DNMT3B Rabbit mAb

Catalog No.: A22658 Recombinant



Basic Information

Observed MW

100kDa

Calculated MW

96kDa

Category

Primary antibody

Applications

WB, ELISA, ChIP, ChIP-seq

Cross-Reactivity

Human, Mouse

CloneNo number

ARC57702

Background

CpG methylation is an epigenetic modification that is important for embryonic development, imprinting, and X-chromosome inactivation. Studies in mice have demonstrated that DNA methylation is required for mammalian development. This gene encodes a DNA methyltransferase which is thought to function in de novo methylation, rather than maintenance methylation. The protein localizes primarily to the nucleus and its expression is developmentally regulated. Mutations in this gene cause the immunodeficiency-centromeric instability-facial anomalies (ICF) syndrome. Eight alternatively spliced transcript variants have been described. The full length sequences of variants 4 and 5 have not been determined.

Recommended Dilutions

WB 1:1000 - 1:5000

ELISA Recommended starting

concentration is 1 µg/mL. Please optimize the concentration based on your specific assay

requirements.

5μg antibody for **ChIP**

10μg-15μg of Chromatin

1:50 - 1:100 ChIP-seq

Immunogen Information

Gene ID Swiss Prot 1789 Q9UBC3

Immunogen

Recombinant protein (or fragment). This information is considered to be commercially sensitive.

Synonyms

ICF; ICF1; FSHD4; M.HsallIB; DNMT3B

Contact

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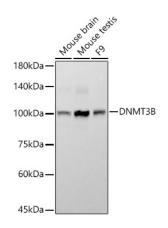
Product Information

Source Isotype **Purification** Rabbit IgG Affinity purification

Storage

Store at -20°C. Avoid freeze / thaw cycles.

Buffer: PBS with 0.05% proclin300,0.05% BSA,50% glycerol,pH7.3.



Western blot analysis of various lysates using DNMT3B Rabbit mAb (A22658) at 1:2000 dilution incubated overnight at 4° C.

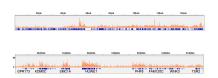
Secondary antibody: HRP-conjugated Goat anti-Rabbit IgG (H+L) (AS014) at 1:10000 dilution.

Lysates/proteins: 25µg per lane.

Blocking buffer: 3% nonfat dry milk in TBST.

Detection: ECL Basic Kit (RM00020).

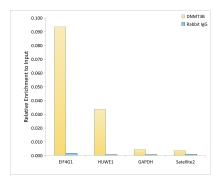
Exposure time: 60s.



Chromatin immunoprecipitation was performed with 14 μ g of cross-linked chromatin from A-549 cells using 5 μ g of DNMT3B Rabbit mAb (A22658). DNA libraries were prepared using Scale ssDNA-seq Lib Prep Kit for Illumina V2 (RK20228). The ChIP sequencing results indicate the enrichment pattern of DNMT3B across chromosome X (upper panel) and the genomic region encompassing HUWE1, a representative gene enriched in DNMT3B (lower panel).



Chromatin immunoprecipitation was performed with 14 μ g of cross-linked chromatin from NCCIT cells using 5 μ g of DNMT3B Rabbit mAb (A22658). DNA libraries were prepared using Scale ssDNA-seq Lib Prep Kit for Illumina V2 (RK20228). The ChIP sequencing results indicate the enrichment pattern of DNMT3B in the representative genomic region surrounding HUWE1 gene.



Chromatin immunoprecipitation was performed with 10 μg of cross-linked chromatin from NCCIT cells, using 5 μg of DNMT3B antibody (A22658) and Rabbit IgG isotype control (AC042). The enrichment of immunoprecipitated DNA at different genomic loci was examined by quantitative PCR. The histogram compares the ratio of the immunoprecipitated DNA to the input at given loci.