

## **Product datasheet for SC204913**

## HAAO (NM 012205) Human 3' UTR Clone

## **Product data:**

**Product Type:** 3' UTR Clones

Product Name: HAAO (NM\_012205) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: HAAO

Synonyms: 3-HAO; h3HAO; HAO; VCRL1

**ACCN:** NM\_012205

**Insert Size:** 378 bp

Insert Sequence: >SC204913 3'UTR clone of NM\_012205

The sequence shown below is from the reference sequence of NM\_012205. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGGCCCTCAATAAAGGCTTCCTGAGGAACGCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**Restriction Sites:** Sgfl-Mlul

OTI Disclaimer: 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.

**RefSeg:** NM 012205.3



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**Summary:** 3-Hydroxyanthranilate 3,4-dioxygenase is a monomeric cytosolic protein belonging to the

> family of intramolecular dioxygenases containing nonheme ferrous iron. It is widely distributed in peripheral organs, such as liver and kidney, and is also present in low amounts in the central nervous system. HAAO catalyzes the synthesis of quinolinic acid (QUIN) from 3hydroxyanthranilic acid. QUIN is an excitotoxin whose toxicity is mediated by its ability to activate glutamate N-methyl-D-aspartate receptors. Increased cerebral levels of QUIN may participate in the pathogenesis of neurologic and inflammatory disorders. HAAO has been suggested to play a role in disorders associated with altered tissue levels of QUIN. [provided

by RefSeq, Jul 2008]

13.1

Locus ID: 23498 MW: