

Product datasheet for **SC206707**

SNAP45 (SNAPC2) (NM_003083) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: SNAP45 (SNAPC2) (NM_003083) Human 3' UTR Clone
Symbol: SNAP45
Synonyms: PTFDELTA; SNAP45
Mammalian Cell Selection: Neomycin
Vector: pMirTarget (PS100062)
ACCN: NM_003083
Insert Size: 516 bp
Insert Sequence: >SC206707 3'UTR clone of NM_003083
 The sequence shown below is from the reference sequence of NM_003083. 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
CTGGGTCGGGCAGCCACCCCTGCCAGTGAGGGGCATGGCGGCAGGAGGCCACACCAGGCCCCCGCC
CTGCCCCTCGGTTCTGCTCGGCTGGCCCTGGCTCTTTCTGAGGATCCCGTCATGGGGGAAGTCTTGA
GATGATGCTCAGCTGTGGGGCGGCCTTAAGATGCCCCATACTTTGGGGTCTCAGAAATGGAACCCC
CGTTGTACAGGGTTGGGTGGGGTTGCAGGACTCCACTCACAAGCCTCCTGATGTCAAGGACAGGCGG
ACAGGGCTGGCCTCCCCAGTCCCAAGCCCACTGTGCCTTGTGTCTGCTGGGGGCCATAGCTGGC
ACTGCCACCGTAAAGGCCCTCGCACATTTTCCCCTTCTGTACACCTCGGGGCCAGCATCCTCACCT
TCTTAACTGACCAGTCGTGGTTACTCCCTGCTGCCAGGTCTTCCCCTTCCCGGGGTATTCTGTGAC
CATGAATAAAGTTATCATTCTCTTTCTTTCA
ACGCGTAAGCGGCCGCGGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
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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).



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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:	NM_003083.4
Summary:	This gene encodes a subunit of the snRNA-activating protein complex which is associated with the TATA box-binding protein. The encoded protein is necessary for RNA polymerase II and III dependent small-nuclear RNA gene transcription. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Nov 2009]
Locus ID:	6618
MW:	18.3