

Product datasheet for SC206065

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Rad51L1 (RAD51B) (NM_133510) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Rad51L1 (RAD51B) (NM_133510) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: RAD51B

Synonyms: R51H2; RAD51L1; REC2

ACCN: NM_133510

Insert Size: 459 bp

Insert Sequence: >SC206065 3'UTR clone of NM_133510

The sequence shown below is from the reference sequence of NM_133510. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TAATTACAGGAGGAACATTTCCGAATAAAGTATTGTCTACCAGA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

RefSeq: <u>NM 133510.4</u>





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Summary: The protein encoded by this gene is a member of the RAD51 protein family. RAD51 family

members are evolutionarily conserved proteins essential for DNA repair by homologous recombination. This protein has been shown to form a stable heterodimer with the family member RAD51C, which further interacts with the other family members, such as RAD51, XRCC2, and XRCC3. Overexpression of this gene was found to cause cell cycle G1 delay and

cell apoptosis, which suggested a role of this protein in sensing DNA damage.

Rearrangements between this locus and high mobility group AT-hook 2 (HMGA2, GeneID 8091) have been observed in uterine leiomyomata. [provided by RefSeq, Mar 2016]

Locus ID: 5890

MW: 17.2