

Product datasheet for **SC214840**

ARMC8 (NM_015396) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	ARMC8 (NM_015396) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	ARMC8
Synonyms:	GID5; HSPC056; S863-2; VID28
ACCN:	NM_015396
Insert Size:	2000 bp



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Insert Sequence: >SC214840 3'UTR clone of NM_015396
 The sequence shown below is from the reference sequence of NM_015396. 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
AAGATGGCACTGCAGCAGTACCTGGCATGATGGGAGTGCCCTGGGCACCTGCGAGCATCCACCTCCT
TGTTTAAGGAAGTACAGGAACCAGCCTCATTGATTCTTCTATTGCACAAGTACCTTGACTGCAG
GGAGCTGTTTTGCAAAAGCAGTTTAGTAGGCTTAGATCTCAAATTCATCTTGAGAACATTTTTTTGAGG
TAGTAATTTCTCAGAAGACTCTGTGTTTTGTTTTGTTTTTTTTCTGAGCTACTCGGACTCTTTAT
TAGAACAAATCAATCATTTTCTTTGGACCTACAATTTTTGCTATGCTGCAGCCACTTTGTGAGTGAGA
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TTAATGCAATTTTTCTGCTCTCAACATGACCACCAAGATTCAAACACAGAGACTTCCAAGTTACTA
TTTTTTGAGAACATCACCAGATTTAAAACTGTCAACACAGGACTTGGTTTTGTGAGATATCAGAGTG
CTAATGTGAGAAACCCAGTTAAAAGAAAACCTAGTTTATTGGGAAGAAAGCCTCAGGCCACTCGTTTT
TAGAACTGTGAAAATATCCTTGGTTGGCATTTTAACACAAAACACTATAGTGTGGTATTAAATAGT
ACGCGT AAGCGGCCGCGGCATCTAGATTGAAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCACCGCCGCCTTCTATGAAAGG
  
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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).

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_015396.6](#)

Summary: Component of the CTLH E3 ubiquitin-protein ligase complex that selectively accepts ubiquitin from UBE2H and mediates ubiquitination and subsequent proteasomal degradation of the transcription factor HBP1.[UniProtKB/Swiss-Prot Function]

Locus ID: 25852

MW: 76.6