

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://www.fdi-lab.org) on Apr 23, 2025

## MV4-11

RRID:CVCL\_0064

Type: Cell Line

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### Proper Citation

(RRID:CVCL\_0064)

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### Cell Line Information

**URL:** [https://web.expasy.org/cellosaurus/CVCL\\_0064](https://web.expasy.org/cellosaurus/CVCL_0064)

**Proper Citation:** (RRID:CVCL\_0064)

**Sex:** Male

**Defining Citation:** [PMID:1423625](#), [PMID:2656885](#), [PMID:3496132](#), [PMID:8353274](#), [PMID:8358709](#), [PMID:9195772](#), [PMID:12529668](#), [PMID:14504097](#), [PMID:14671638](#), [PMID:15843827](#), [PMID:16408098](#), [PMID:19608861](#), [PMID:20215515](#), [PMID:20922763](#), [PMID:21552520](#), [PMID:22460905](#), [PMID:25485619](#), [PMID:25877200](#), [PMID:25984343](#), [PMID:26589293](#), [PMID:27397505](#), [PMID:30285677](#), [PMID:30629668](#), [PMID:30894373](#), [PMID:31068700](#), [PMID:35839778](#)

**Comments:** Omics: Transcriptome analysis by RNAseq., Omics: Transcriptome analysis by microarray., Omics: SNP array analysis., Omics: shRNA library screening., Omics: DNA methylation analysis., Omics: Deep quantitative phosphoproteome analysis., Omics: Deep exome analysis., Omics: Acetylation analysis by proteomics., Karyotypic information: 48,XY,t(4;11)(q21;q23),+8,+19 (ATCC=CRL-9591)., Characteristics: CSF2 and IL3 dependent., Population: Caucasian., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

**Category:** Cancer cell line

**Name:** MV4-11

**Synonyms:** MV-4-11, MV-4:11, MV4:11, MV 4;11, MV4;11, MV411, MV(4;11), MV4II

**Cross References:** BTO:BTO\_0006413, CLO:CLO\_0007906, EFO:EFO\_0002242, CLDB:cl3610, CLDB:cl3611, AddexBio:C0003025/4955, ArrayExpress:E-MTAB-38, ArrayExpress:E-MTAB-2706, ArrayExpress:E-MTAB-2770, ArrayExpress:E-MTAB-3610,

ATCC:CRL-9591, ATCC:HTB-189, BioGRID\_ORCS\_Cell\_line:138, BioSample:SAMN01821582, BioSample:SAMN01821649, BioSample:SAMN03473104, BioSample:SAMN10988366, cancercellines:CVCL\_0064, CCRID:3101HUMSCSP5031, Cell\_Model\_Passport:SIDM00657, ChEMBL-Cells:CHEMBL3308063, ChEMBL-Targets:CHEMBL613835, CLS:300295, Cosmic:787416, Cosmic:798673, Cosmic:908156, Cosmic:975284, Cosmic:994182, Cosmic:996315, Cosmic:1012104, Cosmic:1037736, Cosmic:1078732, Cosmic:1089516, Cosmic:1127256, Cosmic:1150899, Cosmic:1152712, Cosmic:1181603, Cosmic:1278781, Cosmic:1281340, Cosmic:1308221, Cosmic:1319550, Cosmic:1451846, Cosmic:1476424, Cosmic:1516631, Cosmic:1524837, Cosmic:1623635, Cosmic:1696137, Cosmic:1779133, Cosmic:2131568, Cosmic:2306226, Cosmic:2392908, Cosmic:2393011, Cosmic:2542841, Cosmic:2750867, Cosmic-CLP:908156, DepMap:ACH-000045, DSMZ:ACC-102, DSMZCellDive:ACC-102, EGA:EGAS00001000610, EGA:EGAS00001000978, EGA:EGAS00001002554, GDSC:908156, GEO:GSM236796, GEO:GSM236832, GEO:GSM482560, GEO:GSM887344, GEO:GSM888420, GEO:GSM1374694, GEO:GSM1446746, GEO:GSM1670137, IARC\_TP53:21521, IGRhCellID:MV4II, LiGeA:CCELE\_055, LINCS\_LDP:LCL-1092, Lonza:996, PharmacDB:MV411\_972\_2019, PRIDE:PXD030304, Progenetix:CVCL\_0064, PubChem\_Cell\_line:CVCL\_0064, TOKU-E:2536, Ubigen:YC-C060, Wikidata:Q54907109

**ID:** CVCL\_0064

**Record Creation Time:** 20250131T201438+0000

**Record Last Update:** 20250131T203107+0000

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## Ratings and Alerts

No rating or validation information has been found for MV4-11.

**Warning:** Discontinued: ATCC; HTB-189

Omics: Transcriptome analysis by RNAseq., Omics: Transcriptome analysis by microarray., Omics: SNP array analysis., Omics: shRNA library screening., Omics: DNA methylation analysis., Omics: Deep quantitative phosphoproteome analysis., Omics: Deep exome analysis., Omics: Acetylation analysis by proteomics., Karyotypic information: 48,XY,t(4;11)(q21;q23),+8,+19 (ATCC=CRL-9591)., Characteristics: CSF2 and IL3 dependent., Population: Caucasian., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

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## Data and Source Information

**Source:** [Cellosaurus](#)

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## Usage and Citation Metrics

We found 448 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](https://www.fdi-lab.com/sci-crunch).

Melo Garcia L, et al. (2025) Overcoming CD226-related immune evasion in acute myeloid leukemia with CD38 CAR-engineered NK cells. *Cell reports*, 44(1), 115122.

Lu MJ, et al. (2024) SLC25A51 decouples the mitochondrial NAD<sup>+</sup>/NADH ratio to control proliferation of AML cells. *Cell metabolism*, 36(4), 808.

Howard GC, et al. (2024) Ribosome subunit attrition and activation of the p53-MDM4 axis dominate the response of MLL-rearranged cancer cells to WDR5 WIN site inhibition. *eLife*, 12.

Chen M, et al. (2024) Metformin synergizes with gilteritinib in treating FLT3-mutated leukemia via targeting PLK1 signaling. *Cell reports. Medicine*, 5(7), 101645.

Qiao X, et al. (2024) Diversifying the anthracycline class of anti-cancer drugs identifies aclarubicin for superior survival of acute myeloid leukemia patients. *Molecular cancer*, 23(1), 120.

Caulier B, et al. (2024) CD37 is a safe chimeric antigen receptor target to treat acute myeloid leukemia. *Cell reports. Medicine*, 5(6), 101572.

Coleman DJL, et al. (2024) Pharmacological inhibition of RAS overcomes FLT3 inhibitor resistance in FLT3-ITD+ AML through AP-1 and RUNX1. *iScience*, 27(4), 109576.

Lee JK, et al. (2024) Pim Kinase Inhibitors Increase Gilteritinib Cytotoxicity in FLT3-ITD Acute Myeloid Leukemia Through GSK-3 $\alpha$  Activation and c-Myc and Mcl-1 Proteasomal Degradation. *Cancer research communications*, 4(2), 431.

Diepstraten ST, et al. (2024) Putting the STING back into BH3-mimetic drugs for TP53-mutant blood cancers. *Cancer cell*, 42(5), 850.

Wang X, et al. (2024) LncRNA IRAIN overcomes imatinib resistance in chronic myeloid leukemia via NF- $\kappa$ B/CD44 pathway inhibition. *iScience*, 27(6), 109851.

Howard GC, et al. (2024) Ribosome subunit attrition and activation of the p53-MDM4 axis dominate the response of MLL-rearranged cancer cells to WDR5 WIN site inhibition. *bioRxiv* : the preprint server for biology.

Li Y, et al. (2024) The predictive value of BTG1 for the response of newly diagnosed acute myeloid leukemia to decitabine. *Clinical epigenetics*, 16(1), 16.

Kansy AG, et al. (2024) Pharmacological degradation of ATR induces antiproliferative DNA replication stress in leukemic cells. *Molecular oncology*, 18(8), 1958.

Guo HZ, et al. (2024) A CD36-dependent non-canonical lipid metabolism program promotes

immune escape and resistance to hypomethylating agent therapy in AML. *Cell reports. Medicine*, 5(6), 101592.

Sun YM, et al. (2024) lncRNAs maintain the functional phase state of nucleolar prion-like protein to facilitate rRNA processing. *Molecular cell*, 84(24), 4878.

Do BT, et al. (2024) Nucleotide depletion promotes cell fate transitions by inducing DNA replication stress. *Developmental cell*, 59(16), 2203.

Lai J, et al. (2024) ULK2 Is a Key Pro-Autophagy Protein That Contributes to the High Chemoresistance and Disease Relapse in FLT3-Mutated Acute Myeloid Leukemia. *International journal of molecular sciences*, 25(1).

Bianchi M, et al. (2024) The CD33xCD123xCD70 Multispecific CD3-Engaging DARPin MP0533 Induces Selective T Cell-Mediated Killing of AML Leukemic Stem Cells. *Cancer immunology research*, 12(7), 921.

Lee S, et al. (2024) B7H6 is the predominant activating ligand driving natural killer cell-mediated killing in patients with liquid tumours: evidence from clinical, in silico, in vitro, and in vivo studies. *EBioMedicine*, 110, 105459.

Wang P, et al. (2024) Foretinib Is Effective in Acute Myeloid Leukemia by Inhibiting FLT3 and Overcoming Secondary Mutations That Drive Resistance to Quizartinib and Gilteritinib. *Cancer research*, 84(6), 905.