

AN3 Ca Cells | 300119

General information

Description

The An3 Ca cell line is derived from a human endometrial adenocarcinoma, a type of cancer originating from the lining of the uterus. This cell line is estrogen receptor negative (ER-) and exhibits aggressive tumorigenic potential when assessed in vivo. An3 Ca cells are used extensively in research focused on understanding the molecular and cellular mechanisms underlying endometrial cancer progression, including studies on cancer cell proliferation, metastasis, and response to therapeutic agents.

Characteristically, An3 Ca cells display an epithelial morphology and have been utilized to study the impact of various genetic and environmental factors on cancer cell behavior. Research using this cell line has contributed to identifying potential therapeutic targets and understanding the resistance mechanisms against conventional treatments. They serve as a valuable model for evaluating new drugs or treatment strategies that could be effective against aggressive forms of endometrial cancer.

Overall, the An3 Ca cell line is instrumental in advancing the scientific knowledge of endometrial adenocarcinoma, offering insights that might lead to more effective interventions for this challenging and often lethal disease.

Organism Human

Tissue Uterus, Endometrium

Disease Adenocarcinoma

Synonyms AN3_CA, AN3-CA, AN3 Ca, AN3CA, AN-3, AN3, Acanthosis Nigricans 3rd attempt-Carcinoma

Characteristics

Age 55 years

Gender Female

Ethnicity Caucasian

Morphology Epithelial-like

Cell type Epithelial

Growth properties Adherent

Regulatory Data

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Citation AN3 Ca (Cytion catalog number 300119)

Biosafety level 1

NCBI_TaxID 9606

CellosaurusAccession CVCL_0028

Biomolecular Data

Isoenzymes PGM3, 1-2, PGM1, 1, ES-D, 1, AK-1, 1-2, GLO-1, 2, G6PD, B,

Tumorigenic Yes, in nude mice. Produces undifferentiated malignant tumor, also at low frequency (22%) in the cheek pouch of cortisone treated hamsters

Ploidy status Aneuploid, Phenotype Frequency Product: 0.0054

Handling

Culture Medium EMEM (MEM Eagle), w: 2 mM L-Glutamine, w: 2.2 g/L NaHCO₃, w: EBSS (Cytion article number 820100a)

Doubling time 45 to 50 hours

Subculturing Remove the old medium from the adherent cells and wash them with PBS that lacks calcium and magnesium. For T25 flasks, use 3-5 ml of PBS, and for T75 flasks, use 5-10 ml. Then, cover the cells completely with Accutase, using 1-2 ml for T25 flasks and 2.5 ml for T75 flasks. Let the cells incubate at room temperature for 8-10 minutes to detach them. After incubation, gently mix the cells with 10 ml of medium to resuspend them, then centrifuge at 300xg for 3 minutes. Discard the supernatant, resuspend the cells in fresh medium, and transfer them into new flasks that already contain fresh medium.

Split ratio A ratio of 1:3 to 1:6 is recommended

Seeding density An initial seeding density of 3 to 4 x 10⁴ cells/cm² is recommended. Later on, 2 x 10⁴ cells/cm² will yield a confluent layer in 4 to 5 days.

Fluid renewal 2 to 3 times per week

Freeze medium As a cryopreservation medium, use complete growth medium (including FBS) + 10% DMSO for adequate post-thaw viability, or CM-1 (Cytion catalog number 800100), which includes optimized osmoprotectants and metabolic stabilizers to enhance recovery and reduce cryo-induced stress.

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Quality Control & Molecular Analysis

Sterility

Mycoplasma contamination is excluded using both PCR-based assays and luminescence-based mycoplasma detection methods.

To ensure there is no bacterial, fungal, or yeast contamination, cell cultures are subjected to daily visual inspections.

STR profile

Amelogenin: x,x
CSF1PO: 12,14,15
D13S317: 12,14
D16S539: 10,14,15
D5S818: 11,14
D7S820: 7.1,10
TH01: 9.3,10
TPOX: 8,1
vWA: 14,19,20,21
D3S1358: 17
D21S11: 29,3
D18S51: 15,17,18
Penta E: 9,16
Penta D: 9,16
D8S1179: 12,14
FGA: 23
D1S1656: 13,18.3
D6S1043: 12,13,14,15,18
D2S1338: 20,23
D12S391: 20,21,23,24,25
D19S433: 14

HLA alleles

A*: '03:01:01
B*: '44:02:01, '57:01:01
C*: '05:01:01, '06:02:01
DRB1*: '04:01:01G, '16:01:01
DQA1*: '01:02:02, '03:01:01
DQB1*: '03:02:01, '05:02:01
DPB1*: '05:01:01G, '13:01:01G
E: '01:03:02