

ORGANIZATION

To register for the Symposium, please use the registration form (link below) or contact us (Registration is free):

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Venue:

Alte Mensa der Georg-August-Universität zu Göttingen
Wilhelmsplatz 3, 37073 Goettingen



UNIVERSITÄTSMEDIZIN
GÖTTINGEN **UMG**

DKG DEUTSCHE KREBSGESELLSCHAFT Zertifiziertes
Onkologisches Zentrum

UCC
UniversitätsKrebszentrum
Göttingen

15 YEARS MMML

INTERNATIONAL ANNIVERSARY SYMPOSIUM

Organized by MMML-consortium and WP Biology of the GLA e.V.
Conference Chairs: Reiner Siebert, Ulm; Lorenz Trümper, Goettingen

Georg August University Goettingen, Alte Mensa Conference Center

November 7, 2018; 11 AM - 7 PM

“Molecular Mechanisms of Malignant Lymphoma”

Confirmed Speakers:

Chapuy, B., Boston/Göttingen • Duyster, J., Freiburg • Klapper, W., Kiel • Koch, R., Göttingen • Lenz, G., Münster • Lichter, P., Heidelberg • Loeffler, M., Leipzig • Möller, P., Ulm • Mottok, A., Ulm • Oefner, P., Regensburg • Oellerich, T., Frankfurt • Ott, G., Stuttgart • Reinhardt, H.C., Köln • Ruland, J., München • Schlesner, M., Heidelberg • Siebert, R., Ulm • Spang, R., Regensburg • Steidl, C., Vancouver • Stein, H., Berlin • Wagener, R., Ulm • Weigert, O., München • Weniger, M., Essen



Unter der Schirmherrschaft der



ICGC MMML-Seq (2010-2016)
ICGC DE-Mining (2016-2017)

MMML Demonstrators

Program

11:00-11:30

Molecular Mechanisms in Malignant Lymphoma – Where did we start and where do we stand 2018?

Harald Stein, Berlin

*The first years of the consortium:
Molecular Mechanisms of Lymphoma*

German Ott, Stuttgart

Correlative science and translational research in the consortium: A story of success

11:30-12:15

Key Note Lecture

Peter Lichter, Heidelberg

*From Genome Sequencing to Translational Oncology:
The Heidelberg Experience*

12:15-13.30

B-cell lymphomas - Genomes, Exomes, Transcriptomes, Methylomes

Matthias Schlesner, Heidelberg

Mutational Mechanisms in Malignant Lymphoma

Björn Chapuy, Goettingen

Actionable Genetic Signatures in Large B-cell Lymphoma

Rabea Wagener, Ulm

Molecular pathogenesis of pediatric B-cell lymphomas - on Burkitt lymphoma and its relatives

Anja Mottok, Ulm

Towards a molecular classification of primary mediastinal large B-cell lymphoma

13:30-14:00

Lunch break

14:00-15:15

Molecular Heterogeneity of Malignant B-cell Lymphoma

Wolfram Klapper, Kiel

DLBCL in children and young adults – what we have learned from our Kids

Marc Weniger, Essen

Of Normal and Malignant CD30⁺ B-cells

Peter Möller, Ulm

SOCS-Mutations in Large B-cell Lymphoma

Oliver Weigert, Munich

Translational Genetics in Follicular lymphoma: Towards Risk- and Biology-adapted Treatment strategies

15:15-16:00

Molecular Heterogeneity of T-cell Lymphoma

Justus Duyster, Freiburg

Molecular Genetics of ALK positive Lymphomas

Jürgen Ruland, Munich

Immunreceptor Signals in T-cell Lymphoma: Implications for Therapy

Raphael Koch, Goettingen

Targetable vulnerabilities in T-cell lymphoma

16:00-16:15

Coffee break

16:15-17:30

Oncogenic Signaling in Lymphoma

Georg Lenz, Muenster

Oncogenic PI3K signaling in aggressive lymphoma

Thomas Oellerich, Frankfurt am Main

Mechanisms of response to targeted therapies in lymphoma

Peter Oefner, Regensburg

Molecular classification of B-cell lymphoma by label-free, data-independent acquisition proteomics

Christian Steidl, Vancouver

Molecular and Genetic Characterization of MHC Deficiency in Diffuse Large B-cell Lymphoma

17:30-19:00

Modeling of Disease

Markus Löffler, Leipzig

Evolution of Follicular lymphomas

Rainer Spang, Regensburg

Modeling of lymphoma treatment response

H. Christian Reinhardt, Cologne

Autochthonous mouse models as preclinical tools for the development of therapeutic strategies for aggressive lymphomas

Reiner Siebert, Ulm

Goettingen lymphoma festivals – from playing with data via pathogenetic concepts towards stable structures