

## Product datasheet for **RC227975**

### Ki67 (MKI67) (NM\_001145966) Human Tagged ORF Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Ki67 (MKI67) (NM\_001145966) Human Tagged ORF Clone  
**Tag:** Myc-DDK  
**Symbol:** MKI67  
**Synonyms:** KIA; MIB-; MIB-1; PPP1R105  
**Vector:** pCMV6-Entry (PS100001)  
**E. coli Selection:** Kanamycin (25 ug/mL)  
**Cell Selection:** Neomycin  
**ORF Nucleotide Sequence:** >RC227975 representing NM\_001145966  
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
 GCCCGCATCGCC

ATGTGGCCACGAGACGCTGGTTACTATCAAAGGAGCGGGTTCGACGGTCCCCACTTCCCCTGAGCC  
 TCAGCACCTGCTTGTGGGAAGGGTATTGAATGTGACATCCGTATCCAGCTTCCTGTTGTGTCAAAACA  
 ACATTGCAAAATTGAAATCCATGAGCAGGAGGCAATATTACATAATTTTCAGTCCACAAATCCAACACAA  
 GTAAATGGGTCTGTTATTGATGAGCCTGTACGGCTAAAACATGGAGATGTAATAACTATTATTGATCGTT  
 CCTTCAGGTATGAAAATGAAAAGTCTTCAGAATGGAAGGAAGTCAACTGAATTTCCAAGAAAAATACGTGA  
 ACAGGAGCCAGCACGTGCTCAAGATCTAGCTTCTCTTCTGACCCTGATGAGAGTGAGGGAATACCT  
 TTGAAAAGAAGGCGTGTGTCCTTTGGTGGGCACCTAAGACCTGAACTATTTGATGAAAACCTGCCTCCTA  
 ATACGCCTCTCAAAGGGGAGAAGCCCCAACCAAAAAGAAAGTCTCTGGTAATGCACACTCCACCTGTCCT  
 GAAGAAAATCATCAAGGAACAGCCTCAACCATCAGGAAAACAAGAGTCAGGTTCCAGAAATCCATGTGGAA  
 GTGAAGGCACAAAGCTTGGTTATAAGCCCTCCAGCTCCTAGTCCTAGGAAAACCCAGTTGCCAGTGATC  
 AACGCCGTAGGTCCTGCAAAACAGCCCCTGCTCCAGCAGCAAATCTCAGACAGAGGTTCCCTAAGAGAGG  
 AGGGAGAAAGAGTGGCAACCTGCCTTCAAAGAGAGTGTCTATCAGCCGAAGTCAACATGATATTTTACAG  
 ATGATATGTTCCAAAAGAAGAAGTGGTCTCGGAAGCAAATCTGATTGTTGCAAAATCATGGGCAGATG  
 TAGTAAAACCTGGTGCAAAACAAACACAACTAAAGTCATAAAACATGGTCTCAAAGGTCAATGAACAA  
 AAGGCAAGAAGACCTGCTACTCAAAGAAGCCTGTGGCGAAGTTCACAGTCAATTTAGTACAGGCCAC  
 GCAAACCTCCTTGTACCATAATAATAGGGAAGCTCATACTGAAAAAGTACATGTGCCTGCTCGACCCT  
 ACAGAGTGCTCAACAACCTCATTTCACCAAAAAAATGGACTTTAAGGAAGATCTTTCAGGAATAGCTGA  
 AATGTTCAAGACCCAGTGAAGGAGCAACCGCAGTTGACAAGCACATGTCACATCGCTATTTCAAATTC  
 GAGAATTTGCTTGGAAAACAGTTTCAAGGAAGTATTGAGGAGAAAGAACCTCTGCTCCCCACCTCAGAGA  
 GTTTTGGAGGAAATGTGTTCTTCAGTGCACAGAATGCAGCAAAACAGCCATCTGATAAATGCTCTGCAAG  
 CCCTCCCTTAAGACGGCAGTGTATTAGAGAAAATGGAACGTAGCAAAAACGCCAGGAACACCTACAAA  
 ATGACTTCTCGGAGACAAAACCTCAGATACTGAGACAGAGCCTTCAAAAACAGTATCCACTGCAAAACA



GGTCAGGAAGGTCTACAGAGTTCAGGAATATACAGAAGCTACCTGTGGAAAGTAAGAGTGAAGAAACAAA  
 TACAGAAATTGTTGAGTGCATCCTAAAAAGAGGTGAGAAGGCAACACTACTACAACAAAGGAGAGAAGGA  
 GAGATGAAGGAAATAGAAAGACCTTTTGGAGACATATAAGGAAAAATTGAATTAAGAAACCGATGAAA  
 AGATGAAAGCAATGAAGAGATCAAGAAGCTTGGGGCAGAAATGTGCACCAATGTCTGACCTGACAGACCT  
 CAAGAGCTTGCCTGATACAGAACTCATGAAAGACACGGCAGCTGGCCAGAATCTCCTCCAAACCCAAGAT  
 CATGCCAAGGCCAAAGAGTGAAGAAAGGCAAAATCACTAAAAAGCCCTGCCAGTATTACAACCCAGAAC  
 CAATAAACACCCCAACACACACAAAACACAGTTGAAGGCATCCCTGGGAAAGTAGGTGTGAAAGAAGA  
 GCTCCTAGCAGTCGGCAAGTTCACACGGACGTACAGGGGAGACCACGCACACGCACAGAGAGCCAGCAGGA  
 GATGGCAAGAGCATCAGAACGTTTAAAGGAGTCTCCAAAGCAGATCCTGGACCCAGCAGCCGTTAACTG  
 GAATGAAGAAGTGGCAAGAAGCCCTAAGGAAGAGGCCAGTCACTAGAAGACCTGGCTGGCTTCAAAGA  
 GCTCTTCCAGACACCAGTCCCTCTGAGGAATCAATGACTGATGAGAAAACCTACAAAATAGCCTGCAAA  
 TCTCCACCACCAGAATCAGTGGACACTCCAACAAGCACAAAGCAATGGCCTAAGAGAAGTCTCAGAAAAG  
 CAGATGTAGAGGAAGAATTTAGCACTCAGGAACTAACACCATCAGCAGGGAAAGCCATGCTTACGCC  
 CAAACCAGCAGGAGGTGATGAGAAAGACATTAAGCATTATGGGAACTCCAGTGCAGAACTGGACCTG  
 GCAGGAACTTTACCTGGCAGCAAAAGACAGCTACAGACTCCTAAGGAAAAGGCCAGGCTCTAGAAGACC  
 TGGCTGGCTTTAAAGAGCTCTCCAGACTCCTGGTCACACCGAGGAATTAGTGGCTGCTGGTAAAACCA  
 TAAAAACCTGCGACTCTCCACAGTCAGACCCAGTGGACACCCCAACAAGCACAAAGCAACGACCCAAG  
 AGAAGTATCAGGAAAGCAGATGTAGAGGGAGAAGTCTTAGCGTGCAGGAATCTAATGCCATCAGCAGGCA  
 AAGCCATGCACACGCCTAAACCATCAGTAGGTGAAGAGAAAGACATCATATTTGTGGGAACTCCAGT  
 GCAGAACTGGACCTGACAGAGAAGTAAACCGGCAGCAAGAGACGGCCACAAACTCCTAAGGAAGAGGCC  
 CAGGCTCTGGAAGACCTGACTGGCTTTAAAGAGCTCTCCAGACCCCTGGTCACTACTGAAGAAGCAGTGG  
 CTGCTGGCAAACTACTAAAAAGCCCTGCGAATCTTCCACCAGAATCAGCAGACACCCCAACAAGCAC  
 AAGAGGCAGCCCAAGACACCTTTGGAGAAAAGGGACGTACAGAAGGAGCTCTCAGCCCTGAAGAGCTC  
 ACAAGCATCAGGGGAAACACACACACAGATAAAGTACCAGGAGGTGAGGATAAAAGCATCAAGCGCT  
 TTAGGGAAGTGCAAAACAGAACTGGACCCAGCAGCAAGTGTAACTGGTAGCAAGAGGCCCAAAAAAC  
 TAAGGAAAAGGCCAACCCTAGAAAGCCTGGCTGGCTTGAAGAGCTCTCCAGACACCAGTATGCACT  
 GACAAGCCACGACTCACGAGAAAACCTACAAAATAGCCTGCAGATCACAACCAGACCCAGTGGACACAC  
 CAACAAGCTCCAAGCCACAGTCCAAGAGAAGTCTCAGGAAAGTGGACGTAGAAGAAGAATTTCCGCACT  
 CAGGAAACGAACACCATCAGCAGGCAAGCCATGCACACACCCAAACCAGCAGTAAAGTGGTGAAGAAAAC  
 ATCTACGATTTATGGGAACTCCAGTGCAGAACTGGACCTGACAGAGAAGTAACTGGCAGCAAGAGAC  
 GGCTACAACTCCTAAGGAAAAGGCCAGGCTCTAGAAGACCTGGCTGGCTTTAAAGAGCTCTCCAGAC  
 ACGAGGTACACTGAGGAATCAATGACTAACGATAAAACTGCCAAAGTAGCCTGCAAACTCTCACAAACCA  
 GACCCAGACAAAAACCCAGCAAGCTCCAAGCGACGGCTCAAGACATCCCTGGGGAAGTGGGCGTGAAAAG  
 AAGAGCTCCTAGCAGTTGGCAAGCTCACACAGACATCAGGAGAGACTACACACACACACAGAGCCAAC  
 AGGAGATGGTAAGAGCATGAAAGCATTATGGAGTCTCCAAAGCAGATCTTAGACTCAGCAGCAAGTCTA  
 ACTGGCAGCAAGAGGCAGCTGAGAACTCCTAAGGGAAGTCTGAAGTCCCTGAAGACCTGGCCGGCTTCA  
 TCGAGCTCTCCAGACACCAAGTCACTAAGGAATCAATGACTAACGAAAAAACTACCAAAGTATCCTA  
 CAGAGCTTACAGCCAGACCTAGTGGACACCCCAACAAGCTCCAAGCCACAGCCCAAGAGAAGTCTCAG  
 AAAGCAGACACTGAAGAAGAATTTTAGCATTAGGAAACAAACGCCATCAGCAGGCAAGCCATGCACA  
 CACCCAAACAGCAGTAGGTGAAGAGAAAGACATCAACACGTTTTTGGGAACTCCAGTGCAGAAAAGTGA  
 CCAGCCAGGAAATTTACCTGGCAGCAATAGACGGCTACAACTCGTAAGGAAAAGGCCAGGCTCTAGAA  
 GAAGTACTGGCTTCCAGAGAGCTTTCCAGACACCATGCACTGATAACCCACGACTGATGAGAAAACCTA  
 CAAAAAATACTCTGCAATCTCCGCAATCAGACCCAGCGGACACCCCAACAAACACAAAGCAACGGCC  
 CAAGAGAAGCCTCAAGAAAGCAGACGTAGAGGAAGAATTTTAGCATTAGGAAACTAACACCATCAGCA  
 GGCAAGCCATGCACACGCCTAAAGCAGCAGTAGGTGAAGAGAAAGACATCAACACATTTGTGGGGACTC  
 CAGTGGAGAACTGGACCTGCTAGGAAATTTACCTGGCAGCAAGAGACGGCCACAACTCCTAAAGAAA  
 GGCCAAGGCTCTAGAAGATCTGGCTGGCTTCAAAGAGCTCTCCAGACACCAGGTCACTGAGGATCA  
 ATGACCGATGACAAAATCACAGAAGTATCCTGCAATCTCCACAACCAGACCCAGTCAAACCCCAACAA  
 GCTCCAAGCAACGACTCAAGATATCCTTGGGGAAGTAGGTGTGAAAGAAGAGGTCTACCAGTCGGCAA  
 GCTCACACAGACGTGAGGGAAGACCACACAGACACACAGAGAGACAGCAGGAGATGGAAGAGCATCAAA  
 GCGTTTAAAGGAATCTGCAAGCAGATGCTGGACCCAGCAAACTATGGAAGTGGATGGAGAGGTGGCCAA  
 GAACACCTAAGGAAGAGGCCAATCACTAGAAGACCTGGCCGGCTTCAAAGAGCTCTCCAGACACCAGA

CCACACTGAGGAATCAACAACCTGATGACAAAACCTACAAAATAGCCTGCAAATCTCCACCACCAGAATCA  
 ATGGACTCTCAAACAAGCACAAGGAGGCGGCCAAAACACCTTTGGGAAAAGGGATATAGTGGAAAGAGC  
 TCTCAGCCCTGAAGCAGCTCACACAGACCACACACAGACAAAAGTACCAGGAGATGAGGATAAAGGCAT  
 CAACGTGTTCAAGGAACTGCAAAAACAGAACTGGACCCAGCAGCAAGTGTAACTGGTAGCAAGAGGCAG  
 CCAAGAACTCCTAAGGAAAAGCCCAACCCCTAGAAGACTTGGCTGGCTTGAAGAGCTCTCCAGACAC  
 CAATATGCACTGACAAGCCCAGACTCATGAGAAAACCTACAAAATAGCCTGCAGATCTCCACAACCAGA  
 CCCAGTGGGTACCCCAACAATCTTCAAGCCACAGTCCAAGAGAAGTCTCAGGAAAAGCAGACGTAGAGGAA  
 GAATCCTTAGCACTCAGGAAAACGAACACCATCAGTAGGAAAAGCTATGGACACACCCAAACAGCAGGAG  
 GTGATGAGAAAAGACATGAAAGCATTATGGAACTCCAGTGCAGAAAATTGGACCTGCCAGGAAAATTTACC  
 TGGCAGCAAAGATGGCCACAACTCCTAAGGAAAAGGCCAGGCTCTAGAAGACCTGGCTGGCTTCAA  
 GAGCTCTTCCAGACACCAGGCACTGACAAGCCACGACTGATGAGAAAACCTACAAAATAGCCTGCAAAT  
 CTCCACAACCAGACCCAGTGGACACCCAGCAAGCACAAGCAACGGCCCAAGAGAACTCAGGAAAGC  
 AGACGTAGAGGAAGAATTTTAGCACTCAGGAAAACGAACCATCAGCAGGCAAAGCCATGGACACACCA  
 AAACCAGCAGTAAGTGTGAGAAAAATATCAACACATTTGTGAAAACCTCAGTGCAGAAAACCTGGACTGC  
 TAGGAAAATTTACCTGGCAGCAAGAGACGCCACAGACTCCTAAGGAAAAGGCTGAGGCTCTAGAGGACCT  
 GTTTGGCTTCAAAGAACTCTCCAGACACCAGGTCACACTGAGGAATCAATGACTGATGACAAAATCACA  
 GAAGTATCCTGTAAATCTCCACAGCCAGAGTCATTCAAAAACCTCAAGAAGCTCCAAGCAAAGGCTCAAGA  
 TACCCCTGGTGAAGTGGACATGAAAGAAGAGCCCTAGCAGTCAAGCTCACACGGACATCAGGGGA  
 GACTACGCAAACACACACAGAGCCAACAGGAGATAGTAAGAGCATCAAAGCGTTTAAAGAGTCTCCAAAG  
 CAGATCCTGGACCCAGCAGCAAGTGTAACTGGTAGCAGGAGGCACTGAGAACTCGTAAGGAAAAGGCC  
 GTGCTCTAGAAGACCTGGTTGACTTCAAAGAGCTTCTCAGCACCAGGTCACACTGAAGAGTCAATGAC  
 TATTGACAAAAACAAAAATTCCTGCAAACTCCCCACCAGAATAACAGACTGCCACGAGCACA  
 AAGAGATGCCCAAGACACGTCACAGGAAAAGTAAGGAGGAGCTCTCAGCAGTTGAGAGGCTCAGC  
 AAACATCAGGGCAAAGCACACACACACAAAGAACCAGCAAGCGGTGATGAGGGCATCAAAGTATTGAA  
 GCAACGTGCAAAGAAGAAACCAACCCAGTAGAAGAGGAAACCCAGCAGGAGAAGGCCAAGACACCTAAG  
 GAAAAGGCCCAACCCCTGGAAGACCTGGCCGGCTTACAGAGCTCTGAAAACATCAGGTCACTCAGG  
 AATCACTGACTGCTGGCAAAGCCACTAAAATACCCTGCGAATCTCCCCACTAGAAGTGGTAGACACCAC  
 AGCAAGCACAAGAGGCATCTCAGGACACGTGTGAGAAGGTACAAGTAAAAGAAGAGCCTTTCAGCAGTC  
 AAGTTCACACAAAACATCAGGGGAAACACGGATGCAGACAAAGAACCAGCAGGTGAAGATAAAGGCATCA  
 AAGCATTGAAGGAATCTGAAAAACAGACACCGGCTCCAGCAGCAAGTGTAACTGGCAGCAGGAGAGGCC  
 AAGAGCACCCAGGAAAAGTGCCCAAGCCATAGAAGACCTAGCTGGCTTCAAAGACCCAGCAGCAGGTAC  
 ACTGAAGAATCAATGACTGATGACAAAACCACTAAAATACCCTGCAAATCATCACCAGAACTAGAAGACA  
 CCGCAACAAGCTCAAAGAGACGGCCAGGACACGTGCCAGAAAAGTAGAAGTGAAGGAGGAGCTGTTAGC  
 AGTTGGCAAGCTCACACAAAACCTCAGGGGAGACCACGCACACCACAAAGAGCCGGTAGGTGAGGGCAA  
 GGCACGAAAGCATTTAAGCAACCTGCAAAGCGGAAGCTGGACGCAAGAATGTAATTGGCAGCAGGAGAC  
 AGCCAAGAGCACCTAAGGAAAAGGCCCAACCCCTGGAAGATCTGGCCAGCTTCCAAGAGCTCTCTCAAAC  
 ACCAGGCCACTGAGGAACTGGCAATGGTGTGCTGATAGCTTTACAAGCGCTCAAAGCAAACACCT  
 GACAGTGGAAAACCTCTAAAAATATCCAGAAGAGTTCTTCGGGCCCTAAAGTAGAACCCGTGGGAGAGC  
 TGTAAGCACCAGAGACCCTGTAAAATCACAAAGCAAAAAGCAACTTCCCTGCCCCACTGCCCTTCAA  
 GAGGGGAGGTGGCAAAGATGGAAGCGTACGGGAACCAAGAGGCTGCGCTGCATGCCAGCACAGAGGAA  
 ATTGTGGAGGAGCTGCCAGCAGCAAGAAGCAGAGGGTTGCTCCAGGGCAAGAGGCAAATCATCCGAAC  
 CCGTGGTTCATCATGAAGAGAAGTTTGGAGACTTCTGAAAAAGAATTGAACCTGCGGAAGAGCTGAACAG  
 CAACGACATGAAAACCAAAAGAGGAACACAAATTACAAGACTCGGTCCCTGAAAAAAGGGAATATCC  
 CTGCGCTCCAGACGCCAAAATAAGACTGAGGCAGAACAGCAAATAACTGAGGTCTTTGTATTAGCAGAAA  
 GAATAGAAATAAACAGAAATGAAAAGAAGCCCATGAAGACCTCCCAGAGATGGACATTCAGAATCCAGA  
 TGATGGAGCCCGAAAACCCATACCTAGAGACAAAGTCACTGAGAACAAAAGGTGCTTGGAGTCTGCTAGA  
 CAGAATGAGAGCTCCAGCCTAAGGTGGCAGAGGAGAGCGGAGGCAGAAGAGTGCAGAGTTCTCATGC  
 AGAATCAGAAAAGGAAAAGGAGAAGCAGGAAATTCAGACTCCATGTGCCTGAGATCAAGAAAGACAAAAAG  
 CCAGCCTGCAGCAAGCACTTTGGAGAGCAAATCTGTGCAGAGAGTAACCGCGGAGTGTCAAGAGGTGTGCA  
 GAAAATCCAAAGAAGGCTGAGGACAATGTGTGTGCAAGAAAATAAGAACCAGAAGTCATAGGGACAGTG  
 AAGATATT

AGCGGACCGACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC  
TGGATTACAAGGATGACGACGATAAGGTTAA

**Protein Sequence:**

>RC227975 representing NM\_001145966  
Red=Cloning site Green=Tags(s)

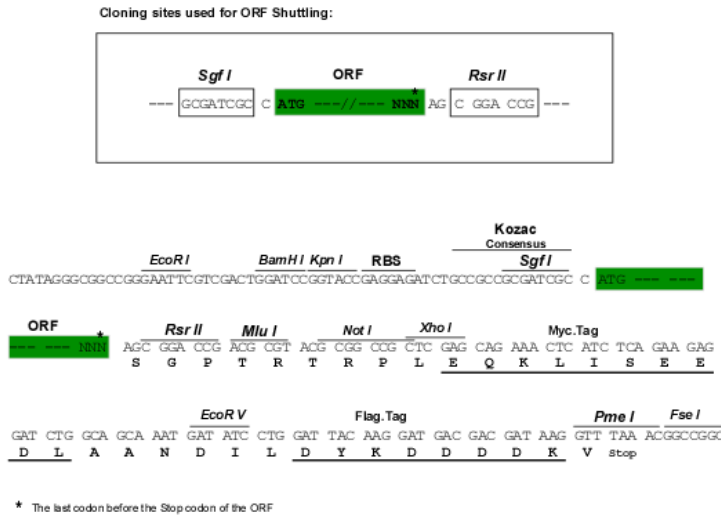
MWPTRRLVTIKRSGVDGPHFPLSLSTCLFGRGIECDIRIQLPVVSKQHCKIEIHEQEAILHNFSSTNPTQ  
VNGSVIDEPVRLKHGDVITIIDRSFRYENESLQNGRKSTEFPRKIREQEPARRVSRSSFSSDPDESEGIP  
LKRRRVVSGGHLRPELFDENLPNTPKLRGEAPTKRKSVMHTPPVLKKIIEKEQPQPSGKQESGSEIHVE  
VKAQSLVISPPAPSPRKTPVASDQRRRSCKTAPASSSSQTEVPKRGGRKSGNLSKRVVSRSQHDILQ  
MICKRRSGASEANLIVAKSWADVVKLGAQTQTKVIKHGPPQRSMNKRQRRPATPKKPVGEVHSQFSTGH  
ANSPCTIIIGKAHTEKVHVPARPYRVLNFIISNQKMDFKEDLSGIAEMFKTPVKEQPQLTSTCHIAISNS  
ENLLGKQFQGTDSGEEPLLPTSESGFNVFFSAQNAAKQPSDKCSASPPLRRQCIRENGNVAKTPRNTYK  
MTSLETKTSDTETEPSKTVSTANRSRSTEFRNIQKLPVESKSEETNTEIVECILKRGQKATLLQORREG  
EMKEIERPFETYKENIELKENDEKMKAMKRSRTWQKCAPMSDLTDLKSLPDELMKDTARGQNLQTQD  
HAKAPKSEKGIITKMPQCQSLQPEPINTPTHTKQQLKASLGKVGVEELLAVGKFRTRSGETTHTHREPAG  
DGKSIRTFKESPKQILDPAARVTGMKKWPRTPKEEAQSLLEDLAGFKELFQTPGPSEESMTDEKTTIACK  
SPPPEVDTPSTKQWPKRSLRKADVEEEFLALRKLTPSAGKAMLTPKAGGDEKDIKAFMGTPVQKLDL  
AGTLPGSKRQLQTPKEKAQALEDLAGFKELFQTPGHTEELVAAGTKTKIPCDSPQSDPVDTPSTKQRPK  
RSIRKADVEGELLACRNLMPGKAMHTPKPSVGEEDKIIIFVGTVPVQKLDL TENLTGSKRRPQTPKEEA  
QALEDLTGFKELFQTPGHTEEEVAAGTKTKMPCESSPESADTPTSTRRQPKTLEKRDVQKEL SALKKL  
TQTSGETTHTDKVPGGEDKSINAFRETAQKLDPAASVTGSKRHPKTKEKAQPLEDLAGLKFQTPVCT  
DKPTTHEKTTKIACRSQDPVDTPSSKQSKRSLRKVDVEEEFFALRKRTPSAGKAMHTPKPAVSGEKN  
IYAFMGTPVQKLDL TENLTGSKRRLQTPKEKAQALEDLAGFKELFQTRGHTEESMTNDKTAKVACKSSQP  
DPDKNPASSKRRLKTS LGKVGVEELLAVGKLTQTSGETTHTHTEPTGDGKSMKAFMESPKQILDSAASL  
TGSKRQLRTPKGVSEVPEDLAGFIELFQTPSHTKESMTNEKTTKVSYRASQPDLDVTPSSKQPKRSLR  
KADTEEEFLAFRKTQPSAGKAMHTPKPAVGEEDINTFLGTPVQKLDQPGNLPGSNRRLQTRKEKAQALE  
ELTGRELQTPCTDNPTTDEKTTKILCKSPQSDPADTPTNTKQRPKRLKADVEEEFLAFRKLTPSA  
GKAMHTPKAAVGEEDINTFVGTVPVEKLDLLGNLPGSKRRPQTPKEKAKALEDLAGFKELFQTPGHTEES  
MTDDKI TEVSCKSPQDPVKTPSSKQRLKISLGKVGVEEVLVPGKLTQTSKTTQTHRETAGDGKSIK  
AFKESAKQMLDPANYGTGMERWPRTPKEEAQSLLEDLAGFKELFQTPDHTEESTDDKTTIACKSPPPE  
MDTPTSTRRRPKTPLGKRDIVEEL SALKQLTQTHTDKVPGEDEKGINVFRETAQKLDPAASVTGSKRQ  
PRTPKGKAQPLEDLAGLKFQTPICTDKPTTHEKTTKIACRSQDPVGTPTIFKQSKRSLRKADVEE  
ESLALRKRTPSVGKAMDTPKAGGDEKDMKAFMGTPVQKLDLPGNLPGSKRWPQTPKEKAQALEDLAGFK  
ELFQTPGTDKPTTDEKTTIACKSPQDPVDTPASTKQRPKRNLRKADVEEEFLALRKRTPSAGKAMDTP  
KPAVSDEKNINTFVETPVQKLDLLGNLPGSKRQPTPKEKAEALDVGFKELFQTPGHTEESMTDDKIT  
EVSCKSPQESFKTSRSSKQRLKIPLVKVDMKEEPLAVSKL TRTSGETTQTHTEPTGDSKSIKAFKESPK  
QILDPAASVTGSRRLRTRKEKARALEDLVDFKELF SAPGHTEESMTIDKNTKIPCKSPPPEL TDTATST  
KRCPKTRPRKEVKEELSAVERLTQTSQSSTHHEKPEASGDEGIKVLKQRAKKKPNPVEEESRRRPRAPK  
EKAQPLEDLAGFTELSETSGHTQESL TAGKATKIPCESPPLEVVDTTASTKRHLRTRVQKVVQVKEEPSAV  
KFTQTSGETTDADKEPAGEDKGIKALKESAKQTPAAPASVTGSRRRRPRAPRESAQAIEDLAGFKDPAAGH  
TEESMTDDKTTKIPCKSSPELEDATSSKRRPRTRAQKVEVKEELLAVGKLTQTSGETTHTDKEPVGEGK  
GTAKFKQPAKRKLD AEDVIGSRROPAPKEKAQPLEDLASFQELSQTPGHTEEL ANGAADSFTSAPKQTP  
DSGKPLKISRRVLRAPKVEPVGDVVSTRDPVKSQSKSNTSLPPLPFKRGGGKDGSVTGKRLRCMPAPEE  
IVEELPASKKQRVAPRARGKSSEPVVIMKRLRTSAKRIEPAEELNSNDMKTNKEEHKLDQSVPENKGIS  
LRSRRQNKTEAEQQITEVFVLAERIEINRNEKKPMKTSPEMDIQNPDDGARKPIPRDKVTENKRLRSAR  
QNESSQKVAEESGGQKSAKVLMLQNKQKGEAGNSDSMCLRSRKTQSPAASTLESKSVQVRVTRSVKRCR  
ENPKKAEDNVCKKIRTRSHRDEDI

SGP TRTRRLEQKLI SEEDLAANDILDYKDDDDKV

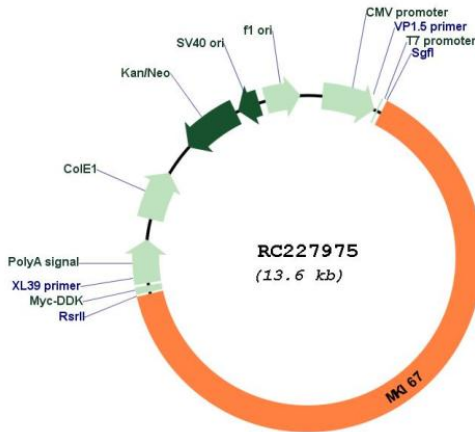
**Restriction Sites:**

Sgfl-RsrII

Cloning Scheme:



Plasmid Map:



ACCN: NM\_001145966

ORF Size: 8688 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](#)

<b>OTI Annotation:</b>	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
<b>Components:</b>	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).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<a href="#">NM_001145966.2</a>
<b>RefSeq ORF:</b>	8691 bp
<b>Locus ID:</b>	4288
<b>UniProt ID:</b>	<a href="#">P46013</a>
<b>Cytogenetics:</b>	10q26.2
<b>Protein Families:</b>	Druggable Genome, ES Cell Differentiation/IPS
<b>MW:</b>	319.3 kDa
<b>Gene Summary:</b>	This gene encodes a nuclear protein that is associated with and may be necessary for cellular proliferation. Alternatively spliced transcript variants have been described. A related pseudogene exists on chromosome X. [provided by RefSeq, Mar 2009]