

Product datasheet for RC221667L4

OGG1 (NM_016820) Human Tagged Lenti ORF Clone

Product data:

OriGene Technologies, Inc.

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Expression Plasmids
OGG1 (NM_016820) Human Tagged Lenti ORF Clone
mGFP
OGG1
HMMH; HOGG1; MUTM; OGH1
Puromycin
pLenti-C-mGFP-P2A-Puro (PS100093)
Chloramphenicol (34 ug/mL)
The ORF insert of this clone is exactly the same as(RC221667).
Sgfl-Mlul
Cloning sites used for ORF Shuttling:
Sgf i ORF Miu i [GCG ATC GC]C <mark>ATG // NNŇ</mark> [ACG CGT]

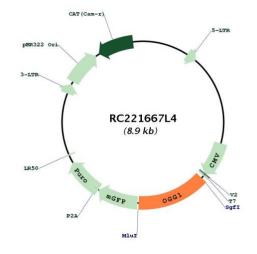


* The last codon before the Stop codon of the ORF.



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Plasmid Map:



ACCN:	NM_016820
ORF Size:	1230 bp
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 clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <u>More info</u>
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:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM 016820.3, NP 058213.1</u>
RefSeq Size:	2574 bp
RefSeq ORF:	1233 bp

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ORIGENE OGG1	(NM_016820) Human Tagged Lenti ORF Clone - RC221667L4
Locus ID:	4968
UniProt ID:	<u>015527</u>
Cytogenetics:	3p25.3
Domains:	HHH, ENDO3c
Protein Families:	Druggable Genome
Protein Pathways:	Base excision repair
MW:	45.6 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]

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