

Product datasheet for RC218559

ZNF292 (NM_015021) Human Tagged ORF Clone

Product data:

Product Type: Expression Plasmids
Product Name: ZNF292 (NM_015021) Human Tagged ORF Clone
Tag: Myc-DDK
Symbol: ZNF292
Synonyms: bA393I2.3; Nbla00365; ZFP292; Zn-15; ZN-16
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
ORF Nucleotide Sequence: >RC218559 representing NM_015021
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGC**C

ATGGCGGACGAAGAGGCCGAGCAGGAGAGGTTGAGTTGCGGGCAAGCGGCTGCGTCGCGGAGCTGCAGC
GCCTGGGCGAGCGGCTCCAGGAGCTGGAGCTACAGCTGCGGGAGAGCCGGTACCGGCCGTGGAAGCGGC
CACCGACTACTGTCAGCAGCTGTGCCAGACACTCTAGAATATGCAGAGAAATGGAAAACCTCAGAAGAT
CCTTTACCTTTATTGGAGGTATACACAGTGGCTATCCAAAGTTATGTTAAAGCCCGACCTTATCTTACCT
CTGAATGTGAAAATGTAGCCTTGTTCTGGAACGCTTGGCATTAAAGCTGTGTTGAACTTTTACTGTGTCT
GCCTGTTGAGTTATCAGATAAACAGTGGGAACAATTTAGACACTGGTGCAGGTAGCTCATGAAAAGCTG
ATGGAGAATGGCAGCTGTGAATTGCATTTTTAGCTACTCTAGCTCAAGAGACTGGGGTGTGAAAAACC
CGGTACTGTGCACTATTCTTCCAGGAACCATGGATAAAGATAAAGTGAATGAATTTTTAGCTTTTGA
GGTCCCATCTTGTGGATATGAGAATTAACATCTAATCAAAACAAATCAGTTAAGTCAAGCAACTGCT
CTAGCAAAGCTGTGTTCTGACCATCCAGAGATTGGCATAAAAGGTAGTTTTAAGCAAACCTACCTTGTCT
GTCTTTGTACATCATACCAAATGGAAAGTTAATCGAAGAGATTTGAAAGTTGATTGCAAAAGTGCAT
GGAAATGATCTGTAACCTAGAATCTGAGGGTGTGAAAAAAGCGCTCTTGTTTTATGACTGCGTTTTTG
TCACGTGAGCTCCAACAAGGAGATATGACTGCGCTTGGGAACCTACTCTCTTTTGGAGTAAATTACAAC
AAAGAGTGAACCATCTATAAAGTGTACCTTGAGAGGTGTCGCAACTTTCTTTGTTAACGAAAACAGT
ATATCACATTTCTTCTGATTAAGTTATTAATTCAGAGACTGAAGGGGCTGGACTTGCTACCTGTATA
GAACTGTGTGTAAGGCTCTTCGCTTGGAGTCTACAGAAAATACTGAAGTAAAAATCTATTTGCAAGA
CCATTTTATGTTTGTGCTGATGATCTGGAAGTTAACGTGCTTGTCAACTGAGTGAATTTCTTATTGA
GCCTACAGTAGATGCGTATTATGCTGTGAAATGTTGTATAATCAGCCAGACCAGAAATATGATGAAGAG
AATCTTCAATACCAAATCTTTACGCTGTGAGCTGTTACTTGTATTGAAAACCTCAATGGCCCTTTGATC
CAGAATTTCTGGGATTGAAAAACCTTGAACGACAATGTCTTGCATTAATGGGAGAGAAGCATCCATTGT
GTCTTCAATAGATGAACTAAATGACAGTGAAGTATGAAAAAGTGGTAGACTACCAAGAAGAGAGTAAA
GAAACTTCTATGAATGGGCTTTCTGGTGGAGTTGGTGCTAATCTGGCCTTCTTAAAGACATTGGTGATG



AAAAGCAGAAGAAGAGAGAGATAAAACAGTTAAGAGAGAGGGGATTTATATCTGCTCGGTTTAGGAATTG
GCAAGCCTACATGCAGTATTGTGTGTTGTGTGACAAAGAATTCCTTGGTCACAGAATAGTACGACATGCT
CAGAAACATTACAAAGATGGAATTTATAGTTGCCCATATGTGCAAAGAATTTAATTCTAAAGAAACTT
TTGTCCCTCATGTCACACTGCATGTTAAACAATCTAGTAAAGAGAGACTAGCAGCTATGAAACCATTAAG
AAGATTGGGAAGGCCTCCAAAGATCACAACACCAATGAAAATCAGAAGACTAATACTGTGGCTAAACAG
GAGCAGCGACCTATAAAAAAGAATAGTCTCTATTCAACAGATTTTATAGTGTTTAATGACAATGATGGTT
CAGATGATGAGAATGATGACAAAGATAAATCCTATGAGCCAGAAGTGATCCAGTCCAGAACCAGTACC
TGTTAATGAATTTAATTGCCCTGTAACCTTTTGTAAAAAGGGCTTTAAGTACTTTAAAAATTTAATTGCT
CATGTGAAGGGGCATAAAGATAATGAAGACGCCAAGCGCTTTCTTGAATGCAGAGCAAAAAAGTTATTT
GCCAGTACTGTAGGCGGCATTTTGTGAGTGTTACTCATCTCAATGATCACTTACAGATGCACTGTGGCAG
TAAACCATATATCTGTATACAGATGAAATGTAAGCTGGTTTTAATAGTTACGCCGAGCTTTTAAACCCAC
CGAAAGGAGCATCAAGTCTTTAGAGCAAAATGTATGTTTCCTAAATGTGGAAGAATTTTTTCGGAAGCTT
ATTTACTATATGATCATGAAGCACAAACATTATAATACGTACACTTGAAGTTCACAGGTTGTGGTAAAGT
TTATCGTTCTCAGGGTGAGCTGGAAAAGCATCTGGATGATCACAGTACTCCTCCTGAAAAAGTGTCTGCT
CCTGAAGCCCACTTAATTCATCTGGAGATTCCATTACGCTTCTGAAGTGAATCAGAACACAGCAGAGA
ATATTGAGAAAAGAAAGATCTATGCTTCCTTCAGAAAATAACATTGAAAACAGCTTACTAGCAGATAGAAG
TGATGCTTGGGATAAAAGCAAAGCAGAATCAGCTGTGACCAAAACAAGACCAGATTTCTGCCTCTGAGCTC
AGGCAAGCTAATGGACCATTGTCAAATGGTTTGGAAAACCTGCTACTACTCCTCTACTTCAATCCAGTG
AAGTAGCTGTGCCATTAAGGTGTCTCTCAATCAGGGGATTGAGGATAACTTTGGAAAGCAAGAAAACCTC
AACTGTGGAAGGCAGTGGTGAAGCACTGGTCACAGACTTACATACGCCAGTTGAAGATACTTGTAAATGAT
TTGTGTCATCCAGGTTTCCAGGAGAGAAAAGAACAAGATTGCTTTAATGATGCCCATGTTACTCAGAATT
CTTTAGTAAATTCAGAACTCTCAAAATAGGTGACCTTACCCACAAAACCTTGAAGACAAAGTGAACAA
CTTGATGACCTTTTCTGTGCAAAATCAGGCAGCATTTCAAACAATTTACCAACTTCAAAATTTGAATGT
GGAGATAATGTTAAACATCATCCAATCTTTATAATTTACCTCTTAAGACATTAGAAAAGTATTGCATTTG
TTCCACCAGCAGTCCGACCTAAGTAATTCATTAGGAACCTCATCAGTGCCTCCTCAAAAGCTCCAGTTCAGAA
ATTCAGCTGCCAGGTCGAGGGATGACTCGAACCTATAATTCTTACAGAGTATTGGGAAACACATGAAG
ACAGCACACCCTGACCAATATGCTGCATTTAAATGCAGCGCAAAAGTAAAAAGGTGAGAAAGCTAACA
ACTTAAATACACCAAATAATGGAAAGTTTGTATTTTTTCCATCACCAGTGAACAGCTCAAATCCATT
TTTTACATCACAGACCAAAGCCAATGGGAATCTGCTTGTTCGGCCAGTTGCAGCATGTCTCGCCACCC
ATTTTTCCAGCTCATTTAGCAAGTGTGCAACTCCATTGTTGCTCCTCAATGGAAGTGCATAAAATCCAA
ATATAACTTCTCAGGATAAAAAAGAACAAGGTGGTATGTTATGTTCCCAATGGAATAATACCTAGTAC
TGCTTTCAGCAGCAAAATGGAAGATCTAACCAAAACAGTTCTGCCTTTGAATATTGACAGTGGCTCAGAT
CCTTTCCTTCTTACCTGCAGAAAGTAGTTCAATGTCTCTCTTCCCTTACCAGCAGATAGTGGGACTA
ATTCTGTTTTTTCCCACTGGAAAATAATACAAATCATTATTCCTCACAGATTGAAGGAAACACTAATTC
CTCCTTCTAAAGGGGGTAAATGGTGAATAATGCAGTTTTTCTTCCACAAGTGAATGTTGCAATAACTTC
AGTAGCACCAATGCCAACAGTCTGCACCTGAAAAAGTAAAAAAGACCGTGGGCGGGGCCAAATGGGA
AGGAAAGAAAACCTAAGCACAAACAAAAGGGCTAAATGGCTGCAATATCAGAGATGGGAAATTTATCTG
TAGCAGGTGTTACAGGGCTTTTACTAATCCAGATCACTGGGTGGCAGTTATCCAAGCGATCTTACTGT
AAACCACTGGATGGAGCCGAAATGCTCAAGAACTTCTACAGAGTAATGGACAGCCTTCTTCTTGGCA
GCATGATTCTCTCCACAAATGCAGTAAATTTGCAGCAGCCACAACAATCTACCTTCAATCCAGAAGCATG
TTTTAAAGATCCATCATTCTACAGCTTCTGCTGAAAATCGCTCGCCAGCATTTTTACCAATACATTT
CCTCGATCTGGTGTGACTAATTTAATACCAGTGTGAGTCAAGAAGGTAGTGAATTTAATAACAGGCTT
TGGAAACTGCTGGCATTCCCAGTACATTTGAGGGTGCCGAAATGCTTTCTCATGTTTCAACAGGTTGTGT
CTCTGATGCATACAAGTAAATGCAACGGTGATGCCAAATCCAAGTGTACCACCCTGTTGCACACTGTA
TGCCATCCAAACACCTTGTGACCAACCAGAATAGGACGTCAAACCTCAAACCTCCTCCATTGAGGAAT
GTAGCAGCTTGCCTGTTTTTCCAACGAATGACTTACTACTGAAGACTGTTGAAAATGGTTTGTGCTCTAG
TTCATTTCTAATCTGGTGGGCCATCACAATAATTTACCAGTAAAGTCTCGTGTCTGTTATAAGT
GGTCTCAGAACACAAGATCCAGTCAATTTAAATAAAAAAGGAAACAGTGTCTTAAAGAGAAGAAAGAAAG
TTGCTCCTCCACTAATTCACCTAACGCTTCCCAAACTTGGTAACAAGTGAATTAACAACAATGGGACT
CATAGCAAAGAGTGTGAAATCCCAACTACTAACCTTCAATCAATGTAATTCCAACTTGTGAACCTCAG
AGTTTGGTGGAAAATCTAACACAGAAATTAATAATGTTAACAATCAGTTATTTATGACTGATGTAAG
AGAATTTCAAACAGTCTTGTGCTCCATACAGTGTAGCCCTTTAACATTAACAACTGAAAATGGTGA

TTCCCAAATGATGGCTTTGAATTCATGCACAACCTCAATAAATTCTGATTTGCAGATTTCTGAAGACAAT
GTTATACAAAACCTTTGAAAAGACTCTTGAAATTATTAATAACTGCATGAATTTCTCAAATCTTGAGGTAA
AAAGTGGATCTCAGGGTCTGGTGAAACTTCACAAAATGCCTCAAATAAATTATAACATTCAGCTTCTCTC
AGTAAACACTGTGCAAAAATACAAAATTACCCGATTCTTCTCCGTTTTCTCTTTATAAGTGTATGCCA
ACAAAAAGTAACATTCCTCAGTCTGAAGTATCACATAAGGAGGATCAAATACAGGAAATTTAGAAAGCT
TACAGAAATTAATAATAGAAAATGACCTATCCACTCCAGCATCCCAATGTGTACTGATAAATACATCAGT
GACACTGACTCCACCGCTGTTAAATCAACTGCAGATATCACAGTTATTCAGCCAGTTTCTGAAATGATA
AACATTCATTTAATGACAAAAGTTAATAAACCCCTTGTGTGTCAAACCAAGGCTGTAACCTACAGTGCTA
TGACAAAAGGATGCACTATTTAAGCACTATGGTAAAATTCATCAATACACTCCAGAAATGATTCTTGAAAT
TAAGAAGAATCAATTGAAATTTGCTCCCTTTAAATGTGTAGTACCTACATGTACAAAAACATTTACAAGA
AATTCTAACCTCCGGGCACACTGTCAGTTGGTGCATCATTTTACAACCTGAAGAAATGGTAAAGTTAAAAA
TTAAAAGGCCCTTATGGAAGAAAATCTCAGAGTGAAGTGTGCCGGCTCACGAAGTACACAAGTGAAGAAA
ACAGCTAGCTATGACAGAGGAAAATAAAAAGGAATCTCAGCCTGCTTTAGAATTGAGAGCAGAGACCCAA
AATACCCACAGTAATGTAGCAGTATCCAGAAAAACAACCTGTAGAAAAAAAAGTCTGACAAAAACAG
AAAGTTCTTTACAAGTATTACAGTTACTTCAGAACAATGTAATACAAATGCACTCACAAACACACAAC
CAAAGGACGGAAGATTAGGAGGCATAAAAAAGAAAAGGAGGAGAAAAACGAAAGAAGCCAGTTTCCCAA
TCCCTTGAGTTTCCAACAAGATACAGTCTTACAGACCTTATCGATGTGTTACCAGGGATGCTTTGCTG
CCTTTACGATACAGCAAACTTGATTCTCCATTACCAGGCTGTACACAAATCAGATCTACCTGCATTTTC
AGCAGAGGTGCAAGAGGAAAAGTGAAGCTGGTAAAGAAAAGTGAAGAACTGAAACTAAACAACTTTGAAA
GAATTCGATGTCAGGTAAAGTACTGTTCTCGAATTTTCCAAGCAATTACTGGCCTAATACAACACTACA
TGAACTTCATGAAATGACTCCTGAAGAAATGAAAGTATGACTGCTTCAGTGGATGTTGGGAAGTTTCC
ATGTGACCAGTTAGAGTGTAAATCTTCTTACTACATATTTGAACTATGTTGTTATCATCTAGAGGCAGAC
CACGGGATTGGACTAAGGGCAAGTAAAACAGAAGAAGTGGTGTATACAAATGTGATTGTGAAGGCTGTG
ACCGTATATATGCAACCCGGTCAATCTCCTCCGACACATTTTTAATAAGCATAATGACAAACATAAGGC
TCATTTGATTCGTCCAAGAAGTTAACACCAGGCCAGGAAAATATGTCAAGCAAGGCAAAACCAAGAAAA
TCAAAGCTAAACATCGGGGGACCAAGCACAGCAGATGTGAAAGGAAGGAATAAAAATGCCAAGACCA
AACGAAAGAAAAAATAATTTAGAAAACAAGAATGCAAAGATTGTGCAGATTGAAGAAAATAAGCCTTA
TTCTCTGAAACGTGGGAAGCATGTATTTCTATAAAGGCTAGAAATGATGCCCTGTCTGAGTGTACAAGC
AGATTTGTAACCCAGTATCCATGTATGATAAAGGGATGACTTCAAGTGTACAAGTGAAGCAATATAA
TTAGACATTATAAGTGCCATAAATTATCTAAGGCATTTACATCACAAACCCGAAATCTTCTTATTGTATT
CAAACGGTGTGCAACTCACAAAGTAAAGGAAACGTCTGAGCAAGAAGGTGCTAAGATGATGTGAAGAT
TCTGACACGTGTATCAGAGAGCAATGATAATTCAAGAACAACAGCTACAGTTTCAAAAAGGAAGTTG
AAAAAATGAAAAAGATGAAATGGATGAACTAACAGAATTGTTTATTACAAAATTAATAATGAAGATAG
CACAAGTGTAGAGACCAAGCTAATACTTCTTCAAATGTAAGTAAATGATTTTTCAGGAAGATAACCTCTGC
CAGTCAGAAAGACAAAAAGCAAGTAATTTGAAGAGAGTTAATAAGGAAAAAATGTCTCACAAAAATAAA
AAAGGAAAGTTGAAAAAGCTGAACCAGCATCAGCAGCTGAGTTAAGTAGCGTGCCTAAAGAAGAAGAAAC
TGCTGTTGCCATTCAAACCATTGAGGAGCATCCTGCATCTTTTACTGGAGCTCTTTTAAAGCAATGGGA
TTTGAAGTATCATTTCTGAAGTTTCTTGAGGAGTCTGCAGTGAAGCAGAAGAAAAATACTGACAAAGACC
ATCCGAACTGGAACAAAAAAGGATCCCATTCAAATTCAAAGAAAAAATATTGATAAAGACTGCTGTGAC
TAGTGGAAATCATGTATGCTTGTAAAGAAAGCGAAACGTTTGTACAGTTTGCCAATCCATCACAGCTT
CAGTGCAGTGATAATGTAAAAATTGTTTTAGACAAGAATCTTAAAGATTGCACTGAGCTTGTCTTAAAGC
AACTTCAGGAAATGAAACCTACCGTCAGTCTGAAAAAAGTGAAGTACATTCAAATGATCCAGATATGTC
TGTTATGAAAGATATCAGTATAGGTAAGCCACAGGCAGAGGTCAGTAC

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
ACAAGGATGACGACGATAAAGTTTAA

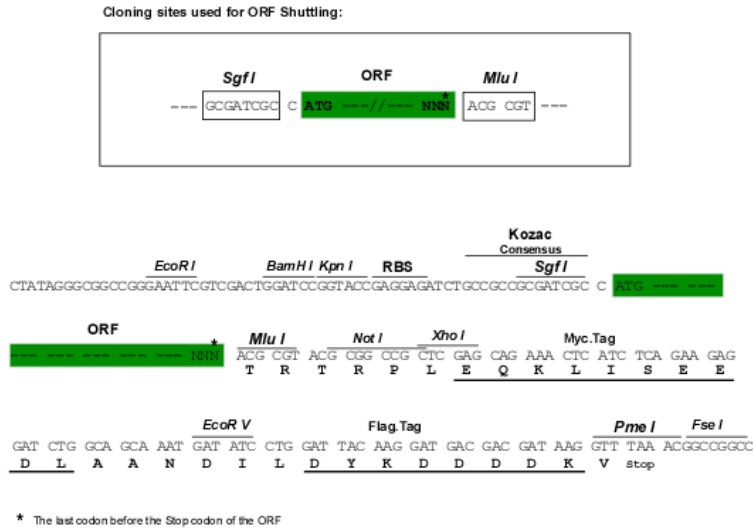
Protein Sequence: >RC218559 representing NM_015021
 Red=Cloning site Green=Tags(s)

MADEEAEQERLSCGEGGCVAELQRLGERLQELELQLRESRVPAAEATDYCQQLCQTLLEAYAEKWKTS
 EDPLPLLEVYTVAIQSYVKARPYLTSECNVALVLERLALSCVELLLCLPVELSDKQWEQFQTLVQVAHEKL
 MENGSCELHFLATLAQETGVWKNPVLCTILSQEPLDKDKVNEFLAFEGPILLDMRIKHLIKTNQLSQATA
 LAKLCSDHPEIGIKGSFKQTYLVCLCTSSPNGKLIIEEISEVDCKDALEMICNLESEGDEKSAVLCTAFL
 SRQLQQGDMYCAWELTLFWSKLLQQRVEPSIQVYLERCRQLSLLTKTVYHIFFLIKVINSETEGAGLATCI
 ELCVKALRLESTENTEVIKISICKTISCLLPDDLEVKRACQLSEFLIEPTVDAYYAVEMLYNQPDQKYDEE
 NLPINSLRCELLLVLKTQWPFDPFWDWTKLRQCLALMGEEASIVSSIDELNDSEVYEKVVVDYQEEESK
 ETSMNGLSGGVGANGLLKDIDGDEKQKREIKQLRERGFISARFRNWQAYMYQYCVLCKEFLGHRIVRHA
 QKHYKDGIIYSCPICAKNFNSKETFPVPHVTLHVQSSKERLAAMKPLRRLGRPPKITTTNENQKTNTVAKQ
 EQRPICKNSLYSTDFIVFNDNDGSDDENDDKDKSYEPEVIVQKPVVNEFNCPVTFCKGFKYFNKLI
 AVHVKGHKDNEDAKRFLMQSKKVICQYCRHFVSVTHLNDHLQMHCGSKPYICIQMKCKAGFNSYAE
 LLTHRKEHQVFRACKMFPKCGRIFSEAYLLYDHEAQHYNTYTCKFTGCGKVYRSQGELEKHLDDHSTP
 PEKVLPEAQLNSSGDSIQPSEVNQNTAENIEKERSMLPSENNIENSLADRSDAWDKSKAESAVTKQDQI
 SASELROANGPLSNGLENPATPLLQSSEVAVSIVKSLNQGIEDNFGKQENSTVEGSGEALVTDLHTP
 VEDTCNDLCHPGFQERKEQDCFNDAAHTQNSLVNSETLKIIGDLTPQNLERQVNNLMTFSVQNA
 AAFQNNLPTSKFECGDNVKTSSNLNPLKTLIESIAFVPPQSDLSNSLGTSPVPPKAPVQKFSQV
 EGCTRTYNSSQSIGKHMKTAAHPDQYAAAFKMQRKSKKQKANNLNTPNNGKFVYFLPSVNSSNPF
 FTSQTKANGNPACSAQLQHVSPPIFPAHLASVSTPLSSMESVINPNITSQDKNEQGGMLCSQ
 MENLPSTALPAQMEDLTKTVLPLNIDSGSDPFLPLPAESSMSLFPSPADSGTNSVFSQ
 LENNTNHYSSQIEGNTNSSFLKGGNGENAVFSPQVNVANNSSTNAQSAPEKVKKDRGRG
 PNGKERKPKHNKRAKWPAAIIRDGKFCISRCYRAF TNPRSLGGHLKSRSYCKPLDGAET
 IAQELLQSNQPSLLASMILSTNAVNLQQPQSTFNPEACFKDPSFLQLLAENRSPAFLPNTF
 PRSGVTNFNTSVSQEGSEI IKQALETAGIPSTFEGAEMLSHVSTGCVSDASQVNAVMPNPTV
 PPLLHTVCHPNTLLTNQRTSNSKTSIEECSSLPVFPTNDLLLKTVENGLCSSSFPNSGGPSQ
 NFTSNSSRVSVISGPQNTRSSHLNKKGNSASKRRKKVAPPLIAPNASQNLVTSDLTMMGLIAK
 SVEIPTTNLHNSVIPTCEPQSLVENLTQKLNNVNNQLFMTDVKENFKTSLESHTVLAPLTLK
 TENGDSQMMALNSCTTSINSDLQISEDNVIQNFKLEI IKTAMNSQILEVKSQSGAGETSQNAQ
 INYNIQLPSVNTVQNNKLPDSSPSSFSISVMPTKSNIPQSEVSHKEDQIQEILEGLQKL
 KLENDLSTPASQCVLINTSVTLTPTPVKSTADITVIQPVSEMINIQFNDKVNKPFV
 CQNQGCNYSAMTKDALFKHYGKIHQYTPEMILEIKKNQLKFAFPKCVVPTCTKTFTR
 NSNLRAHCQLVHHFTTEEMVKLIKRPYGRKSQSENVASRSTQVKKQLAMTEENKKESE
 QPALELRAETQNTHSNVAVIPEKQLVEKKSPDKTESSLQVITVTSQCNTNALTNTQTKGRK
 IRRHKKEKEEKKRKKPVSQSLEFPTRYSPYRYPYRVHQQGCAAFITIQNLILHYQAVHKS
 DLPFAFSAEVEESEEAGKESEETETKQTLKEFRQCQVSDCSRIFQAITGLIQHYMKLHE
 MTPPEIESMTASVDVGKFPDQLECKSSFTTYLNVVHLEADHGI GLRASKTEEDGVYK
 DCEGCDRIYATRSNLLRHIFNKHNDKHKHAHLIRPRRLTPGQENMSSKANQEKSKSK
 HRGTHKSRGCKEGIKMPKTKRKKNNLENKNAKIVQIEENKPYSLKRGKHVYSIKARNDAL
 SECTSRFVTQYPCMIKGTSSVVTSESNIRHYKCHKLSKAFTSQHRNLLIVFKRCCNSQV
 KETSEQEGAKNDVKDSDTCVSESNDNSRTTATVSQKEVEKNEKDEMDDELTELFITKLIN
 EDSTSVETQANTSSNVSNDFQEDNLCQSERQKASNLKRVNKEKNVSQNKRRKVEKAEP
 ASAAELSSVRKEEETAVAIQTIEEHPASFDWSSFKPMGFEVSFLKFLSESAVKQKNTD
 KHPNTGNKKGSHSNSRKNIDKTAVTSGNHVCPCKESETFFVQFANPSQLQCSDNVKIVL
 DKNLKDCTELVLKQLQEMKPTVSLKKLEVHSNDPDMVMKDISISGKATGRGQY

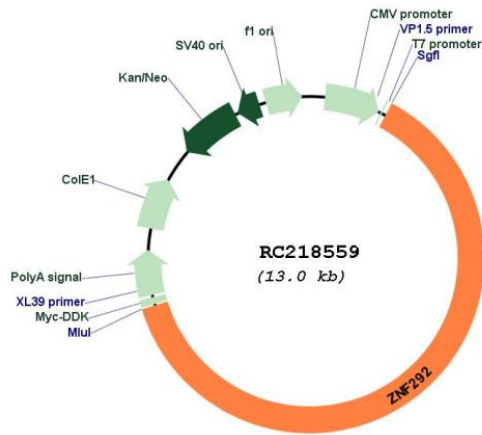
TRTRPLEQKLI SEEDLAANDILDYKDDDDKV

Restriction Sites: Sgfl-MluI

Cloning Scheme:



Plasmid Map:



ACCN:	NM_015021
ORF Size:	8169 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.
RefSeq:	NM_015021.3
RefSeq Size:	10100 bp
RefSeq ORF:	8172 bp
Locus ID:	23036
MW:	304.6 kDa
Gene Summary:	This gene encodes a growth hormone-dependent, zinc finger transcription factor that functions as a tumor suppressor. Naturally occurring mutations in this gene are associated with gastric cancer, colorectal cancer, and chronic lymphocytic leukemia. [provided by RefSeq, May 2017]