

Product datasheet for SC310637

BCL2L2 (NM_004050) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: BCL2L2 (NM_004050) Human Untagged Clone

Tag: Tag Free Symbol: BCL2L2

Synonyms: BCL-W; BCL2-L-2; BCLW; PPP1R51

Mammalian Cell None

Selection:

Vector: pCMV6-XL5

E. coli Selection: Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM_004050 edited

GGCCCCAGCTGGGGGCTTTCATCCGCCTGACCCGGCTCCACGCTGGCCTTTCATCCTCCT GGCCAGCGTGGAGCCGGGTCAGGCCGGGAGGATGAAAGGCCCCAGCTGGGGGCTCCTTGC CACCAGTGCTGTCTTAAGAGCTGCCATCCCGGCTGGCCGCCCGGATGGCGACCCCAGC CTCGGCCCCAGACACACGGGCTCTGGTGGCAGACTTTGTAGGTTATAAGCTGAGGCAGAA GGGTTATGTCTGTGGAGCTGGCCCCGGGGAGGGCCCAGCAGCTGACCCACTGCACCAAGC CATGCGGGCAGCTGGAGATGAGTTCGAGACCCGCTTCCGGCGCACCTTCTCTGATCTGGC GGCTCAGCTGCATGTGACCCCAGGCTCAGCCCAACAACGCTTCACCCAGGTCTCCGATGA ACTTTTCAAGGGGCCCCAACTGGGGCCGCCTTGTAGCCTTCTTTGTCTTTGGGGCTGC ACTGTGTGCTGAGAGTGTCAACAAGGAGATGGAACCACTGGTGGGACAAGTGCAGGAGTG GATGGTGGCCTACCTGGAGACGCGGCTGGCTGACTGGATCCACAGCAGTGGGGGCTGGGC GGAGTTCACAGCTCTATACGGGGACGGGGCCCTGGAGGAGGCGCGCGTCTGCGGGAGGG GAACTGGGCATCAGTGAGGACAGTGCTGACGGGGGCCCTGGCACTGGGGGCCCTGGTAAC TGTAGGGGCCTTTTTTGCTAGCAAGTGAAAGTCCAGGGCCAGGTGGGGCTAGGTGTGGCT GGGGGCCAGGAGCAGGAACAGAACAGAGAAATGCCCTTGGAAGAAGTGGAGTTGGTGG ATGGGTGGGCATGGAACAGGATGGGCAGAGAAAGGGTAGTGTGAGGGAGCTGAGTAGG CCAGGTAGGCGATTGGAAGAGTGAGCAGGACACAGAGGGGAGGGGAATGTTTTGGCAAGT TAGGTGTGGGCACATGAAACGACCTGGAACTTGCTTCACAGCCCTGAGGAAGGTGGACTT ACATAAGCAGCTGTATTCCATTAGATGAGTGGGATTTAGGGAACGCAGAAGGCACATCCC TTTGGAATGGAAGCTTAGGGGTTCTCAGGTGATAGGGAGAGGTGGCTGTTAACAGTGGGC TGCTTGGACACGCGTGTGCATGTGCACGCATGCTGGTGTGCATGCTGGGCTGCCTGGCAA ATCTGGTGGTGGTGGGATTCCTCAAGGAGAAAACATTCCCTCTTGCAATGGCAAGAACTA GGGGCAGTTCTCTGTCCCTCCCAACCCCTCCTTTCCCCTGCCCTTGTCCTGATGCCT

Restriction Sites: Please inquire



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BCL2L2 (NM_004050) Human Untagged Clone - SC310637

ACCN: NM_004050

Insert Size: 1400 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: The ORF of this clone has been fully sequenced and found to be a perfect match to

NM_004050.2.

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: <u>NM 004050.2</u>, <u>NP 004041.1</u>

 RefSeq Size:
 3542 bp

 RefSeq ORF:
 582 bp

 Locus ID:
 599

 UniProt ID:
 Q92843

Cytogenetics: 14q11.2

Domains: Bcl-2, BH4

Protein Families: Druggable Genome, Transmembrane



Gene Summary:

This gene encodes a member of the BCL-2 protein family. The proteins of this family form hetero- or homodimers and act as anti- and pro-apoptotic regulators. Expression of this gene in cells has been shown to contribute to reduced cell apoptosis under cytotoxic conditions. Studies of the related gene in mice indicated a role in the survival of NGF- and BDNF-dependent neurons. Mutation and knockout studies of the mouse gene demonstrated an essential role in adult spermatogenesis. Alternative splicing results in multiple transcript variants. Read-through transcription also exists between this gene and the neighboring downstream PABPN1 (poly(A) binding protein, nuclear 1) gene. [provided by RefSeq, Dec 2010]

Transcript Variant: This variant (1) represents the longer transcript. Both variants 1 and 2 encode the same protein. 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.