

Product datasheet for SC209238

SHFM3 (FBXW4) (NM 022039) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: SHFM3 (FBXW4) (NM_022039) Human 3' UTR Clone

Symbol: SHFM3

Synonyms: DAC; FBW4; FBWD4; SHFM3; SHSF3

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_022039

Insert Size: 655 bp

Insert Sequence: >SC209238 3'UTR clone of NM_022039

The sequence shown below is from the reference sequence of NM_022039. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

TCATTTTGTAAAAATAAAGCTTGATTGTTCACA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

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).



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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.

RefSeq: <u>NM 022039.4</u>

Summary: This gene is a member of the F-box/WD-40 gene family, which recruit specific target proteins

through their WD-40 protein-protein binding domains for ubiquitin mediated degradation. In

mouse, a highly similar protein is thought to be responsible for maintaining the apical ectodermal ridge of developing limb buds; disruption of the mouse gene results in the absence of central digits, underdeveloped or absent metacarpal/metatarsal bones and syndactyly. This phenotype is remarkably similar to split hand-split foot malformation in humans, a clinically heterogeneous condition with a variety of modes of transmission. An autosomal recessive form has been mapped to the chromosomal region where this gene is located, and complex rearrangements involving duplications of this gene and others have been associated with the condition. A pseudogene of this locus has been mapped to one of

the introns of the BCR gene on chromosome 22. [provided by RefSeq, Jul 2008]

Locus ID: 6468 MW: 23.6