

## Product datasheet for SC205616

## MHF1 (CENPS) (NM\_199294) Human 3' UTR Clone

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	MHF1 (CENPS) (NM_199294) Human 3' UTR Clone
Symbol:	MHF1
Synonyms:	APITD1; CENP-S; FAAP16; MHF1
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_199294
Insert Size:	423 bp
Insert Sequence:	>SC205616 3' UTR clone of NM_199294 The sequence shown below is from the reference sequence of NM_199294. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site Blue=Stop Codon
	CAATTGGCAGAGCTCAGAATTCAA <mark>GCGATCGC</mark>
	TTCAAGGCAGCCAGCAGAGGCTGGAGTGGTGGAAAGTGAGAAT <b>TAA</b> AGTCCCTCGCCGCTTGGAAAGTGC AGCCTTCTACAGGTAGAGCCACCTAGAAATGCATATGGCTGCAAAGGAAACTTTGAAGGGTTAAATAGAG ATTTAAAAAAATAAAAT
	ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCG
<b>Restriction Sites:</b>	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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	MHF1 (CENPS) (NM_199294) Human 3' UTR Clone – SC205616
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 199294.1</u>
Summary:	This gene was identified in the neuroblastoma tumor suppressor candidate region on chromosome 1p36. It contains a TFIID-31 domain, similar to that found in TATA box-binding protein-associated factor, TAF(II)31, which is required for p53-mediated transcription activation. This gene was expressed at very low levels in neuroblastoma tumors, and was shown to reduce cell growth in neuroblastoma cells, suggesting that it may have a role in a cell death pathway. The protein is a component of multiple complexes, including the Fanconi anemia (FA) core complex, the APITD1/CENPS complex, and the CENPA-CAD (nucleosome distal) complex. Known functions include an involvement with chromatin associations of the FA core complex, and a role in the stable assembly of the outer kinetochore. Alternative splicing of this gene results in multiple transcript variants. Naturally occurring read-through transcripts also exist between this gene and the downstream cortistatin (CORT) gene, as represented in GeneID:100526739. An APITD1-related pseudogene has been identified on chromosome 7. [provided by RefSeq, Nov 2010]
Locus ID:	378708

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