

Product datasheet for RC218369

LRP1 (NM_002332) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	LRP1 (NM_002332) Human Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	LRP1
Synonyms:	A2MR; APOER; APR; CD91; IGFBP-3R; IGFBP3R; IGFBP3R1; KPA; LRP; LRP1A; TGFBR5
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>RC218369 representing NM_002332. Blue=ORF Red=Cloning site Green=Tag(s)

```
GCTCGTTTGTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC
ATGCTGACCCCGCGTTGCTCCTGCTGCTGCCCTGCTCTCAGCTCTGGTCGCGCGCGCTATCGACGCC
CCTAAGACTTGCAGCCCCAAGCAGTTTGCCTGCAGAGATCAAATAACCTGTATCTCAAAGGGCTGGCGG
TGCGACGGTGAGAGGGACTGCCAGACGGATCTGACGAGGCCCTGAGATTTGCCACAGAGTAAGGCC
CAGCGATGCCAGCCAAACGAGCATAACTGCCTGGTACTGAGCTGTGTGCCATGTCCCGCTCTGC
AATGGGGTCCAGGACTGCATGGACGGCTCAGATGAGGGGCCCACTGCCGAGAGCTCAAGGCAACTGC
TCTCGCTGGGCTGCCAGCACCATTGTGTCCACACTCGATGGGCCACCTGCTACTGCAACAGCAGC
TTTCAGCTTCAGGCAGATGGCAAGACCTGCAAAGATTTTGTAGTGCTCAGTGTACGGCACCTGCAGC
CAGCTATGCACCAACACAGACGGCTCCTTCATATGTGGCTGTGTTGAAGGATACCTCCTGCAGCCGGAT
AACCGCTCCTGCAAGGCCAAGAACGAGCCAGTAGACCGGCCCTGTGCTGTTGATAGCCAACTCCCAG
AACATCTTGGCCACGTACCTGAGTGGGGCCAGGTGTCTACCATCACACCTACGAGCACGGCCAGACC
ACAGCCATGGACTTCAGCTATGCCAACGAGACCGTATGCTGGGTGCATGTTGGGGACAGTGTGCTCAG
ACGCAGCTCAAGTGTGCCCGCATGCCTGGCCTAAAGGGCTTCGTGGATGAGCACACCATCAACATCTCC
CTCAGTCTGCACCAGTGAACAGATGGCCATCGACTGGCTGACAGGCAACTTCTACTTTGTGGATGAC
ATCGATGATAGGATCTTTGTCTGCAACAGAAATGGGGACACATGTGTACATTGCTAGACCTGGAACCTC
TACAACCCCAAGGCATTGCCCTGGACCCTGCCATGGGGAAGGTGTTTTTCACTGACTATGGGCAGATC
CCAAAGGTGGAACACTGTGACATGGATGGGCAGAACCCGACCAAGCTCGTCGACAGCAAGATTGTGTTT
CCTCATGGCATCACGCTGGACCTGGTCAGCCGCTTGTCTACTGGGCAGATGCCTATCTGGACTATATT
GAAGTGGTGGACTATGAGGGCAAGGGCCGACACCATCATCCAGGGCATCCTGATTGAGCACCTGTAC
GGCCTGACTGTGTTTGAATTATCTCTATGCCACCAACTCGGACAATGCCAATGCCAGCAGAAGACG
AGTGTGATCCGTGTGAACCGCTTAACAGCACCGAGTACCAGTTGTACCCGGTGGACAAGGGTGGT
GCCCTCCACATCTACCACCAGAGGCGTCAGCCCCGAGTGAGGAGCCATGCCTGTGAAAACGACAGTAT
GGGAAGCCGGTGGCTGCTGACATCTGCCTGCTGGCCAACAGCCACAAGGCGCGGACCTGCCGCTGC
```



View online »

CGTTCCGGCTTCAGCCTGGGCAGTGACGGGAAGTCATGCAAGAAGCCGGAGCATGAGCTGTTCTCGTG
TATGGCAAGGGCCGGCCAGGCATCATCCGGGCGATGGATATGGGGCCAAAGTCCCGGATGAGCACATG
ATCCCCATTGAAAACCTCATGAACCCCCGAGCCCTGGACTTCCACGCTGAGACCGGCTTCATCTACTTT
GCCGACACCAGCTACCTCATTGGCCGCCAGAAGATTGATGGCACTGAGCGGGAGACCATCCTGAAG
GACGGCATCCACATTGTGGAGGGTGTGGCCGTGGACTGGATGGGAGACAATCTGTACTGGACGGACGAT
GGGCCCCAAAAGACAATCAGCGTGGCCAGGCTGGAGAAAGCTGCTCAGACCCGCAAGACTTAATCGAG
GGCAAAATGACACACCCAGGGCTATTGTGGTGGATCCACTCAATGGGTGGATGTACTGGACAGACTGG
GAGGAGGACCCCAAGGACAGTCGGCGTGGGCGGCTGGAGAGGGCGTGGATGGATGGCTCACACCGAGAC
ATCTTTGTACCTCCAAGACAGTGCTTTGGCCCAATGGGCTAAGCCTGGACATCCCGGCTGGGCGCCTC
TACTGGGTGGATGCCTTCTACGACCCGATCGAGACGATACTGCTCAATGGCACAGACCGGAAGATTGTG
TATGAAGTCTGAGCTGAACCACGCTTTGGCCTGTGTCACCATGGCAACTACCTTTCTGGACTGAG
TATCGGAGTGGCAGTGTCTACCGCTTGGAACGGGGTGTAGGAGGCGCACCCCCACTGTGACCCTTCTG
CGCAGTGAGCGGCCCCCATCTTTGAGATCCGAATGTATGATGCCAGCAGCAGCAAGTTGGACCAAC
AAATGCCGGTGAACAATGGCGGCTGCAGCAGCCTGTGCTTGGCCACCCCTGGGAGCCGACGTCGCC
TGTGCTGAGGACCAGGTGTTGGACGACAGCGGCTCACTTGTGGCGAACCCATCCTACGTGCCTCCA
CCCCAGTGCCAGCCAGGCGAGTTTGCCTGTGCCAACAGCCGCTGCATCCAGGAGCGCTGGAAGTGTGAC
GGAGACAACGATTGCCTGGACAACAGTGATGAGGCCCCAGCCCTCTGCCATCAGCACACCTGCCCTCG
GACCGATTCAAGTGCAGAACACCCGGTGCATCCCAACCCGCTGGCTCTGCGACGGGGACAATGACTGT
GGGAACAGTGAAGATGAGTCCAATGCCACTTGTTCAGCCCGCACCTGCCCCCAACAGTTCTCCTGT
GCCAGTGGCCGCTGCATCCCATCTCCTGGACGTGTGATCTGGATGACGACTGTGGGGACCGCTCTGAT
GAGTCTGCTTCGTGTGCATCCACCTGCTTCCCCCTGACTCAGTTTACCTGCAACAATGGCAGATGT
ATCAACATCAACTGGAGATGCGACAATGACAATGACTGTGGGACAACAGTGACGAAGCCGGCTGCAGC
CACTCCTGTTCTAGCACCCAGTCAAGTGAACAGCGGGCGTTCATCCCGAGCATGGACCTCCGAT
GGGGACAATGACTGCGGAGACTACAGTGATGAGACACACGCCAACTGCACCAACAGGCCACGAGGCC
CCTGGTGGCTGCCACACTGATGAGTTCAGTGCCGGCTGGATGGACTATGCATCCCCCTGCGGTGGCGC
TGCGATGGGGACACTGACTGCATGGACTCCAGCGATGAGAAGAGCTGTGAGGGAGTGACCACGTCTGC
GATCCCAGTGTCAAGTTTGGCTGCAAGGACTCAGCTCGGTGCATCAGCAAAGCGTGGGTGTGTATGGC
GACAATGACTGTGAGGATAACTCGGACGAGGAGAAGTGCAGTCCCTGGCTGCAGGCCACCCTCGCAC
CCTTGTGCCAACACACCTCAGTCTGCCTGCCCTGACAAGCTGTGTATGGCAACGACGACTGTGGC
GACGGCTCAGATGAGGGCAGCTCTGTGACCAGTCTCTGAATAACGGTGGCTGCAGCCACAAGTGC
TCAGTGGCACCTGGCGAAGGCATTGTGTGTTCTGCCCTCTGGCATGGAGCTGGGGCCGACAACCAC
ACCTGCCAGATCCAGAGCTACTGTGCCAAGCATCTCAAATGCAGCCAAAAGTGCAGCCAGAACAAAGTTC
AGCGTGAAGTGTCTGTACGAGGGCTGGGTCTGGAACCTGACGGCGAGAGCTGCCGACGCTGGAC
CCCTTCAAGCCGTTTCATATTTTCTCCAACCGCCATGAAATCCGGCGCATCGATCTTACAAAGGAGAC
TACAGCGTCTGGTGCCCGGCTGCGCAACACCATCGCCCTGGACTTCCACCTCAGCCAGAGCGCCCTC
TACTGGACCGACGTGGTGGAGGACAAGATCTACCGCGGAAGCTGCTGGACAACGGAGCCCTGACTAGT
TTCGAGGTGGTGATTTCAGTATGGCTGGCCACACCCGAGGGCCTGGCTGTAGACTGGATTGCAGGCAAC
ATCTACTGGGTGGAGGTAACCTGGATCAGATCGAGGTGGCCAAGCTGGATGGGACCTCCGACACC
CTGTGGCCGGTGACATTGAGCACCCAAGGGCAATCGCACTGGATCCCGGGATGGGATCCTGTTTTGG
ACAGACTGGGATGCCAGCTGCCCGCATTGAGGCAGCCTCCATGAGTGGGGCTGGGCGCCGACCGTG
CACCGGGAGACCGGCTCTGGGGCTGGCCCAACGGGCTCACCGTGGACTACCTGGAGAAGCGCATCCTT
TGGATTGACGCCAGTTCAGATGCCATTTACTCAGCCCGTTACGACGGCTCTGGCCACATGGAGGTGCTT
CGGGGACACGAGTTCTGTGCGACCCGTTTGCAGTGACGCTGTACGGGGGGAGGTCTACTGGACTGAC
TGGCGAACAAACACACTGGCTAAGGCCAACAGTGGACCGGCCACAATGTACCCTGGTACAGAGGACC
AACACCCAGCCCTTTGACCTGCAGGTGTACCACCCCTCCCGCCAGCCATGGCTCCCAATCCCTGTGAG
GCCAATGGGGCCAGGGCCCTGCTCCACCTGTGTCTCATCACTACAACCGGACCGTGTCTGCGCC
TGCCCCACCTCATGAAGCTCCACAAGGACAACACCACCTGCTATGAGTTAAGAAGTTCCTGCTGTAC
GCACGTACAGATGGAGATCCGAGGTGTGGACCTGGATGCTCCCTACTACAACACTACATCTCCTTACG
GTGCCCGACATCGACAACGTACAGTGCTAGACTACGATGCCCGGAGCAGCGTGTGTAAGTGTGAC
GTGCGGACACAGGCCATCAAGCGGGCTTCATCAACGGCACAGGCGTGGAGACAGTCGTCTCTGCAGAC
TTGCCAAATGCCACGGGCTGGCTGTGGACTGGTCTCCCGAAACCTGTTCTGGACAAGCTATGACACC
AATAAGAAGCAGATCAATGTGGCCCGGCTGGATGGCTCCTTCAAGAAGCAGTGGTGCAGGGCCTGGAG

CAGCCCCATGGCCTTGTGTCGCCACCTCTGCGTGGGAAGCTCTACTGGACCGATGGTGACAACATCAGC
 ATGGCCAAACATGGATGGCAGCAATCGCACCCCTGCTCTTCAAGTGGCCAGAAGGGCCCCGTGGGCTGGCT
 ATTGACTTCCCTGAAAGCAAACCTCTACTGGATCAGCTCCGGGAACCATAACCATCAACCGCTGCAACCTG
 GATGGGAGTGGGCTGGAGGTTCATCGATGCCATGCGGAGCCAGCTGGGCAAGGCCACCGCCCTGGCCATC
 ATGGGGGACAAGCTGTGGTGGGCTGATCAGGTGTGCGAAAAGATGGGCACATGCAGCAAGGCTGACGGC
 TCGGGCTCCGTGGTCTTCGGAACAGCACCCCTGGTGTGCACATGAAGGTCTATGACGAGAGCATC
 CAGCTGGACCATAAGGGCACCAACCCCTGCAGTGTCAACAACGGTGACTGCTCCCAGCTCTGCCTGCC
 ACGTCAGAGACGACCCGCTCCTGCATGTGCACAGCCGGCTATAGCCTCCGGAGTGGCCAGCAGGCCCTGC
 GAGGGCGTAGGTTCTTTCTCCTGACTCTGTGCATGAGGGAATCAGGGGAATCCCTGGATCCCAAT
 GACAAGTCAGATGCCCTGGTCCCAGTGTCCGGACCTCGCTGGCTGTCCGCATCGACTTCCACGCTGAA
 AATGACACCATCTACTGGGTGGACATGGGCTGAGCACGATCAGCCGGGCAAGCGGGACCAGACGTGG
 CGTGAAGACGTGGTACCATGGCATTGGCCGTGTGGAGGGCATTGCAGTGGACTGGATCGCAGGCAAC
 ATCTACTGGACAGACCAGGGCTTGTATGTCATCGAGGTGCGCCGGCTCAATGGCTCCTCCGCTACGTG
 GTGATCTCCAGGTCTAGACAAGCCCCGGCCATCACCGTCCACCCGGAGAAAGGGTACTTGTCTGG
 ACTGAGTGGGGTCAAGTATCCGCGTATTGAGCGGTCTCGGCTAGATGGCACGGAGCGTGTGGTGTGGTC
 AACGTCAGCATCAGCTGGCCCAACGGCATCTCAGTGGACTACCAGGATGGGAAGCTGTACTGGTGGCAT
 GCACGGACAGACAAGATTGAACGGATCGACCTGGAGACAGGTGAGAACCAGGAGGTGTTCTGTCCAGC
 AACAACATGGACATGTTTTCAAGTGTCTGTGTTGAGGATTTCACTACTGGAGTGACAGGACTCATGCC
 AACGGCTCTATCAAGCGCGGGAGCAAAGACAATGCCACAGACTCCGTGCCCTGCGAACCCGGCATCGGC
 GTCCAGCTTAAAGACATCAAAGTCTTCAACCGGGACCGGAGAAAGGCACCAACGTGTGCCGCGTGGCC
 AATGGCGGGTCCAGCAGCTGTCCCTGTACCGGGCCGTGGGACGCGGCTGCCCTGTGCCACGGG
 ATGCTGGCTGAAGACGGAGCATCGTCCCGGAGTATGCCGGTACCTGCTCTACTCAGAGCGACCACTT
 TCAAGAGTATCCACCTGTCCGATGCGCAACCTCAATGCGCCGTCAGCCCTCGAGGACCCCTGAG
 CACATGAAGAACGTCATCGCCCTGGCCTTTGACTACCGGGCAGGACCTCTCCGGGACCCCAATCGC
 ATCTTCTCAGCGACATCCACTTTGGAACATCCAACAGATCAACGACGATGGCTCCAGGAGGATCACC
 ATTGTGAAAACGTGGGCTCCGTGGAAGGCTGGCCTATCACCGTGGCTGGGACACTCTCTATTGGACA
 AGCTACACGACATCCACCATCACGCGCCACACAGTGGACCAGACCCGCCAGGGGCTTCGAGCGTGAG
 ACCGTCATCACTATGTCTGGAGATGACCACCCACGGGCTTCGTTTTGGACGAGTGCCAGAACCTCATG
 TTCTGGACCAACTGGAATGAGCAGCATCCAGCATCATGCGGGCGGCGCTCTCGGAGCCAATGCTCTG
 ACCCTTATCGAGAAGGACATCCGTACCCCAATGGCTGGCCATCGACCACCGTCCGAGAAAGCTCTAC
 TTCTCTGACGCCACCCTGGACAAGATCGAGCGGTGCGAGTATGACGGCTCCACCGCTATGTGATCCTA
 AAGTCAGAGCCTGTCCACCCCTTCGGGCTGGCCGTGTATGGGGAGCACATTTTCTGGACTGACTGGGTG
 CGGCGGGCAGTGCAGCGGGCAACAAGCACGTGGGACGCAACATGAAGCTGCTGCCGCTGGACATCCCC
 CAGCAGCCATGGGCATCATCGCCGTGGCAACGACACCAACAGCTGTGAACTCTCTCCATGCCGAATC
 AACAACGGTGGCTGCCAGGACCTGTGTCTGCTCACTCACAGGGCCATGTCAACTGCTCATGCCAGGG
 GGCCGAATCCTCCAGGATGACCTCACCTGCCGAGCGGTGAATTCCTTTGCCGAGACAAGATGAGTTT
 GAGTGTGCCAATGGCGAGTGCATCAACTCAGCCTGACCTGCGACGGCGTCCCCACTGCAAGGACAAG
 TCCGATGAGAAGCCATCCTACTGCAACTCCCGCGCTGCAAGAAGACTTCCGGCAGTGCAGCAATGGG
 CGCTGTGTGCCAATGCTGTGGTGAACGGGGCCGACGACTGTGGGATGGCTCTGACGAGATCCCT
 TGCAACAAGACAGCCTGTGGTGTGGGCGAGTTCCGCTGCCGGGACGGGACCTGCATCGGGAATCCAGC
 CGCTGCAACCAAGTTTGTGGATTGTGAGGACGCTCAGATGAGATGAACTGCAAGTGCACCCGACTGCAGC
 AGCTACTTCCGCTGGGCGTGAAGGGCGTGTCTTCCAGCCTGCGAGCGGACCTCACTCTGCTACGCA
 CCCAGCTGGGTGTGATGGCGCAATGACTGTGGGGACTACAGTGTGAGCGGACTGCCAGGTGTG
 AAACGCCCCAGATGCCCTCTGAATTAATTCGCTGCCCTAGTGGGCGCTGCATCCCCATGAGCTGGACG
 TGTGACAAAGAGGATGACTGTGAACATGGCGAGGACGAGACCCACTGCAACAAGTTCTGCTCAGAGGCC
 CAGTTTGTGAGTCCAGAACCATCGCTGCATCTCCAAGCAGTGGCTGTGTGACGGCAGCGATGACTGTGGG
 GATGGCTCAGACGAGGCTGCTCACTGTGAAGGCAAGACGTGCGGCCCTCTCTCTCTCTGCCCTGGC
 ACCCACGTGTGCTCCCCGAGCGCTGGCTCTGTGACGGTGACAAAGACTGTGCTGATGGTGCAGACGAG
 AGCATCGCAGCTGGTTGCTGTACAACAGCACTTGTGACGACCGTGAAGTTCATGTGCCAGAACCCGAG
 TGCATCCCCAAGCACTTCGTGTGTGACCACGACCGTACTGTGCAGATGGCTCTGATGAGTCCCCGAG
 TGTGAGTACCCGACCTGCGGCCAGTGTGAGTCCGCTGTGCCAATGGGCGCTGTCTGAGCTCCCGCCAG
 TGGGAGTGTGATGGCGAGAATGACTGCCACGACCAGAGTACGAGGCTCCAAGAACCACACTGCACC

AGCCAAGAGCACAAAGTGAATGCCTCGTCACAGTTCCTGTGCAGCAGTGGGCGCTGTGTGGCTGAGGCA
CTGCTCTGCAACGGCCAGGATGACTGTGGCGACAGCTCGGACGAGCGTGGCTGCCACATCAATGAGTGT
CTCAGCCGCAAGCTCAGTGGCTGCAGCCAGGACTGTGAGGACCTCAAGATCGGCTTCAAGTGCCGCTGT
CGCCCTGGCTTCCGGCTGAAGGACGACGGCCGGACGTGTGCTGATGTGGACGAGTGCAGCACCACCTTC
CCCTGCAGCCAGCGCTGCATCAACTCATGGCAGCTATAAGTGTCTGTGTGGAGGGCTATGCACCC
CGCGCCGGCAGCCCCACAGCTGCAAGGCTGTGACTGACGAGGAACCGTTTCTGATCTTCGCAACCCGG
TACTACCTGCCAAGCTCAACTGGACGGGTCCAACACACGTTACTTAAGCAGGGCTGAACAACGCC
GTTGCCTTGGATTTGACTACCGAGAGCAGATGATCTACTGGACAGATGTGACCACCCAGGGCAGCATG
ATCCGAAGGATGCACCTAACCGGAGCAATGTGCAGGTCCTACACCGTACAGGCCTCAGCAACCCCGAT
GGGCTGGCTGTGGACTGGGTGGGTGGCAACCTGTACTGGTGCACAAAGGCCGGGACACCATCGAGGTG
TCCAAGCTCAATGGGGCTATCGGACGGTGTGGTGTGAGTCTGGCCTCCGTGAGCCAGGGCTCTGGTG
GTGGATGTGCAGAATGGTACCTGTACTGGACAGACTGGGGTACCATTCACTGATCGGCCGCATCGGC
ATGGATGGGTCCAGCCGACGCTCATCGTGGACACCAAGATCACATGGCCCAATGGCCTGACGCTGGAC
TATGCTACTGAGCGCATCTACTGGGCCGACGCCCGGAGGACTACATTGAATTTGCCAGCCTGGATGGC
TCCAATCGCCACGTTGTGCTGAGCCAGGACATCCCGCACATCTTGCAGTACCCTGTTTGAGGACTAC
GTCTACTGGACCGACTGGGAAACAAAGTCCATTAAACCGAGCCACAAGACCACGGGCACCAACAAACG
CTCCTCATCAGCAGCTGCACCGGCCATGGACCTGCATGTCTTCCATGCCCTGCGCCAGCCAGACGTG
CCCAATCACCCCTGCAAGGTCAACAATGGTGGTGCAGCAACCTGTGCCTGTGTCCCCGGGGGAGGG
CACAAATGTGCCTGCCCCACCACTTCTACCTGGGCAGCGATGGGCGCACCTGTGTGTCAAAGTGCACG
GCTAGCCAGTTTGTATGCAAGAACGACAAGTGCATCCCCCTTGTGGTGAAGTGTGACACCCGAGGACGAC
TGCGGGGACCCTCAGACGAGCCCCGGACTGCCCTGAGTTCAAGTGCCGGCCGGACAGTTCAGTGC
TCCACAGGTATCTGCACAAACCTGCCTTCACTGCGATGGCGACAATGACTGCCAGGACAACAGTGC
GAGGCAACTGTGACATCCACGCTGCTTGCAGTCAGTTCAAATGCACCAACCAACCGCTGTATT
CCCGGCATCTCCGCTGCAATGGGCAGGACAACCTGCGGAGATGGGGAGGATGAGAGGGACTGCCCGGAG
GTGACCTGCGCCCCAACCAAGTCCAGTCTCATTACCAACCGTGCATCCCCGGGTCTGGGTCTGC
GACCGGGACAATGACTGTGTGGATGGCAGTGTGAGCCCGCAACTGCACCCAGATGACCTGTGGTGTG
GACGAGTTCGCTGCAAGGATTCGGGCCGCTGCATCCCAGCGCTTGAAGTGTGACGGAGAGGATGAC
TGTGGGGATGGCTCGGATGAGCCAAAGGAAGAGTGTGATGAACGCACCTGTGAGCCATACCAGTTCGCG
TGCAAGAACAACCGCTGCGTGCCCGGCCGCTGGCAGTGCAGTACGACAACGATTGCGGTGACAACCTC
GATGAAGAGAGCTGCACCCCTCGGCCCTGCTCCGAGAGTGAATTCCTGTGCCAACGGCCGCTGCATC
GCGGGGCGCTGGAAATGCGATGGAGACCAGACTGCGCGGACGGCTCGGACGAGAAGAGTGCACCCCC
CGCTGTGACATGGACAGTTCAGTGAAGAGCGGCCACTGCATCCCCCTGCGCTGGCGCTGTGACGCA
GACGCCGACTGCATGGACGGCAGCGACGAGGAGGCTGCGGCACTGGCGTGCAGCTGCCCCCTGGAC
GAGTTCAGTGAACAACACCTTGTGCAAGCCGCTGGCCTGGAAGTGCATGGCGAGGATGACTGTGGG
GACAACCTCAGATGAGAACCCGAGGAGTGTGCCCGTTCTGTGTCCTCCCAACCGGCCCTTCCGTTGC
AAGAAATGACCGCTGTGTCTGTGGATCGGGCGCAATGCGATGGCACGGACAACCTGTGGGGATGGGACT
GATGAAGAGGACTGTGAGCCCCACAGCCACACCCCACTGCAAAGACAAGAAGGAGTTTCTGTGC
CGGAACCAGCGCTGCCTCTCCTCCTCCCTGCGCTGCAACATGTTGATGACTGCGGGGACGGCTGTGAC
GAGGAGGACTGCAGCATCGACCCCAAGCTGACCAGTGCGCCCAATGCCAGCATCTGTGGGGACGAG
GCACGCTGCGTGCACCGAGAAAAGCGGCCTACTGTGCTGCGCTCGGGCTTCCACACCGTGCCTGGC
CAGCCCCGATGCCAAGACATCAACGAGTGCCTGCGCTTCCGACCTGCTCCCAGCTCTGCAACAACACC
AAGGGCGGCCACCTCTGACGCTGCGCTCGGAACCTCATGAAGACGCACAACACCTGCAAGGCCGAAGGC
TCTGAGTACCAGGCTCTGTACATCGCTGATGACAAATGAGATCCGACGCTGTTCCCGGCCACCCCAT
TCGGCTTACGAGCAGGATTCAGGGTACGAGAGTGTCCGATGATGCTATGGATGTCCATGTCAAG
GCTGGCCGTGTCTATTGGACCAACTGGCACACGGGCACCATCTCCTACCGCAGCTGCCACCTGTGCG
CCTCCTACCACTTCCAACCGCCACCGGCGACAGATTGACCGGGTGTACCCACCTCAACATTTAGGG
CTGAAGATGCCAGAGGATCGCCATCGACTGGGTGGCCGAAACGTGTACTGGACCGACTCGGGCCGA
GATGTGATTGAGGTGGCGCAGATGAAGGGCAGAACCGCAAGACGCTCATCTCGGCATGATTGACGAG
CCCCACGCCATTGTGGTGGACCCACTGAGGGGGACCATGTACTGGTCAGACTGGGGCAACCAACCCCAAG
ATTGAGACGGCAGCGATGGATGGGACGCTTCGGGAGACACTGGTGCAGGACAACATTCAGTGGCCACA
GGCCTGGCCGTGGATTATCACAATGAGCGGCTGTACTGGGACGACGCCAAGCTTTAGTCAATCGGCAGC
ATCCGGCTCAATGGCACGGACCCATTGTGGCTGCTGACAGCAAACGAGGCGCTAAGTACCCTTCAGC

ATCGACGTCTTTGAGGATTACATCTATGGTGTCACCTACATCAATAATCGTGTCTTCAAGATCCATAAG
 TTTGGCCACAGCCCTTGGTCAACCTGACAGGGGGCTGAGCCACGCCTCTGACGTGGTCTTTACCAT
 CAGCACAAGCAGCCCGAAGTGACCAACCCATGTGACCGCAAGAAATGCGAGTGGCTCTGCCTGCTGAGC
 CCCAGTGGGCCTGTCTGCACCTGTCCCAATGGGAAGCGGCTGGACAACGGCACATGCGTGCCTGTGCC
 TCTCCAACGCCCCCAGATGCTCCCCGGCTGGAACCTGTACCTGCAGTGTCAACGGTGGCAGC
 TGTTTCTCAATGCACGGAGGCAGCCCAAGTGGCGTGCCAACCCCGCTACACGGGTGACAAGTGTGAA
 CTGGACAGTGTGGGAGCACTGTCGCAATGGGGCACCTGTGCTGCCTCCCCCTGGCATGCCACG
 TGCCGGTGCCCCACGGGCTTACGGGCCCAAATGCACCCAGCAGGTGTGTGCGGGTACTGTGCCAAC
 AACAGCACCTGCACTGTCAACCAGGGCAACCAGCCCCAGTGCCTGACCTACCCGGCTTCTGGGGGAC
 CGCTGCCAGTACCGGCAGTGTCTGGTACTGTGAGAATTTGGCACATGCCAGATGGTGTGATGGC
 TCCCGACAATGCCGTGCACTGCCTACTTTGAGGGATCGAGGTGTGAGGTGAACAAGTGCAGCCGCTGT
 CTCGAAGGGGCTGTGTGGTCAACAAGCAGAGTGGGGATGTACCTGCAACTGCACGGATGGCCGGGTG
 GCCCCAGCTGTCTGACCTGCGTGGCCACTGCAGCAATGGCGGCTCTGTACCTGAACAGCAAAATG
 ATGCTGAGTGCAGTGCACCCACATGACAGGGCCCCGTGTGAGGAGCACGTCTTCAGCCAGCAG
 CAGCCAGGACATATAGCTCCATCTAATCCCTCTGCTGTGCTGCTGCTGCTGTTCTGGTGGCCGGA
 GTGGTATTCTGGTATAAGCGGCGAGTCCAAGGGGCTAAGGGCTTCCAGCACAACGGATGACCAACGGG
 GCCATGAACGTGGAGATTGGAACCCCACTACAAGATGTACGAAGGCGGAGAGCCTGATGATGTGGGA
 GGCTACTGGACGCTGACTTTGCCCTGGACCCTGACAAGCCCACTTACCAACCCCGTGTATGCC
 ACACTCTACATGGGGGGCATGGCAGTGCACCTCCCTGGCCAGCACGGACGAGAAGCGAGAACTCTG
 GGCCGGGGCCCTGAGGACGAGATAGGGGACCCCTTGCA
ACCGGTACGCGGCCGCTCGAGCAGAAACT**ATCTCAGAAGAGGATCTGGCAGCA**AA**TGATATCTGGAT**
TACAAGGATGACGACGATAAGGTTAAACGGCCGGC

Protein Sequence:

>Peptide sequence encoded by RC218369
 Blue=ORF Red=Cloning site Green=Tag(s)

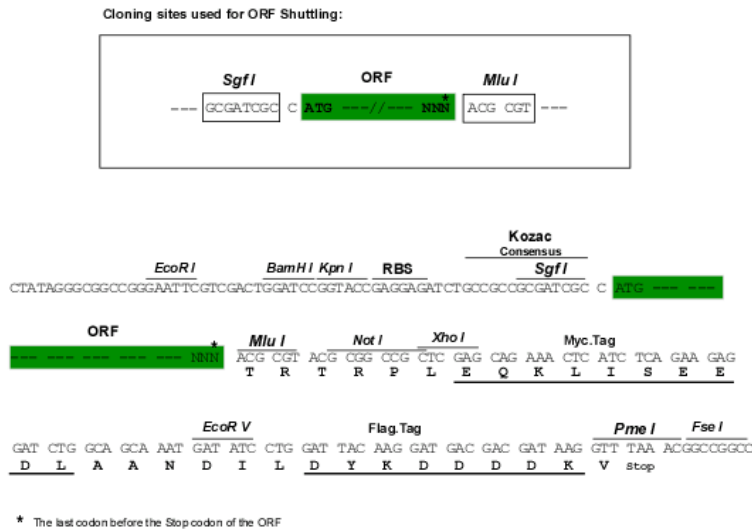
MLTPLLLLLLPLLSALVAAAIDAPKTCSPKQFACRDQITCISKWRCGERDCPDGSDEAPEICPQSKA
 QRCQPNEHNCLGTEL CVPMSRLCNGVQDCMDGSDGPHCRELQGNCSRLGCQHHCVP TLDGPTCYCNSS
 FQLQADGKTKDFDECSVYGTCSQLCTNTDGSFICGCVGYLLQPDNRSCAKNEPVRPPVLLIANSQ
 NILATYLSGAQVSTIPTSTRQTTAMDFSYANETVCWVHVGDSSAAQTQLKCARMPGLKGFVDEHTINIS
 LSLHHVEQMAIDWLTGNFYFVDDIDDRIFVCNRNGDTCVTLLEDLELYNPKGIALDPAMGVFFTDYQGI
 PKVEHCDMDGQNRKLVDSKIVFPHGITL DLVSRLVYWADAYLDYIEVVDYEGKGRQTI IQGILIEHLY
 GLTVFENLYATNSDNANAQKTSVIRVNRFNSTEQVVTRVDKGGALHIYHQRQRPRVRSHACENDQY
 GKPGGSDICLLANSHKARTCRCSGSLGSDGKSCKKPEHELFLVYGGKGRPGIIRGMDMGAKVPDEHM
 IPIENLMNPRALDFHAETGFIFYADTTSYLIGRQKIDGTERETILKDGIIHIVEGVAVDWMGDNLWYTD
 GPKKTISVARLEKAAQTRKTLIEGKMTHPRAIVDPLNGWYWDWEEDPKSRRGRLERAWMDGSHRD
 IFVTSKTVLWPNGLSLDIPAGRLYWVDAFYDRIETILLNGTDRKIVYEGPELNHAFGLCHHGNLFWTE
 YRSGSVYRLERGVGGAPPTVTLR SERPPIFEIRMYDAQQQVGTNKR VNNGGCSSLCLATPGSRQCA
 CAEDQVLDADGVTCLANPSYVPPQCPGEFACANSRCIQERWKCDGDNCLDNSDEAPALCHQHTCPS
 DRFKCENNRIPNRWCDGDNDCGNSSESNATCSARTCPPNQFSCASGRCIPI SWTCDLDDDCGRSD
 ESASCA YPTCFPLTQFTCNNGRCININWRCDNDNDCGNSDEAGCSHSCSSTQFKNSGRCIPEHWTCD
 GDNDGCDYSDETHANCTNQATRPPGGCHTDEFQCRLDGLCIPLRWRCDGDTDCMDSDEKSGEVTHVC
 DPSVKFGCKDSARCI SKAWVCDGNDCEDENSDEENCESLACRPPSHPCANNTSVCLPPDKLCDGNDDCG
 DGSDGELCDQCSL NNGGCSHNCSVAPGEGIVCSCPLGMELGPDNHTCQIQSYCAKHLKCSQKCDQNKF
 SVKCSCEYEGWLEPDGEGSCRSLDPFKPFIIFSNRHEIRRIDLHKGDYSVLPGLRNTIALDFHLSQSAL
 YWTDVVEDKIYRGKLLDNGALTSFEVVIQYGLATPEGLAVDWIAGNIYWVESNLQIEVAKLDGTLRTT
 LLAGDIEHPRAIALDPRDGI LFWTDWDASLPRIEAA SMSGAGRRTVHRE TGSGGWPNGLTVDYLEKRIL
 WIDARSDAIYSARYDGS GHMEVLRGHEFLSHPFAVTL YGGEVYWDWRTNTLAKANKWTHNVTVVQRT
 NTQPFDLQVYHPSRQPMAPNPCEANGGQGPCSHLCLINYNRTVSCACPHMLKHKDNTTCYEFKFLLY
 ARQMEIRGVLDLAPYYNYIISFTVPDIDNVTVDYDAREQRVYWSVVRTQA IKRAFINGTVETVVSAD
 LPNAHGLAVDWVSRNLFWTSYD TNKKQINVARLDGSFKNAV VQGLEQPHGLVVHPLRGKLWYTDGDNIS
 MANMDGSNRTLLFSGQKGPVGLAIDFPESKLYWISSGNHTINRCNL DSGGLEVIDAMRSQLGKATALAI
 MGDKLWWADQVSEKMGTC SKADGSGSVL RNSTL VMHMKVYDESIQLDHKGTNPCSVNNGDCSQLCLP

TSETTRSCMCTAGYSLRSGQACEGVSFLLYSVHEGIRGIPLDPNDKSDALVPVSGTSLAVGIDFHAEN
 NDTIYVWDMGLSTISRARDQTWREDVVTNGIGRVEGIAVDWIAGNIYWDQGFVDIEVARLNGSFRYV
 VISQGLDKPRAITVHPEKGYLFWTEWGYPRIERSRLDGTERTVVLVNVSIWPNGISVDYQDGKLYWCD
 ARTDKIERIDLETGENREVVLSNNMDFSVSVFEDFIYWSDRTHANGSIKRGSKDNATDSVPLRTGIG
 VQLKDIKVFNRDRQKGTNVCAVANGGCQQLCLYRGRGQRACACAHGMLAEDGASCREYAGYLLYSERTI
 LKSIHLSDERNLNAPVQPFEDPEHMKNVIALAFDYRAGTSPGTPNRIFFSDIHFNGIQIINDDGSRRIT
 IVENVGSVEGLAYHRGWDTLWTSYTTSTITRHTVDQTRPGAFERETVITMSGDDHPRFVLDCEQNL
 FWTNWNQHPSPIMRAALSGANVLTLEKDIRTPNGLAIDHRAEKLYFSDATLTKIERCEYDGSYRVIIL
 KSEPVHFPGLAVYGEHIFWTDWVRRRAVQRANKHVGSNMKLLRVDIPQQPMGIIVAVANDTNSCELSPCRI
 NNGGCQDLCLLTHQGHVNCSCRGGRIQLQDDLTCRAVNSSCRAQDEFECANGECINFSLTCDGVPHCKDK
 SDEKPSYCNSRRCKKTFRQCSNGRCVSNMLWCGADDCCGDSDEIPCNKACGVGEFRCDGTICGNSS
 RCNQFVDCEDASDEMNCSDCASSYFRLGVKGVLFQPCERTSLCYAPSWVCDGANDCGDYSYDERDCPGV
 KRPRCPLNYFACPSGRCIPMSWTCDEKDDCEHGEDETHCNKFCSEAQFECQNHRCISKQWLDGSDDCG
 DGSDEAAHCEGKTCGPSSFSCPGTHVCVPERWLDGDKDCADGADESIAAGCLYNSTCDDREFMCCNRQ
 CIPKHFVCDHRCADGSDSPECEYPTCGPSEFRANGRCLSSRQWECGENDCHDQSDAPKNPHCT
 SQEHKCNASSQFLCSSGRCVAEALLCNGQDDCGDSSDERGCHINECLSRKLSGCSQDCEDLKI
 GFKCRCRPGFRLKDDGRTCADVDECSTTFPCSQRICINTHGSYKCLVEGYAPRGGDPHSCAVTDEEPFLIFANR
 YYLRKLNLDGSNYTLKQGLNNAVALDFDYREQMIYWTDVTTQGSIMRRMHLNGSNVQLHRTGLSNPD
 GLAVDWVGGNLWCDKGRDTEVSKLNGAYRTVLVSSGLREPRALVVDVQNGYLWTDWGDHSLIGRIG
 MDGSSRSVIVDTKITWPNGLTLDYVTERIYWADAREDYIEFASLDGNSRHVVLSDIPHIFALTLFEDY
 VYWTDWETKINRAHKTGTNKTLLISTLHRPMDLHVHALRQPDVNPCKVNNGGCSNLCLLSPGGG
 HKCACPTNFYLGSDGRTCVSNCTASQFVCKNDKIPFVWKCDTEDDCGDHSDEPPDCPEFKCRPGQFQC
 STGICTNPAFICDGDNDCCQNSDEANCDIHVCLPSQFKCTNTNRCIPGIFRCNGQDNCGDGEDRDCPE
 VTCAPNFQCSITKRCIPRVWVCDRDNDVDSDEPANCTQMTCGVDEFRCCKDSGRCIPARWCKDGEDD
 CGDGSDEPKKECDERTCEPYQFRCKNNRCPGRWQCDYDNDCGDNDSEESCTPRPCESEFSCANGRCI
 AGRWCKDGDHDCADGSDKEDCTPRCDMDQFQCKSGHCIPLRWRCDADADCMDGSDEEACGTGVRTCLPD
 EFQCNNTLCKPLAWKCDGEDDCGDNSENPEECARFVCPNRPFRCKNDRVCLWIGRQCDGTNCGDGT
 DEEDCEPPTAHTTHCKDKKEFLCRNQRLSSSLRCNMFDCCGDSDEEDCSIDPKLTSCATNASICGDE
 ARCVRTEKAAYCACRSGFHTVPGQPGCQDINECLRFGTCSQLCNNTKGGHLCSCARNFMKTHNTCKAEG
 SEYQVLYIADDNEIRSLFPGHPSAYEQAFQGDSEVRIDAMDVHVKAGRVYWTNWHTGTISYRSLPPAA
 PPTTSNRHRRQIDRGVTHLNIISGLKMPRGIAIDWVAGNVYWTDSGRDVIEVAQMKGENRKTLSGMIDE
 PHAIVVDPLRGTMYSWSDWGNHPKIETAAMDGTLRETLVQDNIQWPTGLAVDYHNERLYWADAKLSVIGS
 IRLNGTDPIVAADSKRGLSHPFSIDVFEDYIYGVTYINNRVFKIHKFGHSPLVNLTGGLSHASDVVLYH
 QHKQPEVTNPCRKKCEWLCLLSPSGPVCTCPNGKRLDNGTCVVPVSPPTPPDAPRPGTCNLQCFNGGS
 CFLNARRQPKCRCQPRYTGDCKELDQCWEHCRNGGTCAASPSGMPTCRCPTGFTGPKCTQVQVACAYCAN
 NSTCTVNQGNQPCRCPLPGFLGDRQCQYRQCSGYCENFGTCQMAADGSRQCRCTAYFEGSRCEVNKCSRC
 LEGACVVNKQSGDVTNCNTDGRVAPSCLTCVGHCSNGGCTMNSKMMPECCQPPHMTGPRCEEHVFSQQ
 QPGHIASILIPLLLLLLLLVLAGVVFVYKRRVQAKGFQHQRMNTGAMNVEIGNPTYKMYEGGEPDDVG
 GLLDADFALDPDKPTNFTNPVYATLYMGHGHSRSLASTDEKRELLGRGPEDEIGDPLA
 TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Chromatograms: https://cdn.origene.com/chromatograms/mg2663_f01.zip

Restriction Sites: Sgfl-MluI

Cloning Scheme:



ACCN: NM_002332

ORF Size: 13632 bp

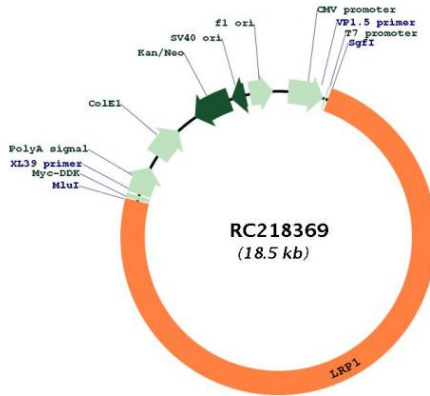
OTI Disclaimer: Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at custsupport@origene.com or by calling 301.340.3188 option 3 for pricing and delivery.

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](#)

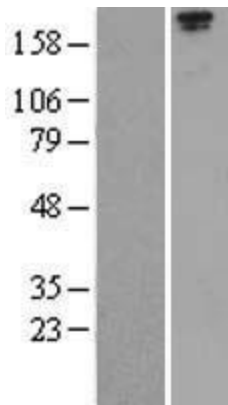
OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

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:	<ol style="list-style-type: none">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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_002332.3
RefSeq Size:	14896 bp
RefSeq ORF:	13635 bp
Locus ID:	4035
UniProt ID:	Q07954
Cytogenetics:	12q13.3
Protein Families:	Druggable Genome, Transmembrane
Protein Pathways:	Alzheimer's disease
MW:	504.6 kDa
Gene Summary:	This gene encodes a member of the low-density lipoprotein receptor family of proteins. The encoded preproprotein is proteolytically processed by furin to generate 515 kDa and 85 kDa subunits that form the mature receptor (PMID: 8546712). This receptor is involved in several cellular processes, including intracellular signaling, lipid homeostasis, and clearance of apoptotic cells. In addition, the encoded protein is necessary for the alpha 2-macroglobulin-mediated clearance of secreted amyloid precursor protein and beta-amyloid, the main component of amyloid plaques found in Alzheimer patients. Expression of this gene decreases with age and has been found to be lower than controls in brain tissue from Alzheimer's disease patients. [provided by RefSeq, Oct 2015]

Product images:



Circular map for RC218369



Western blot validation of overexpression lysate (Cat# [LY400839]) using anti-DDK antibody (Cat# [TA50011-100]). Left: Cell lysates from untransfected HEK293T cells; Right: Cell lysates from HEK293T cells transfected with RC218369 using transfection reagent MegaTran 2.0 (Cat# [TT210002]).