

Ku70 Antibody

Rabbit mAb

Catalog # AP91239

Product Information

Application	WB, IHC, IF, FC, ICC, IP, IHF
Primary Accession	P12956
Reactivity	Human
Clonality	Monoclonal
Other Names	CTC75; CTCBF; DNA repair protein XRCC6; G22P1; Ku 70; Ku autoantigen 70kDa; Ku autoantigen, 70kDa; Ku p70; Ku70; Kup70; TLAA; XRCC6;
Isotype	Rabbit IgG
Host	Rabbit
Calculated MW	69843

Additional Information

Dilution	WB 1:500~1:2000 IHC 1:50~1:200 ICC/IF 1:50~1:200 IP 1:50 FC 1:50
Purification	Affinity-chromatography
Immunogen	A synthesized peptide derived from human Ku70
Description	It works in the 3'-5' direction. Binding to DNA may be mediated by XRCC6. Involved in DNA non-homologous end joining (NHEJ) required for double-strand break repair and V(D)J recombination. The XRCC5/6 dimer acts as regulatory subunit of the DNA-dependent protein kinase complex DNA-PK by increasing the affinity of the catalytic subunit PRKDC to DNA by 100-fold. The XRCC5/6 dimer is probably involved in stabilizing broken DNA ends and bringing them together.
Storage Condition and Buffer	Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid freeze / thaw cycle.

Protein Information

Name	XRCC6
Synonyms	G22P1
Function	Single-stranded DNA-dependent ATP-dependent helicase that plays a key role in DNA non-homologous end joining (NHEJ) by recruiting DNA-PK to DNA (PubMed: 11493912 , PubMed: 12145306 , PubMed: 20493174 , PubMed: 2466842 , PubMed: 7957065 , PubMed: 8621488 , PubMed: 9742108). Required for double-strand break repair and V(D)J recombination (PubMed: 11493912 , PubMed: 12145306 , PubMed: 20493174 , PubMed: 2466842 , PubMed: 7957065 , PubMed: 8621488 , PubMed: 9742108). Also has a role in chromosome translocation (PubMed: 11493912 , PubMed: 12145306 , PubMed: 20493174 , PubMed: 2466842 , PubMed: 7957065 ,

PubMed:[8621488](#), PubMed:[9742108](#)). Has a role in chromosome translocation (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). The DNA helicase II complex binds preferentially to fork-like ends of double-stranded DNA in a cell cycle-dependent manner (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). It works in the 3'-5' direction (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). During NHEJ, the XRCC5-XRRC6 dimer performs the recognition step: it recognizes and binds to the broken ends of the DNA and protects them from further resection (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). Binding to DNA may be mediated by XRCC6 (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). The XRCC5-XRRC6 dimer acts as a regulatory subunit of the DNA-dependent protein kinase complex DNA-PK by increasing the affinity of the catalytic subunit PRKDC to DNA by 100-fold (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). The XRCC5-XRRC6 dimer is probably involved in stabilizing broken DNA ends and bringing them together (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). The assembly of the DNA-PK complex to DNA ends is required for the NHEJ ligation step (PubMed:[11493912](#), PubMed:[12145306](#), PubMed:[20493174](#), PubMed:[2466842](#), PubMed:[7957065](#), PubMed:[8621488](#), PubMed:[9742108](#)). Probably also acts as a 5'-deoxyribose-5-phosphate lyase (5'-dRP lyase), by catalyzing the beta-elimination of the 5' deoxyribose-5-phosphate at an abasic site near double-strand breaks (PubMed:[20383123](#)). 5'-dRP lyase activity allows to 'clean' the termini of abasic sites, a class of nucleotide damage commonly associated with strand breaks, before such broken ends can be joined (PubMed:[20383123](#)). The XRCC5-XRRC6 dimer together with APEX1 acts as a negative regulator of transcription (PubMed:[8621488](#)). In association with NAA15, the XRCC5-XRRC6 dimer binds to the osteocalcin promoter and activates osteocalcin expression (PubMed:[12145306](#)). 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](#)). Negatively regulates apoptosis by interacting with BAX and sequestering it from the mitochondria (PubMed:[15023334](#)). Might have deubiquitination activity, acting on BAX (PubMed:[18362350](#)).

Cellular Location

Nucleus. Chromosome. Cytoplasm. Note=When trimethylated, localizes in the cytoplasm.

Images

Western blot analysis of Ku70 expression in HeLa cell lysate.

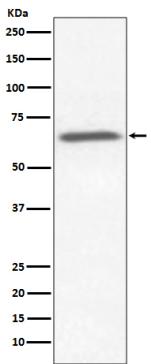


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Immunohistochemical analysis of paraffin-embedded human tonsil, using Ku70 Antibody.

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