

## Product datasheet for **MG221567**

### Rad54l (NM\_001122959) Mouse Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Rad54l (NM_001122959) Mouse Tagged ORF Clone
Tag:	TurboGFP
Symbol:	Rad54l
Synonyms:	RAD54
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)



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**ORF Nucleotide Sequence:**

>MG221567 representing NM\_001122959  
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCC**CGGATCGCC**

ATGAGGAGGAGCTTAGCTCCCAGCCAGTTGGCCAGGAGGAAACCAGAAGACAGATCATCAGATGATGAAG  
 ACTGGCAGCCTGGGACAGTAACCTCTAAGAAACGCAAGTCCAGCAGTGAGACCCAGGTCCAGGAGTGTTT  
 CCTGTCTCCTTTTCGGAAGCCCTTGACTCAGCTACTCAACCGGCCACCTTGTCTGGATAGCAGTCAACAT  
 GAAGCATTTATTGCAAGTATTTTGTCAAAGCCTTTCAAGGTCCCATCCCAAATTATCAAGGTCCCTGG  
 GCTCTCGTGCATTGGGCCTGAAAAGGGTTGGAGTTCGTCGTGCCCTTCATGACCCCTGGAAGAAGGTGC  
 CTTGGTTCTCTATGAGCCTCCCCACTCAGCGCCCATGACCAACTGAAGCTTGACAAGGAAAACTCCCT  
 GTTCATGTGGTTGTTGATCCTATTCTCAGTAAGGTATTGCGGCCTCATCAGAGAGAGGGAGTGAAGTTCC  
 TATGGGAGTGTGCACCAGTCGTCGAATTCCTGGAAGCCATGGCTGCATCATGGCTGATGAGATGGCCT  
 GGGGAAGACACTGCAGTGCATCACATTGATGTGGACACTTTTACGCCAGAGCCCAGAATGCAAGCCAGAA  
 ATCGAGAAGGCAGTGGTGGTGTACCTTAGCCTGGTGAAGAAGTGGTACAATGAGGTTGAGAAATGGC  
 TTGGAGGGAGGATTCAACCTCTGGCCATCGATGGAGGCTCGAAGGACGAGATAGACCCGAAAACTGGAAGG  
 ATTCATGAACCAGCGTGGAGCTAGAGTGCCCTTCCCCATCCTCATCATTTCTATGAGACTTTCCGCCTG  
 CATGTTGGAGTCTTAAAAAAGGAAATGTTGGACTGGTCAATGTGACGAGGGCCACAGGCTAAAGAACT  
 CTGAGAATCAGACTTACCAGGCTCTGGACAGCTTGAATACCAGCCGGCGGGTGTAACTCTCCGGGACCCC  
 CATCCAAAATGATTTGCTGGAGTATTTAGCTTGGTGCCTTTGTAATTTCTGGCATTGTTGGAACTGCC  
 CATGAATTCAGAAGCATTGTTGAGTTGCCAATTTGAAGAGTCGAGATGCAGTGCAGTGCAGTGCAGTGCAG  
 GGCAGCGTGGGAGGAGCGTCTGCGGGAGCTCATCGGTATTGTGAACAGGTGCCTGATACGGAGAACATC  
 TGATATCCTCTAAGTATCTGCCCGTGAAGATTGAGCAGGTGGTTTGTGTAGGCTGACACCCCTTCAA  
 ACTGAGCTATACAAGAGATTTCTGAGACAGGCTAAGCCTGAAGAAGAATTGCGTGAGGGCAAGATGAGTG  
 TGTCTTCCCTGTCTTCTATCACCTCTCTAAAGAAGCTGTGTAATCATCCAGCTCTAATCTATGACAAGTG  
 TGTGGCAGAGGAGGATGGCTTTGAGGGCACTTTGGGTATCTTCCACCTGGTTATAACTCTAAAGCTGTA  
 GAGCCACAGTTGTCAGGTAAGATGCTGGTCCTTGATTACATTCTGGCCGTGACTCGAAGCCGTAGCAGTG  
 ACAAAGTCGTGCTGGTGTCTAATTATACTCAGACATTGGATCTCTTTGAAAAGCTGTGCCGGTTTGAAG  
 GTACTTGTATGTTCCCTGGATGGCAGCATGTCCATTAAGAAGCGAGCCAAGGTTGTGGAGCGCTTCAAT  
 AGCCCATCGAGCCCTGATTTTGTTCATGCTGAGCAGCAAAGCTGGGGCTGTGGTCTTAACTCATTG  
 GTGCTAACCGACTGGTCAATGTTGATCCTGACTGGAATCCAGCCAATGATGAACAAGCTATGGCCCGAGT  
 CTGGCGTGATGGTCAAAAGAAGATCTGCTATATCTACCGACTGCTATCTGCAGGAACAATTGAGGAGAAG  
 ATCTTTACGCGCAGAGCCACAAGAAGGCTCTGAGCAGCTGTGTGGTGGACGAGGAGCAGGATGTGGAGC  
 GCCACTTTTCTTGGTGGAGCTTAAAGAGCTGTTACTCTGGATGAAGCAAGCCTCAGTGACACACATGA  
 CAGATTGCATTGCCCGCCTTGTGTAACAACCGTCAGGTCTGGCCACCCCTGATGGTTCTGACTGCACT  
 TCAGACCTGGCTCAGTGAACCACAGCACAGATAAACGAGGGCTCCAGGATGAGGTACTCCAGGCTGCCT  
 GGGATGCTTCTATACAGCCATCACCTTCGCTTCCACCAGCGTTCATGAGGAGCAGCGCGGTCTTCA  
 C

**ACGCGT**ACGCGGCCGCTCGAG – GFP Tag – GTTTAA

Protein Sequence: >MG221567 representing NM\_001122959  
 Red=Cloning site Green=Tags(s)

MRRSLAPSQLARRKPEDRSSDDEDWQPGTVTPKKRKSSSETQVQECFLSPFRKPLTQLLNRPCLDSSQH  
 EAFIRSILSKPFKVPINYPQGLSRALGLKRVGVRRALHDPLEEGALVLYEPPPLSAHDQLKLDKEKLP  
 VHVVDPIILSKVLRPHQREGVKFLWECVTSRRIPGSHGCIMADEMGLGKTLQCITLMWTLRLRQSPECKPE  
 IEKAVVSPSSLVKNWYNEVEKWLGGRIQPLAIDGGSKDEIDRKLEGFMNQRGARVPSILIIISYETFRL  
 HVGVLKKGNVGLVICDEGHRLKNSNQTYQALDSLNTSRRVLSGTPIQNDLLEYFSLVHFVNSGILGTA  
 HEFKKHFEPLILKSRDAAASEADRQRGEERLRELIGIVNRCLIRRTSDILSKYLPVKIEQVVCRLTPLQ  
 TELYKRFRLQAKPEEELREGKMSVSSLSITSLLKLCNHPALIYDKVAEEDGFEGTLGIFPPGYNSKAV  
 EPQLSGKMLVLDYILAVTRSRSSDKVVLVSNTQTLDLFEKLCRVRRYL YVRLDGTMSIKKRAKVVERFN  
 SPSSPDFVFM LSSKAGGCGNLIGANRLVMFDPDWN PANDEQAMARVWRDGGKKICYIYRLLSAGTIEEK  
 IFQRQSHKKALSSCVDEEQDVERHFSLGELKELFTLDEASLSDTHDRLHCRRCVNNRQVWPPPDGSDCT  
 SDLAQWNHSTDKRGLQDEVLQAAWDASSTAITFVFHQRSHEEQRGLH

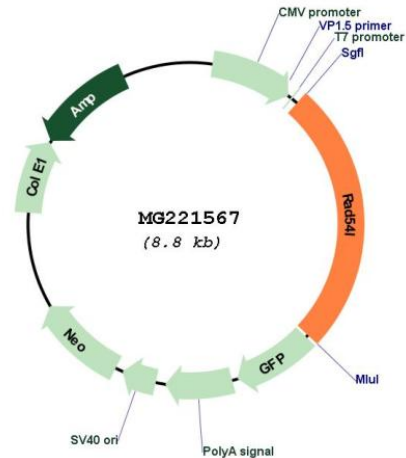
TRTRPLE - GFP Tag - V

Restriction Sites: SgfI-MluI

Cloning Scheme:

Cloning sites used for ORF Shutting:



**Plasmid Map:**


**ACCN:** NM\_001122959

**ORF Size:** 2241 bp

**OTI Disclaimer:** The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

**OTI Annotation:** This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

**Reconstitution Method:**

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

**RefSeq:** [NM\\_001122959.1](#), [NP\\_001116431.1](#)

**RefSeq Size:** 2742 bp

**RefSeq ORF:** 2244 bp

**Locus ID:** 19366

UniProt ID: [P70270](#)

Cytogenetics: 4 D1

**Gene Summary:** Involved in DNA repair and mitotic recombination. Functions in the recombinational DNA repair (RAD52) pathway. Dissociates RAD51 from nucleoprotein filaments formed on dsDNA. Could be involved in the turnover of RAD51 protein-dsDNA filaments (By similarity). Deficient mice also show significantly shorter telomeres than wild-type controls, indicating that the protein activity plays an essential role in telomere length maintenance in mammals. Deficiency also resulted in an increased frequency of end-to-end chromosome fusions involving telomeres compared to the controls, suggesting a putative role in telomere capping. Non-homologous end joining (NHEJ) and homologous recombination (HR) represent the two major pathways of DNA double-strand break (DSB) repair in eukaryotic cells. LIG4 and RAD54L cooperate to support cellular proliferation, repair spontaneous DSBs, and prevent chromosome and single chromatid aberrations.[UniProtKB/Swiss-Prot Function]