

**Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729]**  
**Purified Mouse Monoclonal Antibody**  
**Catalog # AH10371**

**Specification**

**Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729] - Product Information**

Application	IF, FC
Primary Accession	<a href="#">P12956</a>
Other Accession	<a href="#">P13010</a>
Reactivity	Human
Host	Mouse
Clonality	Monoclonal
Isotype	IgG1, kappa
Calculated MW	70kDa & 80kDa KDa

**Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729] - Additional Information**

**Gene ID 2547**

**Other Names**

Ku (p70): 70kDa subunit of Ku antigen; ATP dependent DNA helicase 2 subunit 1; ATP-dependent DNA helicase II 70kDa subunit; CTC box-binding factor 75kDa subunit; CTC75; CTCBF; DNA repair protein XRCC6; G22P1; Ku autoantigen, 70kDa; Ku70; Kup70; Lupus Ku autoantigen protein p70; ML8; Thyroid autoantigen 70kD (Ku antigen); Thyroid-lupus autoantigen (TLAA); X-ray repair cross-complementing protein 6 (XRCC6) Ku (p80): 86kDa subunit of Ku antigen; ATP dependent DNA helicase 2 subunit 2; ATP dependent DNA helicase II 86kD subunit; ATP-dependent DNA helicase II 80kDa subunit; CTC box-binding factor 85kDa subunit; CTC85; CTCBF; DNA repair protein XRCC5; KARP1; Ku autoantigen 80kDa; Ku80; Ku86 autoantigen related protein 1; KUB2; Lupus Ku autoantigen protein p86; Nuclear factor IV (NFIV); Thyroid-lupus autoantigen (TLAA); X-ray repair cross-complementing protein 5 (XRCC5)

**Target/Specificity**

Nuclear extract of human HL-60 cells

**Application Note**

<span class = "dilution\_IF">IF~~~1:50~200</span><br /><span class = "dilution\_FC">FC~~~1:10~50</span>

**Format**

0.5ml at 100ug/ml with BSA and azide

**Storage**

Store at 2 to 8°C. Antibody is stable for 24 months.

**Precautions**

Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729] is for research use only and not for use in diagnostic or therapeutic procedures.

**Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729] - 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/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</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/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/9742108" target="\_blank">9742108</a>). Required for double-strand break repair and V(D)J recombination (PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</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/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/9742108" target="\_blank">9742108</a>). Also has a role in chromosome translocation (PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</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/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/9742108" target="\_blank">9742108</a>). Has a role in chromosome translocation (PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</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/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/9742108" target="\_blank">9742108</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/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</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/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/9742108" target="\_blank">9742108</a>). It works in the 3'-5' direction (PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">12145306</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/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/9742108" target="\_blank">9742108</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/11493912" target="\_blank">11493912</a>, PubMed:<a

href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</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/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/9742108" target="\_blank">>9742108</a>). Binding to DNA may be mediated by XRCC6 (PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</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/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/9742108" target="\_blank">>9742108</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/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</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/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/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/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</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/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/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/11493912" target="\_blank">>11493912</a>, PubMed:<a href="http://www.uniprot.org/citations/12145306" target="\_blank">>12145306</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/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/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>). Negatively regulates apoptosis by interacting with BAX and sequestering it from the mitochondria (PubMed:<a href="http://www.uniprot.org/citations/15023334" target="\_blank">>15023334</a>). Might have deubiquitination activity, acting on BAX (PubMed:<a

[18362350](http://www.uniprot.org/citations/18362350)).

#### Cellular Location

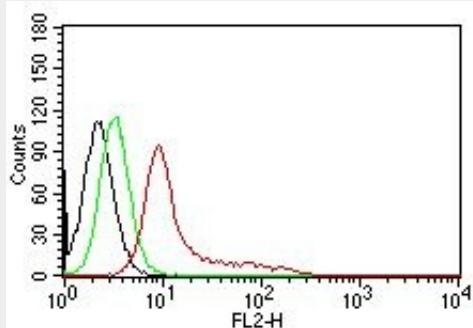
Nucleus. Chromosome. Cytoplasm. Note=When trimethylated, localizes in the cytoplasm.

#### Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729] - 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](#)

#### Ku (p70/p80)Mouse Monoclonal Antibody [Clone KU729] - Images



Flow Cytometric analysis of human Ku (p70/p80) on K562 Cells. Black: Cells alone; Green: Isotype Control; Red: PE-labeled Ku MAb (KU729).