



pHKHK1-2

95693™

Description

Organism: *Homo sapiens*, human

Clone type: Clone

Shipping information: Rehydrate with TE

Storage Conditions

Product format: Dried

Intended Use

This product is intended for laboratory research use only. It is not intended for any animal or human therapeutic use, any human or animal consumption, or any diagnostic use.

BSL 1

ATCC determines the biosafety level of a material based on our risk assessment as guided by the current edition of *Biosafety in Microbiological and Biomedical Laboratories (BMBL)*, U.S. Department of Health and Human Services. It is your responsibility to understand the hazards associated with the material per your organization's policies and procedures as well as any other applicable regulations as enforced by your local or national agencies.

Certificate of Analysis

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For batch-specific test results, refer to the applicable certificate of analysis that can be found at www.atcc.org.

Insert Information

Insert size (kb): 0.9310000000000005

Type of DNA: cDNA

Insert source: Hep G2 cell line

Insert tissue: Hep G2 cell line

Insert information:

DESCRIPTION OF INSERT COMPONENT:

Disease: hepatocellular carcinoma

Cross references: DNA Seq. Acc.: X78677

Nucleotides 1-931 of the insert correspond to

Disease: hepatocellular carcinoma

nucleotides 1-931 of X78677.

Genome: Homo sapiens

Gene name: ketohexokinase (fructokinase)

Gene product: ketohexokinase (fructokinase) [KHK]

Gene symbol: KHK

Contains complete coding sequence: No

Vector Information

Construct size (kb): 5.429999828338623

Intact vector size: 4.356

Vector name: pCDM8

Type of vector: phagemid

Construction: CMV, SV40

Host range: vertebrate cells

Vector end: BstXI

Cloning sites: HindIII; XbaI; XhoI; BstXI; PstI; NotI

Markers: ampR; supF

MCS: HindIII...XbaI, ->, 2198-2655

Polylinker sites: HindIII; XbaI; XhoI; BstXI; --stuffer--BstXI; XhoI; PstI; NotI; XbaI

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Promoters: CMV; T7**Replicon:** pMB1, 1-587; M13, ←, 588-1182; polyoma, 3353-4188; SV40, 4189-4352**Terminator:** SV40 splice site and polyadenylation, →, 2656-3352

Notes

Restriction digests of the clone give the following sizes (kb): XbaI--4.2, 1.2; HindIII--5.4; BamHI--3.8, 1.6; PstI--4.5, 0.5, 0.4; SacI--4.8, 0.5.

- ATCC staff

Derived from the form C variant transcript (alternately spliced third coding exon). The first polyadenylation signal was used.

- Hum. Mol. Genet. 3: 1627-1631, 1994

The insert contains the following restriction sites (approximate kb from the 5' end): BamHI--0.2; PstI--0.24, 0.67; SacI--0.33.

- Hum. Mol. Genet. 3: 1627-1631, 1994

Material Citation

If use of this material results in a scientific publication, please cite the material in the following manner: pHKHK1-2 (ATCC 95693)

References

References and other information relating to this material are available at www.atcc.org.

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