

Product datasheet

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ARG62471 anti-DNA PKcs antibody [18-2]

Package: 100 μl Store at: -20°C

Summary

Product Description Mouse Monoclonal antibody [18-2] recognizes DNA PKcs

Tested Reactivity Hu, Ms, Rat

Tested Application FACS, ICC/IF, IHC-Fr, IHC-P, IP, Inhib, WB

Host Mouse

Clonality Monoclonal

Clone 18-2

Isotype IgG1

Target Name DNA PKcs
Species Human

Immunogen Human DNA-dependent Protein Kinase purified from HeLa cells

Epitope aa 1-2713

Conjugation Un-conjugated

Alternate Names p350; DNAPK; HYRC1; DNA-PKcs; DNA-dependent protein kinase catalytic subunit; DNPK1; IMD26;

HYRC; EC 2.7.11.1; p460; DNA-PK catalytic subunit; XRCC7

Application Instructions

Application table	Application	Dilution
	FACS	Assay-dependent
	ICC/IF	Assay-dependent
	IHC-Fr	Assay-dependent
	IHC-P	1:400
	IP	1:400
	Inhib	Assay-dependent
	WB	1:200
• •	* The dilutions indicate recommended starting dilutions and the optimal dilutions or concentrations should be determined by the scientist.	
Positive Control	LS174T cells. Tonsil.	

Properties

Form	Liquid
Purification	Purified Antibody

Buffer 1X PBS and 0.1% Sodium azide

Preservative 0.1% Sodium azide

Concentration 0.2 mg/ml

Storage instruction For continuous use, store undiluted antibody at 2-8°C for up to a week. For long-term storage, aliquot

and store at -20°C or below. Storage in frost free freezers is not recommended. Avoid repeated

freeze/thaw cycles. Suggest spin the vial prior to opening. The antibody solution should be gently mixed

before use.

Note For laboratory research only, not for drug, diagnostic or other use.

Bioinformation

Database links GenelD: 19090 Mouse

GenelD: 5591 Human

Swiss-port # P78527 Human

Swiss-port # P97313 Mouse

Gene Symbol PRKDC

Gene Full Name protein kinase, DNA-activated, catalytic polypeptide

Background This gene encodes the catalytic subunit of the DNA-dependent protein kinase (DNA-PK). It functions

with the Ku70/Ku80 heterodimer protein in DNA double strand break repair and recombination. The $\,$

protein encoded is a member of the PI3/PI4-kinase family.[provided by RefSeq, Jul 2010]

Function Serine/threonine-protein kinase that acts as a molecular sensor for DNA damage. Involved in DNA non-

homologous 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, XRCC4, XRCC5, XRCC6, WRN, 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 p53/TP53 in the presence of supercoiled DNA is dependent on C1D. 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 machanism. Interacts with CRY1 and CRY2; negatively regulates

CRY1 phosphorylation. [UniProt]

Research Area Gene Regulation antibody

Calculated Mw 469 kDa

PTM Autophosphorylated on Ser-2056, Thr-2609, Thr-2638 and Thr-2647. Ser-2056 and Thr-2609 are DNA

damage-inducible phosphorylation sites (inducible with ionizing radiation, IR) dephosphorylated by PPP5C. Autophosphorylation induces a conformational change that leads to remodeling of the DNA-PK

complex, requisite for efficient end processing and DNA repair.

S-nitrosylated by GAPDH.

Polyubiquitinated by RNF144A, leading to proteasomal degradation.