

WILSEDE XXXII.

Kind-Philipp-Meeting 2019

Tagungsprogramm (Scientific Program)

XXXII. Jahrestagung der Kind-Philipp-Stiftung
für Leukämieforschung
in Wilsede
vom 5. – 7. Juni 2019



Ausrichter (organized by):

Klinik für Kinder- und Jugendmedizin, Universitätsklinikum Halle; Institut für Pharmazeutische Biologie, Goethe Universität Frankfurt am Main; Pädiatrische Hämatologie und Onkologie, Medizinische Hochschule Hannover; Princess Maxima Center for Pediatric Oncology, Utrecht

Supported by:

- Stifterverband für die deutsche Wissenschaft, DSZ Deutsches Stiftungszentrum, Kind-Philipp-Stiftung
- Verein für krebskranke Kinder Hannover e.V.
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- SERVIER Deutschland GmbH, München
- Eurocept International B.V., Ankeveen
- Hilfe für krebskranke Kinder Frankfurt e.V.
- Abteilung für Kinder- und Jugendmedizin des Universitätsklinikums Ulm
- Fördergemeinschaft Kinderkrebs-Zentrum Hamburg e.V.

PROGRAM AT A GLANCE

Wednesday, June 5	Thursday, June 6	Friday, June 7
	9:00-10:45 Molecular mechanisms of disease I	9:00-10:30 Immunotherapy
	11:00-12:45 Drugs I	10:45-12:30 Molecular mechanisms of disease III
	12:45-14:15 Lunch at Heidemuseum	12:30-13:45 Lunch at Heidemuseum
	14:15-15:45 Diagnostics and Biomarkers	13:45-15:00 Molecular mechanisms of disease IV
16:00-17:30 Welcome and Registration	16:15-17:45 Molecular mechanisms of disease II	15:15-16:45 Drugs II
17:30-18:30 Invited Lecture: T. Mercher	18:15-19:00 Invited Lecture: M. Suttorp	16:45-17:15 Wilsede Award, Farewell
18:30-19:45 Omics Studies	19:00 Barbecue	17:15 coaches to Undeloh 18:00 Bus transfer to HH-Dammtor
19:45 Dinner		

WEDNESDAY, June 5, 2019

16:00 - 17:30 h

Welcome and registration

17:30 – 17:45 h

Opening of the meeting – Rolf Marschalek

17:45 – 18:30 h

(1) Opening Lecture

Chair: Jan-Henning Klusmann

Prof. Dr. Thomas Mercher

Ontogeny-dependent susceptibility to fusion oncogenes controls pediatric myeloid leukemia

Institut Gustave Roussy; Villejuif

18:30 – 19:45 h

(2) Omics studies

Chair: Thomas Sternsdorf

2.a Mutational and transcriptomic landscapes in hematopoietic cells of Fanconi anemia patients.

Mittapalli, VR; Freiburg

2.b Mapping the single cell transcriptome reveals the cellular composition of ATRT subgroups

Buellesbach, A; Heidelberg

2.c Giant cell glioblastoma does not represent a distinct entity but stratifies into different genetically defined entities

Wiedey, A; Bonn

2.d WNT Medulloblastoma: More than just β -Catenin Mutations – Rare Genetic Features in the Focus

Goschzik, T; Bonn

2.e Epigenetics profiling for minimal residual disease in paediatric acute myeloid leukaemia

Al Ameri, M; London

19:45

Dinner

THURSDAY, June 6, 2019

09:00 - 10:45 h

(3) Molecular mechanisms of disease I

Chair: Owen Williams

3.a Splicing factors of the SF3b complex PHF5A and SF3B1 regulate the DNA damage response in ALL

Moorthy, S; Newcastle upon Tyne

3.b Inactivation of Nsd1 impairs terminal erythroid maturation and induces erythroleukemia

Tauchmann, S; Basel

3.c TRIM28 haploinsufficiency predisposes to Wilms tumor

Metzler, M; Erlangen

3.d CRISPR-mediated genome editing of t(4;11) in human prenatal and perinatal hematopoietic stem/progenitor cells

Torres-Ruiz, R; Barcelona

3.e Therapeutic application of the tumour suppressive miR-193b in acute myeloid leukaemia

Issa, H; Halle

3.f Genome-wide CRISPR screen in patient derived cells reveals the mechanism of ALL chemoresistance

Szolysek, K; Newcastle upon Tyne

3.g CRISPRi screen to identify functional long noncoding RNAs in pediatric acute myeloid leukemia

Ng, M; Halle

10:45 - 11:00 h

Coffee break

11:00 - 12:45 h

(4) Drugs I

Chair: Torsten Pietsch

4.a Tumour Necrosis Factor receptor (TNFR)-signalling dependent killing in T-cell acute lymphoblastic leukaemia (T-ALL)

Krippner-Heidenreich, A; Utrecht

4.b iPSC Model of Stepwise Leukemia Development in Congenital Neutropenia Reveals BAALC as a Key Mediator of Leukemogenesis

Dannenmann, B; Tübingen

4.c Targeting c-MYB in Acute Leukaemia through Drug Repositioning

Clesham KJ; London

4.d Functional analysis of class I HDAC inhibition in group 3 medulloblastoma to identify synergistic drug combinations

Vollmer, J; Heidelberg

4.e Screening assay to identify potential Taspase1 inhibitors

Luciano, V; Frankfurt

4.f Analyzing the therapeutic efficacy of navitoclax and MCL-1 inhibitors in juvenile myelomonocytic leukemia

Wu, Y; Freiburg

4.g Establishment of new therapeutic options for the treatment of Myeloid Leukemia in Down syndrome (ML-DS)

Samulowski, D; Halle

12:45 - 14:15 h

Lunch (Wilsede, Heidemuseum)

14:15 – 15:45 h

(5) Diagnostics and Biomarkers

Chair: Roland Kappler

5.a Masked hypodiploidy against uniparental disomies in hyperdiploid ALL: Observations from the diagnostics perspective

Schieck, M; Hannover

5.b Large amplicon droplet digital PCR for DNA-based monitoring of pediatric chronic myeloid leukemia (CML)

Krumbholz, M; Erlangen

5.c Stratification and prognosis of IGH-DUX4 positive acute lymphoblastic leukemia in children

Müller, J; Kiel

5.d Multiplex droplet digital PCR-based targeted enrichment NGS for identification of tumor markers in Ewing sarcoma (EWS)

Eiblwieser, J; Erlangen

5.e Machine learning algorithms for the automated classification of pediatric anemia

Zierk, J; Erlangen

5.f Identification of the genetic mechanisms linked to the occurrence of H3.3K27M mutation in pediatric diffuse intrinsic pontine gliomas

Kubiak, K; Göttingen

15:45 – 16:15 h

Coffee Break

16:15 – 17:45 h

(6) Molecular mechanisms of disease II

Chair: Deepali Pal

6.a Deciphering the interactive network of the DLK1-DIO3 locus in hematopoiesis and pediatric acute megakaryoblastic leukemia

Verboon LJ; Halle

6.b Identification of GATA1s interaction partners in Down syndrome-associated myeloid leukemia

Bräuer-Hartmann, D; Halle

6.c Isolation and characterization of tumor-derived exosomes from a patient-derived xenograft mouse model of acute leukemia

Bartholomé, R; München

6.d Interrogating the role of chromatin regulator BRD4 in the DNA damage response in medulloblastoma

Vu-Han, T-L; Hamburg

6.e Analyzing the effects of oncogenic SHP2 on apoptosis signaling during malignant transformation

Koleci, N; Freiburg

6.f Chromosome structure and mitotic defects are major pathogenic mechanisms in hyperdiploid childhood B-ALL.

Molina, O; Barcelona

18:15 - 19:00 h

(7) Invited Lecture:

Senior-Prof. em. Dr. Meinolf Suttorp *Chair: Karl Welte*

Do you know a chimera?

Pediatric Hematology and Oncology

Medical Faculty, Technical University Dresden; Dresden

19:00 h

Barbecue

Starting later

Evening with the famous Mac-Guitar

FRIDAY, June 7, 2019

09:00 - 10:30 h

(8) Immunotherapy

Chair: Olaf Heidenreich

8.a Development of a novel central nervous system (CNS)-targeted CD19-antibody (AB) in acute lymphoblastic leukemia (ALL)

Winterberg, D; Kiel

8.b CD19 and CD22-directed bispecific CAR for B-cell acute lymphoblastic leukemia

Zanetti, SR; Barcelona

8.c Low regulatory T-cells are associated with improved survival of neuroblastoma patients treated with anti-GD2 antibodies

Troschke-Meurer, S; Greifswald

8.d CRISPR/Cas9 Gene-modification Platform of ELANE Mutations in iPSCs and HSPCs of Severe Congenital Neutropenia Patients

Nasri, M; Tübingen

8.e Venetoclax enhances the efficacy of therapeutic antibodies in B-cell malignancies

Heymann, J; Kiel

8.f Blockade of suppressive myeloid cells is effective against neuroblastoma

von Lojewski, L; Tübingen

10:30 – 10:45 h

Coffee break

10:45 - 12:30 h

(9) Molecular mechanisms of disease III

Chair: Jasper de Boer

9.a Too little is too much: Leukemogenesis in the Gata2 syndrome

Weiss, JM; Freiburg

9.b Using acute promyelocytic leukemia to study the H3.3 histone chaperone system and its role in pediatric malignancies

Cui, W; Hamburg

9.c Functional characterization of RUNX1 variants in the context of FPDMM

Decker, M; Hannover

9.d FOS and FOSB are linked with CNS-infiltration and inferior prognosis in childhood T-cell acute lymphoblastic leukemia

Spory, L; Kiel

9.e Deciphering role of lncRNAs in pediatric AML

Bhayadia, R; Halle

9.f Genomic characterisation of lineage switched MLL-rearranged leukemias

Tirtakusuma, R; Newcastle upon Tyne

9.g Downstream effect of CSF3R and RUNX1 mutations that underlie leukemic transformations in congenital neutropenia (CN)

Ritter, MU; Tübingen

12:30 - 13:45 h

Lunch (Wilsede, Heidemuseum)

13:45 - 15:00 h	<p>(10) Molecular mechanisms of disease IV <i>Chair: Markus Metzler</i></p> <p><u>10.a</u> Identification of a new splice variant of the human transcription factor ONECUT2 <i>Lein, P; Halle</i></p> <p><u>10.b</u> NAMPT-mediated LMO2 deacetylation is indispensable for hematopoiesis and T-ALL leukemogenesis <i>Morishima, T; Tübingen</i></p> <p><u>10.c</u> GADD45b plays an essential role in the G-CSF triggered granulocytic differentiation of human hematopoietic cells <i>Mir, P; Tübingen</i></p> <p><u>10.d</u> Tripartite motif-containing 71 (TRIM71) is a major factor of oncogenic activity in human hepatoblastoma <i>Jiang, T; München</i></p> <p><u>10.e</u> Cooperativity between miR-125b and Gata1s in the pathogenesis of Down syndrome-associated myeloid leukemia <i>Alejo-Valle, O; Halle</i></p>
15:00 - 15:15 h	<p>Coffee break</p>
15:15 - 16:45 h	<p>(11) Drugs II <i>Chair: Julia Skokowa</i></p> <p><u>11.a</u> Investigating HDACi and dnTaspase1 for the treatment of (4;11) leukemic cells <i>Wilhelm, A; Frankfurt</i></p> <p><u>11.b</u> Dasatinib and dexamethasone offer a novel therapeutic strategy for T-cell acute lymphoblastic leukaemia <i>Yuzhe, S; Newcastle upon Tyne</i></p> <p><u>11.c</u> Drug repositioning in infant leukaemia <i>Looi-Somoye, R; London</i></p> <p><u>11.d</u> Synthetic lethality of Wnt pathway activation and asparaginase in drug-resistant acute leukemias <i>Hinze, L; Hannover</i></p> <p><u>11.e</u> A human BM-iPSC-derived oncogenic niche identifies CDH2 as therapeutic niche target in leukemia <i>Pal, D; Newcastle upon Tyne</i></p> <p><u>11.f</u> Inhibition of the polycomb repressive complex 1 (PRC1) as a therapeutic option in childhood liver tumors <i>Bentrop, M; München</i></p>
16:45 - 17:00 h	<p>Wilsede Award and subsequent closure of the meeting – Martin Stanulla</p>
16:50 - 17:15 h	<p>Farewell at the Emhoff</p>
17:15 h	<p>Departure of the coaches to Undeloh</p>
18:00 h	<p>Bus transfer to Hamburg Dammtor railway station</p>