

**Anti-Ku70 Antibody**  
**Catalog # ABV11941****Specification****Anti-Ku70 Antibody - Product Information**

Application	WB
Primary Accession	<a href="#">P12956</a>
Reactivity	Human, Mouse, Rat
Host	Rabbit
Isotype	Rabbit IgG
Calculated MW	69843

**Anti-Ku70 Antibody - Additional Information****Gene ID 2547**

Positive Control                            WB; Hela, NIH3T3, PC12 cell lysates  
Application & Usage                WB; 1:500 - 1:2000

**Other Names**

G22P1; X-ray repair cross-complementing protein 6; 5'-deoxyribose-5-phosphate lyase Ku70; 5'-dRP lyase Ku70; 70 kDa subunit of Ku antigen; ATP-dependent DNA helicase 2 subunit 1; ATP-dependent DNA helicase II 70 kDa subunit; CTC box-binding factor 75 kDa subunit; CTC75; CTCBF; DNA repair protein XRCC6; Lupus Ku autoantigen protein p70; Ku70; Thyroid-lupus autoantigen; TLAA; X-ray repair complementing defective repair in Chinese hamster cells 6

**Target/Specificity**  
XRCC6**Antibody Form**  
Liquid**Appearance**  
Colorless liquid**Handling**  
The antibody solution should be gently mixed before use**Reconstitution & Storage**  
-20°C**Background Descriptions****Precautions**

Anti-Ku70 Antibody is for research use only and not for use in diagnostic or therapeutic procedures.

**Anti-Ku70 Antibody - 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:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">9742108</a>). Required for double-strand break repair and V(D)J recombination (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">9742108</a>). Also has a role in chromosome translocation (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">9742108</a>). Has a role in chromosome translocation (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">9742108</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>). The DNA helicase II complex binds preferentially to fork-like ends of double-stranded DNA in a cell cycle-dependent manner (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">9742108</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>). It works in the 3'-5' direction (PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">9742108</a>, PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>). During NHEJ, the XRCC5-XRCC6 dimer performs the recognition step: it recognizes and binds to the broken ends of the DNA and protects them from further resection (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">8621488</a>, PubMed:<a

href="http://www.uniprot.org/citations/20493174" target="\_blank">>20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">>2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">>9742108</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</a>). Binding to DNA may be mediated by XRCC6 (PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">>20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">>2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">>9742108</a>, PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">>7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">>8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</a>). 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:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">>7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">>8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">>20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">>2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">>9742108</a>). The XRCC5-XRRC6 dimer is probably involved in stabilizing broken DNA ends and bringing them together (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">>7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">>8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">>20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">>2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">>9742108</a>). The assembly of the DNA-PK complex to DNA ends is required for the NHEJ ligation step (PubMed:<a href="http://www.uniprot.org/citations/7957065" target="\_blank">>7957065</a>, PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">>8621488</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/20493174" target="\_blank">>20493174</a>, PubMed:<a href="http://www.uniprot.org/citations/2466842" target="\_blank">>2466842</a>, PubMed:<a href="http://www.uniprot.org/citations/9742108" target="\_blank">>9742108</a>). 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:<a href="http://www.uniprot.org/citations/20383123" target="\_blank">>20383123</a>). 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:<a href="http://www.uniprot.org/citations/20383123" target="\_blank">>20383123</a>). The XRCC5-XRRC6 dimer together with APEX1 acts as a negative regulator of transcription (PubMed:<a href="http://www.uniprot.org/citations/8621488" target="\_blank">>8621488</a>). In association with NAA15, the XRCC5-XRRC6 dimer binds to the osteocalcin promoter and activates osteocalcin expression (PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</a>). 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:<a href="http://www.uniprot.org/citations/28712728" target="\_blank">>28712728</a>).

## Cellular Location

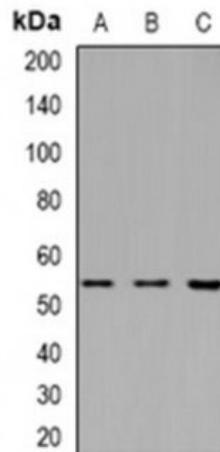
Nucleus. Chromosome

## Anti-Ku70 Antibody - Protocols

Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

## Anti-Ku70 Antibody - Images



WB analysis of Ku70 (AcK338) expression in HeLa (A); NIH3T3 (B); PC12 (C) whole cell lysates