

Resource Summary Report

Generated by PRECISE-TBI on Sep 16, 2026

Encyclopedia of Hepatocellular Carcinoma Genes Online

RRID:SCR_007636

Type: Tool

Proper Citation

Encyclopedia of Hepatocellular Carcinoma Genes Online (RRID:SCR_007636)

Resource Information

URL: <http://ehco.iis.sinica.edu.tw>

Proper Citation: Encyclopedia of Hepatocellular Carcinoma Genes Online (RRID:SCR_007636)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented August 23, 2016. ECHO is a web resource of Hepatocellular Carcinoma genes. The fundamental part of EHCO2 is the collections of thirteen gene sets related to HCC. It also contains tools to search by homology, pathway, or phenotype.

Synonyms: EHCO

Resource Type: data or information resource, database

Keywords: hepatocellular carcinoma, hepatocellular carcinoma gene, oncology

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Encyclopedia of Hepatocellular Carcinoma Genes Online

Resource ID: SCR_007636

Alternate IDs: nif-0000-02791

Alternate URLs: <https://www.hsls.pitt.edu/obrc/index.php?page=URL1151599427>

Record Creation Time: 20220129T080242+0000

Record Last Update: 20260912T200149+0000

Ratings and Alerts

No rating or validation information has been found for Encyclopedia of Hepatocellular Carcinoma Genes Online.

No alerts have been found for Encyclopedia of Hepatocellular Carcinoma Genes Online.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Jhuang HJ, et al. (2015) Gluconeogenesis, lipogenesis, and HBV replication are commonly regulated by PGC-1??-dependent pathway. Oncotarget, 6(10), 7788.

Wang J, et al. (2015) Pathway and network approaches for identification of cancer signature markers from omics data. Journal of Cancer, 6(1), 54.

Zhang Y, et al. (2013) Identification of GRB2 and GAB1 coexpression as an unfavorable prognostic factor for hepatocellular carcinoma by a combination of expression profile and network analysis. PloS one, 8(12), e85170.

Kubrycht J, et al. (2012) Virtual interactomics of proteins from biochemical standpoint. Molecular biology international, 2012, 976385.

Chen MH, et al. (2011) Gene expression-based chemical genomics identifies potential therapeutic drugs in hepatocellular carcinoma. PloS one, 6(11), e27186.

Huang Q, et al. (2011) RNA-Seq analyses generate comprehensive transcriptomic landscape and reveal complex transcript patterns in hepatocellular carcinoma. PloS one, 6(10), e26168.

Hsu CN, et al. (2007) Detection of the inferred interaction network in hepatocellular carcinoma from EHCO (Encyclopedia of Hepatocellular Carcinoma genes Online). BMC bioinformatics, 8, 66.