

SaOS-2 Cells | 300331

General information

Description

Saos-2 cells are an osteosarcoma cell line derived from the primary osteogenic sarcoma of an 11-year-old Caucasian female. These cells are a widely recognized model for studying osteosarcoma and bone biology, due to their osteoblastic characteristics and the ability to produce a bone-like extracellular matrix.

Characterized by their high level of alkaline phosphatase activity and expression of bone-specific proteins such as osteocalcin and osteopontin, Saos-2 cells serve as an effective in vitro system to study bone formation and the pathophysiology of osteosarcoma. They are particularly valuable for investigating cellular responses to various biochemical stimuli and mechanical forces that mimic the bone environment.

Saos-2 cells also exhibit an aneuploid karyotype, lacking several chromosomes but with extra copies of others, typical of many cancer cell lines. They are negative for mycoplasma and possess a robust capacity for calcification, making them suitable for assays related to mineral deposition.

In the context of cancer research, Saos-2 cells are extensively used to explore the molecular mechanisms of tumorigenesis, metastasis, and the effects of anticancer drugs on osteosarcoma. The cells are also employed to study gene expression profiles associated with osteoblastic differentiation and malignancy.

Due to their high transfectability, Saos-2 cells are amenable to genetic manipulation, which allows for the study of gene function and the validation of molecular targets for therapeutic intervention. This adaptability has facilitated significant advancements in understanding the genetic and molecular basis of bone cancer and in developing targeted treatments for osteosarcoma.

Organism Human

Tissue Bone

Disease Osteosarcoma

Synonyms SAOS-2, Saos-2, SAOS 2, Saos 2, Saos2, SaOs2, SAOS2, Sarcoma OSteogenic-2, SaOS, SAOS

Characteristics

Age 11 years

Gender Female

Ethnicity Caucasian

Morphology Epithelial-like

Growth properties Monolayer, adherent

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Identifiers / Biosafety / Citation**Citation** SaOS-2 (Cytion catalog number 300331)**Biosafety level** 1**Expression / Mutation****Receptors expressed** Epidermal growth factor (EGF), transforming growth factor beta (type 1 and type 2)**Antigen expression** Blood Type B, Rh+, HLA A2, A3, Bw16, Bw47**Isoenzymes** Me-2, 1, PGM3, 1-2, PGM1, 1-2, ES-D, 2, AK-1, 1, GLO-1, 2, G6PD, B, Phenotype Frequency Product: 0.0002**Tumorigenic** No**Karyotype** The stemline chromosome number is hypotriploid with the modal number of 56 chromosomes per cell and the 2S component occurring at 13.2%. Over two-thirds of the chromosome complement consisted of structurally rearranged chromosomes. Most marker chromosomes had complex rearrangements. The origin of the segments composing these markers could not be identified. Of the identifiable markers, 6p+/q+, 7p+, 11p+, and 12p+ occasionally were present at 2 copies per cell. The Y chromosome was not detected in the QM stained preparation.**Handling****Culture Medium** DMEM:Ham's F12, w: 3.1 g/L Glucose, w: 1.6 mM L-Glutamine, w: 15 mM HEPES, w: 1.0 mM Sodium pyruvate, w: 1.2 g/L NaHCO₃ (Cytion article number 820400a)**Medium supplements** Supplement the medium with 10% FBS**Passaging solution** Accutase**Doubling time** 35 to 40 hours

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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 1:2 to 1:4 is recommended

Seeding density 1×10^4 cells/cm²

Fluid renewal 2 to 3 times per week

Freezing recovery Fast

Freeze medium CM-1 (Cytion catalog number 800100) or CM-ACF (Cytion catalog number 806100)

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Handling of cryopreserved cultures

1. Confirm that the vial remains deeply frozen upon delivery, as cells are shipped on dry ice to maintain optimal temperatures during transit.
2. Upon receipt, either store the cryovial immediately at temperatures below -150°C to ensure the preservation of cellular integrity, or proceed to step 3 if immediate culturing is required.
3. For immediate culturing, swiftly thaw the vial by immersing it in a 37°C water bath with clean water and an antimicrobial agent, agitating gently for 40-60 seconds until a small ice clump remains.
4. Perform all subsequent steps under sterile conditions in a flow hood, disinfecting the cryovial with 70% ethanol before opening.
5. Carefully open the disinfected vial and transfer the cell suspension into a 15 ml centrifuge tube containing 8 ml of room-temperature culture medium, mixing gently.
6. Centrifuge the mixture at $300 \times g$ for 3 minutes to separate the cells and carefully discard the supernatant containing residual freezing medium. Optionally, skip centrifugation but remove any remaining freezing medium after 24 hours.
7. Gently resuspend the cell pellet in 10 ml of fresh culture medium. For adherent cells, divide the suspension between two T25 culture flasks; for suspension cultures, transfer all the medium into one T25 flask to promote effective cell interaction and growth.
8. Adhere to established subculture protocols for continued growth and maintenance of the cell line, ensuring reliable experimental outcomes.

Quality control / Genetic profile / HLA

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.

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STR profile

CSF1PO: 10
D13S317: 12,13
D16S539: 12,13
D5S818: 12
D7S820: 8,1
TH01: 6,9
TPOX: 8
vWA: 18
D3S1358: 14,18
D21S11: 28,3
D18S51: 15
Penta E: 14,19
Penta D: 11,12
D8S1179: 10,12
FGA: 22,25

HLA alleles

A*: 02:01:01, 24:02:01
B*: 13:02:01, 44:27:01
C*: 06:02:01, 07:04:01
DRB1*: 11:04:01, 12:01:01
DQA1*: 05:05:01
DQB1*: 03:01:01
DPB1*: 02:01:02, 04:01:01
E: 01:01:01, 01:03:01