

Product datasheet for SC207238

OGG1 (NM 016826) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: OGG1 (NM_016826) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: OGG1

Synonyms: HMMH; HOGG1; MUTM; OGH1

ACCN: NM_016826

Insert Size: 545 bp

Insert Sequence: >SC207238 3'UTR clone of NM_016826

The sequence shown below is from the reference sequence of NM_016826. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TTTAAGAAGCAGTGTTTTAGAACAGGTTCTTAAAAAGGAACAAATAAACTCATTTAACTAAA

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 016826.3



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Summary:

This gene encodes the enzyme responsible for the excision of 8-oxoguanine, a mutagenic base byproduct which occurs as a result of exposure to reactive oxygen. The action of this enzyme includes lyase activity for chain cleavage. Alternative splicing of the C-terminal region of this gene classifies splice variants into two major groups, type 1 and type 2, depending on the last exon of the sequence. Type 1 alternative splice variants end with exon 7 and type 2 end with exon 8. All variants share the N-terminal region in common, which contains a mitochondrial targeting signal that is essential for mitochondrial localization. Many alternative splice variants for this gene have been described, but the full-length nature for every variant has not been determined. [provided by RefSeq, Aug 2008]

Locus ID: 4968 MW: 20.7