

Resource Summary Report

Generated by [RRID](#) on Sep 6, 2026

32D

RRID:CVCL_0118

Type: Cell Line

Proper Citation

DSMZ Cat# ACC-411, RRID:CVCL_0118

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_0118

Proper Citation: DSMZ Cat# ACC-411, RRID:CVCL_0118

Description: Cell line 32D is a Factor-dependent cell line with a species of origin Mus musculus (Mouse)

Sex: Male

Defining Citation: [PMID:25277546](#)

Comments: Characteristics: IL3 dependent.

Category: Factor-dependent cell line

Organism: Mus musculus (Mouse)

Name: 32D

Cross References: BTO:BTO_0002882, CLO:CLO_0001296, CLO:CLO_0051044, EFO:EFO_0022796, MCCL:MCC:0000010, DSMZ:ACC-411, DSMZCellDive:ACC-411, Lonza:673, RCB:RCB1145, Wikidata:Q54584854

ID: CVCL_0118

Vendor: DSMZ

Catalog Number: ACC-411

Record Creation Time: 20220427T215048+0000

Record Last Update: 20260905T184311+0000

Ratings and Alerts

No rating or validation information has been found for 32D.

No alerts have been found for 32D.

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Le BV, et al. (2026) Pol γ activity modulates sensitivity to standard therapies in DNMT3A-deficient leukemia. Cell reports. Medicine, 7(3), 102687.

Zwick M, et al. (2026) FLT3-ITD Induces CMTM6 and Enhances Immune Escape in Acute Myeloid Leukemia. Cancer research, 86(2), 367.

Kamimae-Lanning AN, et al. (2026) Metabolite-induced DNA damage drives stochastic stem cell loss and clonal hematopoiesis. Cell stem cell, 33(4), 642.

Schmidt D, et al. (2024) Oncogenic Calreticulin Induces Immune Escape by Stimulating TGF β Expression and Regulatory T-cell Expansion in the Bone Marrow Microenvironment. Cancer research, 84(18), 2985.

Bermes M, et al. (2023) Exploiting Synthetic Lethality between Germline BRCA1 Haploinsufficiency and PARP Inhibition in JAK2V617F-Positive Myeloproliferative Neoplasms. International journal of molecular sciences, 24(24).

Wünsche P, et al. (2018) Mapping Active Gene-Regulatory Regions in Human Repopulating Long-Term HSCs. Cell stem cell, 23(1), 132.