

Product datasheet for **SC329420**

CORO2B (NM_001190456) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: CORO2B (NM_001190456) Human Untagged Clone
Tag: Tag Free
Symbol: CORO2B
Synonyms: CLIPINC
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC329420 representing NM_001190456.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

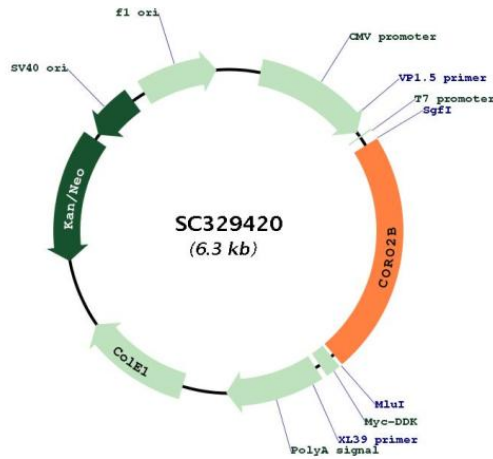
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CTCCAGCTGGAAGTAAAAAAGTTCGCAACAGCCCAAGAAGTGTAG
  
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Restriction Sites: SgfI-MluI



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Plasmid Map:


ACCN: NM_001190456

Insert Size: 1428 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).

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_001190456.1](#)

RefSeq Size: 3344 bp

RefSeq ORF: 1428 bp

Locus ID: 10391

UniProt ID: [Q9UQ03](#)

Cytogenetics: 15q23

MW: 54.4 kDa

Gene Summary: May play a role in the reorganization of neuronal actin structure.[UniProtKB/Swiss-Prot Function]

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 (2) is shorter at the N-terminus, compared to isoform 1. Variants 2 and 3 encode the same isoform. 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.