OriGene Technologies, Inc.
9620 Medical Center Drive, Ste 200
Rockville, MD 20850, US
Phone: +1-888-267-4436
https://www.origene.com
techsupport@origene.com
EU: info-de@origene.com
CN: techsupport@origene.cn

Product datasheet for SC207189

MYBBP1A (NM_014520) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	MYBBP1A (NM_014520) Human 3' UTR Clone
Symbol:	MYBBP1A
Synonyms:	P160; PAP2; Pol5
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_014520
Insert Size:	541 bp
Insert Sequence:	>SC207189 3'UTR clone of NM_014520 The sequence shown below is from the reference sequence of NM_014520. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GCACAGGTGAGGAAGGCAGGGAAGCCCTGAGCACAGGTACGGGCCCCCCTCAGCCCCTGCCTCCATCTG CCTGAGACGCCTATTTTTTTTTTTTTTTTTTAAACCATGATTTTAATACGCAAGCTGTTTCTAAGGCGC TGCCACTGGGGAGGGTGGCTGTTGCCGCCTGCCCGGGCATCCTGCTCTGGCAAGCACAGCCTGAGCCAT TCCTGCAGGGGTCCCAGGGTGCAGAGACCTCCCCACCCCCGGTTCTGGGCTGGGACCCTGGCTCCAGGG CCATGTCCAGGGCTCTGGTGTTTGCCTGGGTTGGTGCAGGTTGATGTGCTGGCTGCAGGCAGGTGTGAC CATCTCTCGTGCCTGCCACCTCTTTGCCCCCAGGCTTTTTTGCTGTGAGGGAGCCACCAGGGGGTGATT TAAATAGGTTTATTTCTTCATTTACAAGAGGAATATATTTGGCTTCTCTCTTAAGACTCTGAGATTCAC AATCAGCAGCTCTAAAAAATAAAGGAGCAGTTTGGCTTCCGGAAGGAAGAGGAGGCAA 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.
RefSeq:	NM_014520.4
Summary:	This gene encodes a nucleolar transcriptional regulator that was first identified by its ability to bind specifically to the Myb proto-oncogene protein. The encoded protein is thought to play a role in many cellular processes including response to nucleolar stress, tumor suppression and synthesis of ribosomal DNA. Alternate splicing results in multiple transcript variants. [provided by RefSeq, Sep 2013]
Locus ID:	10514
MW:	20.3