

## **Product datasheet for SC209155**

## HBS1L (NM\_001145207) Human 3' UTR Clone

**Product data:** 

Product Type: 3' UTR Clones

Symbol: HBS1L

**Synonyms:** EF-1a; eRF3c; ERFS; HBS1; HSPC276

Mammalian Cell Neomycin

Selection:

Vector: pMirTarget (PS100062)

**ACCN:** NM\_001145207

Insert Size: 723 bp

Insert Sequence: >SC209155 3'UTR clone of NM\_001145207

The sequence shown below is from the reference sequence of NM\_001145207. The complete sequence

of this clone may contain minor differences, such as  $\ensuremath{\mathsf{SNPs}}\xspace.$ 

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AGTATACACTTTAAAATAAATATATAAAAAAGTG

**ACGCGT**AAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul



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Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms

(SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**Note:** Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um

filter is required.

**RefSeq:** <u>NM\_001145207.2</u>

Summary: This gene encodes a member of the GTP-binding elongation factor family. It is expressed in

multiple tissues with the highest expression in heart and skeletal muscle. The intergenic region

of this gene and the MYB gene has been identified to be a quantitative trait locus (QTL) controlling fetal hemoglobin level, and this region influnces erythrocyte, platelet, and

monocyte counts as well as erythrocyte volume and hemoglobin content. DNA polymorphisms at this region associate with fetal hemoglobin levels and pain crises in sickle cell disease. A single nucleotide polymorphism in exon 1 of this gene is significantly associated with severity in beta-thalassemia/Hemoglobin E. Multiple alternatively spliced transcript variants encoding

different protein isoforms have been found for this gene. [provided by RefSeq, May 2009]

**Locus ID:** 10767

MW: 29