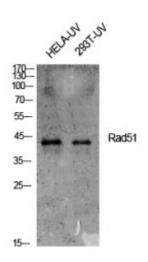


## Anti-Rad51 antibody



**Description** Rabbit polyclonal to Rad51.

Model STJ95330

**Host** Rabbit

**Reactivity** Human, Mouse

**Applications** ELISA, IHC, WB

**Immunogen** Synthesized peptide derived from human Rad51 around the non-

phosphorylation site of Y315.

**Immunogen Region** 250-330 aa

**Gene ID** <u>5888</u>

Gene Symbol RAD51

**Dilution range** WB 1:500-1:2000IHC 1:100-1:300ELISA 1:20000

**Specificity** Rad51 Polyclonal Antibody detects endogenous levels of Rad51 protein.

**Tissue Specificity** Highly expressed in testis and thymus, followed by small intestine, placenta,

colon, pancreas and ovary. Weakly expressed in breast.

**Purification** The antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

**Note** For Research Use Only (RUO).

Protein Name DNA repair protein RAD51 homolog 1 HsRAD51 hRAD51 RAD51 homolog

A

Molecular Weight 42 kDa

**Clonality** Polyclonal

**Conjugation** Unconjugated

**Isotype** IgG

**Formulation** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

**Concentration** 1 mg/ml

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

Database Links HGNC:98170MIM:114480

Alternative Names DNA repair protein RAD51 homolog 1 HsRAD51 hRAD51 RAD51 homolog

A

**Function** Plays an important role in homologous strand exchange, a key step in DNA

repair through homologous recombination (HR). Binds to single and double-stranded DNA and exhibits DNA-dependent ATPase activity. Catalyzes the recognition of homology and strand exchange between homologous DNA partners to form a joint molecule between a processed DNA break and the repair template. Binds to single-stranded DNA in an ATP-dependent manner to form nucleoprotein filaments which are essential for the homology search and strand exchange . Part of a PALB2-scaffolded HR complex containing BRCA2 and RAD51C and which is thought to play a role in DNA repair by HR. Plays a role in regulating mitochondrial DNA copy number under conditions of oxidative stress in the presence of RAD51C and XRCC3. Also

involved in interstrand cross-link repair.

**Sequence and Domain Family** The nuclear localization may reside in the C-terminus (between 259 and 339)

AA).

Cellular Localization Nucleus Cytoplasm Cytoplasm, perinuclear region. Mitochondrion matrix

Cytoplasm, cytoskeleton, microtubule organizing center, centrosome. Colocalizes with RAD51AP1 and RPA2 to multiple nuclear foci upon induction of DNA damage. DNA damage induces an increase in nuclear levels. Together with FIGNL1, redistributed in discrete nuclear DNA damage-induced foci after ionizing radiation (IR) or camptothecin (CPT) treatment. Accumulated at sites of DNA damage in a SPIDR-dependent manner.

**Post-translational**Ubiquitinated by the SCF(FBXO18) E3 ubiquitin ligase complex, regulating **Modifications**RAD51 subcellular location and preventing its association with DNA.

RAD51 subcellular location and preventing its association with DNA. Phosphorylated. Phosphorylation of Thr-309 by CHEK1 may enhance association with chromatin at sites of DNA damage and promote DNA repair by homologous recombination. Phosphorylation by ABL1 inhibits function.