

Product datasheet for RC213611

OGG1 (NM_016827) Human Tagged ORF Clone

Product data:

Product Type: Expression Plasmids
Product Name: OGG1 (NM_016827) Human Tagged ORF Clone
Tag: Myc-DDK
Symbol: OGG1
Synonyms: HMMH; HOGG1; MUTM; OGH1
Mammalian Cell Selection: Neomycin
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
ORF Nucleotide Sequence: >RC213611 representing NM_016827
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGATCGCC**

ATGCTGCCCGCGCTTCTGCCAGGCGCATGGGGCATCGTACTCTAGCCTCCACTCCTGCCCTGTGGG
 CCTCCATCCCGTGCCTCGCTCTGAGCTGCGCCTGGACCTGGTTCTGCCTTCTGGACAATCTTCCGGTG
 GAGGGAGCAAAGTCTGCACACTGGAGTGGTGTACTAGCGGATCAAGTATGGACTGACTCAGACTGAG
 GAGCAGCTCCACTGCACTGTGTACCGAGGAGACAAGGCCAGGCTAGCAGGCCACACCAGACGAGCTGG
 AGGCCGTGCGCAAGTACTTCCAGCTAGATGTTACCTGGCTCACTGTATCACCCTGGGTTCCGTGGA
 CTCCCACTTCCAAGAGGTGGCTCAGAAATCCTCAAGGTGTGCGACTGCTGCGACAAGACCCCATCGAATGC
 CTTTTCTTTTTATCTGTTCTCTCAACAACAACATCGCCCGCATCACTGGCATGGTGGAGCGGCTGTGCC
 AGGCTTTTGGACCTCGGCTCATCCAGCTTGATGATGTCACCTACCATGGCTTCCCAGCCTCGAGGCCCT
 GGCTGGGCTCCTTGGCAATGCATT

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >RC213611 representing NM_016827
 Red=Cloning site Green=Tags(s)

MPARALLPRRMGHRRLASTPALWASIPCRSELRLDLVPSGQSFWRREQSPAHWGVLADQVWTLTQTE
 EQLHCTVYRGDKSQASRPTPDELEAVRKYFQLDVTLAQLYHHWGSVDSHFQEVAQKFQGVRLLRQDP
 IEC LFSFICSSNNNIARITGMVERLCQAFGPRLIQLDVITYHGFP
 SLQALAGPPWQCI

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

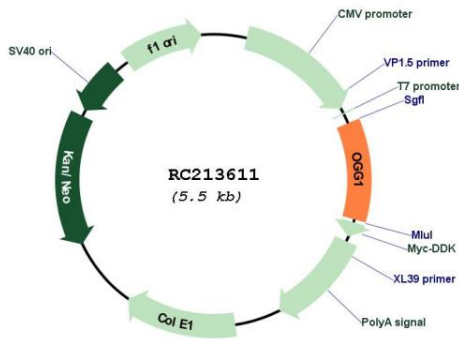


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Cytogenetics: 3p25.3
Protein Families: Druggable Genome
Protein Pathways: Base excision repair
MW: 22.2 kDa

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:



Circular map for RC213611