Elabscience®

(KO Validated) DNMT1 Polyclonal Antibody

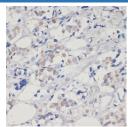
catalog number: E-AB-66916

Note: Centrifuge before opening to ensure complete recovery of vial contents.

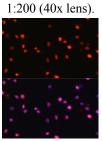
| Description | |
|--------------|--|
| Reactivity | Human;Mouse;Rat |
| Immunogen | Recombinant protein of human DNMT1. |
| Host | Rabbit |
| Is otype | IgG |
| Purification | Affinity purification |
| Buffer | Phosphate buffered solution, pH 7.4, containing 0.05% stabilizer and 50% glycerol. |
| Applications | Recommended Dilution |

IHC 1:50-1:200 IF 1:50-1:200

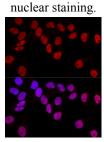
Data



Immunohistochemistry of paraffin-embedded Human breast Immunohistochemistry of paraffin-embedded Human tonsil cancer using DNMT1 Polyclonal Antibody at dilution of



Immunofluorescence analysis of C6 cells using DNMT1 Polyclonal Antibody at dilution of 1:100. Blue: DAPI for

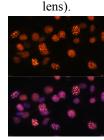


Confocal immunofluorescence analysis of HeLa cells using DNMT1 Polyclonal Antibody at dilution of 1:100. Blue: DAPI for nuclear staining.

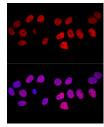
Preparation & Storage

For Research Use Only

Toll-free: 1-888-852-8623 Web:www.elabscience.com using DNMT1 Polyclonal Antibody at dilution of 1:200 (40x



Immunofluorescence analysis of NIH/3T3 cells using DNMT1 Polyclonal Antibody at dilution of 1:100. Blue: DAPI for nuclear staining.



Confocal immunofluorescence analysis of HeLa cells using DNMT1 Polyclonal Antibody at dilution of 1:100. Blue: DAPI for nuclear staining.

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Elabscience Bionovation Inc. A Reliable Research Partner in Life Science and Medicine

Storage Shipping Store at -20°C Valid for 12 months. Avoid freeze / thaw cycles. The product is shipped with ice pack,upon receipt,store it immediately at the temperature recommended.

Background

This gene encodes an enzyme that transfers methyl groups to cytosine nucleotides of genomic DNA. This protein is the major enzyme responsible for maintaining methylation patterns following DNA replication and shows a preference for hemi-methylated DNA. Methylation of DNA is an important component of mammalian epigenetic gene regulation. Aberrant methylation patterns are found in human tumors and associated with developmental abnormalities. Variation in this gene has been associated with cerebellar ataxia, deafness, and narcolepsy, and neuropathy, hereditary sensory, type IE. Alternative splicing results in multiple transcript variants.