

Product datasheet for SC329847

ALS2CR2 (STRADB) (NM 001206864) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: ALS2CR2 (STRADB) (NM_001206864) Human Untagged Clone

Tag: Tag Free Symbol: **STRADB**

Synonyms: ALS2CR2; CALS-21; ILPIP; ILPIPA; PAPK; PRO1038

Vector: pCMV6-Entry (PS100001)

>SC329847 representing NM_001206864. **Fully Sequenced ORF:**

Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGTCTCTTTTGGATTGCTTCTGCACTTCAAGAACACAAGTTGAATCACTCAGACCTGAAAAACAGTCT GAAACCAGTATCCATCAATACTTGGTTGATGAGCCAACCCTTTCCTGGTCACGTCCACTAGAGCC AGTGAAGTACTATGTTCCACCAACGTTTCTCACTATGAGCTCCAAGTAGAAATAGGAAGAGGATTTGAC AACTTGACTTCTGTCCATCTTGCACGGCATACTCCCACAGGAACACTGGTAACTATAAAAAATTACAAAT CTGGAAAACTGCAATGAAGAACGCCTGAAAGCTTTACAGAAAGCCGTGATTCTATCCCACTTTTTCCGG CATCCCAATATTACAACTTATTGGACAGTTTTCACTGTTGGCAGCTGGCTTTGGGTTATTTCTCCATTT ATGGCCTATGGTTCAGCAAGTCAACTCTTGAGGACCTATTTTCCTGAAGGAATGAGTGAAACTTTAATA AGAAACATTCTCTTTGGAGCCGTGAGAGGGTTGAACTATCTGCACCAAAATGGCTGTATTCACAGGAGT ATTAAAGCCAGCCATATCCTCATTTCTGGTGATGGCCTAGTGACCCTCTCTGGCCTGTCCCATCTGCAT AGTTTGGTTAAGCATGGACAGAGGCATAGGGCTGTGTATGATTTCCCACAGTTCAGCACATCAGTGCAG CCGTGGCTGAGTCCAGAACTACTGAGACAGGATTTACATGGGTATAATGTGAAGTCAGATATTTACAGT GTTGGGATTACAGCATGTGAATTAGCCAGTGGGCAGGTGCCTTTCCAGGACATGCATAGAACTCAGATG CTGTTACAGAAACTGAAAGGTCCTCCTTATAGCCCATTGGATATCAGTATTTTCCCTCAATCAGAATCC AGAATGAAAAATTCCCAGTCAGGTGTAGACTCTGGGATTGGAGAAAGTGTGCTTGTCTCCAGTGGAACT CACACAGTAAATAGTGACCGATTACACACACCATCCTCAAAAAACTTTCTCTCCTGCCTTCTTTAGCTTG

TTCAAACAGCCTTATTTTGAGTTTCTTTAA

Restriction Sites: Sgfl-Mlul



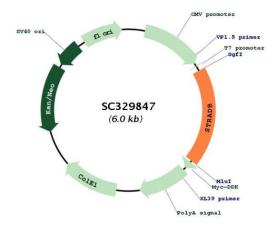
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Plasmid Map:



ACCN: NM_001206864

Insert Size: 1134 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: <u>NM 001206864.1</u>

RefSeq Size: 2328 bp



ALS2CR2 (STRADB) (NM_001206864) Human Untagged Clone - SC329847

 RefSeq ORF:
 1134 bp

 Locus ID:
 55437

 UniProt ID:
 Q9C0K7

 Cytogenetics:
 2q33.1

Protein Families: Druggable Genome, Protein Kinase

MW: 42.4 kDa

Gene Summary: This gene encodes a protein that belongs to the serine/threonine protein kinase STE20

subfamily. One of the active site residues in the protein kinase domain of this protein is altered, and it is thus a pseudokinase. This protein is a component of a complex involved in the activation of serine/threonine kinase 11, a master kinase that regulates cell polarity and energy-generating metabolism. This complex regulates the relocation of this kinase from the nucleus to the cytoplasm, and it is essential for G1 cell cycle arrest mediated by this kinase. The protein encoded by this gene can also interact with the X chromosome-linked inhibitor of apoptosis protein, and this interaction enhances the anti-apoptotic activity of this protein via the JNK1 signal transduction pathway. Two pseudogenes, located on chromosomes 1 and 7, have been found for this gene. Alternatively spliced transcript variants encoding different

isoforms have been found for this gene. [provided by RefSeq, May 2011]

Transcript Variant: This variant (2) has an alternate splice site in the 3' coding region, resulting in a longer transcript and a reading frame-shift, compared to variant 1. The resulting isoform

(2) has a shorter and distinct C-terminus, compared to isoform 1.