

Resource Summary Report

Generated by [ASWG](#) on Aug 31, 2026

CHEBI

RRID:SCR_002088

Type: Tool

Proper Citation

CHEBI (RRID:SCR_002088)

Resource Information

URL: <https://www.ebi.ac.uk/chebi/beta/>

Proper Citation: CHEBI (RRID:SCR_002088)

Description: Collection of chemical compounds and other small molecular entities that incorporates an ontological classification of chemical compounds of biological relevance, whereby the relationships between molecular entities or classes of entities and their parents and/or children are specified. The molecular entities in question are either products of nature or synthetic products used to intervene in the processes of living organisms.

Abbreviations: ChEBI

Synonyms: CHEBI, Chemical Entities of Biological Interest

Resource Type: data or information resource, database

Defining Citation: [PMID:19854951](#), [PMID:19496059](#), [PMID:17932057](#)

Keywords: complex, conformer, ion, ion pair, isotope, molecular entity, molecule, radical, radical ion, small molecule, obo, gold standard, biochemistry, metabolomics, bio.tools

Funding: BBSRC BB/G022747/1

Availability: Freely available

Resource Name: CHEBI

Resource ID: SCR_002088

Alternate IDs: nif-0000-02655, biotools:chebi, r3d100012626

Alternate URLs: <http://bioportal.bioontology.org/ontologies/1007>,
<http://www.obofoundry.org/cgi-bin/detail.cgi?id=chebi>,

ftp://ftp.ebi.ac.uk/pub/databases/chebi/ontology/chebi.obo,
http://chebi.wiki.sourceforge.net/, https://bio.tools/chebi

Old URLs: http://www.ebi.ac.uk/chebi/

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260829T182920+0000

Ratings and Alerts

No rating or validation information has been found for CHEBI.

No alerts have been found for CHEBI.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 129 mentions in open access literature.

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

Petrier M, et al. (2026) Identification of a robust metabolic signature associated with hospital-acquired pneumonia and response to interferon-gamma treatment in critically ill patients. *Critical care* (London, England), 30(1).

Liu Y, et al. (2026) A microbiome catalog of Chinese traditional artisanal cheeses provides insights into functional and microbial diversity. *Nature communications*, 17(1).

Vaz S, et al. (2025) Catalytic degradation of kraft lignin to determine its antioxidant potential for industrial purposes. *Heliyon*, 11(4), e42783.

Karakoltzidis A, et al. (2025) AI-driven parametrization of Michaelis-Menten maximal velocity: Advancing in silico new approach methodologies (NAMs). *NAM journal*, 1, 100012.

Wang S, et al. (2025) ACSS2 coupled with KAT7 regulates histone ??-hydroxybutyrylation to enhance transcription. *Science advances*, 11(33), eadv8448.

Ninzatti L, et al. (2025) Artificial symbiont replacement in a vertically transmitted plant symbiosis reveals a role for microbe-microbe interactions in enforcing specificity. *The ISME journal*, 19(1).

Girija A, et al. (2025) Metabolomic approaches suggest two mechanisms of drought response post-anthesis in Mediterranean oat (*Avena sativa* L.) cultivars. *Physiologia plantarum*, 177(2), e70181.

Milani C, et al. (2025) Functional modulation of the human gut microbiome by bacteria vehicled by cheese. *Applied and environmental microbiology*, 91(3), e0018025.

Latini S, et al. (2024) Unveiling the signaling network of FLT3-ITD AML improves drug sensitivity prediction. *eLife*, 12.

Witting M, et al. (2024) Challenges and perspectives for naming lipids in the context of lipidomics. *Metabolomics : Official journal of the Metabolomic Society*, 20(1), 15.

Mullin S, et al. (2024) Chemical entity normalization for successful translational development of Alzheimer's disease and dementia therapeutics. *Journal of biomedical semantics*, 15(1), 13.

Jordan EN, et al. (2024) Integrated omics of *Saccharomyces cerevisiae* CENPK2-1C reveals pleiotropic drug resistance and lipidomic adaptations to cannabidiol. *NPJ systems biology and applications*, 10(1), 63.

Sternberg PW, et al. (2024) WormBase 2024: status and transitioning to Alliance infrastructure. *Genetics*, 227(1).

Nev OA, et al. (2024) Metabolic modelling as a powerful tool to identify critical components of *Pneumocystis* growth medium. *PLoS computational biology*, 20(10), e1012545.

Hay AE, et al. (2024) Comparative metabolomics reveals how the severity of predation by the invasive insect *Cydalima perspectalis* modulates the metabolism re-orchestration of native *Buxus sempervirens*. *Plant biology (Stuttgart, Germany)*.

Kyoda K, et al. (2024) SSBD: an ecosystem for enhanced sharing and reuse of bioimaging data. *Nucleic acids research*.

Kwoji ID, et al. (2024) Elucidating the Mechanisms of Cell-to-Cell Crosstalk in Probiotics Co-culture: A Proteomics Study of *Limosilactobacillus reuteri* ZJ625 and *Ligilactobacillus salivarius* ZJ614. *Probiotics and antimicrobial proteins*, 16(5), 1817.

Frleta Matas R, et al. (2024) Influence of Nutrient Deprivation on the Antioxidant Capacity and Chemical Profile of Two Diatoms from Genus *Chaetoceros*. *Marine drugs*, 22(2).

Hager MS, et al. (2024) Untargeted metabolomics profiling of oat (*Avena sativa* L.) and wheat (*Triticum aestivum* L.) infested with wheat stem sawfly (*Cephus cinctus* Norton) reveals differences associated with plant defense and insect nutrition. *Frontiers in plant science*, 15, 1327390.

Coudert E, et al. (2023) Annotation of biologically relevant ligands in UniProtKB using ChEBI. *Bioinformatics (Oxford, England)*, 39(1).