrik Kaessmann**,

University of Lausanne, Switzerland

**Philipp Khaitovich**,

Institute for Computational Biology, China

**JONG-IL KIM**,

Seoul National University College of Medicine, KOREA

**Young-Joon Kim**,

Yonsei University, KOREA

**Yingrui Li**, BGI Shenzhen, China

**Partha Majumder**,

Indian Statistical Institute, India

**Naoyuki Takahata**,

Graduate University for Advanced Study, Japan

**Hongyan Wang**,

Fudan University, China

**Wen Wang**,

Chinese Academy of Sciences, China

**Chia-Lin Wei**, Genome Institute of Singapore, Singapore

**Liping Wei**,

Peking University, China

**Kou-Juey Wu**,

National Yang-Ming University, Taiwan