

Product datasheet for **SC207782**

MEPCE (NM_019606) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	MEPCE (NM_019606) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	MEPCE
Synonyms:	BCDIN3
ACCN:	NM_019606
Insert Size:	594 bp
Insert Sequence:	>SC207782 3'UTR clone of NM_019606 The sequence shown below is from the reference sequence of NM_019606. 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
TTCCACAAGGCCGATCCCCAGCCACTAAGTGGCCCCCTAAACAGAAAAGTGTGAAGAGGCTGCCCTCG
CTGCTCATAAGGACCTGGGGGAAGAGGAAAGTGTCCCAAGTCTTTCTTCTGACTCCAAAAATAGTT
TCCTTTCTTGGATCTGCAAAGAAAGCTTTTCTTCCGTCGCTGCCTCAGCCTCCTCCTATGCCTCTGGC
ACCTGCGCAGCAAGGCTGGCTGTGCTGGAGTCACCATCATCTTCTCTCCCCAGCCTCCCAGGCTGGA
TGGCATGGACTGTTTGTGACCTCTGTTCTCTTAGGGCATGGGAGGTGGGAGGATATCAAATCTCTAG
CCCTTTCTCCTATTCTCCCAAGGAGAGAGATTCCCATTCTCCTCGGCCATTGTACCTAGCTCTTGTC
CCTAGCTGCATTTCAAGTGGACCATGGATAGAGGGACTGAGGGTTAGACGGGGAAGACTGGCAGGGAGGC
ACGCAGGTACTGTAAAAATCCTTCCCTTTGCCCTCCCCAGTGGGAGAGGGGTTGGGTTTTCAATGTG
AGAACAGCACAATAAACTTGATGTCTAGGGCAGTGGCCCCCA
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

Restriction Sites:	Sgfl-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.



View online »

RefSeq: [NM_019606.6](#)

Summary: S-adenosyl-L-methionine-dependent methyltransferase that adds a methylphosphate cap at the 5'-end of 7SK snRNA, leading to stabilize it.[UniProtKB/Swiss-Prot Function]

Locus ID: 56257

MW: 21.4