

Product datasheet for SC329621

SEM1 (NM 001201450) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: SEM1 (NM_001201450) Human Untagged Clone

Tag: Tag Free Symbol: SEM1

Synonyms: C7orf76; DSS1; ECD; SHFD1; Shfdg1; SHFM1; SHSF1

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329621 representing NM_001201450.

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

GGTGACCTCTACAATCTGGCCCCAGCAGAAAGAACTTGCTAG

Restriction Sites: Sgfl-Mlul

ACCN: NM_001201450

Insert Size: 387 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.



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RefSeq: NM 001201450.1

RefSeq Size: 2707 bp
RefSeq ORF: 387 bp
Locus ID: 7979
Cytogenetics: 7q21.3

Protein Pathways: Homologous recombination, Proteasome

MW: 14.1 kDa

Gene Summary: The product of this gene has been localized within the split hand/split foot malformation

locus SHFM1 at chromosome 7. It has been proposed to be a candidate gene for the

autosomal dominant form of the heterogeneous limb developmental disorder split hand/split foot malformation type 1. In addition, it has been shown to directly interact with BRCA2. It also may play a role in the completion of the cell cycle. [provided by RefSeq, Jul 2008]

Transcript Variant: This variant (6) represents the use of an alternate promoter, resulting in a different 5' UTR and use of an alternate start codon, compared to variant 1. It encodes

isoform 4, which is longer and has a distinct N-terminus, compared to isoform 1.