

## **Rabbit Anti-SPIDR antibody**

SL17664R

Product Name:	SPIDR
Chinese Name:	
Alias:	DNA repair-scaffolding protein; KIAA0146; Scaffolding protein involved in DNA
	repair; SPIDR; SPIDR_HUMAN.
Organism Species:	Rabbit
Clonality:	Polyclonal
React Species:	Human,Mouse,Rat,Cow,Horse,
Applications:	WB=1:500-2000ELISA=1:500-1000IHC-P=1:400-800IHC-F=1:400-800ICC=1:100-
	500IF=1:100-500 (Paraffin sections need antigen repair)
	not yet tested in other applications.
	optimal dilutions/concentrations should be determined by the end user.
Molecular weight:	100kDa
Cellular localization:	The nucleus
Form:	Lyophilized or Liquid
<b>Concentration:</b>	1mg/ml
immunogen:	KLH conjugated synthetic peptide derived from human SPIDR:131-230/915
Lsotype:	IgG
Purification:	affinity purified by Protein A
Storage Buffer:	0.01M TBS(pH7.4) with 1% BSA, 0.03% Proclin300 and 50% Glycerol.
Storage:	Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. The lyophilized
	antibody is stable at room temperature for at least one month and for greater than a year
	when kept at -20°C. When reconstituted in sterile pH 7.4 0.01M PBS or diluent of
	antibody the antibody is stable for at least two weeks at 2-4 °C.
PubMed:	PubMed
Product Detail:	Plays a role in DNA double-strand break (DBS) repair via homologous recombination
	(HR). Serves as a scaffolding protein that helps to promote the recruitment of DNA-
	processing enzymes like the helicase BLM and recombinase RAD51 to site of DNA
	damage, and hence contributes to maintain genomic integrity.
	Function:

Plays a role in DNA double-strand break (DBS) repair via homologous recombination (HR). Serves as a scaffolding protein that helps to promote the recruitment of DNA-processing enzymes like the helicase BLM and recombinase RAD51 to site of DNA damage, and hence contributes to maintain genomic integrity.

## Subunit:

Found in a complex, at least composed of BLM, RAD51 and SPIDR; the complex formation is mediated by SPIDR. Interacts (via C-terminal region) with BLM; the interaction is direct. Interacts with RAD51; the interaction is direct. Interacts (via the C-terminal region) with FIGNL1 (via N-terminal one-half region);the interaction is direct.

## Subcellular Location:

Nucleus. Together with BLM, is redistributed in discrete nuclear DNA damage-induced foci following hydroxyurea (HU) or camptothecin (CPT) treatment.

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**SWISS:** Q14159

Gene ID: 23514

Database links:

Entrez Gene: 23514 Human

<u>Omim: 615384</u> Human

SwissProt: Q14159 Human

Unigene: 381058 Human

Important Note:

This product as supplied is intended for research use only, not for use in human, therapeutic or diagnostic applications.

