

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on May 18, 2025

HBL-1 [Human diffuse large B-cell lymphoma]

RRID:CVCL_4213

Type: Cell Line

Proper Citation

(RRID:CVCL_4213)

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_4213

Proper Citation: (RRID:CVCL_4213)

Sex: Male

Defining Citation: [PMID:2854303](#), [PMID:3338018](#), [PMID:8558920](#), [PMID:15028022](#), [PMID:20054396](#), [PMID:21179087](#), [PMID:23257783](#), [PMID:26787899](#), [PMID:29416618](#), [PMID:29666304](#), [PMID:30165192](#), [PMID:31510952](#)

Comments: Omics: Transcriptome analysis by microarray., Omics: Array-based CGH., Karyotypic information: Has lost chromosome Y., Population: Japanese.

Category: Cancer cell line

Name: HBL-1 [Human diffuse large B-cell lymphoma]

Synonyms: HBL1

Cross References: BTO:BTO_0002522, CLO:CLO_0037256, EFO:EFO_0022408, BioGRID_ORCS_Cell_line:747, cancercellines:CVCL_4213, Cosmic:1486586, Cosmic:1487545, Cosmic:1517652, Cosmic:2129634, Cosmic:2437308, GEO:GSM1035338, GEO:GSM1059804, GEO:GSM1374489, GEO:GSM1890017, GEO:GSM1890018, GEO:GSM1890019, GEO:GSM1890020, GEO:GSM2037050, GEO:GSM2037051, Progenetix:CVCL_4213, Wikidata:Q54881426

ID: CVCL_4213

Record Creation Time: 20250131T200317+0000

Ratings and Alerts

No rating or validation information has been found for HBL-1 [Human diffuse large B-cell lymphoma].

No alerts have been found for HBL-1 [Human diffuse large B-cell lymphoma].

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 16 mentions in open access literature.

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

Choi J, et al. (2024) Molecular targets of glucocorticoids that elucidate their therapeutic efficacy in aggressive lymphomas. *Cancer cell*, 42(5), 833.

Bolonsky A, et al. (2024) IRF4 requires ARID1A to establish plasma cell identity in multiple myeloma. *Cancer cell*, 42(7), 1185.

Witmond M, et al. (2024) Dynamic hydrogen peroxide levels reveal a rate-dependent sensitivity in B-cell lymphoma signaling. *Scientific reports*, 14(1), 4265.

Phelan JD, et al. (2024) Response to Bruton's tyrosine kinase inhibitors in aggressive lymphomas linked to chronic selective autophagy. *Cancer cell*, 42(2), 238.

Ecker V, et al. (2023) Negative feedback regulation of MAPK signaling is an important driver of chronic lymphocytic leukemia progression. *Cell reports*, 42(10), 113017.

Rodina A, et al. (2023) Systems-level analyses of protein-protein interaction network dysfunctions via epichaperomics identify cancer-specific mechanisms of stress adaptation. *Nature communications*, 14(1), 3742.

Venturutti L, et al. (2023) An Aged/Autoimmune B-cell Program Defines the Early Transformation of Extranodal Lymphomas. *Cancer discovery*, 13(1), 216.

Scheich S, et al. (2023) Targeting N-linked Glycosylation for the Therapy of Aggressive Lymphomas. *Cancer discovery*, 13(8), 1862.

Rana S, et al. (2023) Methotrexate-based PROTACs as DHFR-specific chemical probes. *Cell chemical biology*.

Donati G, et al. (2023) Oxidative stress enhances the therapeutic action of a respiratory inhibitor in MYC-driven lymphoma. *EMBO molecular medicine*, 15(6), e16910.

Wei P, et al. (2022) Mitochondrial pyruvate supports lymphoma proliferation by fueling a glutamate pyruvate transaminase 2-dependent glutaminolysis pathway. *Science advances*, 8(39), eabq0117.

Ruiz Cuevas MV, et al. (2021) Most non-canonical proteins uniquely populate the proteome or immunopeptidome. *Cell reports*, 34(10), 108815.

Yan P, et al. (2020) Molecular Stressors Engender Protein Connectivity Dysfunction through Aberrant N-Glycosylation of a Chaperone. *Cell reports*, 31(13), 107840.

Zhang J, et al. (2020) Assessing IRAK4 Functions in ABC DLBCL by IRAK4 Kinase Inhibition and Protein Degradation. *Cell chemical biology*, 27(12), 1500.

Díaz N, et al. (2018) Chromatin conformation analysis of primary patient tissue using a low input Hi-C method. *Nature communications*, 9(1), 4938.

Reddy A, et al. (2017) Genetic and Functional Drivers of Diffuse Large B Cell Lymphoma. *Cell*, 171(2), 481.