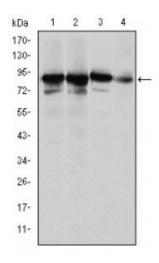


Anti-Ku-80 antibody





Description Mouse monoclonal to Ku-80.

Model STJ98211

Host Mouse

Reactivity Human, Mouse

Applications ELISA, FC, IF, IHC, WB

Immunogen Purified recombinant fragment of human Ku-80 expressed in E. Coli.

Gene ID 7520

Gene Symbol XRCC5

Dilution range WB 1:500-1:2000IHC 1:200-1:1000IF 1:200-1:1000FC 1:200-1:400ELISA

1:10000

Specificity Ku-80 Monoclonal Antibody detects endogenous levels of Ku-80 protein.

Purification Affinity purification

Clone ID 5C5

Note For Research Use Only (RUO).

Protein Name X-ray repair cross-complementing protein 5 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

rep

Clonality Monoclonal

Conjugation Unconjugated

Isotype IgG1

Formulation Ascitic fluid containing 0.03% sodium azide.

Storage Instruction Store at -20°C, and avoid repeat freeze-thaw cycles.

Database Links HGNC:12833OMIM:194364

Alternative Names X-ray repair cross-complementing protein 5 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

rep

Function Single-stranded DNA-dependent ATP-dependent helicase. Has a role in

chromosome translocation. The DNA helicase II complex binds preferentially to fork-like ends of double-stranded DNA in a cell cycle-dependent manner. 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 . The assembly of the DNA-PK complex to DNA ends is required for the NHEJ ligation step. In association with NAA15, the XRCC5/6 dimer binds to the osteocalcin promoter and activates osteocalcin expression . The XRCC5/6 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-

deoxyribose-5-phosphate lyase (5'-dRP lyase), by catalyzing the betaelimination of the 5' deoxyribose-5-phosphate at an abasic site near doublestrand breaks. 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. The XRCC5/6 dimer together with APEX1 acts as a negative

regulator of transcription.

Sequence and Domain Family The EEXXXDDL motif is required for the interaction with catalytic subunit

PRKDC and its recruitment to sites of DNA damage.

Cellular Localization Nucleus Nucleus, nucleolus Chromosome

Post-translational Phosphorylated on serine residues. Phosphorylation by PRKDC may enhance **Modifications** helicase activity. Sumovlated. Ubiquitinated by RNF8 via 'Lys-48'-linked

helicase activity. Sumoylated. Ubiquitinated by RNF8 via 'Lys-48'-linked ubiquitination following DNA damage, leading to its degradation and removal from DNA damage sites . Ubiquitinated by RNF138, leading to remove the

Ku complex from DNA breaks.