

## Product datasheet for **SC204732**

### **SNF5 (SMARCB1) (NM\_003073) Human 3' UTR Clone**

#### **Product data:**

|                           |                                                                                          |
|---------------------------|------------------------------------------------------------------------------------------|
| Product Type:             | 3' UTR Clones                                                                            |
| Product Name:             | SNF5 (SMARCB1) (NM_003073) Human 3' UTR Clone                                            |
| Symbol:                   | SNF5                                                                                     |
| Synonyms:                 | BAF47; CSS3; hSNFS; INI1; MRD15; PPP1R144; RDT; RTPS1; Sfh1p; SNF5; SNF5L1; Snr1; SWNTS1 |
| Mammalian Cell Selection: | Neomycin                                                                                 |
| Vector:                   | pMirTarget (PS100062)                                                                    |
| ACCN:                     | NM_003073                                                                                |
| Insert Size:              | 2000 bp                                                                                  |



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**Insert Sequence:**

>SC204732 3'UTR clone of NM\_003073

The sequence shown below is from the reference sequence of NM\_003073. 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
CGTCTTGCCAACACGGCCCCGGCCTGGTAACAGCCCATCAGCACACGGCTCCCACGGAGCATCTCAGA
AGATTGGGCCGCCTCTCCTCCATCTTCTGGCAAGGACAGAGGCGAGGGGACAGCCAGCGCCATCCTGA
GGATCGGGTGGGGGTGGAGTGGGGGCTTCCAGGTGGCCCTTCCCGGCACACATTCCATTTGTTGAGCCC
CAGTCTGCCCCCACCACCCCTCCCTACCCCTCCCACTCTCTGGGTGAGGAAGAACTTATTTT
AGGTTGTGTTTTGTTTGTATAGGAGCCCCAGGCAGGGCTAGTAACAGTTTTTAAATAAAAGGCAACA
GGTCATGTTCAATTTCTCAACAGGTGATGTTCAATTTCTCAAAGTTTTAACATAAAAAAATAGAGAG
CCAGGAGTGGGGCGGGCCTGGGGGACGAAGGTGGTATGTGAACAAGTTGGCACACAGGCCTCACC
CTCCTCTGCCTCAGATTCCTCAAGTGGGAGGTGGGGGTGAATGGGGCTCCGGGTAGCACCTCAGCTCCT
CTCAGCTCCCCTCAGCCTGTTCTCCTTCCAGACCCAGAGAGCTGAGAAGAGTAGCTGTGAGGCTCAGGG
CAAGAGGCTCTCTGCCTTTCAGGAACAGCCCTAACCTGTCTCCCTTGCTTGGCCTCAGGAAGGTGCCG
CGAGCTCTCCTGCCGTCCCTGGGCCGCCCTGGCTCTGCTGTGTCCAGATGGTCAGGCTACTGCCAGCTG
GGGCCCTTGCTGCTCTGAAGTCCCTGCGGAGGGCCAGTCTGTGTGGGCACTGCTGGGCTGTGCCAG
CCTGGGTGACAGGAGGCTGTTCTAGCTCCAGTGGCACCCATAGCCAGGTGAGTGGGGCCCTTCCAC
CCCAGCAGGTGCTGTGGCCTGGGCCAGTCTGCTTACAAGCCAGCTGTGAGGAATATGGGAATAGCC
CTCCCGGCCTGGTGCCAGCTCTTGGAGTTGACACGGTACAGGGAGGAGACACAGCCAGGGTCCCTTCC
CAGCCCTGCCTCCAAGGAGTTCATGTCCCCTCTGTTCTCATCTGTAATAGGGAGGTGTCCCATTTCTC
AGAATGGACACAGGATCTGGGAGGGCAGCAAACTGGCTCGCAGCTCCAGCCTTACTGAAGAGAATGGGC
ACAGATCCGGGCACAGATCCCAGCACAGACTGCTGCCACCTCAGCTGTTGGCAGTCCCATGCTGCCA
GGGCAGGGCTAGGGTCAAGGCTGCTGTGCTCCCTGGAAGTGGGGTAGGGCCCCATGTGGGCAGAGGC
AGAGCTCTGATTAGGGATTGGGGTCTTGGTCTGCTGAGATGTGAGAGGAGGGCTCCTTTGAGCACATGT
TAGCATGGGACTCTTCCAGGGAGTTTGCCTCAGGGCCTTGCCTCCATCAAAGAGTGGAACTCCCC
AGAGCCCCATGCACAGCAAGGGGACAGTGGGCCTTACTGGAAGGCCTTGAACAAAGGGGAAGATTCCC
AGCCCAGCTGCTCTTAGACATGAACAGGTTTCATTGCTGAGGTGTTTGTCTGTCCATGAGGTAGGAAC
CTCGGCAATGAAAGGGTGAAGCAGCCCTGTGTCTCCCAACTGGGGGATGGAAGGAACCTTGGCTGCC
TACCCCCACAGGTGCGGCAGGGCACCTGGCTGGGAGGTGCCGGGAAGGCTGGGCCCTCACTCCTGACC
GCCAGCTCACACGCCGCAAGCCATCTCCACAAGGTCTGGCTACAACACGGAGGGCAGACTCAACAGA
GAACAGTGTGTTACCATGAAAATGACAACCTGTCTTTGGAGGAGGCCCCGTGCCACTGAGCATCCAGA
AATAAACCAACATGGACAGGCTTAGAACAACAAGGAAAGCTGCCAGGTGAGAAGAGAAAAATGAGCC
ACAGGGGTGCGATAAGGCTCACACAGTCTCAGCTAAAAGGGCAGGAACAGAACCTTCCAGAAGTC
ACGCGTAAGCGGCCGCGGCATCTAGATTGGAAGAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
    
```

**Restriction Sites:**

SgfI-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.

**RefSeq:**

NM\_003073.5

|           |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               |
|-----------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Summary:  | The protein encoded by this gene is part of a complex that relieves repressive chromatin structures, allowing the transcriptional machinery to access its targets more effectively. The encoded nuclear protein may also bind to and enhance the DNA joining activity of HIV-1 integrase. This gene has been found to be a tumor suppressor, and mutations in it have been associated with malignant rhabdoid tumors. Alternatively spliced transcript variants have been found for this gene. [provided by RefSeq, Dec 2015] |
| Locus ID: | 6598                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |
| MW:       | 71.5                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          |