

Product datasheet for **SC204469**

Nectin 2 (NECTIN2) (NM_002856) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: Nectin 2 (NECTIN2) (NM_002856) Human 3' UTR Clone
Vector: pMirTarget (PS100062)
Symbol: NECTIN2
Synonyms: CD112; HVEB; PRR2; PVRL2; PVRR2
ACCN: NM_002856
Insert Size: 348 bp
Insert Sequence: >SC204469 3'UTR clone of NM_002856
The sequence shown below is from the reference sequence of NM_002856. The complete sequence of this clone may contain minor differences, such as SNPs.
Blue=Stop Codon **Red**=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTCATCTCACGGCGGCAGTTTATGTGTGACCTGGACACAGACAGACAGACAGCCAGGCCCGGCCCTCC
CGCCCCGACCTGACCACGCCGGCCTAGGGTCCAGACTGGTTGGACTTGTTCGTCTGGACGACACTGG
AGTGGAACACTGCCTCCCACCTTTCTTGGGACTTGGAGGGAGGTGGAACAGCACACTGGACTTCTCCCGT
CTCTAGGGCTGCATGGGAGCCCCGGGAGCTGAGTAGTGGGGATCCAGAGAGGACCCCGCCCCAGAG
ACTTGGTTTTGGCTCCAGCCTTCCCCTGGCCCCGTGACACTCAGGAGTTAATAAATGCCTTGGAGGAAA
ACA
ACGCGTAAGCGGCCGCGGCATCTAGATTCAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
```

Restriction Sites: Sgfl-MluI

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.

RefSeq: [NM_002856.3](#)



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Summary:

This gene encodes a single-pass type I membrane glycoprotein with two Ig-like C2-type domains and an Ig-like V-type domain. This protein is one of the plasma membrane components of adherens junctions. It also serves as an entry for certain mutant strains of herpes simplex virus and pseudorabies virus, and it is involved in cell to cell spreading of these viruses. Variations in this gene have been associated with differences in the severity of multiple sclerosis. Alternate transcriptional splice variants, encoding different isoforms, have been characterized. [provided by RefSeq, Jul 2008]

Locus ID:

5819

MW:

12.9