

## Product datasheet for RC221784L3V

## OriGene Technologies, Inc.

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## OGG1 (NM\_016826) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** OGG1 (NM\_016826) Human Tagged ORF Clone Lentiviral Particle

Symbol: OGG<sup>\*</sup>

Synonyms: HMMH; HOGG1; MUTM; OGH1

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-Myc-DDK-P2A-Puro (PS100092)

 Tag:
 Myc-DDK

 ACCN:
 NM\_016826

ORF Size: 1071 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC221784).

Sequence:

Cytogenetics:

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This

naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

**RefSeg:** NM 016826.1, NP 058434.1

3p25.3

 RefSeq Size:
 1909 bp

 RefSeq ORF:
 1074 bp

 Locus ID:
 4968

 UniProt ID:
 015527

Protein Families: Druggable Genome
Protein Pathways: Base excision repair





ORIGENE

**MW:** 39.5 kDa

**Gene 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]