

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/>

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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 toolkit, software resource

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: 20260803T040851+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 24 mentions in open access literature.

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

Umei TC, et al. (2025) Serine synthesis pathway regulates cardiac differentiation from human pluripotent stem cells. *iScience*, 28(7), 112843.

An Z, et al. (2025) Identification and verification of immune and oxidative stress-related diagnostic indicators for malignant lung nodules through WGCNA and machine learning. *Scientific reports*, 15(1), 22449.

Ren X, et al. (2025) CRISPR-Cas9 screening identifies a gene signature predictive of prognosis in glioblastoma. *Scientific reports*, 15(1), 21077.

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.

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.

Huang Z, et al. (2025) Mechanistic insights into Sanbi Decoction for osteoarthritis treatment based on network pharmacology and experimental validation. *Scientific reports*, 15(1), 18707.

Mondragon-Estrada E, et al. (2025) Protocol to analyze deep-learning-predicted functional scores for noncoding de novo variants and their correlation with complex brain traits. *STAR protocols*, 6(2), 103738.

Liu HT, et al. (2025) Overexpression of STOML1 is associated with good prognosis in nasopharyngeal carcinoma. *Oncology letters*, 30(4), 461.

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.

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.

Yao XR, et al. (2025) C16orf74 is a novel prognostic biomarker and associates with immune infiltration in head and neck squamous cell carcinoma. *PloS one*, 20(5), e0322701.

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.

Scialò C, et al. (2025) Seeded aggregation of TDP-43 induces its loss of function and reveals early pathological signatures. *Neuron*, 113(10), 1614.

Arce M, et al. (2025) KRIT1 heterozygous mutations are sufficient to induce a pathological phenotype in patient-derived iPSC models of cerebral cavernous malformation. *Cell reports*, 44(5), 115576.

Zamuner FT, et al. (2025) Epigenetic profiling reveals key super-enhancer networks driving oncogenesis in HPV-positive HNSCC. *iScience*, 28(11), 113911.

Li T, et al. (2024) N6-methyladenosine-associated genetic variants in NECTIN2 and HPCAL1 are risk factors for abdominal aortic aneurysm. *iScience*, 27(4), 109419.

Pavon N, et al. (2024) Patterning ganglionic eminences in developing human brain organoids using a morphogen-gradient-inducing device. *Cell reports methods*, 4(1), 100689.

Fowler JL, et al. (2024) Lineage-tracing hematopoietic stem cell origins in vivo to efficiently make human HLF+ HOXA+ hematopoietic progenitors from pluripotent stem cells. *Developmental cell*, 59(9), 1110.

Jessa S, et al. (2024) FOXR2 targets LHX6+/DLX+ neural lineages to drive CNS neuroblastoma. *Cancer research*.

Zhang W, et al. (2024) Prediction and verification of the prognostic biomarker SLC2A2 and its association with immune infiltration in gastric cancer. *Oncology letters*, 27(2), 70.