

Product datasheet for SC210807

OriGene Technologies, Inc.

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Carbonic Anhydrase III (CA3) (NM_005181) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Carbonic Anhydrase III (CA3) (NM_005181) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: CA3

Synonyms: CAIII; Car3
ACCN: NM_005181

Insert Size: 917 bp

Insert Sequence: >SC210807 3'UTR clone of NM_005181

The sequence shown below is from the reference sequence of NM_005181. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

ATAAAGACATAAATGGAAAA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

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).





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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 005181.4</u>

Summary: Carbonic anhydrase III (CAIII) is a member of a multigene family (at least six separate genes

are known) that encodes carbonic anhydrase isozymes. These carbonic anhydrases are a class

of metalloenzymes that catalyze the reversible hydration of carbon dioxide and are

differentially expressed in a number of cell types. The expression of the CA3 gene is strictly tissue specific and present at high levels in skeletal muscle and much lower levels in cardiac and smooth muscle. A proportion of carriers of Duchenne muscle dystrophy have a higher CA3 level than normal. The gene spans 10.3 kb and contains seven exons and six introns.

[provided by RefSeq, Oct 2008]

Locus ID: 761 **MW:** 36.2