

**Subject:** eRundschreiben 27 (6. September 2005)  
**From:** Biometrische Gesellschaft <gernot.wassmer@uni-koeln.de>  
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**To:** "gernot.wassmer@uni-koeln.de" <gernot.wassmer@uni-koeln.de>

Liebe Kolleginnen und Kollegen,

dieses E-Rundschreiben enthaelt zwei Ankuendigungen fuer Workshops:

1. Die AG Multiple Verfahren der DR veranstaltet einen gemeinsamen Workshop mit der Arbeitsgruppe "Adaptive Designs" am 6./7. Oktober in Basel.
2. Die AG Bioinformatik veranstaltet den BCB Workshop on Machine Learning in Bioinformatics am 10. Oktober in Berlin.

Mit freundlichen Gruessen

Gernot Wassmer  
--- Schriftfuehrer ---

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Joint Workshop of the Working Groups  
“Multiple Methods” (German Region / IBS) and “Adaptive Designs”  
with symposium on  
“Adaptive Designs in Drug Development  
Recent Advances and Health Authority Perspectives  
Basel, October 06/07, 2005  
Novartis Ausbildungszentrum Horburg  
Mühlheimerstrasse 195  
CH-4057 Basel

Dear colleagues and friends,

A joint workshop of the Working Groups “Multiple Methods” (German Region of the International Biometric Society (IBS)) and “Adaptive Designs” will be held in Basel at October 6-7, 2005.

The symposium on “Adaptive Designs in Drug Development Recent Advances and Health Authority Perspectives” will form the first day of the workshop. The symposium will feature invited talks from health authorities, academia and industry. The second day of the workshop will consist of an invited talk by Jason Hsu (Ohio State Univ.) on the partitioning principle to control the generalized familywise error rate and contributed presentations on adaptive designs and multiple testing.

For more information visit [http://www.bioinf.uni-hannover.de/mcp\\_home/meetings/Basel05/](http://www.bioinf.uni-hannover.de/mcp_home/meetings/Basel05/)

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BCB Workshop on Machine Learning in Bioinformatics

Dear colleague,

This is to inform you of the upcoming

INTERNATIONAL BCB-WORKSHOP ON MACHINE LEARNING IN BIOINFORMATICS

to be held at the Magnus House in Berlin on October 10, 2005.

Invited Speakers include

David Wild (Claremont, CA)

Gunnar Rätsch (Tübingen)

Alessandro Verri (Genova)

Jason Hsu (Columbus, OH)

Cesare Furlanello (Trento)

Mike Showe (Philadelphia, PA)

James Reid (Milano)

For detailed information please see <http://comdiag.molgen.mpg.de/iccb2005/>

Please circulate this announcement to interested scientists in your environment.

Best regards,

Rainer Spang

Berlin Center for Genome Based Bioinformatics and

Max Planck Institute for Molecular Genetics

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