

Product datasheet for SC108880

ANKRD15 (KANK1) (NM_153186) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ANKRD15 (KANK1) (NM_153186) Human Untagged Clone
Tag:	Tag Free
Symbol:	ANKRD15
Synonyms:	ANKRD15; CPSQ2; KANK
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>OriGene ORF sequence for NM_153186 edited

ATGGAGACCCGGAGAAGACTGGAACAGGAGAGAGCCACCATGCAGATGACACCGGGTGAG
 TTCAGAAGGCCAGGCTGGCCAGTTTGGAGGCATGGGCACCACAAGCTCCCTCCCTTCT
 TTTGTGGGTTCTGAAACCAACAATCCTGCCAAGCACCAGCTTCAGAATGGATACCAAGGT
 AATGGGGATTATGGTAGCTATGCCCCAGCTGCTCCCACCACTTCCTCCATGGGGAGCTCC
 ATCCGCCACAGCCCCCTGAGCTCAGGGATCTCCACCCAGTGACCAACGTGAGCCCCATG
 CACCTGCAGCACATCCGCGAGCAGATGGCCATTGCTCTGAAACGCCTGAAGGAGCTGGAG
 GAGCAGGTGCGAACCATCCCTGTGCTCCAGGTAAGATCTCTGTCTTGCAAGAAGAGAAA
 AGGCAGTTGGTCTCACAGCTGAAAAACCAAGGGCTGCATCCCAGATCAATGTCTGTGGT
 GTGAGGAAGCGGTCCTATAGTGGGGGAACGCCTCCCAGCTGGAACAGCTCTCCCGGGCC
 CGAAGAAGTGGCGGGGAATTATACATTGACTATGAGGAGGAAGAAATGGAGACCGTAGAA
 CAGAGCACGCAGAGGATAAAGGAGTTCCGGCAACTTACAGCAGACATGCAAGCCCTGGAG
 CAGAAGATCCAGGACAGCAGCTGTGAGGCCTCCTCAGAGCTCAGGGAGAATGGAGAGTGC
 CGGTCTGTGGCTGTGGGTGCCGAGGAGAACATGAACGACATCGTCGTGTACCACAGAGGC
 TCCAGGTCCTGTAAGGATGCAGCTGTAGGGACACTTGTTGAGATGAGAAAATTGTGGGGTC
 AGCGTGACAGAGGCCATGCTTGGAGTGATGACTGAAGCTGACAAAGAAATTGAGCTGCAA
 CAGCAGACCATAGAATCCTTGAAGGAAAAGATCTATCGCTAGAAGTACAGCTTAGAGAA
 ACCACCCATGACCGGGAGATGACTAAACTGAAACAAGAGCTGCAGGCTGCTGGATCGAGG
 AAAAAGGTTGACAAAGCCACGATGGCCAGCCGCTTGTTTTCAGTAAGGTGGTGGAGGCA
 GTGGTGACAGACCAGAGACCAAAATGGTCGGCAGTCACATGGACCTGGTGGACAGTGTGTT
 GGGACCTCCGTGGAACAAACAGTGTAGGCATCTCCTGCCAGCCTGAATGTAAGAATAAA
 GTCGTAGGCTGAGCTGCCTATGAATTGGTGGATTGTTAAGGAGAGGGTGGAAATGCAT
 GACCGATGTGCTGGGAGGTCTGTGGAATGTGTGACAAGAGTGTGAGTGTGGAAGTCAGC
 GTCTGCGAAACAGGCAGCAACACAGAGGAGTCTGTGAACGACCTCACACTCCTCAAGACA
 AACTTGAATCTCAAAGAAGTGGGTCTATCGTTGTGGAGATTGTTCTGTTGACGTGACC
 GTCTGCTCTCAAAGGAGTGGCCTCCCGGGGCGTGAACACTGAGGCTGTTAGCCAGGTG
 GAAGCTGCCGTATGGCAGTGCCTCGTACTGCAGACCAGGACACTAGCACAGATTGGAA


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CAGGTGCACCAAGTTCACCAACACCGAGACGGCCACCCTCATAGAGTCTGCACCAACACT
 TGTCTAAGCACTTTGGACAAGCAGACCAGCACCCAGACTGTGGAGACGCGGACAGTAGCT
 GTAGGAGAAGGCCGTGTCAAGGACATCAACTCCTCCACCAAGACGCGTCCATTGGTGTT
 GGAACGTTGCTTTCTGGCCATTCTGGGTTTGACAGGCCATCAGCTGTGAAGACCAAGAG
 TCAGGTGTGGGCGAGATAAATATTAACGACAATATCTGGTTGGTCTCAAAATGAGGACT
 ATAGCTTGTGGGCCACCACAGTTGACTGTGGGGCTGACAGCCAGCAGAAGGAGCGTGGGG
 GTTGGGATGACCCTGTAGGGGAATCTCTGGAGAACCCCAAGCTCAAGTCCACTTGGGA
 ATGATGACTGGCCTGGATCACTACATTGAGCGTATCCAGAAGCTGCTGGCAGAACAGCAG
 ACACTGCTGGCTGAGAACTACAGTGAAGTGGCAGAAGCTTTCGGGGAACCTCACTCACAG
 ATGGGCTCCCTCAACTCTCAGCTCATCAGCACCTGTCTGTCTATCAACTCTGTATGAAA
 TCTGCAAGCACTGAAGAGCTGAGGAACCCTGACTTCCAGAAAACCACTGTGGTAAAAATC
 ACAGGCAATTATTTGGGATATACCTGTAAGTGTGGGGCTTCAGTCAGGAAGTCCCTTA
 AGCTCCCAGACATCCCAGCTGAGCAAGAAGTGGGGACCTCAGAAGGAAAGCCAATCAGC
 AGCCTGGATGCCTTCCCCACTCAGGAAGGTACGCTGTCTCCAGTGAACCTGACAGACGAC
 CAGATCGCCGCTGGCCTCTATGATGTACAAACAATGAAAGTAACTGAAGTCCATCATG
 AAGAAGAAAGATGGTAACAAAGATTCAAATGGCGCAAAAAAGAAATCTTCAGTTTGTGGC
 ATTAATGGAGGGTATGAAACAACCTTCAAGTGATGATTCCAGCTCAGATGAAAGCTCTTCT
 TCCGAGTCAGATGACGAGTGTGATGTCATTGAGTATCCTCTTGAAGAAGAGGAGGAGGAG
 GAGGATGAAGACACTCGGGGAATGGCAGAAGGGCACCATGCAGTTAATATTGAAGTTTG
 AAGTCTGCCAGGGTGAAGATGAAATGCAGGTTCAAGAATGTGAACCTGAGAAGGTGGAA
 ATCAGAGAGAGGTATGAATTAAGTGAAGAGATGTTGTCTGCATGCAACTTACTGAAAAAT
 ACTATAAATGACCCCAAAGCTTTGACCAGCAAAGATATGAGGTTCTGTCTGAACACCGTC
 CAGCACGAGTGGTTCGCGGTGTCCAGTCAGAAGTCAGCCATTCAGCCATGGTGGGGGAC
 TACATAGCTGCTTTTGGAGCCATTTCAGATGTCTCCGCTATGTCATCAACTTGGCA
 GACGGCAACGGCAACACAGCCCTCCATTACAGCGTGTCCCACTCAACTTCGAGATTGTG
 AAGCTGCTGTTAGATGCCGATGTGTGTAATGTGGATCACCAGAACAAGGCAGGCTACACC
 CCCATCATGTTGGCGGCCCTCGCCGCTGTGGAAGCAGAGAAGGACATGCGGATTGTGAA
 GAACTCTCGGCTGTGGGATGTGAATGCCAAAGCTAGTCAGGCGGGACAGACGGCCCTC
 ATGCTGGCGGTCACTCACGGACGGATAGACATGGTGAAGGGCTTCTGGCCTGTGGGGCT
 GATGTCAACATCCAGGATGACGAGGGCTCCACGGCCCTCATGTGTGCCAGCGAGCAGGG
 CACGTGGAGATTGTCAAGCTGCTGCTGGCCAGCCGGCTGCAACGGTCACCTAGAGGAC
 AACGATGGCAGCACTGCGCTCTCAATCGCCCTGGAAGCAGGACACAAGGACATCGCTGTT
 CTTCTGTATGCCCATGTCAACTTTGCAAAAGGCCAGTCTCCGGGACCCCTAGGCTTGA
 AGGAAGACGTCTCCTGGCCCCACCCACCGAGGTTCAATTGATTGA

Restriction Sites:

NotI-NotI

ACCN:

NM_153186

Insert Size:

5100 bp

OTI Disclaimer:

Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

OTI Annotation:

no

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_153186.3](#), [NP_694856.1](#)

RefSeq Size: 5263 bp

RefSeq ORF: 3585 bp

Locus ID: 23189

UniProt ID: [Q14678](#)

Cytogenetics: 9p24.3

Domains: ANK

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]

Transcript Variant: This variant (2) differs in the 5' UTR, lacks a portion of the 5' coding region, and uses a downstream in-frame start codon, compared to variant 1. The resulting isoform (b, also known as Kank-S) is shorter at the N-terminus, compared to isoform a.