

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## MatrixDB

RRID:SCR\_001727

Type: Tool

### Proper Citation

MatrixDB (RRID:SCR\_001727)

### Resource Information

**URL:** <http://matrixdb.univ-lyon1.fr/>

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**Description:** Freely available database focused on interactions established by extracellular proteins and polysaccharides, taking into account the multimeric nature of the extracellular proteins (e.g. collagens, laminins and thrombospondins are multimers). MatrixDB is an active member of the International Molecular Exchange (IMEx) consortium and has adopted the PSI-MI standards for annotating and exchanging interaction data. It includes interaction data extracted from the literature by manual curation, and offers access to relevant data involving extracellular proteins provided by the IMEx partner databases through the PSICQUIC webservice, as well as data from the Human Protein Reference Database. The database reports mammalian protein-protein and protein-carbohydrate interactions involving extracellular molecules. Interactions with lipids and cations are also reported. MatrixDB is focused on mammalian interactions, but aims to integrate interaction datasets of model organisms when available. MatrixDB provides direct links to databases recapitulating mutations in genes encoding extracellular proteins, to UniGene and to the Human Protein Atlas that shows expression and localization of proteins in a large variety of normal human tissues and cells. MatrixDB allows researchers to perform customized queries and to build tissue- and disease-specific interaction networks that can be visualized and analyzed with Cytoscape or Medusa. Statistics (2013): 2283 extracellular matrix interactions including 2095 protein-protein and 169 protein-glycosaminoglycan interactions.

**Abbreviations:** MatrixDB

**Synonyms:** MatrixDB: Extracellular Matrix Interactions Database, Extracellular Matrix Interactions Database

**Resource Type:** data or information resource, database, service resource, production service resource

**Defining Citation:** [PMID:20852260](#), [PMID:19147664](#)

**Keywords:** extracellular, protein fragment, biomolecule, cation, cleavage, collagen, glycosaminoglycan, human, interaction, laminin, lipid, mammalian, matricryptin, matrikin, matrix, molecule, monomer, mulimerization, multimer, polysaccharide, protein, protein-carbohydrate interaction, protein-protein interaction, recognition, thrombospondin, interactome, extracellular protein, protein-polysaccharide interaction, extracellular interaction, molecular interaction, model organism, inorganic, small molecule-protein, small molecule, extracellular matrix protein, protein-glycosaminoglycan interaction, bio.tools, FASEB list

**Funding:** European Union contract FP7-HEALTH-2007-223411

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** MatrixDB

**Resource ID:** SCR\_001727

**Alternate IDs:** biotools:matrixdb, r3d100010672, nif-0000-10226

**Alternate URLs:** <https://bio.tools/matrixdb>, <https://doi.org/10.17616/R3M03H>

**Old URLs:** <http://matrixdb.ibcp.fr/>

**Record Creation Time:** 20220129T080209+0000

**Record Last Update:** 20260731T042622+0000

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## Ratings and Alerts

No rating or validation information has been found for MatrixDB.

No alerts have been found for MatrixDB.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 86 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Rizvi SA, et al. (2026) Scaling Large Language Models for Next-Generation Single-Cell Analysis. bioRxiv : the preprint server for biology.

Farahi N, et al. (2025) Phase-separating fusion proteins drive cancer by upsetting transcription regulation. Genome biology, 26(1), 330.

Ling Z, et al. (2025) Chemoselective Characterization of New Extracellular Matrix Deposition in Bioengineered Tumor Tissues. Advanced materials (Deerfield Beach, Fla.),

37(47), e05445.

Bhutada S, et al. (2025) Degradomics defines proteolysis information flow from human knee osteoarthritis cartilage to matched synovial fluid and the contributions of secreted proteases ADAMTS5, MMP13 and CMA1 to articular cartilage breakdown. *Osteoarthritis and cartilage*, 33(1), 116.

Richard JB, et al. (2025) Synovial matrix turnover controls immune cell spatial patterning in inflammation resolution. *Molecular systems biology*, 21(11), 1638.

Lamba R, et al. (2025) MatriCom, a single-cell RNA-sequencing data mining tool to infer cell-extracellular matrix interactions. *Journal of cell science*, 138(13).

Samarasinghe KW, et al. (2025) MatrixDB 2024: an increased coverage of extracellular matrix interactions, a new Network Explorer and a new web interface. *Nucleic acids research*, 53(D1), D1677.

Wright SN, et al. (2025) State of the interactomes: an evaluation of molecular networks for generating biological insights. *Molecular systems biology*, 21(1), 1.

Okoye GD, et al. (2025) Single-cell map of innate-like lymphocyte response to *Francisella tularensis* infection reveals interleukin-17-dependent protection by MAIT cells. *iScience*, 28(3), 111810.

Enríquez-Rodríguez CJ, et al. (2024) Proteomic Blood Profiles Obtained by Totally Blind Biological Clustering in Stable and Exacerbated COPD Patients. *Cells*, 13(10).

Lamba R, et al. (2024) MatriCom: a scRNA-Seq data mining tool to infer ECM-ECM and cell-ECM communication systems. *bioRxiv : the preprint server for biology*.

Martinez K, et al. (2024) Functional implications of glycans and their curation: insights from the workshop held at the 16th Annual International Biocuration Conference in Padua, Italy. *Database : the journal of biological databases and curation*, 2024.

Li N, et al. (2024) Glycosaminoglycans: Participants in Microvascular Coagulation of Sepsis. *Thrombosis and haemostasis*, 124(7), 599.

Kyriakopoulou K, et al. (2023) Trends in extracellular matrix biology. *Molecular biology reports*, 50(1), 853.

Cheraghi-Shavi T, et al. (2023) TGM2, HMGA2, FXYD3, and LGALS4 genes as biomarkers in acquired oxaliplatin resistance of human colorectal cancer: A systems biology approach. *PloS one*, 18(8), e0289535.

Chaussé AM, et al. (2023) Epithelial cell invasion by salmonella typhimurium induces modulation of genes controlled by aryl hydrocarbon receptor signaling and involved in extracellular matrix biogenesis. *Virulence*, 14(1), 2158663.

Petrov PB, et al. (2023) Matrisome AnalyzeR: A suite of tools to annotate and quantify ECM molecules in big datasets across organisms. *bioRxiv : the preprint server for biology*.

Naba A, et al. (2023) Ten Years of Extracellular Matrix Proteomics: Accomplishments, Challenges, and Future Perspectives. *Molecular & cellular proteomics : MCP*, 22(4),

100528.

Petrov PB, et al. (2023) Matrisome AnalyzeR - a suite of tools to annotate and quantify ECM molecules in big datasets across organisms. *Journal of cell science*, 136(17).

Li X, et al. (2022) dbBIP: a comprehensive bipolar disorder database for genetic research. *Database : the journal of biological databases and curation*, 2022.