

Product datasheet for **SC203639**

HMBS (NM_000190) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	HMBS (NM_000190) Human 3' UTR Clone
Symbol:	HMBS
Synonyms:	PBG-D; PBGD; PORC; UPS
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_000190
Insert Size:	296 bp
Insert Sequence:	>SC203639 3'UTR clone of NM_000190 The sequence shown below is from the reference sequence of NM_000190. The complete sequence of this clone may contain minor differences, such as SNPs. Blue =Stop Codon Red =Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA GCGATCGCC GTTGCACGGCAGCTTAACGATGCCCAT TA ACTGGTTTGTGGGGCACAGATGCCTGGGTTGCTGCTGTCC AGTGCCTACATCCCGGCCCTCAGTGCCCCATTCTCACTGCTATCTGGGGAGTGATTACCCCGGGAGACT GAACTGCAGGGTTCAAGCCTTCCAGGGATTTGCCTCACCTTGGGGCCTTGATGACTGCCTTGCCCTCCTC AGTATGTGGGGGTTTCTCTCTTTAGAGAAGTCCAAGCAACAGCCTTTGAATGTAACCAATCCTACTAA TAAACCAGTTCTGAAGGTG ACGCGT AAGCGGCCGCGCATCTAGATTGGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
Restriction Sites:	Sgfl-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.
RefSeq:	<u>NM_000190.4</u>



[View online »](#)

Summary: This gene encodes a member of the hydroxymethylbilane synthase superfamily. The encoded protein is the third enzyme of the heme biosynthetic pathway and catalyzes the head to tail condensation of four porphobilinogen molecules into the linear hydroxymethylbilane. Mutations in this gene are associated with the autosomal dominant disease acute intermittent porphyria. Alternatively spliced transcript variants encoding different isoforms have been described. [provided by RefSeq, Jul 2008]

Locus ID: 3145

MW: 10.5