Resource Summary Report

Generated by FDI Lab - SciCrunch.org on May 14, 2025

B95-8

RRID:CVCL_1953
Type: Cell Line

Proper Citation

(ECACC Cat# 85011419, RRID:CVCL_1953)

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_1953

Proper Citation: (ECACC Cat# 85011419, RRID:CVCL_1953)

Sex: Female

Defining Citation: PMID:2153236, PMID:2824833, PMID:3017841, PMID:3145248, PMID:4333982, PMID:4346033, PMID:4357683, PMID:6265077, PMID:6294325, PMID:20454443, PMID:24380390

Comments: Omics: YY1 ChIP-seq epigenome analysis., Omics: CTCF ChIP-seq epigenome analysis., Virology: This cell line sheds the Epstein-Barr virus (EBV) strain B95-8 and is often used to transform human or other primates B lymphocytes., Characteristics: Was established by exposure extracts of the 883L cell line (Cellosaurus=CVCL_6A87)., Group: Non-human primate cell line.

Category: Transformed cell line

Name: B95-8

Synonyms: B95.8, B 95.8, B 95-8, B-95-8, B958, GM07404, GM07404A, GM07404D

Cross References: BTO:BTO_0000107, CLO:CLO_0001776, CLO:CLO_0001838, EFO:EFO_0022801, CLDB:cl364, CLDB:cl365, CLDB:cl366, CLDB:cl367, CLDB:cl368, CLDB:cl369, ArrayExpress:E-MTAB-1511, ATCC:CRL-1612, BCRJ:0042, CCRID:3101MONGNO3, CCRID:4201MON-CCTCC00015, CCRID:5301MON-KCB93008YJ, CCTCC:GDC0015, ChEMBL-Cells:CHEMBL3307906, ChEMBL-Targets:CHEMBL614276, CLS:305152, CLS:601102, Coriell:GM07404, DSMZ:ACC-100, DSMZCellDive:ACC-100, ECACC:85011419, IBRC:C10050, ICLC:ATL95004, IZSLER:BS CL 110, JCRB:JCRB9123, KCB:KCB 91004YJ, KCB:KCB 93008YJ, KCLB:21612, NCBI_Iran:C110,

PubChem_Cell_line:CVCL_1953, TKG:TKG 0186, TOKU-E:3559, Wikidata:Q54752921

ID: CVCL_1953

Vendor: ECACC

Catalog Number: 85011419

Record Creation Time: 20250131T193959+0000

Record Last Update: 20250131T194226+0000

Ratings and Alerts

No rating or validation information has been found for B95-8.

Warning: Discontinued: ATCC; CRL-1612
Omics: YY1 ChIP-seq epigenome analysis., Omics: CTCF ChIP-seq epigenome analysis., Virology: This cell line sheds the Epstein-Barr virus (EBV) strain B95-8 and is often used to transform human or other primates B lymphocytes., Characteristics: Was established by exposure extracts of the 883L cell line (Cellosaurus=CVCL_6A87)., Group: Non-human primate cell line. Warning: Discontinued: Coriell; GM07404
Omics: YY1 ChIP-seq epigenome analysis., Omics: CTCF ChIP-seq epigenome analysis., Virology: This cell line sheds the Epstein-Barr virus (EBV) strain B95-8 and is often used to transform human or other primates B lymphocytes., Characteristics: Was established by exposure extracts of the 883L cell line (Cellosaurus=CVCL_6A87)., Group: Non-human primate cell line. Warning: Discontinued: KCLB; 21612
Omics: YY1 ChIP-seq epigenome analysis., Omics: CTCF ChIP-seq epigenome analysis., Virology: This cell line sheds the Epstein-Barr virus (EBV) strain B95-8 and is often used to transform human or other primates B lymphocytes., Characteristics: Was established by exposure extracts of the 883L cell line (Cellosaurus=CVCL_6A87)., Group: Non-human primate cell line.

Data and Source Information

Source: Cellosaurus

Usage and Citation Metrics

We found 245 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

San Roman AK, et al. (2024) The human Y and inactive X chromosomes similarly modulate autosomal gene expression. Cell genomics, 4(1), 100462.

Huang J, et al. (2024) Assembly and activation of EBV latent membrane protein 1. Cell, 187(18), 4996.

Grain A, et al. (2023) Two Ways of Targeting a CD19 Positive Relapse of Acute Lymphoblastic Leukaemia after Anti-CD19 CAR-T Cells. Biomedicines, 11(2).

Castelo Rueda MP, et al. (2023) Molecular phenotypes of mitochondrial dysfunction in clinically non-manifesting heterozygous PRKN variant carriers. NPJ Parkinson's disease, 9(1), 65.

Ho JWY, et al. (2023) Comprehensive Profiling of EBV Gene Expression and Promoter Methylation Reveals Latency II Viral Infection and Sporadic Abortive Lytic Activation in Peripheral T-Cell Lymphomas. Viruses, 15(2).

El Kenz B, et al. (2023) Definition of a New HLA B*52-Restricted Rev CTL Epitope Targeted by an HIV-1-Infected Controller. Viruses, 15(2).

Chen S, et al. (2023) LMP1 mediates tumorigenesis through persistent epigenetic modifications and PGC1? upregulation. Oncology reports, 49(3).

Uddin MK, et al. (2023) Epstein-Barr Virus BBLF1 Mediates Secretory Vesicle Transport to Facilitate Mature Virion Release. Journal of virology, 97(6), e0043723.

Reguraman N, et al. (2023) Assessing the Efficacy of VLP-Based Vaccine against Epstein-Barr Virus Using a Rabbit Model. Vaccines, 11(3).

Chao TY, et al. (2023) Subcellular Distribution of BALF2 and the Role of Rab1 in the Formation of Epstein-Barr Virus Cytoplasmic Assembly Compartment and Virion Release. Microbiology spectrum, 11(1), e0436922.

Kumar R, et al. (2023) Reduced G protein signaling despite impaired internalization and ?-arrestin recruitment in patients carrying a CXCR4Leu317fsX3 mutation causing WHIM syndrome. JCI insight, 8(5).

Chen H, et al. (2023) CTCF variant begets to short stature by down-regulation of IGF1. Journal of molecular endocrinology, 70(4).

Fu X, et al. (2023) Impacts of plasma microbial lipopolysaccharide translocation on B cell perturbations and anti-CD4 autoantibody production in people with HIV on suppressive antiretroviral therapy. Cell & bioscience, 13(1), 78.

Li Z, et al. (2023) Construction of a Hantaan Virus Phage Antibody Library and Screening for Potential Neutralizing Activity. Viruses, 15(5).

Ai Q, et al. (2023) Targeting KRASG12V mutations with HLA class II-restricted TCR for the immunotherapy in solid tumors. Frontiers in immunology, 14, 1161538.

Zhou X, et al. (2023) Novel biallelic mutations in TMEM126B cause splicing defects and lead

to Leigh-like syndrome with severe complex I deficiency. Journal of human genetics, 68(4), 239.

Zhang X, et al. (2023) GMP development and preclinical validation of CAR-T cells targeting a lytic EBV antigen for therapy of EBV-associated malignancies. Frontiers in immunology, 14, 1103695.

Lee SH, et al. (2023) Characterization of a new CCCTC-binding factor binding site as a dual regulator of Epstein-Barr virus latent infection. PLoS pathogens, 19(1), e1011078.

San Roman AK, et al. (2023) The human inactive X chromosome modulates expression of the active X chromosome. Cell genomics, 3(2), 100259.

Vietzen H, et al. (2023) Ineffective control of Epstein-Barr-virus-induced autoimmunity increases the risk for multiple sclerosis. Cell, 186(26), 5705.