GORIIGene
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## Product datasheet for RC200522

## OGG1 (NM_002542) Human Tagged ORF Clone

## Product data:

## Product Type: Expression Plasmids

Product Name:
OGG1 (NM_002542) Human Tagged ORF Clone

## Tag:

Symbol:
Synonyms:
Mammalian Cell
Selection:
Vector:
E. coli Selection:

Restriction Sites:
Cloning Scheme:

Myc-DDK
OGG1
HMMH; HOGG1; MUTM; OGH1
Neomycin
pCMV6-Entry (PS100001)
Kanamycin ( $25 \mathrm{ug} / \mathrm{mL}$ )
Sgfl-Mlul
ACCN:
ORF Size:
NM_002542
1035 bp


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## OTI Disclaimer:

| OTI Annotation: | This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene. |
| :---: | :---: |
| Components: | The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water). |
| Reconstitution Method: | 1. Centrifuge at $5,000 \mathrm{xg}$ for 5 min . <br> 2. Carefully open the tube and add 100 ul of sterile water to dissolve the DNA. <br> 3. Close the tube and incubate for 10 minutes at room temperature. <br> 4. Briefly vortex the tube and then do a quick spin (less than 5000 xg ) to concentrate the liquid at the bottom. <br> 5. Store the suspended plasmid at $-20^{\circ} \mathrm{C}$. The DNA is stable for at least one year from date of shipping when stored at $-20^{\circ} \mathrm{C}$. |
| RefSeq: | NM 002542.5 NP 002533.1 |
| RefSeq Size: | 1652 bp |
| RefSeq ORF: | 1038 bp |
| Locus ID: | 4968 |
| UniProt ID: | 015527 |
| Cytogenetics: | 3 p 25.3 |
| Domains: | HHH, ENDO3c |
| Protein Families: | Druggable Genome |
| Protein Pathways: | Base excision repair |
| MW: | 38.8 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]

## Product images:



