

U2OS-CRISPR-SNAPf-SEH1 | 300664

U2OS-CRISPR-SNAPf-SEH1

Description U2OS-CRISPR-SNAPf-SEH1 is a cell line derived from U2OS cells, which are a human osteosarcoma cell line. The cells are stably transfected with a CRISPR-Cas9 system targeting the SEH1 gene. The CRISPR-Cas9 system is used to generate a stable cell line with a targeted mutation in the SEH1 gene. The SEH1 gene is a member of the SNAPf family of proteins, which are involved in the regulation of gene expression. The SEH1 protein is a transcription factor that binds to the SEH1 promoter and activates the transcription of the SEH1 gene. The SEH1 protein is also involved in the regulation of the cell cycle and cell growth. The SEH1 protein is a member of the SNAPf family of proteins, which are involved in the regulation of gene expression. The SEH1 protein is a transcription factor that binds to the SEH1 promoter and activates the transcription of the SEH1 gene. The SEH1 protein is also involved in the regulation of the cell cycle and cell growth.

Organism Homo sapiens

Tissue Bone marrow

Disease Osteosarcoma

Characteristics

Age 15 days

Gender Male

Ethnicity Caucasian

Morphology Adherent

Growth properties High

References

Citation U2OS-CRISPR-SNAPf-SEH1 (ATCC CCL-219) Cytion 300664

Biosafety level 1

NCBI_TaxID 9606

Depositor EMBL

GMO Status GMO-S1: U2OS-CRISPR-SNAPf-SEH1 (U2OS-CRISPR-SNAPf-SEH1) CRISPR-Cas9 system targeting the SEH1 gene

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Flask Coating

Freezing Procedure **-78°C**

Shipping Conditions **-78°C**

Storage Conditions **-150** **196**

/ / **HLA**

Sterility