

## **Product datasheet for SC205164**

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## BubR1 (BUB1B) (NM\_001211) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: BubR1 (BUB1B) (NM\_001211) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: BUB1B

Synonyms: Bub1A; BUB1beta; BUBR1; hBUBR1; MAD3L; MVA1; SSK1

ACCN: NM\_001211

**Insert Size:** 394 bp

Insert Sequence: >SC205164 3'UTR clone of NM\_001211

The sequence shown below is from the reference sequence of NM\_001211. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

ACTAGTCCTGGGGCTTTGCTCTTTCAGTGAGCTAGGCAATCAAGTCTCACAGATTGCTGCCTCAGAGCA ATGGTTGTATTGTGGAACACTGAAACTGTATGTGCTGTAATTTAATTTAGGACACACTTTAGATGCACTA CCATTGCTGTTCTACTTTTTGGTACAGGTATATTTTTGACGTCACTGATATTTTTATACAGTGATATAC TTACTCATGGCCTTGTCTAACTTTTGTGAAGAACTATTTTATTCTAAACAGACTCATTACAAATGGTTA CCTTGTTATTTAACCCATTTGTAAATTGTCCCTGTACTTTTCCCTGTACTTTTCCCATTTTGTAAATTTGTAAAATGTTC

TCTTATGATCACCATGTATTTTGTAAATAATAAAAATAGTATCTGTTAAA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

**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.

**RefSeg:** NM 001211.6





## BubR1 (BUB1B) (NM\_001211) Human 3' UTR Clone - SC205164

Summary: This gene encodes a kinase involved in spindle checkpoint function. The protein has been

localized to the kinetochore and plays a role in the inhibition of the anaphase-promoting

complex/cyclosome (APC/C), delaying the onset of anaphase and ensuring proper  $% \left( A_{1}\right) =\left( A_{1}\right) +\left( A_{2}\right) +\left( A_{3}\right) +\left( A_{3}$ 

chromosome segregation. Impaired spindle checkpoint function has been found in many

forms of cancer. [provided by RefSeq, Jul 2008]

Locus ID: 701

MW: 15.3