

Product datasheet for SC204706

SH2D2A (NM_003975) Human 3' UTR Clone

Product data:

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| Product Type: | 3' UTR Clones |
| Product Name: | SH2D2A (NM_003975) Human 3' UTR Clone |
| Symbol: | SH2D2A |
| Synonyms: | F2771; SCAP; TSAD; VRAP |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pMirTarget (PS100062) |
| ACCN: | NM_003975 |
| Insert Size: | 364 bp |
| Insert Sequence: | >SC204706 3'UTR clone of NM_003975 The sequence shown below is from the reference sequence of NM_003975. 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
GCATGGCTTCCCCTTGGCCTCCTCAGTAGGCGGTCTGGCCTGACCCCCAACAAAGAAGCCTGGAGGTC
AGAGAAGCAAATGCGGAGCCTGCTCCCTCCTAAGAAGATCCCAAGAATCCAATGGCTCAGTCCTTGGTG
ATCTAAGACAGCAAAGAAGTGCAAGGAGGGCCCTGTTAGCTCCCACTGTCCTGGTTTCTCCTCCTGG
AGTCTAATTTCTTGGCCCTCTGAGCCTTTTGAGTCTGGGCCCTGGTCCAATGCTGCTGTTGTCTGAGG
AATGGTTTGGTGAGAACAGATGTTAGAACTTGTTGTTGATTCTGTCTGGCTAATAAATCATCACCAA
CTGCCTTCTCCTACAGGGA
ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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| 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. |



[View online »](#)

RefSeq: [NM_003975.4](#)

Summary: This gene encodes an adaptor protein thought to function in T-cell signal transduction. A related protein in mouse is responsible for the activation of lymphocyte-specific protein-tyrosine kinase and functions in downstream signaling. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2010]

Locus ID: 9047

MW: 13.3