

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://www.fdi-lab.com) on Apr 5, 2025

ECC10

RRID:CVCL_1188

Type: Cell Line

Proper Citation

(RRID:CVCL_1188)

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_1188

Proper Citation: (RRID:CVCL_1188)

Sex: Male

Defining Citation: [PMID:8392984](#), [PMID:20164919](#), [PMID:22460905](#), [PMID:26589293](#), [PMID:27397505](#), [PMID:30894373](#), [PMID:31068700](#), [PMID:35839778](#)

Comments: Omics: Transcriptome analysis by RNAseq., Omics: Transcriptome analysis by microarray., Omics: SNP array analysis., Omics: DNA methylation analysis., Omics: Deep quantitative proteome analysis., Omics: Deep exome analysis., Population: Japanese., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

Category: Cancer cell line

Name: ECC10

Synonyms: ECC-10, ECC 10

Cross References: CLO:CLO_0050796, ArrayExpress:E-MTAB-2770, ArrayExpress:E-MTAB-3610, BioSample:SAMN03472196, BioSample:SAMN10987724, cancercellines:CVCL_1188, Cell_Model_Passport:SIDM00280, ChEMBL-Cells:ChEMBL3308191, ChEMBL-Targets:ChEMBL2366117, Cosmic:848376, Cosmic:906848, Cosmic:1187269, Cosmic-CLP:906848, DepMap:ACH-000560, EGA:EGAS00001000978, GDSC:906848, GEO:GSM886994, GEO:GSM888063, GEO:GSM1669750, IARC_TP53:21318, LiGeA:CCL_472, LINCS_LDP:LCL-1897, PharmacDB:ECC10_317_2019, PRIDE:PXD030304, Progenetix:CVCL_1188, PubChem_Cell_line:CVCL_1188, RCB:RCB0983, Wikidata:Q54831922

ID: CVCL_1188

Record Creation Time: 20250131T195408+0000

Record Last Update: 20250131T200219+0000

Ratings and Alerts

No rating or validation information has been found for ECC10.

No alerts have been found for ECC10.

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Wang Z, et al. (2024) Molecular subtypes of neuroendocrine carcinomas: A cross-tissue classification framework based on five transcriptional regulators. *Cancer cell*, 42(6), 1106.