

Product datasheet for SC318742

KALRN (NM_001024660) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	KALRN (NM_001024660) Human Untagged Clone
Tag:	Tag Free
Symbol:	KALRN
Synonyms:	ARHGEF24; CHD5; CHDS5; DUET; DUO; HAPIP; TRAD
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>SC318742 representing NM_001024660. Blue=Insert sequence Red=Cloning site Green=Tag(s)

```
GCTCGTTT TAGTGAACCGTCAGAATTTTGT AATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG
GATCCGGTACCGAGGAGATCTGCCGCC GCGATCGCC
ATGACGGACCGCTTCTGGGACCAAGTGGTATCTCTGGTATCTCCGCTTGCTCCGGCTGCTGGATCGAGGG
TCTTTTCGGAATGATGGTTTGAAGCTTCTGATGTCCTTCTATCCTAAAGGAAAAGGTGGCCTTCGTG
TCTGGGGTCTGATAAGCGAGGCGGACCCATCCTGACCTCCCTGCTCGCAGCAATCATGACAGAATA
AGACAGGAAGACCTGCGGAACTCGTGACGATTTGGCCAGCGTGCCAAGTGAGGACGTGTGCAAACGT
GGCTTCACTGTCATCATCGACATGCGGGGCTCCAAGTGGGACCTCATCAAGCCCCCTCTCAAACCGTG
CAGGAAGCCTTTCCAGCTGAGATCCATGTGGCCCTCATCATTAAACCCGACAACCTCTGGCAGAAACAG
AAGACCAACTTTGGCAGCTCAAATTCATCTTTGAGACGAGCATGGTATCTGTGGAGGGCTCACAAAG
CTGGTGGACCCCTCCCAGCTGACGGAGGAGTTTGATGGCTCCCTGGACTACAACCATGAGGAGTGGATC
GAACTGCGGCTCTCCCTGGAGGAGTTTCTCAACAGCGCGGTGCACCTGCTCTCGCGCTCGAGGACCTC
CAGGAGATGCTAGCCCGAAGGAGTTTCTGTGGATGTGGAGGGCTCTCGCGGGCTCATTGACGAACAC
ACACAGCTCAAGAAAAGGTGCTGAAGGCCCTGTGGAGGAGCTGGACCGGGAGGGGCGAGCGGCTGCTG
CAGTGCATCCGCTGCAGCGACGGCTTCTCAGGACGCAACTGCATCCCGGGCAGTGTGACTTCCAGAGC
CTGGTGCCCAAGATCACCAGTCTCCTGGACAAGCTGCACTCCACCCGGCAGCACCTGCACCAGATGTGG
CATGTGCGCAAGCTCAAGCTGGACCAAGCTGCTTTCAGCTGCGGCTCTTCGAGCAGGATGCTGAGAAGATG
TTTGACTGGATAAGCCACAACAAGGAGTTATTCTCCAGAGCCACACGGAGATCGGAGTACGATACCAG
TACGCCCTTGACCTCCAGACGCAGCACAATCACTTTGCCATGAACTCCATGAATGCCTATGTCAACATC
AACCAGCATCATGTCGTTGCCCTCCCGCTCTCTGAGGCCGGTTCATTATGCCTCACAACAAATCAAGCAG
ATCTCCACCCAGCTGGACCAGGAGTGGAAGAGCTTCGCTGCTGCCCTGGATGAACGCAGCACCATCCTC
GCCATGTCTGCTGTGTTCCACCAGAAGGCTGAGCAGTTCTGTGCGGAGTGGATGCCTGGTGCAAGATG
TGCAGTGAAGGTGGTCTGCCATCCGAGATGCAAGACCTAGAGCTGGCAATCCACCACCACCAGACCTTG
TATGAGCAGGTGACCCAAGCCTACACAGAGGTGAGCCAGGATGGCAAAGCACTACTTGATGTGCTGCAG
CGGCCCTGAGCCCTGGAACTCCGAATCCCTCACGGCCACAGCAACTACTCCAAGGCAGTGCACCAG
```



[View online »](#)

GTGCTGGACGTGGTGCATGAGGTGTTACATCACCAGCGACGGCTGGAGAGCATCTGGCAGCACCGCAAG
 GTGCGGCTCCACCAGCGGCTGCAGCTCTGCGTCTTCCAGCAGGATGTACAGCAGGTGTTGGACTGGATT
 GAAAACCATGGTGAAGCCCTTCTCAGCAAAACACTGGAGTTGGGAAGTCCCTACATCGAGCCCGGGCC
 CTGCAGAAGAGGCATGATGACTTTGAAGAGGTGGCTCAGAATACGTACACCAATGCGGACAAGCTCCTA
 GAAGCAGCAGAGCAGTTGGCTCAGACGGGGAAATGTGACCCCGAGGAGATCTACAAGGCAGCTCGACAC
 CTGGAGGTGCGCATCAAAGACTTCGTGCCAGGGTGGAGCAGCGGAAGCTTCTCTGGACATGTCTGTT
 TCCTTCCACACACACCAAGAGTTGTGGACATGGATGGAAGACCTTCAAGAGGATGTTGGAGGAT
 GTCTGTGCAGATTCTGTGGATGCAGTCCAGGAAGTCAAGCAGTTCCAGCAGCAGCAGACCGCCACT
 CTAGATGCCACACTCAATGTCATCAAGGAAGGCGAAGACCTTATCCAGCAGCTCAGGTACAGCCTCCC
 TCCCTCGGGGAGCCAGCGAGGCCAGGGACTCGGCTGTGTCCAACAACAAAAACCCACAGCAGCTCC
 ATCAGCCACATCGAGTCGGTCTGCAGCAGCTTGTATGATGCCAGGTGCAGATGGAGGAGCTGTTCCAC
 GAGCGGAAGTCAAGCTGGACATCTTCTGCAACTGCGCATCTTTGAGCAGTACACCATCGAGGTGACA
 GCAGAGCTAGACGCTGGAATGAAGACTTGCTTCGGCAGATGAATGACTTCAACACAGAGGACCTAACC
 CTGGCAGAACAGCGGCTGCAGCGCCACACAGAACGGAAGCTAGCCATGAACAACATGACCTTTGAGGTT
 ATCCAGCAGGGACAGGATCTGCACCAAGTACATCACGGAGGTCCAGGCATCAGGAATTGAGTTGATCTGT
 GAAAAAGACATTGATCTGGCAGCCAGGTGCAAGAGTTATTGGAATTTCTCCATGAGAAGCAGCATGAA
 TTGGAGCTCAATGCAGAGCAGACTCATAAGCGGCTAGAGCAGTGCCTCCAATTACGTCACCTCCAGGCT
 GAAGTCAAACAGGTTCTGGGATGGATCCGCAATGGAGAGTCAATGCTCAACGCCAGCCTGGTCAATGCC
 AGCTCTTTGTGCGAAGCAGAGCAGCTGCAGCGGGAGCAGCAGCAGTTCCTCAACTGGCCATCGAGTCCCTC
 TTTTATGCCACTTCTTTCAGAGAAGCAGCAGGAGTGCCTGCAGGTACAGCAGAAAAGCCGAGGTGCTG
 CTCCAGGCCGGCCACTACGATGCCGATGCCATCCGGGAATGTGCTGAGAAGGTGGCCCTCCACTGGCAG
 CAGCTCATGCTGAAGATGGAAGACCGGCTAAAAATTGGTCAATGCCTCTGTGGCCTTTTAAAAAAGTTCT
 GAACAGGTGTGTAGTGTCTGGAGAGCTTAGAGCAAGAATACCGGAGAGATGAGGACTGTGTGGTGA
 CGAGATAAGCTGGGGCCAGCAGCAGAGATCGACCATGTCAATCCCTCCTCAGCAAAACATTTGGAACAA
 AAGGAGGCCTTTCTTAAGCCTGCACCCTGGCTCGGCGGAATGCTGAGGTGTTTCTCAAGTACATCCAC
 AGGAACAACGTCAGCATGCCAGTGTGCCAGCCACACTCGGGGACCCGAGCAACAAGTAAAGCCATC
 CTGAGTGAGCTCCTGCAGAGGGAGAATCGCGTGTGCATTTCTGGACCTTGAAGAAGCGGCGTTAGAC
 CAATGCCAGCAATATGTGGTGTTCGAGCGCAGCGCTAAGCAGGCGCTTGACTGGATCCAAGAAACAGGT
 GAATTTTACCTCTCAACACATACCTCCACTGGAGAGACCACAGAGGAGACTCAGGAAGTGTGAAAGAA
 TATGGGAATTCAGGGTGCCTGCCAAGCAAAACAAAGGAGAAGGTGAAGCTTCTGATTGAGTGGCCGAT
 AGCTTTGTGAAAAAGGCCACATTCATGCCACGGAGATAAGGAAATGGGTGACCACGGTGGACAAGCAC
 TACAGAGATTTCTCCCTGAGGATGGGAAAGTACCGATACTCACTGGAGAAAGCCCTAGGAGTCAACACA
 GAGGATAATAAGGACCTGGAGCTGGATATTATCCAGCAAGCCTTTCCGGATCGGGAGGTCAAGTCCGG
 GACGCCAACACGAAGTCAATGAAGAGAAGCGGAAGTCAAGCCCGAAGAAAGAAATTTATTATGGCTGAA
 CTACTCCAGACAGAGAAGGCTTATGTAAGGGATTTGCATGAGTGCCTTAGAGACCTACCTGTGGGAAATG
 ACCAGTGGTGTGGAGGAGATCCCCCTGGGATCCTCAATAAAGAGCATATCATCTTTGGCAACATCCAA
 GAGATCTACGATTTCCATAACAACATCTTCTCAAAGAGCTGGAGAAGTACGAGCAACTGCCTGAGGAT
 GTGGGCACTGCTTTGTTACCTGGGCAGACAAATTTAGATGTATGTACCTACTGTAAAAACAAGCCT
 GATTTCAACCAGCTTATCCTGGAGCATGCGGGCACCTTTTGTGAGATACAACAGCGGCATGGTCTG
 GCCAACTCCATCTTCTACCTAATTAAGCCTGTCAAAGGATACCAAAATATCAACTGCTCCTGAAG
 GAACCTTTAAGTGTGTGAAGAAGGAAAGGGGAGCTCAAGGATGGCCTGGAGGTGATGCTCAGTGTC
 CCAAAGAAAGCCAATGATGCCATGCATGTGAGCATGCTGGAAGGGTTCGACGAGAACCTGGATGTGCAG
 GGGGAGTTGATTCTCAGGATGCCTTTCAAGTGTGGGACCCGAAGTCGCTGATCCGGAAGGGGCGGGAG
 CGGCACCTGTTCTCTTTGAGATCTCCTTGTTTTAGCAAGGAGATCAAAGATTCTCAGGACACAG
 AAATATGTTTACAAGAACAAGCTACTGACCTCAGAGCTGGGTGTGACCGAGCACGTGGAGGGCGATCCC
 TGCAAATTCGCTTGTGGTCTGGGCGCACCCATCCTCAGACAATAAAACAGTGTGAAAGCCTCCAAC
 ATTGAAACCAAGCAGGAGTGGATCAAGAACATTCGAGAAGTATTCAAGAAAGGATCATTACCTGAAA
 GGAGCTTTAAAGGAGCCACTTCAGCTCCCCAAAACACCAGCCAAACAGAGGAACAATAGTAAGAGGGAT
 GGAGTGGAGGATATTGACAGCCAGGGGGATGGGAGCAGCCAACCAGACACCATCTCCATTGCTTCTAGG
 ACCTCTCAGAACACAGTGGACAGTGACAAGCTCTCTGGTGGATGTGAGCTGACAGTGGTCTCCAGGAC
 TTCAGTGGCGGCCACAGCAGTGAAGTACCATCCAGGTGGGGCAGACGGTAGAGCTGCTGGAGCGGCC
 AGCGAGCGCCTGGTGGTGTCTGGTCCGTACCACCGAACGGAGCCCGCCTTGGAGGTCTGGTCCCC

AGCAGCGCCCTGTGCATCTCACACTCCCGAAGCAGCGTGGAGATGGACTGCTTCTCCCTTGGTGAAA
 GATGCATACTCTCATTCCCTCAAGCGAGAATGGAGGCAAGTCCGAGTCCGTGGCCAACCTGCAGGCCAG
 CCCTCCCTGAACTCCATCCACAGTTCCTCCGGGTCCCAAGCGCTCCACCAACTCTTAAGAAGTGGCTG
 ACGAGTCTGTGCGTCCGCTTAACACGGGGAAGGCAGATGGAAACATCAAAAAGCAGAAGAAAGTTCGC
 GATGGTCGGAAGAGCTTTGACCTGGGATCTCCCAAGCCTGGGGATGAAACAACCCCTCAGGGAGACAGC
 GCTGATGAGAAGCAAGAAAGGTTGGGGTGAAGATGAGCCGGATGAAGAGTACACACACCCCTCCCA
 CCACCTATGAAGATTTTTGACAACGACCCTACACAGGATGAAATGTCTCTCTTTGCTAGCAGCCCGG
 CAGGCTTCCACTGAAGTACCTACTGCTGCAGACCTTGCAATGCAATAGAAAAGTTGGTCAAAAACAAG
 CTGAGTCTAGAAGGAAGCTCATACCGGGGAGCTTGAAGACCCTGCAGGCTGCCTGAATGAGGGATG
 GCCCACCCACACCTCTAAAAACCAGAAGAAGAACAGAAAAGCCAAGGCCCTGAGAGGCAGGATGTTT
 GTCCTGAATGAGCTGGTACAGACAGAGAAAGACTATGTCAAGGATCTGGGCATTGTGGTGGAGGGCTTC
 ATGAAGAGAATAGAAGAAAAGGGTGTCCCTGAGGATATGCGAGGAAAGGACAAAACTGTTTGGAAAT
 ATTCATCAGATTTATGACTGGCATAAGGATTTTTCTGGCGAACTGGAAAAGTGTATCCAGGAGCAA
 GACAGATTGGCACAGCTTTTAAAGCACGAGCGGAAGCTGCACATCTACGTGTGGTATTGTCAGAA
 AAGCCCGCTCAGAGTACATCGTTGCTGAGTATGACGCCTACTTTGAGGAGGTAAAACAGGAGATAAT
 CAGAGGCTGACACTGAGTGACTTCTCATCAAGCCCATTCAGAGAATAACAAAATACCAGTTGCTCCTC
 AAGGACTTCTGAGATACAGTGAGAAGGCTGGTTTGGAGTGTTCAGATATTGAGAAAAGCAGTGGAGTTA
 ATGTGCCTTGTCCCAAACGCTGCAATGACATGATGAATCTAGGACGCTGCAGGGCTTTGAGGGCACT
 CTGACTGCTCAGGGGAAGCTGCTGCAGCAGGACACATTCTATGTGATCGAGCTGGATGCAGGCATGCAG
 TCCCGGACCAAAGAGAGGGCGCTGTTCTCTTCGAGCAGATTGTCATCTTCAGTGAATGCTCAGGAAG
 GGATCCCTCACCCCTGGTACATGTTCAAAGGAGCATCAAGTGAATTACTTGGTCTGGAGGAGAA
 GTGGACAATGATCCCTGCAAGTTTGCACTCATGAACAGAGAGACTTCTGAGAGGGTTGTTCTGCAAGCC
 CCAACGCTGACATCCAGCAGGCTGGGTGCAGGACATCAATCAAGTCTTAGAAAACACAGCGAGACTTT
 TTGAATGCACTGCAATCGCCATTGAGTATCAACGGAAAAGAAAGGAGCACAGCTGTGATGAGGTTCAA
 CCTGCCAGGCTTCCCAAGCCAGCCCAAGGCCCTACTCCTCTGTTCTGCGGGCTCAGAGAAGCCCCCA
 AAGGGCTCCAGCTATAACCCACCTCTGCCTCCCTGAAGATATCTACCTCCAATGGCAGTCCAGGGTTT
 GAATACCACCAGCCTGGGGACAAGTTCGAAGCCAGCAAGCAGAACGACCTGGGAGGCTGCAATGGGACC
 TCGTCCATGGCCGTGATCAAAGATTACTATGCACTGAAGGAGAATGAAATCTGTGTGAGCCAAGGTGAG
 GTGGTCCAGTCTCGCCGTCAACCAGCAGAACATGTGTCTGGTGTACCAGCCTGCCAGCGACCATTCC
 CCCGCCCGGAGGCTGGGTCCCAGGCAGCATCCTGGCGCCCTCACAAAGCCACAGCAGCAGAAAAGT
 AGTGACGGGAGCATCAAGAAGTCATGTTTATGGCATACTCTACGCATGAGAAAAGCGGGCGGAAGTGGAG
 AACACGGTAAAAATGAAGCCACAGGGCTCGTAAACCCAAGGATATTCTGGGCAACAAAGTCTCTGTT
 AAAGAGACGAACAGTTCGAGGAATCAGAGTGTGATGATCTTGACCCATAACTAGCATGGAGATCTTA
 AATCCAATTTTCAAGAAAGTGGCCCCAGAATTCTTGTGCCCTTGGTGGATGTGACCTGCTTGTCTT
 GGGGACACAGTGATACTGCAGTGCAAAGTCTGTGGGCGGCCAAAGCCACCATCACTTGGAAAGGTCCA
 GACCAGAACATCCTTGACACTGATAACAGCTCAGCCACATACACGGTCTCCTCTTGTGATTCTGGAGAA
 ATCACCCTGAAGATCTGTAATCTGATGCCCCAAGCAGTGGGATTTATACCTGCATAGCAACAAATGAC
 CACGGGACCACATCAACGTCTGCAACAGTCAAAGTGAAGGTGTTCCAGCAGCCCTAACCGCCCCATT
 GCCCAGGAGAGAAGCTGCACCTCCGTGATTCTCCGCTGGCTGCCCCCTCCAGCACAGAAAAGTGCAC
 ATTTCTGGTTACACTGTGGAGTACAGAGAGGAAGGTTCTCAGATCTGGCAGCAGTCAAGTGGCTTCGACC
 TTGGACACTTACCTCGTCATCGAAGACCTTAGTCCCGGTGTCCTTATCAGTTCAGAGTCAAGTGGCAGT
 AACCCCTGGGGAATCAGCCTTCCAGCAGGACCCTCGGAGTTTGTGCGACTTCCAGAATATGATGCTGCT
 GCTGATGGTGCCACCATTTCTTGAAGGAAAAATTTGACTCAGCTTACACTGAGCTGAATGAAATTTGGA
 AGAGGCCGTTTCTCTATAGTAAAGAAATGCATTCAAAAGCTACCCGCAAAGATGTGGCTGTGAAATTT
 GTTAGCAAAAAATGAAGAAGAAAGAACAGGCTGCCACGAGGCTGCCCTGCTTACAGCACCTACAGCAC
 CCCCAGTACATCACTCTCCATGACACCTATGAGTCCCCACATCCTACATCCTGATCTTGAAGTATGATG
 GATGATGGCCGGCTTTAGACTACCTTATGAATCATGATGAACTGATGGAGGAAAAAGTAGCTTTCTAT
 ATCCGAGACATCATGGAGGCTCTGCAGTACCTTCAAACTGCAGGTTGCACATTTGGACATAAAGCCT
 GAAAACCTGCTCATTGACCTACGGATTCCAGTGCCTCGAGTGAAGCTCATTGACTTGGAGGATGCTGTC
 CAGATCTCGGGTCACTTCCACATTCACCACCTGCTGGGGAACCCTGAGTTTGTGCCCCAGAAGTCAAT
 CAAGGCATCCCCGTCTCCTGGGGACAGACATCTGGAGCATCGGGGTTCTGACATATGTCATGCTGAGT
 GGGGCTCCCCCTTCTGGATGAGAGCAAAGAGGAGACATGTATCAACGTATGCAGGTTGGATTTTCAGC

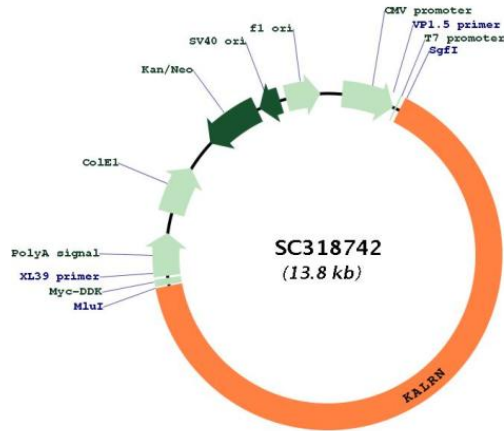
```

TCCCCCATGAATACTTCTGTGGTGTGAGCAATGCTGCCAGAGATTTTCATCAATGTGATCTTACAGGAA
GATTTTCGGAGGCGGCCACAGCAGCCACATGCTTGCAGCATCCATGGCTGCAGCCCATAATGGCAGC
TACTCTAAGATCCCCCTGGACACCTCCCGCCTAGCATGCTTCATAGAACGTGCAAGCACCAGAATGAT
GTGCGGCTATTCCCAATGTCAAGAGCTACATTGTCAACCGGGTGAACCAAGGGACGTAG
ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT
TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
    
```

Restriction Sites:

Sgfl-MluI

Plasmid Map:



ACCN:

NM_001024660

Insert Size:

8961 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:

This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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_001024660.4](#)

RefSeq Size:	15982 bp
RefSeq ORF:	8961 bp
Locus ID:	8997
UniProt ID:	O60229
Cytogenetics:	3q21.1-q21.2
Protein Families:	Druggable Genome, Protein Kinase
MW:	340.3 kDa
Gene Summary:	<p>Huntington's disease (HD), a neurodegenerative disorder characterized by loss of striatal neurons, is caused by an expansion of a polyglutamine tract in the HD protein huntingtin. This gene encodes a protein that interacts with the huntingtin-associated protein 1, which is a huntingtin binding protein that may function in vesicle trafficking. [provided by RefSeq, Apr 2016]</p> <p>Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1). There are no publicly available transcripts representing the full-length splicing pattern of this variant; it is inferred from data in PMID:11891045 and from the rat homolog, NM_032062.1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.</p>