

## Product datasheet for SC205295

### IDH3G (NM\_174869) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	IDH3G (NM_174869) Human 3' UTR Clone
Symbol:	IDH3G
Synonyms:	H-IDHG
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_174869
Insert Size:	411 bp
Insert Sequence:	<p>&gt;SC205295 3'UTR clone of NM_174869 The sequence shown below is from the reference sequence of NM_174869. The complete sequence of this clone may contain minor differences, such as SNPs. <b>Blue</b>=Stop Codon <b>Red</b>=Cloning site</p> <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA<b>GCGATCGCC</b> CGCCAGTCTCCCCTGCAGCCTCCTGT<b>AG</b>CCCGCCTCATGCACTGACCACAGCCCCAGGGGAGGATG GTACTGGATCTGGCCCCATTTCTGGCCCTTAGCTGGTGCCACTCCCTCCTGTGGCAATCCCAGTCCT GCACGTACAGGAGACAGCGGGGGTGGCAGGGCCTGGTGTGTGCCGAGCCTGCCTGTGCCTACAGAT GCACACTCCGGACATCGGGGGCCAGGGACAACATCTGAAGCCATCCAGGACGTATCCGCCACATCCG CGTCATCAACGGCCGGGCGGTGGAGGCCTAGGCTGGCCCTAGGACCTTCTTGTTTGCTCCTTGGATTCC CCCTTCCCACTCCAGCACCCAGCCAGCCTGGTACGCAGATCCCAGAATAAAGCACCTTCTCCCTA <b>ACGCGT</b>AAGCGGCCGCGGCATCTAGATTCAAGAAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG </pre>
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.



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RefSeq: [NM\\_174869.3](#)

**Summary:** Isocitrate dehydrogenases catalyze the oxidative decarboxylation of isocitrate to 2-oxoglutarate. These enzymes belong to two distinct subclasses, one of which utilizes NAD(+) as the electron acceptor and the other NADP(+). Five isocitrate dehydrogenases have been reported: three NAD(+)-dependent isocitrate dehydrogenases, which localize to the mitochondrial matrix, and two NADP(+)-dependent isocitrate dehydrogenases, one of which is mitochondrial and the other predominantly cytosolic. NAD(+)-dependent isocitrate dehydrogenases catalyze the allosterically regulated rate-limiting step of the tricarboxylic acid cycle. Each isozyme is a heterotetramer that is composed of two alpha subunits, one beta subunit, and one gamma subunit. The protein encoded by this gene is the gamma subunit of one isozyme of NAD(+)-dependent isocitrate dehydrogenase. This gene is a candidate gene for periventricular heterotopia. Several alternatively spliced transcript variants of this gene have been described, but only some of their full length natures have been determined. [provided by RefSeq, Jul 2008]

Locus ID: 3421

MW: 14.5