

**XRCC5 Antibody (Center K439) Blocking peptide**  
**Synthetic peptide**  
**Catalog # BP11960c**

**Specification**

**XRCC5 Antibody (Center K439) Blocking peptide - Product Information**

Primary Accession [P13010](#)

**XRCC5 Antibody (Center K439) Blocking peptide - Additional Information**

**Gene ID** 7520

**Other Names**

X-ray repair cross-complementing protein 5, 364-, 86 kDa subunit of Ku antigen, ATP-dependent DNA helicase 2 subunit 2, ATP-dependent DNA helicase II 80 kDa subunit, CTC box-binding factor 85 kDa subunit, CTC85, CTCBF, DNA repair protein XRCC5, Ku80, Ku86, Lupus Ku autoantigen protein p86, Nuclear factor IV, Thyroid-lupus autoantigen, TLAA, X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining), XRCC5, G22P2

**Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

**XRCC5 Antibody (Center K439) Blocking peptide - Protein Information**

**Name** XRCC5

**Synonyms** G22P2

**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>). 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>). 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>). 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/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/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-XRRC6 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/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/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/20383123" target="\_blank">>20383123</a>, PubMed:<a href="http://www.uniprot.org/citations/11493912" target="\_blank">>11493912</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/20383123" target="\_blank">>20383123</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/20383123" target="\_blank">>20383123</a>). The XRCC5-XRRC6 dimer 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>). XRCC5 probably acts as the catalytic subunit of 5'- dRP activity, and 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>). As part of the DNA-PK complex, involved in the early steps of ribosome assembly by promoting the processing of precursor rRNA into mature 18S rRNA in the small-subunit processome (PubMed:<a href="http://www.uniprot.org/citations/32103174" target="\_blank">>32103174</a>). Binding to U3 small nucleolar RNA, recruits PRKDC and XRCC5/Ku86 to the small-subunit processome

(PubMed:<a href="http://www.uniprot.org/citations/32103174" target="\_blank">32103174</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. Nucleus, nucleolus. Chromosome

#### **XRCC5 Antibody (Center K439) Blocking peptide - Protocols**

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

- [Blocking Peptides](#)

#### **XRCC5 Antibody (Center K439) Blocking peptide - Images**

#### **XRCC5 Antibody (Center K439) Blocking peptide - Background**

The protein encoded by this gene is the 80-kilodalton subunit of the Ku heterodimer protein which is also known as ATP-dependant DNA helicase II or DNA repair protein XRCC5. Ku is the DNA-binding component of the DNA-dependent protein kinase, and it functions together with the DNA ligase IV-XRCC4 complex in the repair of DNA double-strand break by non-homologous end joining and the completion of V(D)J recombination events. This gene functionally complements Chinese hamster xrs-6, a mutant defective in DNA double-strand break repair and in ability to undergo V(D)J recombination. A rare microsatellite polymorphism in this gene is associated with cancer in patients of varying radiosensitivity.

#### **XRCC5 Antibody (Center K439) Blocking peptide - References**

Gomes, B.C., et al. Oncol. Rep. 24(4):1079-1085(2010) Liu, Y., et al. Carcinogenesis 31(10):1762-1769(2010) Ho-Pun-Cheung, A., et al. Pharmacogenomics J. (2010) In press : Briggs, F.B., et al. Am. J. Epidemiol. 172(2):217-224(2010) Monsees, G.M., et al. Breast Cancer Res. Treat. (2010) In press :