

Resource Summary Report

Generated by [dkNET](#) on Sep 12, 2026

ES-E14

RRID:CVCL_C320

Type: Cell Line

Proper Citation

RRID:CVCL_C320

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_C320

Proper Citation: RRID:CVCL_C320

Description: Cell line ES-E14 is a Embryonic stem cell with a species of origin Mus musculus (Mouse)

Sex: Male

Defining Citation: [PMID:3821905](#), [PMID:19581487](#), [PMID:23850244](#), [PMID:25004115](#), [PMID:26484140](#)

Comments: Part of: ENCODE project mouse cell lines.

Category: Embryonic stem cell

Organism: Mus musculus (Mouse)

Name: ES-E14

Synonyms: E14, E-14, ES (E14)

Cross References: BTO:BTO_0005136, EFO:EFO_0007075, 4DN:4DNSRR6CIPYP, BioSamples:SAMEA2155660, ENCODE:ENCBS033ENC, ENCODE:ENCBS042ENC, ENCODE:ENCBS043ENC, ENCODE:ENCBS054SMP, ENCODE:ENCBS324NLT, GEO:GSE53149, GEO:GSM344759, GEO:GSM344760, GEO:GSM397410, GEO:GSM397411, GEO:GSM413111, GEO:GSM413112, GEO:GSM413113, GEO:GSM413117, GEO:GSM413118, GEO:GSM413119, GEO:GSM413123, GEO:GSM413124, GEO:GSM413125, GEO:GSM413129, GEO:GSM413130, GEO:GSM413131, GEO:GSM413135, GEO:GSM413136, GEO:GSM413137, GEO:GSM413141, GEO:GSM413142, GEO:GSM413143, GEO:GSM413147, GEO:GSM413148, GEO:GSM413149, GEO:GSM413153, GEO:GSM413154, GEO:GSM413155, GEO:GSM1000121, GEO:GSM1000123, GEO:GSM1000124,

GEO:GSM1000125, GEO:GSM1000126, GEO:GSM1003750, GEO:GSM1003751, GEO:GSM1003756, GEO:GSM1003799, GEO:GSM1003807, GEO:GSM1003809, GEO:GSM1003810, GEO:GSM1014154, Wikidata:Q54832448

ID: CVCL_C320

Record Creation Time: 20220427T215835+0000

Record Last Update: 20260913T003043+0000

Ratings and Alerts

No rating or validation information has been found for ES-E14.

No alerts have been found for ES-E14.

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 44 mentions in open access literature.

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

Olmeda F, et al. (2026) Scaling and self-similarity in the formation of the embryonic epigenome. Nature physics, 22(6), 931.

Tran AJ, et al. (2026) Loss of SPECC1L in cranial neural crest cells results in increased hedgehog signaling and frontonasal dysplasia. Frontiers in physiology, 17, 1751758.

Charlton SJ, et al. (2024) The fork protection complex promotes parental histone recycling and epigenetic memory. Cell, 187(18), 5029.

Iyer DP, et al. (2024) mTOR activity paces human blastocyst stage developmental progression. Cell, 187(23), 6566.

Sandovici I, et al. (2024) Overexpression of Igf2-derived Mir483 inhibits Igf1 expression and leads to developmental growth restriction and metabolic dysfunction in mice. Cell reports, 43(9), 114750.

Sun Z, et al. (2023) Chromatin regulation of transcriptional enhancers and cell fate by the Sotos syndrome gene NSD1. Molecular cell, 83(14), 2398.

Vainorius G, et al. (2023) Ascl1 and Ngn2 convert mouse embryonic stem cells to neurons via functionally distinct paths. Nature communications, 14(1), 5341.

Carraro M, et al. (2023) DAXX adds a de novo H3.3K9me3 deposition pathway to the histone chaperone network. Molecular cell, 83(7), 1075.

Flury V, et al. (2023) Recycling of modified H2A-H2B provides short-term memory of chromatin states. *Cell*, 186(5), 1050.

Xu X, et al. (2022) Mcm2 promotes stem cell differentiation via its ability to bind H3-H4. *eLife*, 11.

Liang X, et al. (2022) E-cadherin mediates apical membrane initiation site localisation during de novo polarisation of epithelial cavities. *The EMBO journal*, 41(24), e111021.

Gopalan S, et al. (2022) Multi-CUT&Tag to simultaneously profile multiple chromatin factors. *STAR protocols*, 3(1), 101100.

Pernaute B, et al. (2022) DRP1 levels determine the apoptotic threshold during embryonic differentiation through a mitophagy-dependent mechanism. *Developmental cell*, 57(11), 1316.

Abe K, et al. (2022) Distinct patterns of RNA polymerase II and transcriptional elongation characterize mammalian genome activation. *Cell reports*, 41(13), 111865.

Tavares M, et al. (2022) JAZF1-SUZ12 dysregulates PRC2 function and gene expression during cell differentiation. *Cell reports*, 39(9), 110889.

Li R, et al. (2021) CUT&Tag-BS for simultaneous profiling of histone modification and DNA methylation with high efficiency and low cost. *Cell reports methods*, 1(8).

Houlard M, et al. (2021) MCPH1 inhibits Condensin II during interphase by regulating its SMC2-Kleisin interface. *eLife*, 10.

Li R, et al. (2021) A simple and robust method for simultaneous dual-omics profiling with limited numbers of cells. *Cell reports methods*, 1(3).

Lima A, et al. (2021) Cell competition acts as a purifying selection to eliminate cells with mitochondrial defects during early mouse development. *Nature metabolism*, 3(8), 1091.

Petryk N, et al. (2021) Genome-wide and sister chromatid-resolved profiling of protein occupancy in replicated chromatin with ChOR-seq and SCAR-seq. *Nature protocols*, 16(9), 4446.