



# OGG1 Rabbit mAb

<b>Catalog No</b>	YP-rAb-17777
<b>Isotype</b>	IgG
<b>Reactivity</b>	Human
<b>Applications</b>	WB,IHC,IF,ELISA
<b>Gene Name</b>	OGG1 MMH MUTM OGH1
<b>Protein Name</b>	N-glycosylase/DNA lyase [Includes: 8-oxoguanine DNA glycosylase lyase (AP lyase) (EC 4.2.99.18)]
<b>Purification Process</b>	Protein A
<b>Specificity</b>	Endogenous
<b>Formulation</b>	PBS, 50% glycerol, 0.05% Proclin 300, 0.05%BSA
<b>Source</b>	Monoclonal, Rabbit,IgG
<b>Dilution</b>	IHC 1:200-1:1000; WB 1:10000-1:50000; IF 1:200-1:1000; ELISA 1:5000-1:20000; Note: For IHC, we suggest antigen retrieval with TE buffer pH 9.0
<b>Concentration</b>	0.5 mg/ml
<b>Purity</b>	≥90%
<b>Storage Stability</b>	-15° C to -25° C/1 year(Do not lower than -25° C)
<b>Synonyms</b>	
<b>Observed Band</b>	39kD
<b>Calculated Molecular Weight</b>	39kD
<b>Cell Pathway</b>	Nucleus, nucleoplasm . Nucleus speckle . Nucleus matrix . Together with APEX1 is recruited to nuclear speckles in UVA-irradiated cells.; [Isoform 1A]: Nucleus.; [Isoform 2A]: Mitochondrion.
<b>Tissue Specificity</b>	Ubiquitous.
<b>Function</b>	Catalytic activity:The C-O-P bond 3' to the apurinic or apyrimidinic site in DNA is broken by a beta-elimination reaction, leaving a 3'-terminal unsaturated sugar and a product with a terminal 5'-phosphate.; Disease:Defects in OGG1 are a cause of renal cell carcinoma (RCC1) [MIM:144700].; Disease:Defects in OGG1 are associated with tumor formation.; Function:DNA repair enzyme that incises DNA at 8-oxoG residues. Excises 7,8-dihydro-8-oxoguanine and 2,6-diamino-4-hydroxy-5-N-methylformamidopyrimidine (FAPY) from damaged DNA. Has a beta-lyase activity that nicks DNA 3' to the lesion.; similarity:Belongs to the type-1 OGG1 family.; tissue specificity:Ubiquitous.;





## Background

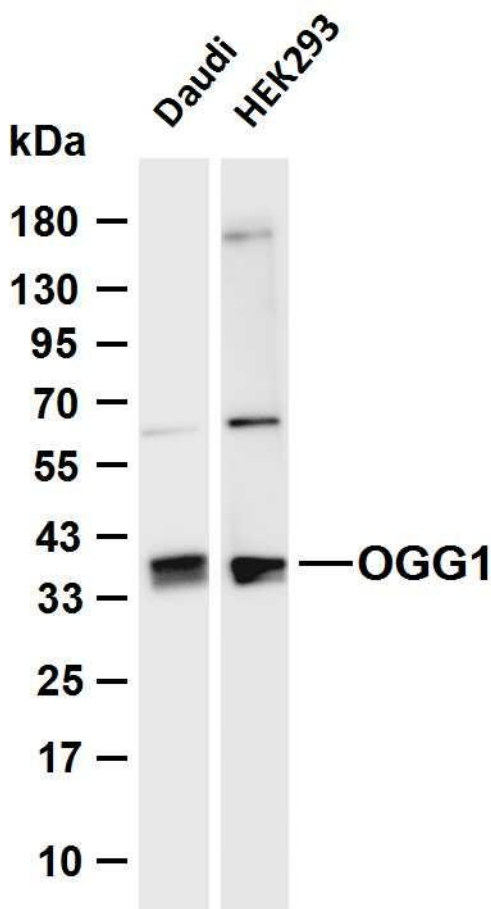
This gene encodes the enzyme responsible for the excision of 8-oxoguanine, a mutagenic base byproduct which occurs as a result of exposure to reactive oxygen. The action of this enzyme includes lyase activity for chain cleavage. Alternative splicing of the C-terminal region of this gene classifies splice 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],

## matters needing attention

Avoid repeated freezing and thawing!

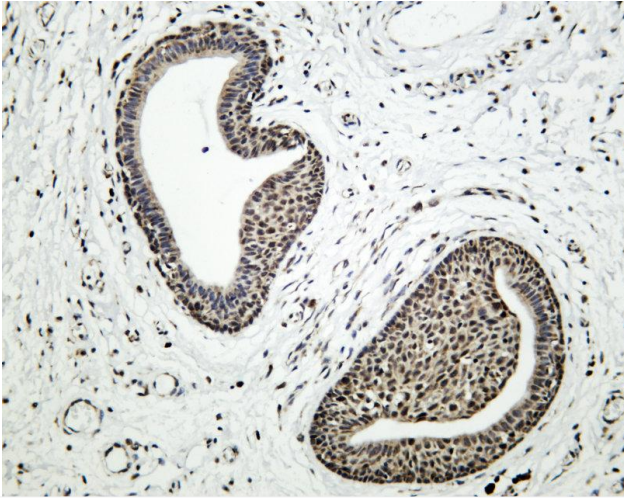
## Usage suggestions

This product can be used in immunological reaction related experiments. For more information, please consult technical personnel.



Various whole cell lysates were separated by 4-20% SDS-PAGE, and the membrane was blotted with anti-OGG1 antibody. The HRP-conjugated Goat anti-Rabbit IgG (H + L) antibody was used to detect the antibody. Lane 1: Daudi Lane 2: HEK293 Predicted band size: 39kDa Observed band size: 39kDa





Human tonsil was stained with anti-OGG1 Rabbit antibody

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