

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

Max Delbrück Center for Molecular Medicine; Berlin; Germany

RRID:SCR_011369

Type: Tool

Proper Citation

Max Delbrück Center for Molecular Medicine; Berlin; Germany (RRID:SCR_011369)

Resource Information

URL: <https://www.mdc-berlin.de/>

Proper Citation: Max Delbrück Center for Molecular Medicine; Berlin; Germany (RRID:SCR_011369)

Description: A major biomedical research institute located in the northeastern corner of Berlin, Germany, with a mission of translating discoveries from molecular research into applications to improve the prevention, diagnosis, and treatment of major human diseases. The work of the MDC's 57 research groups centers around three classes of disease which have an enormous impact on society: * Cardiovascular and metabolic diseases * Cancer * Nervous system disorders

Abbreviations: MDC

Synonyms: Max Delbrück Center for Molecular Medicine, Max Delbrück Center for Molecular Medicine

Resource Type: institution

Resource Name: Max Delbrück Center for Molecular Medicine; Berlin; Germany

Resource ID: SCR_011369

Alternate IDs: Wikidata: Q1912019, ISNI: 0000 0001 1014 0849, grid.419491.0, nlx_156718

Alternate URLs: <https://ror.org/04p5ggc03>

Record Creation Time: 20220129T080304+0000

Record Last Update: 20260905T132702+0000

Ratings and Alerts

No rating or validation information has been found for Max Delbrück Center for Molecular Medicine; Berlin; Germany.

No alerts have been found for Max Delbrück Center for Molecular Medicine; Berlin; Germany.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#).

Qin D, et al. (2023) Exosomal miR-23a-3p derived from human umbilical cord mesenchymal stem cells promotes remyelination in central nervous system demyelinating diseases by targeting Tbr1/Wnt pathway. The Journal of biological chemistry, 300(1), 105487.

Chen C, et al. (2023) ALKBH5-mediated CHAC1 depletion promotes malignant progression and decreases cisplatin-induced oxidative stress in gastric cancer. Cancer cell international, 23(1), 293.

Coronel-Córdoba P, et al. (2022) FoxK1 is Required for Ectodermal Cell Differentiation During Planarian Regeneration. Frontiers in cell and developmental biology, 10, 808045.

Tack A, et al. (2021) Towards novel osteoarthritis biomarkers: Multi-criteria evaluation of 46,996 segmented knee MRI data from the Osteoarthritis Initiative. PloS one, 16(10), e0258855.

Wibberg D, et al. (2019) The de.NBI / ELIXIR-DE training platform - Bioinformatics training in Germany and across Europe within ELIXIR. F1000Research, 8.

Xie S, et al. (2019) Identification of miRNAs Involved in Bacillus velezensis FZB42-Activated Induced Systemic Resistance in Maize. International journal of molecular sciences, 20(20).

Zhang Y, et al. (2018) Upregulation of HOXA1 promotes tumorigenesis and development of non-small cell lung cancer: A comprehensive investigation based on reverse transcription-quantitative polymerase chain reaction and bioinformatics analysis. International journal of oncology, 53(1), 73.

Li MF, et al. (2018) Investigation of miR-490-3p Expression in Hepatocellular Carcinoma Based on Reverse Transcription-Polymerase Chain Reaction (RT-qPCR) and a Meta-Analysis of 749 Cases. Medical science monitor : international medical journal of experimental and clinical research, 24, 4914.

Zhang Y, et al. (2018) Microarray-based bioinformatics analysis of the prospective target gene network of key miRNAs influenced by long non-coding RNA PVT1 in HCC.

Oncology reports, 40(1), 226.