

Product datasheet for SC204773

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

Exonuclease 1 (EXO1) (NM_006027) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Exonuclease 1 (EXO1) (NM_006027) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: EXO1

Synonyms: HEX1; hExol ACCN: NM_006027

Insert Size: 385 bp

Insert Sequence: >SC204773 3'UTR clone of NM_006027

The sequence shown below is from the reference sequence of NM_006027. 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 TAATTGGGAAAATCCTCTGGAGGTTTATAAAAAGTCTACTCTAAATATTTCTGTAATGTTGTCAAGTAGAA

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.

RefSeq: <u>NM_006027.4</u>





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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: 15.5