

Product datasheet for SC207572

BRMS1 (NM_015399) Human 3' UTR Clone

Product data:

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

Froduct data.	
Product Type:	3' UTR Clones
Product Name:	BRMS1 (NM_015399) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	BRMS1
ACCN:	NM_015399
Insert Size:	573 bp
Insert Sequence:	<pre>>SC207572 3'UTR clone of NM_015399 The sequence shown below is from the reference sequence of NM_015399. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC</pre>
	CCTCAGAAGAGAAAATCGGATGGACCTTGACCCTGCTGTTCACAGCCAGGGGGACCCTCAGAGCAGCTG GCACTGCACCCAGGATTCTCGTCTTCCTCCTGCAGACAGGCGGACCCACAGGCCCCTCAGGGTCTGCCC AGCCAGGCTCCTGTGGTGCTGCTGGGCCCTCCCACTCCATCTGGCACTGGCCTGGACTCCTCCTCTGCC CTCCTCGAGGCCTGCACAGCTGTGGCCGTGGAGCTGACCTGACCAGGCAAGGCTGCTGTCTCCATCCCT GAGCCGCCTGCCACCTCCCACTCCTGAAGATCCATCTCTTGGGGCTCCCCTGACAGAGAAGACAGCCGA AGTCAAAGCCACATCCTCTTGCTGATGTTGGATGCAGGCTGTCCGGCCTCAGGGCCAGGGAAGCCAGTTT CCACTGTGCGGGGAACTCTGAGTCAGACGTGATTATCTGGGGGTCTGTCCACCCTGGCAGGGAGCCAGTTT CCACTGTGCGGGGAACTCTGAGTCAGACGTGATTATCTGGGGGTCTGCTCCCCTGGCTGG
Restriction Sites:	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).
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 015399.4</u>



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This gene reduces the metastatic potential, but not the tumorogenicity, of human breast cancer and melanoma cell lines. The protein encoded by this gene localizes primarily to the nucleus and is a component of the mSin3a family of histone deacetylase complexes (HDAC).
The protein contains two coiled-coil motifs and several imperfect leucine zipper motifs. Alternative splicing results in two transcript variants encoding different isoforms. [provided by RefSeq, Jul 2008]
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