

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://www.fdi-lab.org) on Apr 3, 2025

HEL 92.1.7

RRID:CVCL_2481

Type: Cell Line

Proper Citation

(RRID:CVCL_2481)

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_2481

Proper Citation: (RRID:CVCL_2481)

Sex: Male

Defining Citation: [PMID:20215515](#), [PMID:22460905](#), [PMID:30285677](#), [PMID:30894373](#), [PMID:31068700](#), [PMID:31978347](#)

Comments: Donor information: Originally the patient was suffering from Hodgkin lymphoma., Omics: Transcriptome analysis by RNAseq., Omics: Transcriptome analysis by microarray., Omics: SNP array analysis., Omics: Deep quantitative proteome analysis., Omics: Deep exome analysis., Population: Caucasian., Part of: Tumor Immunology Bank (TIB) collection from Salk (transferred to ATCC in 1981)., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

Category: Cancer cell line

Name: HEL 92.1.7

Synonyms: HEL92.1.7, HEL-92.1.7, HEL-92-1-7, HEL-92_1_7, HEL-92, HEL9217

Cross References: BTO:BTO_0003320, CLO:CLO_0003682, CLO:CLO_0003683, EFO:EFO_0002193, CLDB:cl5067, CLDB:cl7132, ArrayExpress:E-MTAB-38, ArrayExpress:E-MTAB-2770, ATCC:TIB-180, BCRC:60370, BCRJ:0308, BioSample:SAMN10987842, cancercellines:CVCL_2481, Cell_Model_Passport:SIDM00593, ChEMBL-Cells:ChEMBL4295457, ChEMBL-Targets:ChEMBL4296425, CLS:300462, Cosmic:683543, Cosmic:976723, DepMap:ACH-000005, ECACC:92111706, GEO:GSM887076, GEO:GSM888146, GEO:GSM1374526, GEO:GSM1374527, IARC_TP53:21369, ICLC:HTL03002, IGRhCellID:HEL9217,

LiGeA:CCLE_672, LINCS_LDP:LCL-1076, Lonza:1042,
PharmacDB:HEL92_1_7_535_2019, Progenetix:CVCL_2481,
PubChem_Cell_line:CVCL_2481, Wikidata:Q54882504

ID: CVCL_2481

Record Creation Time: 20250131T200356+0000

Record Last Update: 20250131T201611+0000

Ratings and Alerts

No rating or validation information has been found for HEL 92.1.7.

No alerts have been found for HEL 92.1.7.

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 21 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Graham K, et al. (2024) Discovery of YAP1/TAZ pathway inhibitors through phenotypic screening with potent anti-tumor activity via blockade of Rho-GTPase signaling. *Cell chemical biology*, 31(7), 1247.

Blöchl C, et al. (2023) Transcriptionally imprinted glycomic signatures of acute myeloid leukemia. *Cell & bioscience*, 13(1), 31.

Nicosia L, et al. (2023) Therapeutic targeting of EP300/CBP by bromodomain inhibition in hematologic malignancies. *Cancer cell*, 41(12), 2136.

Saleiro D, et al. (2023) Targeting CHAF1B Enhances IFN Activity against Myeloproliferative Neoplasm Cells. *Cancer research communications*, 3(5), 943.

Wei TT, et al. (2022) Cannabinoid receptor 1 antagonist genistein attenuates marijuana-induced vascular inflammation. *Cell*, 185(10), 1676.

Roversi FM, et al. (2022) Novel inhibitor of hematopoietic cell kinase as a potential therapeutic agent for acute myeloid leukemia. *Cancer immunology, immunotherapy : CII*, 71(8), 1909.

Yang BH, et al. (2022) Epigenetics-Associated Risk Reduction of Hematologic Neoplasms in

a Nationwide Cohort Study: The Chemopreventive and Therapeutic Efficacy of Hydralazine. *Frontiers in oncology*, 12, 809014.

Polyanskaya SA, et al. (2022) SCP4-STK35/PDIK1L complex is a dual phospho-catalytic signaling dependency in acute myeloid leukemia. *Cell reports*, 38(2), 110233.

Cao Z, et al. (2021) ZMYND8-regulated IRF8 transcription axis is an acute myeloid leukemia dependency. *Molecular cell*, 81(17), 3604.

Takao S, et al. (2021) Convergent organization of aberrant MYB complex controls oncogenic gene expression in acute myeloid leukemia. *eLife*, 10.

Jin S, et al. (2020) 5-Azacytidine Induces NOXA to Prime AML Cells for Venetoclax-Mediated Apoptosis. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 26(13), 3371.

Morita K, et al. (2020) Allosteric Activators of Protein Phosphatase 2A Display Broad Antitumor Activity Mediated by Dephosphorylation of MYBL2. *Cell*, 181(3), 702.

Gier RA, et al. (2020) High-performance CRISPR-Cas12a genome editing for combinatorial genetic screening. *Nature communications*, 11(1), 3455.

Saenz DT, et al. (2020) Mechanistic basis and efficacy of targeting the β -catenin-TCF7L2-JMJD6-c-Myc axis to overcome resistance to BET inhibitors. *Blood*, 135(15), 1255.

Prigent A, et al. (2019) Acute inflammation down-regulates alpha-synuclein expression in enteric neurons. *Journal of neurochemistry*, 148(6), 746.

Lin KH, et al. (2019) Systematic Dissection of the Metabolic-Apoptotic Interface in AML Reveals Heme Biosynthesis to Be a Regulator of Drug Sensitivity. *Cell metabolism*, 29(5), 1217.

Celik H, et al. (2018) JARID2 Functions as a Tumor Suppressor in Myeloid Neoplasms by Repressing Self-Renewal in Hematopoietic Progenitor Cells. *Cancer cell*, 34(5), 741.

Tarumoto Y, et al. (2018) LKB1, Salt-Inducible Kinases, and MEF2C Are Linked Dependencies in Acute Myeloid Leukemia. *Molecular cell*, 69(6), 1017.

Jin L, et al. (2018) MAST1 Drives Cisplatin Resistance in Human Cancers by Rewiring cRaf-Independent MEK Activation. *Cancer cell*, 34(2), 315.

Chung J, et al. (2017) Erythropoietin signaling regulates heme biosynthesis. *eLife*, 6.