

Resource Summary Report

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[org.Hs.eg.db](#)

RRID:SCR_024739

Type: Tool

Proper Citation

org.Hs.eg.db (RRID:SCR_024739)

Resource Information

URL: <https://bioconductor.org/packages/org.Hs.eg.db/>

Proper Citation: org.Hs.eg.db (RRID:SCR_024739)

Description: Software R package as object for simple mapping of Entrez Gene identifiers <https://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=gene> to all possible GenBank accession numbers.

Resource Type: software resource, software toolkit

Keywords: simple mapping of Entrez Gene identifiers, GenBank accession numbers,

Availability: Free, Available for download, Freely available

Resource Name: org.Hs.eg.db

Resource ID: SCR_024739

License: Artistic v2.0

Record Creation Time: 20231204T190628+0000

Record Last Update: 20260829T183330+0000

Ratings and Alerts

No rating or validation information has been found for org.Hs.eg.db.

No alerts have been found for org.Hs.eg.db.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 45 mentions in open access literature.

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

Song Y, et al. (2026) Variations of structural-functional coupling in post-traumatic stress disorder are associated with underlying molecular and transcriptional features. *Psychological medicine*, 56, e196.

Xun Y, et al. (2026) Integrative analysis of a novel snoRNA-based prognostic signature in patients with breast cancer. *Frontiers in oncology*, 16, 1779697.

Subba P, et al. (2026) Prenatal acoustic communication triggers adaptive vascular programming in the developing avian brain. *The Journal of experimental biology*, 229(11).

Sævarsson T, et al. (2026) Enhanced IFN γ response in dedifferentiated melanoma cells is due to chromatin remodeling as revealed by ATAC-seq. *Cell communication and signaling : CCS*, 24(1).

Zhao J, et al. (2026) Identification and Validation of Mitophagy-Related Biomarkers in Colorectal Cancer: An Integrated Analysis of Single-Cell Transcriptome and Mendelian Randomization. *Genetics research*, 2026(1), e5579542.

Shen P, et al. (2026) High RMI1 expression is associated with cancer cell progression and poor prognosis in prostate cancer. *Oncology reports*, 55(4).

Ni W, et al. (2026) Integrative bioinformatics and experimental analysis reveals FRA1 as a key mediator of tubulointerstitial inflammation in lupus nephritis. *Molecular medicine reports*, 33(4).

Yan X, et al. (2026) NDC80 kinetochore complex component is a potential molecular target of adenoid cystic carcinoma. *Oncology letters*, 31(3), 114.

Xia Z, et al. (2026) PI3K and MAPK Signaling Nodes Serve as Divergent Drivers of Phenotypic Plasticity in Cancer-Associated Fibroblasts in Colorectal Cancer. *Cancer research*, 86(12), 3040.

Cai Y, et al. (2026) Identification of STMN1 as a lactylation-related driver of lung cancer progression using Mendelian randomization. *Molecular medicine reports*, 33(5).

Liu B, et al. (2026) Shared molecular features and candidate pathways underlying gastric cancer-depression comorbidity: a systems biology analysis. *Frontiers in bioinformatics*, 6, 1836419.

Zink A, et al. (2026) Pluripotent stem-cell-based screening uncovers sildenafil as a mitochondrial disease therapy. *Cell*, 189(6), 1656.

Ramsey AC, et al. (2026) TGF- β signaling promotes astroglial activation and TDP-43 proteinopathy in organoid models of frontotemporal lobar degeneration. *The Journal of clinical investigation*, 136(14).

Li P, et al. (2025) ADORA2B promotes proliferation and migration in head and neck squamous cell carcinoma and is associated with immune infiltration. BMC cancer, 25(1), 673.

Huang CQ, et al. (2025) Elevated POSTN expression predicts poor prognosis and is associated with radioresistance in cervical cancer patients treated with radical radiotherapy. Scientific reports, 15(1), 4174.

Shen T, et al. (2025) Integrative machine learning frameworks to uncover specific protein signature in neuroendocrine cervical carcinoma. BMC cancer, 25(1), 57.

Tian D, et al. (2025) Exploring the evolving role of apolipoproteins in oncology: global trends and emerging frontiers. Journal of cancer research and clinical oncology, 151(12), 325.

Wang H, et al. (2025) Phenotypic plasticity and increased infiltration of peripheral blood-derived TREM1+ mono-macrophages following radiotherapy in rectal cancer. Cell reports. Medicine, 6(1), 101887.

He X, et al. (2025) CDT1 is a Potential Therapeutic Target for the Progression of NAFLD to HCC and the Exacerbation of Cancer. Current genomics, 26(3), 225.

Pang KL, et al. (2025) Deciphering a proliferation-essential gene signature based on CRISPR-Cas9 screening to predict prognosis and characterize the immune microenvironment in HNSCC. BMC cancer, 25(1), 756.