

Resource Summary Report

Generated by [NIF](#) on Sep 16, 2026

SNG-M

RRID:CVCL_1707

Type: Cell Line

Proper Citation

RRID:CVCL_1707

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_1707

Proper Citation: RRID:CVCL_1707

Description: Cell line SNG-M is a Cancer cell line with a species of origin Homo sapiens (Human)

Sex: Female

Disease: Endometrial adenocarcinoma

Defining Citation: [PMID:858140](#), [PMID:2429643](#), [PMID:2448191](#), [PMID:9436037](#), [PMID:10457904](#), [PMID:12893190](#), [PMID:20164919](#), [PMID:20215515](#), [PMID:22460905](#), [PMID:25877200](#), [PMID:26589293](#), [PMID:27397505](#), [PMID:28196595](#), [PMID:30894373](#), [PMID:30971826](#), [PMID:31068700](#), [PMID:31978347](#), [PMID:35839778](#)

Comments: Population: Japanese., Part of: MD Anderson Cell Lines Project., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE)., Problematic cell line: Possibly contaminated. HHUA (distributed by RCB) and SNG-M (distributed by JCRB) share similar STR profiles and have been established by the same research group. While HHUA was described in 1984 and SNG-M in 1977 it has not been possible to establish which of the two cell lines are derived from each other. Thus it has been decided to consider them as being sister cell lines originating from the same donor..

Category: Cancer cell line

Organism: Homo sapiens (Human)

Name: SNG-M

Synonyms: SNGM

Cross References: CLO:CLO_0037165, EFO:EFO_0022497, ArrayExpress:E-MTAB-783, ArrayExpress:E-MTAB-2770, ArrayExpress:E-MTAB-3610, BioGRID_ORCS_Cell_line:460, BioSample:SAMN03470892, BioSample:SAMN03472782, BioSample:SAMN10987627, cancercellines:CVCL_1707, Cell_Model_Passport:SIDM00373, ChEMBL-Cells:ChEMBL3308466, ChEMBL-Targets:ChEMBL1075582, Cosmic:909735, Cosmic:1152552, Cosmic:1241347, Cosmic:2030483, Cosmic-CLP:909735, DepMap:ACH-000974, EGA:EGAS00001000978, GDSC:909735, GEO:GSM827242, GEO:GSM887601, GEO:GSM888684, GEO:GSM1670461, IARC_TP53:27580, JCRB:FDSC0030, JCRB:IFO50313, JCRB:JCRB0179, LiGeA:CCLE_345, LINCS_LDP:LCL-1503, PharmacDB:SNGM_1431_2019, PRIDE:PXD030304, Progenetix:CVCL_1707, PubChem_Cell_line:CVCL_1707, Wikidata:Q54955061

ID: CVCL_1707

Record Creation Time: 20220427T221550+0000

Record Last Update: 20260913T010054+0000

Ratings and Alerts

No rating or validation information has been found for SNG-M.

Warning: Discontinued: JCRB; FDSC0030

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Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Hu X, et al. (2026) Immune gene diversity and STING1 variants in shaping cancer immunity across different genetic ancestry populations. Cell reports, 45(2), 116882.

Lum MA, et al. (2025) Small-molecule modulators of B56-PP2A restore 4E-BP function to suppress eIF4E-dependent translation in cancer cells. The Journal of clinical investigation, 135(4).

Graham K, et al. (2024) Discovery of YAP1/TAZ pathway inhibitors through phenotypic screening with potent anti-tumor activity via blockade of Rho-GTPase signaling. *Cell chemical biology*, 31(7), 1247.

Corr BR, et al. (2023) Combination CDC-like kinase inhibition (CLK)/Dual-specificity tyrosine-regulated kinase (DYRK) and taxane therapy in CTNNB1 -mutated endometrial cancer. *bioRxiv* : the preprint server for biology.