

Ausgewählte Aktivitäten in der klinischen und translationalen Epigenetik	Publikationen	Aktuelle Labor-basierte Projekte	Aktuelle klinische Studien	Link
Prof. Dr. Heiko Becker	<p>Epigenetics in Pancreatic Ductal Adenocarcinoma: Impact on Biology and Utilization in Diagnostics and Treatment. Cancers 2022 DOI 10.3390/cancers14235926.</p> <p>Bisulfite-free epigenomics and genomics of single cells through methylation-sensitive restriction. Commun Biol 2021 DOI 10.1038/s42003-021-01661-w.</p> <p>Monosomal karyotype and chromosome 17p loss or TP53 mutations in decitabine-treated patients with acute myeloid leukemia. Ann Hematol 2020 DOI 10.1007/s00277-020-04082-7.</p>	<p>The DNA methylome as a therapeutic target in elderly patients with acute myeloid leukemia: Development of combination treatments Förderorganisation: DFG (FOR2674)</p> <p>Predictive epigenetic markers in immunonological and epigenetically active treatments Förderorganisation: BMBF (NanodiagBW)</p>	<p>Implementation of Epigenetically active treatment in the care of patients with unresectable or metastasized Pancreatic ductal adenoCarcinoma (EPIC) (IT, DRKS00031481)</p>	<p>https://www.uniklinik-freiburg.de/medizin1/forschung/experimentelle-forschung/becker-laboratory.html</p>
Dr. Michael Daskalakis	<p>Reactivation of endogenous retroviral elements via treatment with DNMT- and HDAC-inhibitors. Cell Cycle 2018 DOI: 10.1080/15384101.2018.1442623</p> <p>DNMT and HDAC inhibitors induce cryptic transcription start sites encoded in long terminal repeats. Nat Genet 2017 DOI 10.1038/ng.3889</p>	<p>Immune cellular therapy (ICT) platform for liquid and solid cancer types and systemic autoimmune diseases; Förderorganisation: University of Bern</p> <p>Translational study on viral mimicry: Marker for prediction of epigenetic treatment response or prognosis in patients with AML or MDS Förderorganisation: Bernese Cancer League</p> <p>Characterization of Oncogenic Protein Isoforms as Targets for Myelodysplastic Syndromes Strategic Focus Area of Personalized Health and Related Technologies (PHRT) funded by six Swiss research institutions: ETHZ, EPFL, PSI, Empa, Eawag and WSL</p>		<p>http://www.hzi.unsel.ch/de/lehre-und-forschung/blood-research-program/research-group-daskalakis</p>
Prof. Dr. Florian Heide	<p>Dynamic DNA methylation reveals novel cis-regulatory elements in mouse hematopoiesis. Exp Hematol 2022 DOI: S0301-472X(22)00805-0</p> <p>Histone demethylase KDM4C is a functional dependency in JAK2-mutated neoplasms. Leukemia 2022 DOI: 10.1038/s41375-022-01611-3</p> <p>YBX1 mediates translation of oncogenic transcripts to control cell competition in AML. Leukemia 2022 DOI: 10.1038/s41375-021-01393-0</p>	<p>Targeted combination therapies exploiting (de-) stabilization of oncogenic mRNAs Förderorganisation: Hector Stiftung</p> <p>Pre-clinical assessment of pharmacologic YBX1 inhibition in myeloid cancers Förderorganisation: Else Kröner-Fresenius Stiftung</p> <p>Macrohistone H2AFY Function in AML Förderorganisation: DFG</p> <p>YBX1 Function in AML Förderorganisation: DFG</p> <p>Tick-Tock: Linking pathology and lifestyle to epigenetic determinants of biological vs. chronological stem cell aging Förderorganisation: Leibniz-Gemeinschaft</p>	<p>Pelabresib Monotherapy in Patients with Myelofibrosis (IST, NCT02158858)</p>	<p>https://www.mhh.de/haematologie</p>
Prof. Dr. Michael Kühn	<p>Menin inhibitor ziftomenib (KO-539) synergizes with drugs targeting chromatin regulation or apoptosis and sensitizes acute myeloid leukemia with MLL rearrangement or NPM1 mutation to venetoclax. Haematologica 2023. doi: 10.3324/haematol.2022.282160.</p> <p>Synergistic targeting of FLT3 mutations in AML via combined menin-MLL and FLT3 inhibition. Blood 2020. doi: 10.1182/blood.2020005037.</p> <p>Targeting Chromatin Regulators Inhibits Leukemogenic Gene Expression in NPM1 Mutant Leukemia. Cancer Discovery 2016. doi: 10.1158/2159-8290.CD-16-0237.</p>	<p>Epigenetische Kontrolle leukämogener Genexpression durch Menin und MLL1-Komplexpartner-Proteine. Förderorganisation: DFG (Emmy Noether-Programm)</p> <p>Therapeutische Manipulation von Chromatin-Komplexen zur Verstärkung der anti-leukämischen Effizienz zellulärer Immuntherapien. Förderorganisation: DFG (SFB1292)</p>	<p>A Study of SNDX-5613 in R/R Leukemias Including Those With an MLLr/KMT2A Gene Rearrangement or NPM1 Mutation (IST, NCT04065399)</p> <p>Ziftomenib in Relapsed or Refractory Acute Myeloid Leukemia (IST, NCT04067336)</p>	<p>https://www.unimedizin-mainz.de/2-forschung/blood-research-program/med/arbeitsgruppen/emmy-noether-ag-kuehn/ag-kuehn/startseite-home.html</p>
PD Dr. Daniel Lipka	<p>Hotspot DNMT3A mutations in clonal hematopoiesis in acute myeloid leukemia sensitize cells to azacytidine via viral mimicry response. Nat Cancer 2021 DOI: 10.1038/s43018-021-00213-9</p> <p>International Consensus Definition of DNA Methylation Subgroups in Juvenile Myelomonocytic Leukemia. Clin Cancer Res 2021 DOI: 10.1158/1078-0432.CCR-20-3184</p>	<p>Etablierung präklinischer Modelle der juvenilen myelomonocyären Leukämie zur Entwicklung neuer therapeutischer Ansätze für Hochrisikopatienten. Förderorganisation: Deutsche José Carreras Leukämie Stiftung</p> <p>Heterogeneity, Evolution, and Resistance in Oncogenic fusion gene-Expressing Sarcomas affecting Adolescents and Young Adults (HEROES-AYA), SP3 - Non-invasive characterization, WP 3.1: Liquid biopsy-based assessment of intratumor heterogeneity. Förderorganisation: BMBF</p>	N/A	<p>www.translational-cancer-epigenomics.de/</p>
Prof. Dr. Michael Lübbert	<p>10-day decitabine versus 3 + 7 chemotherapy followed by allografting in older patients with acute myeloid leukaemia: an open-label, randomised, controlled, phase 3 trial. Lancet Haematol 2023 DOI 10.1016/S2352-3026(23)00273-9.</p> <p>Decitabine Induces Gene Derepression on Monosomic Chromosomes: In Vitro and In Vivo Effects in Adverse-Risk Cytogenetics AML. Cancer Res 2021 DOI 10.1158/0008-5472.CAN-20-1430.</p>	<p>The DNA methylome as a therapeutic target in elderly patients with acute myeloid leukemia: Development of combination treatments Förderorganisation: DFG (FOR2674)</p>	<p>Decitabine + venetoclax +/- ATRA in AML (DECIDER-2) (IT, EudraCT 2020-005495-36)</p> <p>"InDACTION" vs "3+7" Induction in AML (AML21/EORTC1301) (IT, NCT02172872)</p>	<p>https://www.uniklinik-freiburg.de/medizin1/forschung/experimentelle-forschung/luebber-laboratory.html</p>
Dr. Florian Perner	<p>MEN1 mutations mediate clinical resistance to Menin inhibition. 2023. Nature 2023 DOI: 10.1038/s41586-023-05755-9.</p> <p>Novel Inhibitors of the Histone-Methyltransferase DOT1L Show Potent Antileukemic Activity in Patient-derived Xenografts. Blood 2020. DOI: 10.1182/blood.202006113.</p>	<p>Deciphering epigenetic mechanisms underlying disease persistence under Menin-inhibitor treatment in Acute Myeloid Leukemia. Förderorganisation: Else Kröner-Fresenius Stiftung</p> <p>Targeting aberrant chromatin modifiers in clonal hematopoiesis and pre-leukemia. Förderorganisation: Deutsche Forschungsgemeinschaft, Emmy Noether-Programme</p>	<p>Ziftomenib in patients with R/R AML (IST, NCT04067336)</p>	<p>https://www.linkedin.com/in/florian-perner</p>
Prof. Dr. Christoph Plass	<p>DNMT and HDAC inhibitors induce cryptic transcription start sites encoded in long terminal repeats. Nat Genetics 2017 DOI: 10.1038/ng.1117-1661c</p> <p>DNMT and HDAC inhibition induces immunogenic neoantigens from human endogenous retroviral element-derived transcripts. Nat Commun doi: 10.1038/s41467-023-42417-w</p>	<p>Veränderungen im Methylohm hämatopoetischer Stammzellen in der altersungs-assoziierten klonalen Hämatopoese und myeloidischen Leukämogenese Förderorganisation: DFG</p>	N/A	<p>https://www.dkfz.de/en/CanEp/index.php</p>