

## **Product datasheet for SC204772**

## OriGene Technologies, Inc.

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## Exonuclease 1 (EXO1) (NM\_130398) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

Product Name: Exonuclease 1 (EXO1) (NM\_130398) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: EXO1

Synonyms: HEX1; hExol ACCN: NM\_130398

**Insert Size:** 369 bp

Insert Sequence: >SC204772 3'UTR clone of NM\_130398

The sequence shown below is from the reference sequence of NM\_130398. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

GGCCGTGTTCAAAGAGCAATATTCCAGTAAATGCAGACTGCTGCAAAGCTTTTGCCTGCAAGAGAATCT
GATCAATTTGAAGTCCCTGTTTGGGAATGAGGCACTTATCAGCATGAAGAATTTTTTCTCATTCTGTGC
CATTTTAAAAATAGAATACATTTTGTATATTAACTTTATAATTGGGTTGTGGTTTTTTTGCTCAGCTTT
TTATATTTTTATAAGAAGCTAAATAGAAGAATAATTGTATCTCTGACAGGTTTTTGGAGGTTTTAGTGT
TAATTGGGAAAATCCTCTGGAGTTTATAAAAGTCTACTCTAAATATTTCTGTAATGTTGTCAAGTAGAA

AGATAGTAAATGGAGAAACTACAA

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 130398.4





## Exonuclease 1 (EXO1) (NM\_130398) Human 3' UTR Clone - SC204772

Summary: This gene encodes a protein with 5' to 3' exonuclease activity as well as an RNase H activity. It

is similar to the Saccharomyces cerevisiae protein Exo1 which interacts with Msh2 and which is involved in mismatch repair and recombination. Alternative splicing of this gene results in three transcript variants encoding two different isoforms. [provided by RefSeq, Jul 2008]

**Locus ID:** 9156

**MW:** 14.8