

## DNA-PKCS (Phospho Ser2056) Rabbit pAb

CatalogNo: YP0987

### Key Features

#### Host Species

- Rabbit

#### Reactivity

- Human,Mouse

#### Applications

- WB,IHC,IF,ELISA

#### MW

- 469kD (Calculated)

#### Isotype

- IgG

### Storage

**Storage\*** -15°C to -25°C/1 year (Do not lower than -25°C)

**Formulation** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

### Recommended Dilution Ratios

**IHC 1:100-1:300**

**ELISA 1:20000**

**WB 1:500-2000**

**IF 1:50-200**

### Basic Information

**Clonality** Polyclonal

### Immunogen Information

**Immunogen** The antiserum was produced against synthesized peptide derived from human DNA-PK around the phosphorylation site of Ser2056. AA range:2022-2071

**Specificity**

Phospho-DNA-PKCS (S2056) Polyclonal Antibody detects endogenous levels of DNA-PKCS protein only when phosphorylated at S2056. The name of modified sites may be influenced by many factors, such as species (the modified site was not originally found in human samples) and the change of protein sequence (the previous protein sequence is incomplete, and the protein sequence may be prolonged with the development of protein sequencing technology). When naming, we will use the "numbers" in historical reference to keep the sites consistent with the reports. The antibody binds to the following modification sequence (lowercase letters are modification sites): YSsQD

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**| Target Information****Gene name** PRKDC**Protein Name** DNA-dependent protein kinase catalytic subunit

Organism	Gene ID	UniProt ID
Human	<a href="#">5591</a> ;	<a href="#">P78527</a> ;
Mouse	<a href="#">19090</a> ;	<a href="#">P97313</a> ;

**Cellular Localization** Nucleus . Nucleus , nucleolus .**Tissue specificity** Brain , Cervix carcinoma , Epithelium , Fetal lung , Placen

## Function

Catalytic activity:ATP + a protein = ADP + a phosphoprotein. ,enzyme regulation:Inhibited by wortmannin. Activity of the enzyme seems to be attenuated by autophosphorylation. ,Function:Serine/threonine-protein kinase that acts as a molecular sensor for DNA damage. Involved in DNA nonhomologous end joining (NHEJ) required for double-strand break (DSB) repair and V (D) J recombination. Must be bound to DNA to express its catalytic properties. Promotes processing of hairpin DNA structures in V (D) J recombination by activation of the hairpin endonuclease artemis (DCLRE1C) . The assembly of the DNA-PK complex at DNA ends is also required for the NHEJ ligation step. Required to protect and align broken ends of DNA. May also act as a scaffold protein to aid the localization of DNA repair proteins to the site of damage. Found at the ends of chromosomes , suggesting a further role in the maintenance of telomeric stability and the prevention of chromosomal end fusion. Also involved in modulation of transcription. Recognizes the substrate consensus sequence [ST]-Q. Phosphorylates 'Ser-139' of histone variant H2AX/H2AFX , thereby regulating DNA damage response mechanism. Phosphorylates DCLRE1C , c-Abl/ABL1 , histone H1 , HSPCA , c-jun/JUN , p53/TP53 , PARP1 , POU2F1 , DHX9 , SRF , XRCC1 , XRCC1 , XRCC4 , XRCC5 , XRCC6 , WRN , c-myc/MYC and RFA2. Can phosphorylate C1D not only in the presence of linear DNA but also in the presence of supercoiled DNA. Ability to phosphorylate TP53/p53 in the presence of supercoiled DNA is dependent on C1D. ,PTM:Phosphorylated upon DNA damage , probably by ATM or ATR. Autophosphorylated on Thr-2609 , Thr-2638 and Thr-2647. Thr-2609 is a DNA damage-inducible phosphorylation site (inducible with ionizing radiation , IR) . Autophosphorylation induces a conformational change that leads to remodeling of the DNA-PK complex , requisite for efficient end processing and DNA repair. ,similarity:Belongs to the PI3/PI4-kinase family. ,similarity:Contains 1 FAT domain. ,similarity:Contains 1 FATC domain. ,similarity:Contains 1 PI3K/PI4K domain. ,similarity:Contains 2 HEAT repeats. ,similarity:Contains 3 TPR repeats. ,subunit:DNA-PK is a heterotrimer of PRKDC and the Ku p70-p86 (XRCC6-XRCC5) dimer. Formation of this complex may be promoted by interaction with ILF3. Associates with the DNA-bound Ku heterodimer , but it can also bind to and be activated by free DNA. Interacts with DNA-PKcs-interacting protein (KIP) with the region upstream the kinase domain. PRKDC alone also interacts with and phosphorylates DCLRE1C , thereby activating the latent endonuclease activity of this protein. Interacts with C1D. ,

## Validation Data

## Contact information

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