

KD-Validated Anti-Uracil DNA Glycosylase Rabbit Monoclonal Antibody

Rabbit monoclonal antibody Catalog # AGI1748

Specification

KD-Validated Anti-Uracil DNA Glycosylase Rabbit Monoclonal Antibody - Product Information

Application WB, FC, ICC
Primary Accession P13051
Reactivity Human
Clonality Monoclonal
Isotype Rabbit IgG

Calculated MW Predicted, 35 kDa , observed , 32 kDa KDa

Gene Name UN

Aliases UNG; Uracil DNA Glycosylase; UNG1; UDG;

HIGM4; UNG2; DGU; Uracil-DNA

Glycosylase 1, Uracil-DNA Glycosylase 2;

Uracil-DNA Glycosylase; UNG15; EC

3.2.2.27; EC 3.2.2; HIGM5

Immunogen A synthesized peptide derived from human

UNG

KD-Validated Anti-Uracil DNA Glycosylase Rabbit Monoclonal Antibody - Additional Information

Gene ID 7374

Other Names

KD-Validated Anti-Uracil DNA Glycosylase Rabbit Monoclonal Antibody - Protein Information

Name UNG {ECO:0000255|HAMAP-Rule:MF 03166}

Function

Uracil-DNA glycosylase that hydrolyzes the N-glycosidic bond between uracil and deoxyribose in single- and double-stranded DNA (ssDNA and dsDNA) to release a free uracil residue and form an abasic (apurinic/apyrimidinic; AP) site. Excises uracil residues arising as a result of misincorporation of dUMP residues by DNA polymerase during replication or due to spontaneous or enzymatic deamination of cytosine (PubMed:12958596, PubMed:15967827, PubMed:17101234, PubMed:22521144, PubMed:7671300, PubMed:8900285, PubMed:<a href="http://www.uniprot.org/citations/9016624"



target=" blank">9016624, PubMed:9776759). Mediates error-free base excision repair (BER) of uracil at replication forks. According to the model, it is recruited by PCNA to S-phase replication forks to remove misincorporated uracil at U:A base mispairs in nascent DNA strands. Via trimeric RPA it is recruited to ssDNA stretches ahead of the polymerase to allow detection and excision of deaminated cytosines prior to replication. The resultant AP sites temporarily stall replication, allowing time to repair the lesion (PubMed: 22521144). Mediates mutagenic uracil processing involved in antibody affinity maturation. Processes AICDA-induced U:G base mispairs at variable immunoglobulin (Ig) regions leading to the generation of transversion mutations (PubMed:12958596). Operates at switch sites of Ig constant regions where it mediates Ig isotype class switch recombination. Excises AICDA-induced uracil residues forming AP sites that are subsequently nicked by APEX1 endonuclease. The accumulation of staggered nicks in opposite strands results in double strand DNA breaks that are finally resolved via non-homologous end joining repair pathway (By similarity) (PubMed:12958596).

Cellular Location

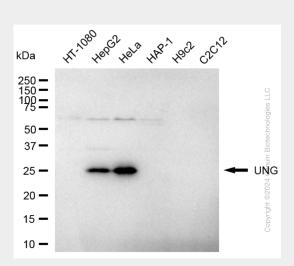
[Isoform 1]: Mitochondrion

KD-Validated Anti-Uracil DNA Glycosylase Rabbit Monoclonal Antibody - Protocols

Provided below are standard protocols that you may find useful for product applications.

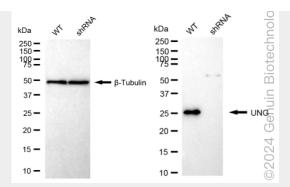
- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- Immunoprecipitation
- Flow Cytomety
- Cell Culture

KD-Validated Anti-Uracil DNA Glycosylase Rabbit Monoclonal Antibody - Images

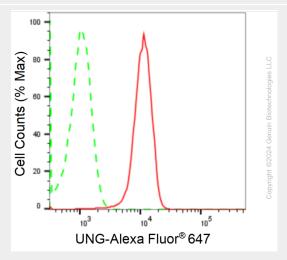


Western blotting analysis using anti-UNG antibody (Cat#AGI1748). Total cell lysates (30 μ g) from various cell lines were loaded and separated by SDS-PAGE. The blot was incubated with anti-UNG antibody (Cat#AGI1748, 1:5,000) and HRP-conjugated goat anti-rabbit secondary antibody respectively.

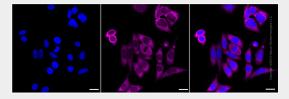




Western blotting analysis using anti-UNG antibody (Cat#AGI1748). UNG expression in wild type (WT) and UNG shRNA knockdown (KD) HeLa cells with 20 μ g of total cell lysates. β -Tubulin serves as a loading control. The blot was incubated with anti-UNG antibody (Cat#AGI1748, 1:5,000) and HRP-conjugated goat anti-rabbit secondary antibody respectively.



Flow cytometric analysis of UNG expression in HepG2 cells using anti-UNG antibody (Cat#AGI1748, 1:2,000). Green, isotype control; red, UNG.



Immunocytochemical staining of HepG2 cells with anti-UNG antibody (Cat#AGI1748, 1:1,000). Nuclei were stained blue with DAPI; UNG was stained magenta with Alexa Fluor® 647. Images were taken using Leica stellaris 5. Protein abundance based on laser Intensity and smart gain: Medium. Scale bar: $20~\mu m$.