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Product datasheet for SC207495

MFAP1 (NM_005926) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Symbol:	MFAP1
Synonyms:	AMP
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_005926
Insert Size:	632 bp
Insert Sequence:	>SC207495 3'UTR clone of NM_005926 The sequence shown below is from the reference sequence of NM_005926. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC CCATCTGCCAAGAAGCGGAAAACTACCTAGGGTCCAACTGCTTATTCTTCCAACTGTGGAACACAAGGG GAGTCTCAGCATCTGGTCCTTGATTTGGTTTTTTCATTGTTTCCTTGGCCCCTGTATCCAGATATTGGA CTTACTGCTATACTTGTGATACTGGGTAGCCCAGACTTTGAAGGTGCTTTGTGAGGTTTGGACTCATGC TGAGAAACCCACAGGAAAGCACTGTCCAGGTAGGATTAGAGGCTTCCCACTTAAAACTATTTCTGAGAA ATCTTAGGTTTTATCACTGCTATGGTTTCCCATATTTACTTGGGACTGTTCTGACTTTCTTTTTCCAGC CCTTAGCTTGGGTTAGAAAAGTGGACATGTAAGTGAACAATGCATTACTTCTACCTTAGGTTTAGGAGT AATATACCCGGAAATCTAAGCTCATGGAAACATGTTTTCCATTTTTGCTTGGAGTCCGTTTTTCTAGTT GTACATACTTTTTGATCCATATATGTGTGCATGTCAAGAAATAAAAGAATCACACAACAAGGAGATTTT TAGCATGATACTGGAAATAAAATGTCCAAGCTGACAAGTATTAATTGCCTATTAAAGTAAAAGATAACA ATTGTCAGGCA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG
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.
Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_005926.3
Summary:	Involved in pre-mRNA splicing as a component of the spliceosome.[UniProtKB/Swiss-Prot Function]
Locus ID:	4236
MW:	23.9