

Product datasheet for **SC207636**

HAGHL (NM_207112) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: HAGHL (NM_207112) Human 3' UTR Clone
Vector: pMirTarget (PS100062)
Symbol: HAGHL
Synonyms: GLO2-like/ RJD12; hydroxyacylglutathione hydrolase-like; MGC2605
ACCN: NM_207112
Insert Size: 604 bp
Insert Sequence: >SC207636 3'UTR clone of NM_207112
 The sequence shown below is from the reference sequence of NM_207112. The complete sequence of this clone may contain minor differences, such as SNPs.
 Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
AAGGCATCTGGGACTGCGTGTGGGCTGAGTGAGCATCTCTGGCTTGGGGAGGCTGCTCATTAAAGTG
CCTGCCTGCCCGCCACCCTCGGCGCCATGCTCCCGCTGGGCAGCGGGCCCTGCGCCTCACTGCACC
CCTCCCTGCAGAGAGGAGCCGGTGCACAAAGTTCACGGGCAAGGGCGTCCCGCCGACGTCCTGGAGGCG
CTATGCAAGGAGCGGGCGCGCTTGAACAGGCGGGCGAGCCGCGGCAGCCACAGGCGGGCCCTCCTT
GCGCTGCAGTGGGGCTCCTGAGTGCAGCCCCACGACTGAGCCACCCAGACCCTCACAGGGCTGGGG
CCTGCGTCCCTCCTCGTGACCTCGGCCAGCTGGACCCACATGAGGGCCACCTCTGGAACCTTCTTCGAG
GCCCTGGCCAGCCATCTGCCAGCCTCGGAGGGTGGGCAACCTGGTGCTCCCGGGTGGACACACAGGA
CCACTCAGTGGGGCTGTGGGCGCCGAGACCTGGGTGTCTGGGAAGTGGGGCACAGGGGCTCCGA
ACTATGAATAAAGCTTTGAAAGGCCAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

Restriction Sites: SgfI-MluI

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.



[View online »](#)

RefSeq: [NM_207112.1](#)

Summary: Hydrolase acting on ester bonds.[UniProtKB/Swiss-Prot Function]

Locus ID: 84264

MW: 21.2