

Product Name: Phospho-DNA PKcs (S2056) (5N13)
Rabbit Monoclonal Antibody
Catalog #: AMRe05886

Summary

Production Name	Phospho-DNA PKcs (S2056) (5N13) Rabbit Monoclonal Antibody
Description	Rabbit Monoclonal Antibody
Host	Rabbit
Application	WB,IHC-P,ICC/IF,IF-P
Reactivity	Human

Performance

Conjugation	Unconjugated
Modification	Phospho Antibody
Isotype	IgG
Clonality	Monoclonal
Form	Liquid
Storage	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles. Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.02% New type
Buffer	preservative N and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid freeze / thaw cycle.
Purification	Affinity purification

Immunogen

Gene Name	PRKDC
Alternative Names	DNA- PKcs, DNA-dependent protein kinase catalytic subunit, DNPK1, EC 2.7.11.1, P460, PRKD, PRKDC, XRCC7, kinase DNA-PK
Gene ID	5591.0
SwissProt ID	P78527.

Application

Dilution Ratio	WB 1:1000-1:2000, IHC-P/IF-P 1:100, ICC/IF 1:20-1:100
Molecular Weight	469kDa

Background

The PRKDC gene encodes the catalytic subunit of a nuclear DNA-dependent serine/threonine protein kinase (DNA-PK). The second component is the autoimmune antigen Ku (MIM 152690), which is encoded by the G22P1 gene on chromosome 22q. On its own, the catalytic subunit of DNA-PK is inactive and relies on the G22P1 component to direct it to the DNA and trigger its kinase activity; PRKDC must be bound to DNA to express its catalytic properties. Serine/threonine-protein kinase that acts as a molecular sensor for DNA damage (PubMed: [11955432](http://www.uniprot.org/citations/11955432)), PubMed: [12649176](http://www.uniprot.org/citations/12649176)), PubMed: [14734805](http://www.uniprot.org/citations/14734805)), PubMed: [33854234](http://www.uniprot.org/citations/33854234)). Involved in DNA non-homologous end joining (NHEJ) required for double-strand break (DSB) repair and V(D)J recombination (PubMed: [11955432](http://www.uniprot.org/citations/11955432)), PubMed: [12649176](http://www.uniprot.org/citations/12649176)), PubMed: [14734805](http://www.uniprot.org/citations/14734805)), PubMed: [33854234](http://www.uniprot.org/citations/33854234)). Must be bound to DNA to express its catalytic properties (PubMed: [11955432](http://www.uniprot.org/citations/11955432)). Promotes processing of hairpin DNA structures in V(D)J recombination by activation of the hairpin endonuclease artemis (DCLRE1C) (PubMed: [11955432](http://www.uniprot.org/citations/11955432)). Recruited by XRCC5 and XRCC6 to DNA ends and is required to (1) protect and align broken ends of DNA, thereby preventing their degradation, (2) and sequester the DSB for repair by NHEJ (PubMed: [15574326](http://www.uniprot.org/citations/15574326)), PubMed: [11955432](http://www.uniprot.org/citations/11955432)), PubMed: [12649176](http://www.uniprot.org/citations/12649176)), PubMed: [14734805](http://www.uniprot.org/citations/14734805)), PubMed: [33854234](http://www.uniprot.org/citations/33854234)). Act as a scaffold protein to aid the localization of DNA repair proteins to the site of damage (PubMed: [15574326](http://www.uniprot.org/citations/15574326)), PubMed: [11955432](http://www.uniprot.org/citations/11955432)), PubMed: [12649176](http://www.uniprot.org/citations/12649176)), PubMed: [14734805](http://www.uniprot.org/citations/14734805)). The assembly of the DNA-PK complex at DNA ends is also required for the NHEJ ligation step (PubMed: [15574326](http://www.uniprot.org/citations/15574326)), PubMed: [11955432](http://www.uniprot.org/citations/11955432)), PubMed: [12649176](http://www.uniprot.org/citations/12649176)), PubMed: [14734805](http://www.uniprot.org/citations/14734805)). Found at the ends of chromosomes, suggesting a further role in the maintenance of

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telomeric stability and the prevention of chromosomal end fusion (By similarity). Also involved in modulation of transcription (PubMed:15574326, PubMed:11955432, PubMed:12649176, PubMed:14734805). As part of the DNA-PK complex, involved in the early steps of ribosome assembly by promoting the processing of precursor rRNA into mature 18S rRNA in the small-subunit processome (PubMed:32103174). Binding to U3 small nucleolar RNA, recruits PRKDC and XRCC5/Ku86 to the small-subunit processome (PubMed:32103174). Recognizes the substrate consensus sequence [ST]-Q (PubMed:15574326, PubMed:11955432, PubMed:12649176, PubMed:14734805). Phosphorylates 'Ser-139' of histone variant H2AX, thereby regulating DNA damage response mechanism (PubMed:14627815, PubMed:16046194). Phosphorylates DCLRE1C, c-Abl/ABL1, histone H1, HSPCA, c-jun/JUN, p53/TP53, PARP1, POU2F1, DHX9, FH, SRF, NHEJ1/XLF, XRCC1, XRCC4, XRCC5, XRCC6, WRN, MYC and RFA2 (PubMed:2507541, PubMed:2247066, PubMed:1597196, PubMed:8407951, PubMed:8464713, PubMed:9362500, PubMed:9139719, PubMed:10026262, PubMed:10467406, PubMed:12509254, PubMed:11889123, PubMed:14612514, PubMed:14599745, PubMed:15177042, PubMed:18644470, PubMed:26666690, PubMed:30247612, PubMed:14704337

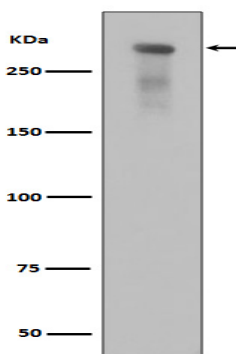
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target="_blank">14704337, PubMed:16397295, PubMed:26237645, PubMed:28712728). Can phosphorylate C1D not only in the presence of linear DNA but also in the presence of supercoiled DNA (PubMed:9679063). Ability to phosphorylate p53/TP53 in the presence of supercoiled DNA is dependent on C1D (PubMed:9363941). Contributes to the determination of the circadian period length by antagonizing phosphorylation of CRY1 'Ser- 588' and increasing CRY1 protein stability, most likely through an indirect mechanism (By similarity). Plays a role in the regulation of DNA virus-mediated innate immune response by assembling into the HDP- RNP complex, a complex that serves as a platform for IRF3 phosphorylation and subsequent innate immune response activation through the cGAS-STING pathway (PubMed:28712728).

Research Area

Image Data



Western blot analysis of Phospho-DNA PKcs (Ser2056) expression in alkaline treated Jurkat cell lysate.

Note

For research use only.