

Nibrin Rabbit mAb

Catalog # AP75860

Product Information

Application WB, IP, ICC
Primary Accession O60934
Host Rabbit

Clonality Monoclonal Antibody

Calculated MW 84959

Additional Information

Gene ID 4683

Other Names NBN

Dilution WB~~1/500-1/1000 IP~~N/A ICC~~N/A

Format 50mM Tris-Glycine(pH 7.4), 0.15M NaCl, 40%Glycerol, 0.01% sodium azide and

0.05% BSA.

Storage Store at 4°C short term. Aliquot and store at -20°C long term. Avoid

freeze/thaw cycles.

Protein Information

Name NBN (HGNC:7652)

Function Component of the MRN complex, which plays a central role in double-strand

break (DSB) repair, DNA recombination, maintenance of telomere integrity and meiosis (PubMed:10888888, PubMed:15616588, PubMed:18411307,

PubMed: <u>18583988</u>, PubMed: <u>18678890</u>, PubMed: <u>19759395</u>, PubMed: <u>23115235</u>, PubMed: <u>28216226</u>, PubMed: <u>28867292</u>,

PubMed:<u>23115235</u>, PubMed:<u>28216226</u>, PubMed:<u>28867292</u>, PubMed:<u>9705271</u>). The MRN complex is involved in the repair of DNA

double-strand breaks (DSBs) via homologous recombination (HR), an error-free mechanism which primarily occurs during S and G2 phases (PubMed:19759395, PubMed:28867292, PubMed:9705271). The complex (1) mediates the end resection of damaged DNA, which generates proper single-stranded DNA, a key initial steps in HR, and is (2) required for the recruitment of other repair factors and efficient activation of ATM and ATR upon DNA damage (PubMed:19759395, PubMed:9705271). The MRN complex possesses single-strand endonuclease activity and double-strand-specific 3'-5' exonuclease activity, which are provided by MRE11, to initiate end resection, which is required for single-strand invasion and recombination

(PubMed: 19759395, PubMed: 28867292, PubMed: 9705271). Within the MRN complex, NBN acts as a protein-protein adapter, which specifically recognizes and binds phosphorylated proteins, promoting their recruitment to DNA

damage sites (PubMed: 12419185, PubMed: 15616588, PubMed: 18411307, PubMed:18582474, PubMed:18583988, PubMed:18678890, PubMed: 19759395, PubMed: 19804756, PubMed: 23762398, PubMed:24534091, PubMed:27814491, PubMed:27889449, PubMed:33836577). Recruits MRE11 and RAD50 components of the MRN complex to DSBs in response to DNA damage (PubMed:12419185, PubMed: 18411307, PubMed: 18583988, PubMed: 18678890, PubMed: 24534091, PubMed: 26438602). Promotes the recruitment of PI3/PI4-kinase family members ATM, ATR, and probably DNA-PKcs to the DNA damage sites, activating their functions (PubMed: 15064416, PubMed: 15616588, PubMed: 15790808, PubMed: 16622404, PubMed: <u>22464731</u>, PubMed: <u>30952868</u>, PubMed: <u>35076389</u>). Mediates the recruitment of phosphorylated RBBP8/CtIP to DSBs, leading to cooperation between the MRN complex and RBBP8/CtIP to initiate end resection (PubMed: 19759395, PubMed: 27814491, PubMed: 27889449, PubMed:33836577). RBBP8/CtIP specifically promotes the endonuclease activity of the MRN complex to clear DNA ends containing protein adducts (PubMed: 27814491, PubMed: 27889449, PubMed: 30787182, PubMed:33836577). The MRN complex is also required for the processing of R-loops (PubMed:31537797). NBN also functions in telomere length maintenance via its interaction with TERF2: interaction with TERF2 during G1 phase preventing recruitment of DCLRE1B/Apollo to telomeres (PubMed: 10888888, PubMed: 28216226). NBN also promotes DNA repair choice at dysfunctional telomeres: NBN phosphorylation by CDK2 promotes non- homologous end joining repair at telomeres, while unphosphorylated NBN promotes microhomology-mediated end-joining (MMEJ) repair (PubMed: <u>28216226</u>). Enhances AKT1 phosphorylation possibly by association with the mTORC2 complex (PubMed:23762398).

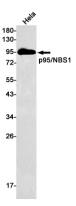
Cellular Location

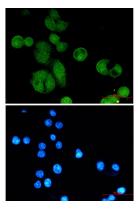
Nucleus. Chromosome. Nucleus, PML body. Chromosome, telomere Note=Localizes to discrete nuclear foci after treatment with genotoxic agents (PubMed:10783165, PubMed:26215093, PubMed:26438602). Localizes to DNA double-strand breaks (DSBs); recruited to DNA damage sites via association with phosphorylated proteins, such as phosphorylated H2AX, phosphorylated MDC1 and phosphorylated RAD17 (PubMed:12419185, PubMed:18411307, PubMed:18582474, PubMed:18583988, PubMed:18678890, PubMed:19338747, PubMed:23115235, PubMed:24534091, PubMed:26438602) Acetylation of 'Lys-5' of histone H2AX (H2AXK5ac) promotes NBN/NBS1 assembly at the sites of DNA damage (PubMed:26438602)

Tissue Location

Ubiquitous (PubMed:9590180). Expressed at high levels in testis (PubMed:9590180).

Images





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