

Product datasheet for SC321392

OriGene Technologies, Inc.

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SM22 alpha (TAGLN) (NM_001001522) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: SM22 alpha (TAGLN) (NM_001001522) Human Untagged Clone

Tag: Tag Free

Symbol: SM22 alpha

Synonyms: SM22; SM22-alpha; SMCC; TAGLN1; WS3-10

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-AC (PS100020)E. coli Selection:Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM_001001522.1

CAAGTCTTCACTCCTTCCTGCGAGCCCTGAGGAAGCCTTGTGAGTGCATTGGCTGGGGCT TGGGGGCCCAAGCTCCAGGCAGGGTGGGCTGGATCACTAGCGTCCTGGATCTCTCAGA CTGGGCAGCCCGGGCTCATTGAAATGCCCCGGATGACTTGGCTAGTGCAGAGGAATTGA CCGCGAAGTGCAGTCCAAAATCGAGAAGAAGTATGACGAGGAGCTGGAGGAGCGGCTGGT GGAGTGGATCATAGTGCAGTGTGGCCCTGATGTGGGCCGCCCAGACCGTGGGCGCTTGGG CTTCCAGGTCTGGCTGAAGAATGGCGTGATTCTGAGCAAGCTGGTGAACAGCCTGTACCC GGAGCAGGTGGCTCAGTTCCTGAAGGCGGCTGAGGACTATGGGGTCATCAAGACTGACAT GTTCCAGACTGTTGACCTCTTTGAAGGCAAAGACATGGCAGCAGTGCAGAGGACCCTGAT GGCTTTGGGCAGCTTGGCAGTGACCAAGAATGATGGGCACTACCGTGGAGATCCCAACTG GTTTATGAAGAAAGCGCAGGAGCATAAGAGGGAATTCACAGAGAGCCAGCTGCAGGAGGG AAAGCATGTCATTGGCCTTCAGATGGGCAGCAACAGAGGGGCCTCCCAGGCCGGCATGAC AGGCTACGGACGACCTCGGCAGATCATCAGTTAGAGCGGAGAGGGCTAGCCCTGAGCCCG TGGTACCTTCAGCCCTGGCCAAGCTTTGAGGCTCTGTCACTGAGCAATGGTAACTGCACC TGGGCAGCTCCTCCTGTGCCCCCAGCCTCAGCCCAACTTCTTACCCGAAAGCATCACTG AGGGGAGAAGCGGGCTGGGGTAGCCTGGATGTGGGCCAAGTCCACTGTCCTCCTTGGCG GCAAAAGCCCATTGAAGAAGAACCAGCCCAGCCTGCCCCCTATCTTGTCCTGGAATATTT



Restriction Sites: Please inquire

ACCN: NM_001001522

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: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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 001001522.1</u>, <u>NP 001001522.1</u>

RefSeq Size: 1574 bp
RefSeq ORF: 606 bp
Locus ID: 6876
UniProt ID: Q01995

Cytogenetics: 11q23.3



Gene Summary:

This gene encodes a shape change and transformation sensitive actin-binding protein which belongs to the calponin family. It is ubiquitously expressed in vascular and visceral smooth muscle, and is an early marker of smooth muscle differentiation. The encoded protein is thought to be involved in calcium-independent smooth muscle contraction. It acts as a tumor suppressor, and the loss of its expression is an early event in cell transformation and the development of some tumors, coinciding with cellular plasticity. The encoded protein has a domain architecture consisting of an N-terminal calponin homology (CH) domain and a C-terminal calponin-like (CLIK) domain. Mice with a knockout of the orthologous gene are viable and fertile but their vascular smooth muscle cells exhibit alterations in the distribution of the actin filament and changes in cytoskeletal organization. [provided by RefSeq, Aug 2017] Transcript Variant: This variant (1) represents the longer transcript. Variants 1 and 2 both 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.