

# Resource Summary Report

Generated by RRID on Aug 4, 2026

## MetaMap

RRID:SCR\_015031

Type: Tool

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### Proper Citation

MetaMap (RRID:SCR\_015031)

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### Resource Information

**URL:** <https://metamap.nlm.nih.gov/>

**Proper Citation:** MetaMap (RRID:SCR\_015031)

**Description:** Program to map biomedical text to the UMLS Metathesaurus and to discover Metathesaurus concepts referred to in text based on symbolic, natural-language processing and computational-linguistic techniques.

**Synonyms:** MetaMap 2016, MetaMap 2016v2

**Resource Type:** text-mining software, software resource, software application

**Keywords:** text mining, biomedical text

**Funding:** NLM

**Availability:** Free, Account required

**Resource Name:** MetaMap

**Resource ID:** SCR\_015031

**License:** UMLS Metathesaurus License

**License URLs:** <https://metamap.nlm.nih.gov/MMTnCs.shtml>

**Record Creation Time:** 20220129T080323+0000

**Record Last Update:** 20260804T043543+0000

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### Ratings and Alerts

No rating or validation information has been found for MetaMap.

No alerts have been found for MetaMap.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 337 mentions in open access literature.

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

Forrest N, et al. (2025) A Dynamic Time Warping Extension to Consensus Weight-Based Cachexia Criteria Improves Prediction of Cancer Patient Outcomes. JCSM communications, 8(1), e107.

Wang L, et al. (2025) DrugCombo: an informatics bridge for anticancer drug combination Phase I trial design. Database : the journal of biological databases and curation, 2025.

Huang H, et al. (2025) An open-source multi-semantic annotation dataset and automated recognition tool for viral carcinogenesis factors. Database : the journal of biological databases and curation, 2025.

Nejat P, et al. (2025) Optimizing Digital Management of Research and Collaboration With Academic Information Manager. Mayo Clinic proceedings. Digital health, 3(2), 100222.

Hu YH, et al. (2025) An Intelligent Trial Eligibility Screening Tool Using Natural Language Processing With a Block-Based Visual Programming Interface: Development and Usability Study. JMIR medical informatics, 13, e80072.

Xin Y, et al. (2025) Improving topic modeling performance on social media through semantic relationships within biomedical terminology. PloS one, 20(2), e0318702.

Forrest N, et al. (2025) Natural language processing algorithms identify wild-type isocitrate dehydrogenase gliomas in electronic health records. Neuro-oncology advances, 7(1), vda111.

Haaker TS, et al. (2025) Approaches for extracting daily dosage from free-text prescription signatures in heart failure with reduced ejection fraction: a comparative study. JAMIA open, 8(1), ooae153.

Tayebi Arasteh S, et al. (2024) Enhancing diagnostic deep learning via self-supervised pretraining on large-scale, unlabeled non-medical images. European radiology experimental, 8(1), 10.

Roberts K, et al. (2024) Natural language processing of clinical notes enables early inborn error of immunity risk ascertainment. The journal of allergy and clinical immunology. Global, 3(2), 100224.

Gutiérrez-Sacristán A, et al. (2024) Development and validation of an open-source pipeline for automatic population of case report forms from electronic health records: a

pediatric multi-center prospective study. *EBioMedicine*, 108, 105337.

Kamp H, et al. (2024) Utility of in vivo metabolomics to support read-across for UVCB substances under REACH. *Archives of toxicology*, 98(3), 755.

Nédellec C, et al. (2024) TaeC: A manually annotated text dataset for trait and phenotype extraction and entity linking in wheat breeding literature. *PloS one*, 19(6), e0305475.

Zhou H, et al. (2024) Complementary and Integrative Health Information in the literature: its lexicon and named entity recognition. *Journal of the American Medical Informatics Association : JAMIA*, 31(2), 426.

Lee HJ, et al. (2024) StrokeClassifier: ischemic stroke etiology classification by ensemble consensus modeling using electronic health records. *NPJ digital medicine*, 7(1), 130.

Song L, et al. (2024) Centrifuger: lossless compression of microbial genomes for efficient and accurate metagenomic sequence classification. *Genome biology*, 25(1), 106.

Xiao Y, et al. (2024) Repurposing non-pharmacological interventions for Alzheimer's disease through link prediction on biomedical literature. *Scientific reports*, 14(1), 8693.

Kim K, et al. (2024) Multifaceted Natural Language Processing Task-Based Evaluation of Bidirectional Encoder Representations From Transformers Models for Bilingual (Korean and English) Clinical Notes: Algorithm Development and Validation. *JMIR medical informatics*, 12, e52897.

Yada S, et al. (2024) Cross-lingual Natural Language Processing on Limited Annotated Case/Radiology Reports in English and Japanese: Insights from the Real-MedNLP Workshop. *Methods of information in medicine*, 63(5-06), 145.

Deng Y, et al. (2024) Natural language processing to identify lupus nephritis phenotype in electronic health records. *BMC medical informatics and decision making*, 22(Suppl 2), 348.