

Product datasheet for **RG207331**

ANKRD15 (KANK1) (NM_015158) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Tag:	TurboGFP
Symbol:	ANKRD15
Synonyms:	ANKRD15; CPSQ2; KANK
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)

ORF Nucleotide Sequence: >RG207331 representing NM_015158
Red=Cloning site Blue=ORF Green=Tags(s)

TTTGTGAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCCGCGATCGCC

ATGGCTCACACCACAAAGGTTAACGGCAGTGCCTCAGGAAAAGCAGGTGATATTCTCAGTGGAGACCAGG
ACAAGGAACAGAAAGACCTTACTTTGTGGAGACCCCTATGGTTATCAACTAGACTTAGATTTCTCTCAA
ATATGTGGATGACATACAGAAGGGAATACCATCAAAAGACTGAACATCCAGAAGAGCGGAAGCCGTCC
GTGCCATGCCAGAACCCAGGACCACATCTGGTCAGCAAGGTATATGGACTTCACTGAATCCCTCTCAT
CCTCCAACAGTGATGACAACAAGCAGTGCCCAACTTCTCATAGCCAGAAGTCAAGTTACATCAACTCC
AATCTCAAAGCCACCTCCCCCTCTGGAGACCTCACTCCCTTTTCTTACCATCCAGAAAATCGACAGCTG
CCACCTCCCTCACCACAACCTCCCAAGCATAACTTCATGTCACCAAGACACTGATGGAGACCCGGAGAA
GACTGGAACAGGAGAGAGCCACCATGCAGATGACACCGGTGAGTTCAGAAGGCCAGGCTGGCCAGTTT
TGGAGGCATGGGCAACCACAAGCTCCCTCCCTTCTTTGTGGTTCTGGAACCAACAAGCCTGCCAAGCAC
CAGCTTCAGAATGGATACCAAGGTAATGGGGATTATGGTAGCTATGCCCCAGCTGCTCCACCACTTCTCT
CCATGGGGAGCTCCATCCGCCACAGCCCCCTGAGCTCAGGGATCTCCACCCAGTGACCAACGTGAGCCC
CATGCACCTGCAGCACATCCGCGAGCAGATGGCCATTGCTCTGAAACGCCTGAAGGAGCTGGAGGAGCAG
GTGCGAACCATCCCTGTGCTCCAGGTAAGATCTCTGTCTTGCAAGAAGAGAAAAGGCAAGTTGGTCTCAC
AGCTGAAAAACCAAGGGCTGCATCCAGATCAATGTCTGTGGTGTGAGGAAGCGGTCTATAGTGGGG
GAACGCCTCCAGCTGGAACAGCTCTCCCGGGCCGAAGAAGTGGCGGGGAATTATACATTGACTATGAG
GAGGAAGAAATGGAGACCGTGAACAGAGCAGCAGAGGATAAAGGAGTTCCGGCACTTACAGCAGACA
TGCAAGCCCTGGAGCAGAAGATCCAGGACAGCAGCTGTGAGGCCTCCTCAGAGCTCAGGGAGAATGGAGA
GTGCCGTCTGTGGCTGTGGGTGCCGAGGAGAACATGAACGACATCGTCGTGTACACAGAGGCTCCAGG
TCCTGTAAGGATGACAGTGTAGGGACACTTGTTTCAGATGAGAAATTGTGGGGTCAGCGTGACAGAGGCCA
TGCTTGGAGTGATGACTGAAGCTGACAAAGAAATTGAGCTGCAACAGCAGACCATAGAATCCTTGAAGGA
AAAGATCTATCGCCTAGAAGTACAGCTTAGAGAAACCAACCATGACCGGGAGATGACTAAACTGAAACAA
GAGCTGCAGGCTGTGGATCGAGGAAAAAGGTTGACAAAGCCACGATGGCCAGCCGCTGTTTTCACTA



AGGTGGTGGAGGCAGTGGTGCAGACCAGAGACCAATGGTCGGCAGTCACATGGACCTGGTGGACACGTG
TGTTGGGACCTCCGTGGAACAAACAGTGTAGGCATCTCCTGCCAGCCTGAATGTAAGAATAAAGTCGTA
GGGCTGAGCTGCCTATGAATTGGTGGATTGTTAAGGAGAGGGTGGAAATGCATGACCGATGTGCTGGGA
GGTCTGTGGAAATGTGTGACAAGAGTGTGAGTGTGGAAGTCAGCGTCTGCGAAACAGGCAGCAACACAGA
GGAGTCTGTGAATGACCTCACACTCCTCAAGACAACTTGAATCTCAAAGAAGTGCAGTCTATCGTTGT
GGAGATTGTTCTGTTGACGTGACCGTCTGCTCTCAAAGGAGTGCCTCCCGGGCGTGAACACTGAGG
CTGTTAGCCAGGTGGAAGCTGCCGTATGGCAGTGCCTCGTACTGCAGACCAGGACACTAGCACAGATTT
GGAACAGGTGCACCAAGTTCACCAACACCGAGACGGCCACCCTCATAGAGTCTGCACCAACACTTGTCTA
AGCACTTTGGACAAGCAGACCAGCACCAGACTGTGGAGACGCGGACAGTAGCTGTAGGAGAAGGCCGTG
TCAAGGACATCAACTCTCCACCAAGACGCGGTCCATTGGTGTGGAACGTTGCTTTCTGGCCATTCTGG
GTTTGACAGGCCATCAGCTGTGAAGACCAAGAGTCAGGTGTGGGGCAGATAAATATTAACGACAACAT
CTGGTTGGTCTCAAAATGAGGACTATAGCTTGTGGGCCACCACAGTTGACTGTGGGGCTGACAGCCAGCA
GAAGGAGCGTGGGGTTGGGGATGACCCTGTAGGGGAATCTCTGGAGAACCCCGAGCCTCAAGCTCCACT
TGGAATGATGACTGGCCTGGATCACTACATTGAGCGTATCCAGAAGCTGCTGGCAGAACAGCAGACTG
CTGGCTGAGAACTACAGTGAAGTGGCAGAAGCTTTCGGGGAACCTCACTCACAGATGGGCTCCCTCAACT
CTCAGTCTATCAGCACCTGTGCTCTATCAACTCTGTATGAAATCTGCAAGCACTGAAGAGCTGAGGAA
CCCTGACTTCCAGAAAACAGTCTGGGTAAATCACAGGCAATTATTTGGGATATACCTGTAAGTGTGGG
GGCCTTCAGTCAGGAAGTCCCTTAAGTCCCAGACATCCAGCCTGAGCAAGAAGTGGGGACCTCAGAAG
GAAAGCCAATCAGCAGCCTGGATGCCCTCCCACTCAGGAAGGTACGCTGTCTCCAGTGAACCTGACAGA
CGACCAGATCGCCGCTGGCCTCTATGCATGTACAAACAATGAAAGTACACTGAAGTCCATCATGAAGAAG
AAAGATGGTAACAAAGATTCAAATGGCGCAAAAAAGAATCTTCAGTTTGTGGCATTAAATGGAGGGTATG
AAACAACCTCAAGTGATGATTCCAGCTCAGATGAAAGCTCTTCTCCGAGTCAGATGACGAGTGTGATGT
CATTGAGTATCCTCTTGAAGAAGAGGAGGAGGAGGATGAAGACTCGGGGAATGGCAGAAGGGCAC
CATGCAGTTAATATTGAAGTTTGAAGTCTGCCAGGGTGGAAAGATGAAATGCAGGTTCAAGAATGTGAAC
CTGAGAAGGTGGAATCAGAGAGAGGTATGAATTAAGTGAAAGATGTTGTCTGCATGCAACTTACTGAA
AAATACTATAAATGACCCAAAGCTTTGACCAGCAAAGATATGAGGTTCTGTCTGAACACCTCCAGCAC
GAGTGGTTCGCGGTGTCCAGTCAGAAGTCAGCCATTCCAGCCATGGTGGGGACTACATAGCTGCTTTTG
AGGCCATTTCCCAAGATGTCCTCCGCTATGTCATCAACTTGGCAGACGGCAACGGCAACACAGCCCTCCA
TTACAGCGTGTCCCACTCCAACCTTCGAGATTGTGAAGCTGCTGTTAGATGCCGATGTGTGAATGTGGAT
CACCAGAACAAGGCAGGCTACACCCCATCATGTTGGCGGCCCTCGCCGCTGTGGAAGCAGAGAAGGACA
TGCGGATTGTGGAAGAAGCTTTGGCTGTGGGGATGTGAATGCCAAAGCTAGTCAGGCGGGACAGACGGC
CCTCATGCTGGCGGTGAGTCACGGACGGATAGACATGGTGAAGGGCCTTCTGGCCTGTGGGGCTGATGTC
AACATCCAGGATGACGAGGGCTCCACGGCCCTCATGTGTGCCAGCGAGCACGGGCACGTGGAGATTGTCA
AGCTGCTGCTGGCCAGCCGGCTGCAACGGTCACCTAGAGGACAACGATGGCAGCACTGCGCTCTCAAT
CGCCCTGGAAGCAGGACACAAGGACATCGCTGTTCTTCTGTATGCCATGTCAACTTTGAAAAGCCAG
TCTCCGGGCACCCCTAGGCTTGAAGGAAGACGCTCTCTGGCCCCACCCACCGAGTTCAATTGAT

ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

Protein Sequence:

>RG207331 representing NM_015158

Red=Cloning site Green=Tags(s)

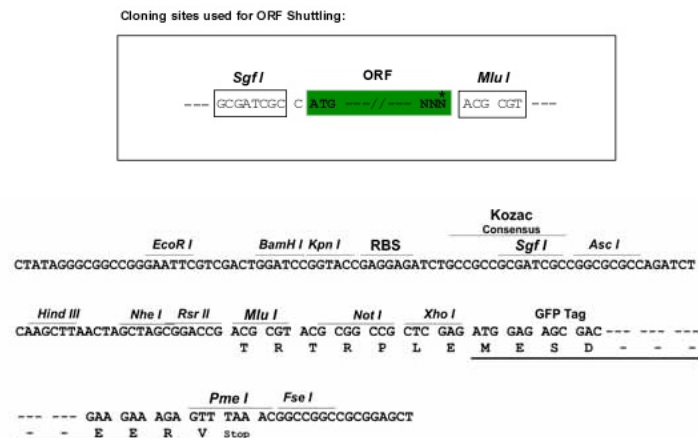
MAHTTKVNGSASGKAGDILSGDQDKEQKDPYFVETPYGYQLDLDFLKYVDDIQKNTIKRLNIQKRRKPS
VPCPEPRRTSGQQGIWSTESLSSNSDDNKQCPNFLIARSQVTSTPIKPPPPLETSLPFLTIPENRQL
PPSPQLPKHNLHVTKLTMETRRRLEQERATMQMTPGEFRRPRLASFVGGMTTSSLPSFVSGNHNKPAKH
QLQNGYQNGDYGSYAAPPTSSMGSSIRHSPLSSGISTPVTNVSPMHLQHIREQMAIALKRLKELEEQ
VRTIPVLQVKISVLQEEKRQLVSQKNQRAASQINVCGVRKRSYSAGNASQLEQLSRARRSGGELYIDYE
EEMETVEQSTQRIKEFRQLTADMQALEQKIQDSSCEASSELRENGECRSVAVGAENMNDIVVYHRGSR
SCKDAAVGTLVQMRNCGVSVTEAMLGVMTEADKEIELQQQTIESLKEKIYRLEVQLRETTTHDREMTKLKQ
ELQAAGSRKKVDKATMAQPLVFSKVVEAVVQTRDQMGVSHMDLVDTCVGTSVETNSVGISCQPECKNKVV
GPPELPMNWWIVKERVEMHDCAGRSVEMCDKSVSVEVSVCTGSNTEESVNDLTLLKTNLNLKEVRSIGC
GDCSDVTVCSPEKASRGVNTEAVSQVEAAVMAVPRTADQDSTDLEQVHQFTNTETATLIESCNTNCL
STLDKQTSTQTVETRTVAVGEGRVKDINSSTKTRSIGVGTLLSGHSGFDRPSAVKTKESGVGQININDNY
LVGLKMRTIACGPPQLTVGLTASRRSVGVGDDPVGESLENPQPAPLGMTGLDHYIERIQKLLAEQQT
LAENYSELAEAFGEPHSQMGLNSQLISTLSSINSVMKSASTEELRNPDFQKTSLGKITGNLYGYTCKCG
GLQSGSPLSSQTSQPEQEVGTSEGKPISSLDAPFTQEGTLPVNLTDQIAAGLYACTNNESTLKSIMKK
KDGKNSNGAKKNLQFVGINGGYETTSSDSSSDESSSESDDCEDVIEYPLEEEEEDEDEDTRGMAEGH
HAVNIEGLKSARVEDEMQVQECEPEKVEIRERYELSEKMLSACNLLKNTINDPKALTSKDMRFCLNTLQH
EWFVRVSSQKSAIPAMVGDIYIAFEATSPDVLRYVINLADGNGNTALHYSVSHSNFEIVKLLLDADV CNVD
HQNKAGYTPIMLAALAAVEAEKDMRIVEELFGCGDVNAKASQAGQTALMLAVSHGRIDMVKGLLACGADV
NIQDDEGSTALMCASEHGHVEIVKLLLAQPGCNGHLEDNDGSTALSI ALEAGHKDIAVLLYAHVNF AKAQ
SPGTPRLGRKTSPPGTHRG SFD

TRTRPLE - GFP Tag - V

Restriction Sites:

SgfI-MluI

Cloning Scheme:



ACCN:

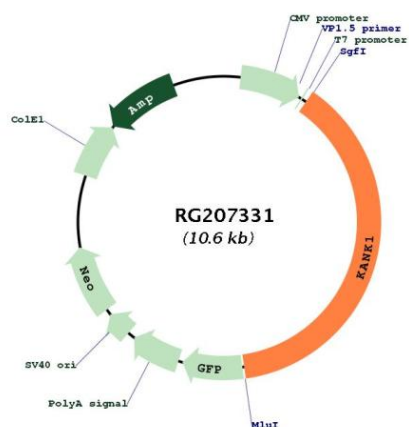
NM_015158

ORF Size:

4056 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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_015158.2 , NP_055973.2
RefSeq Size:	5083 bp
RefSeq ORF:	4059 bp
Locus ID:	23189
UniProt ID:	Q14678
Cytogenetics:	9p24.3
Gene Summary:	The protein encoded by this gene belongs to the Kank family of proteins, which contain multiple ankyrin repeat domains. This family member functions in cytoskeleton formation by regulating actin polymerization. This gene is a candidate tumor suppressor for renal cell carcinoma. Mutations in this gene cause cerebral palsy spastic quadriplegic type 2, a central nervous system development disorder. A t(5;9) translocation results in fusion of the platelet-derived growth factor receptor beta gene (PDGFRB) on chromosome 5 with this gene in a myeloproliferative neoplasm featuring severe thrombocythemia. Alternative splicing of this gene results in multiple transcript variants. A related pseudogene has been identified on chromosome 20. [provided by RefSeq, Dec 2014]

Product images:



Circular map for RG207331