

Product datasheet for **SC117199**

MTA1 (NM_004689) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	MTA1 (NM_004689) Human Untagged Clone
Tag:	Tag Free
Symbol:	MTA1
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL5</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >OriGene ORF within SC117199 sequence for NM_004689 edited (data generated by NextGen Sequencing)

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ATGGCCGCAACATGTACAGGGTCGGAGACTACGTCTACTTTGAGAACTCCTCCAGCAAC
CCATACCTGATCCGGAGAATCGAGGAGCTCAACAAGACGGCCAATGGGAACGTGGAGGCC
AAAGTGGTGTGCTTCTACCGGAGGCGGGACATCTCCAGCACCTCATCGCCCTGGCCGAC
AAGCACGCAACCTGTCACTGTCTATAAGGCCGACCGGGGGCGGACAACGGCGAGGAA
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CACAGCTGCGGCATCGGGAGCTGTTCTCTCCGGCAGCTGGAGTCTCTGCCCGCCACG
CACATCAGGGGCAAGTGCAGCGTACCCTGCTCAACGAGACCGAGTCTGCTCAAGTCTAC
CTGGAGCGGGAGGATTTCTTCTTCTATTCTCTAGTCTACGACCCACAGCAGAAGACCCTG
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ATGCCCAGTAGGGTCTGGCAAACACGGACAGACCAGGCACATGGGACCAAGCCGGAAC
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CCCCAGTCAAGCGGGCGGGATGAACTGGATCGACGCCCGGATGACGTGTTCTACATG
GCCACAGAGGAGACCAGGAAGATCCGCAAGCTGCTCTCATCTCGGAAACCAAGCGTGCT
GCCCGCCGGCCCTACAAGCCATCGCCCTGCGCCAGAGCCAGGCCCTGCCCGCGGCCA
CCGCCACCTGCGCCGTAACGACGAGCCCATCGTCATCGAGGACTAG

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Clone variation with respect to NM_004689.3
 1114 g=>a;1834 g=>a

Reconstitution Method:	<ol style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM_004689.2, NP_004680.1</u>
RefSeq Size:	2662 bp
RefSeq ORF:	2148 bp
Locus ID:	9112
UniProt ID:	<u>Q13330</u>
Cytogenetics:	14q32.33
Domains:	GATA, ELM2, myb_DNA-binding, BAH
Protein Families:	Druggable Genome, Transcription Factors
Gene Summary:	<p>This gene encodes a protein that was identified in a screen for genes expressed in metastatic cells, specifically, mammary adenocarcinoma cell lines. Expression of this gene has been correlated with the metastatic potential of at least two types of carcinomas although it is also expressed in many normal tissues. The role it plays in metastasis is unclear. It was initially thought to be the 70kD component of a nucleosome remodeling deacetylase complex, NuRD, but it is more likely that this component is a different but very similar protein. These two proteins are so closely related, though, that they share the same types of domains. These domains include two DNA binding domains, a dimerization domain, and a domain commonly found in proteins that methylate DNA. The profile and activity of this gene product suggest that it is involved in regulating transcription and that this may be accomplished by chromatin remodeling. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Feb 2011]</p> <p>Transcript Variant: This variant (1) represents the longer transcript and encodes the longer isoform (MTA1).</p>