

Product data sheet

Human SLX4 Knockout A549 cell line (clone 14)

Catalog Number: A549KO-18633

Storage: Liquid nitrogen

Components: One vial of gene knockout cells (1x10^6 cells) and one vial of control parental

cells (1x10^6)

Product description

Engineered clonal A549 cells with SLX4 gene knockout, sequence confirmed.

Cell line description

Parental cell line: A549

Genotype: SLX4-/-

Organism: Homo sapiens (human)

Tissue: Lung

Morphology: epithelial-like Culture Properties: Adherent

Biosafety Level: 2

Medium

1. Complete culture medium: DMEM with 10% fetal bovine serum (FBS)

2. Freeze medium: FBS with 6% DMSO

Culture procedure

Thawing of frozen cells

- 1. Thaw the frozen cryovial by gentle agitation in a 37 °C water bath in 1-2 minutes.
- 2. Remove the cryovial from the water bath as soon as the contents are thawed, and decontaminate by wiping with 70% ethanol.
- 3. Transfer the thawed cell suspension to a centrifuge tube containing 10 ml of Complete culture medium, centrifuge at 500 g for 5 minutes.
- 4. Remove the medium by aspiration, resuspend the cells with 10 ml of the Complete culture medium by gently pipetting up and down.
- 5. Transfer the cells to a 10 cm cell culture dish.
- 6. Place the cells in a 37°C incubator with 5% CO2.

Sub-culturing

Volumes are given for a 10 cm cell culture dish. Increase or decrease the amount of dissociation medium needed proportionally.

- 1. Remove the medium by aspiration.
- 2. Briefly rinse the cell layer with 1xDPBS to remove all traces of serum that contains trypsin inhibitor.
- 3. Add 1 ml of Trypsin-EDTA (0.05%) solution to the dish and observe cells under an inverted microscope until cell layer is dispersed.
- 4. Add 4 ml of complete growth medium and aspirate cells by gently pipetting.
- 5. Add appropriate aliquots of the cell suspension to new culture vessels. Incubate cultures at 37°C with 5% CO2.

Sequence Verification

Three different alleles were identified.	
Type I:	
KO	
GCCCAACAGCGACTCCCAGCCTCCTCCT GCCCAACAGCGACTCCCAGCCTCCTCCTTCCTGTTTGACAACAGCAGTGCCAAG	<mark>agca caattggtcctacag</mark> cgaatgca <mark>g cagttca</mark> CCCTCCA a accccgca ca <mark>g</mark> ca ca attggtcctacagcgaatgcag cagttca
WT	
There is a 44 bp deletion in exon 2.	
Type II:	
KO	
AGAGAATGTGCCCAACAGCGACTCCCCAGCCTCCTT AGAGAATGTGCCCAACAGCGACTCCCAGCCTCCTCCTTCCT	AG TOCCTCCAAACCCCGCACAG CACAATTGGTCCTACAGGGAATGCAGCAGTTCAAGA AG TOCCTCCAAACCCCGCACAG CACAATTGGTCCTACAGCGAATGCAGCAG TTCAAGA
WT	
There is a 43 bp deletion in exon 2.	
Type III:	
KO	
GAGAANT: TSGCCAACAGGGACTCCCAGGCTCCTCCT GAGAATSTGCCCAACAGGGACTCCCAGCCTCCTTCDTGTTTTGACAACAGGATTGCCAAGTCCCTCCAAACCCCGCA	
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There is a 96 bp deletion in exon 2, leading to a deletion of 32 amino acids (191-222).