

Resource Summary Report

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European Molecular Biology Laboratory

RRID:SCR_004473

Type: Tool

Proper Citation

European Molecular Biology Laboratory (RRID:SCR_004473)

Resource Information

URL: <http://www.embl.de/>

Proper Citation: European Molecular Biology Laboratory (RRID:SCR_004473)

Description: Intergovernmental organisation funded by public research money from its member states in Europe. Groups and laboratories perform basic research in molecular biology and molecular medicine, training for scientists, students and visitors. Provides development of services, new instruments and methods, data and technology in its member states.

Abbreviations: EMBL

Synonyms: European Molecular Biology Laboratory, EMBL

Resource Type: organization portal, data or information resource, training resource, portal, postdoctoral program resource, graduate program resource

Keywords: molecular, biology, medicine, data, training, service, organization, data, technology, academic

Funding:

Resource Name: European Molecular Biology Laboratory

Resource ID: SCR_004473

Alternate IDs: nlx_46173

Record Creation Time: 20220129T080224+0000

Record Last Update: 20250421T053439+0000

Ratings and Alerts

No rating or validation information has been found for European Molecular Biology Laboratory.

No alerts have been found for European Molecular Biology Laboratory.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5121 mentions in open access literature.

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

Cassol I, et al. (2025) Key features and guidelines for the application of microbial alpha diversity metrics. *Scientific reports*, 15(1), 622.

Dolgin V, et al. (2025) Severe neonatal hypotonia due to SLC30A5 variant affecting function of ZnT5 zinc transporter. *JIMD reports*, 66(1), e12465.

Montero F, et al. (2025) Exploring the druggability of the UEV domain of human TSG101 in search for broad-spectrum antivirals. *Protein science : a publication of the Protein Society*, 34(1), e70005.

Horsman S, et al. (2025) Molecular Epidemiological Characteristics of *Staphylococcus pseudintermedius*, *Staphylococcus coagulans*, and Coagulase-Negative *Staphylococci* Cultured from Clinical Canine Skin and Ear Samples in Queensland. *Antibiotics (Basel, Switzerland)*, 14(1).

Prattico C, et al. (2025) Identification of novel fructo-oligosaccharide bacterial consumers by pulse metatranscriptomics in a human stool sample. *mSphere*, 10(1), e0066824.

Berardi A, et al. (2025) The C-terminal PHDVC5HCH tandem domain of NSD2 is a combinatorial reader of unmodified H3K4 and tri-methylated H3K27 that regulates transcription of cell adhesion genes in multiple myeloma. *Nucleic acids research*, 53(1).

Krawczyk K, et al. (2025) Chromosome-scale telomere to telomere genome assembly of common crystalwort (*Riccia sorocarpa* Bisch.). *Scientific data*, 12(1), 77.

Bernhard H, et al. (2025) Structural basis of Spliced Leader RNA recognition by the *Trypanosoma brucei* cap-binding complex. *Nature communications*, 16(1), 685.

Thiry V, et al. (2025) Using DNA metabarcoding and direct behavioural observations to identify the diet of proboscis monkeys (*Nasalis larvatus*) in the Kinabatangan Floodplain, Sabah. *PLoS one*, 20(1), e0316752.

Selzer AM, et al. (2025) An SH3-binding allosteric modulator stabilizes the global conformation of the AML-associated Src-family kinase, Hck. *The Journal of biological chemistry*, 301(1), 108088.

Baldari S, et al. (2025) Expression pattern and prognostic significance of aldehyde dehydrogenase 2 in lung adenocarcinoma as a potential predictor of immunotherapy efficacy. *Cancer innovation*, 4(1), e149.

Su S, et al. (2025) Analysis of the CHS Gene Family Reveals Its Functional Responses to Hormones, Salinity, and Drought Stress in Moso Bamboo (*Phyllostachys edulis*). *Plants (Basel, Switzerland)*, 14(2).

Dobrzyński J, et al. (2025) Non-native PGPB consortium consisting of *Pseudomonas* sp. G31 and *Azotobacter* sp. PBC2 promoted winter wheat growth and slightly altered the native bacterial community. *Scientific reports*, 15(1), 3248.

Tan I, et al. (2025) Alu-Sc-mediated exonization generated a mitochondrial LKB1 gene variant found only in higher order primates. *Scientific reports*, 15(1), 3360.

Brezak M, et al. (2025) Defective Mammary Epithelial Outgrowth in Transgenic EKAREV-NLS Mice: Correction via Estrogen Supplementation and Genetic Background Modification. *Journal of mammary gland biology and neoplasia*, 30(1), 1.

Coquille S, et al. (2025) Allostery and Evolution: A Molecular Journey Through the Structural and Dynamical Landscape of an Enzyme Super Family. *Molecular biology and evolution*, 42(1).

Yakovenko I, et al. (2025) Telomemore enables single-cell analysis of cell cycle and chromatin condensation. *Nucleic acids research*, 53(3).

Jiang Z, et al. (2025) The RodentGPOmics Atlas: a comprehensive database of rodent biology for genomes and pathogens. *Nucleic acids research*, 53(D1), D1144.

Nurhidayat L, et al. (2025) Tokay gecko tail regeneration involves temporally collinear expression of HOXC genes and early expression of satellite cell markers. *BMC biology*, 23(1), 6.

Formichetti S, et al. (2025) Genetic gradual reduction of OGT activity unveils the essential role of O-GlcNAc in the mouse embryo. *PLoS genetics*, 21(1), e1011507.