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 SC204759

BUD23 (NM_017528) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	BUD23 (NM_017528) Human 3' UTR Clone
Symbol:	BUD23
Synonyms:	HASJ4442; HUSSY-3; MERM1; PP3381; WBMT; WBSCR22
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_017528
Insert Size:	356 bp
Insert Sequence:	>SC204759 3'UTR clone of NM_017528 The sequence shown below is from the reference sequence of NM_017528. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC ACCGGCCGCAAGCGCAAGCCCCGCTTCTAAGTCACCACGCGGTTCTGGAAAGGCACTTGCCTCTGCACT TTTCTATATTGTTCAGCTGACAAAGTAGTATTTTAGAAAAGTTCTAAAGTTATAAAAATGTTTTCTGCA GTAAAAAAAAAGTTCTCTGGGCCGGGCGTGGTGGCTCACACCTGTAATCCCAGCACCTTGGGAGGCTGA GGTGGGAGGATCATTTGAGGCCAGGAGTTTGAGACCTGCCTGGGCAACATAATGAAACTTCCTTTCCAG GGAGGAAAAAAAAAAAAAAAAAAAGCTCTGAGAGCATCTTATTTTGTTTAAAGGCAAGAAATAAAATTT CCTTTTGTGGA 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).
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.

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RefSeq:	NM_017528.5
Summary:	This gene encodes a protein containing a nuclear localization signal and an S-adenosyl-L-methionine binding motif typical of methyltransferases, suggesting that the encoded protein may act on DNA methylation. This gene is deleted in Williams syndrome, a multisystem developmental disorder caused by the deletion of contiguous genes at 7q11.23. Alternatively spliced transcript variants have been found. [provided by RefSeq, Feb 2011]
Locus ID:	114049
MW:	13.3