

Product datasheet for SC208055

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

Apc7 (ANAPC7) (NM_001137664) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Apc7 (ANAPC7) (NM_001137664) Human 3' UTR Clone

Symbol: Apc7
Synonyms: APC7

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_001137664

Insert Size: 618 bp

Insert Sequence: >SC208055 3'UTR clone of NM_001137664

The sequence shown below is from the reference sequence of NM_001137664. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

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).





Apc7 (ANAPC7) (NM_001137664) Human 3' UTR Clone - SC208055

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 001137664.2</u>

Summary: This gene encodes a tetratricopeptide repeat containing component of the anaphase

promoting complex/cyclosome (APC/C), a large E3 ubiquitin ligase that controls cell cycle progression by targeting a number of cell cycle regulators such as B-type cyclins for 26S proteasome-mediated degradation through ubiquitination. The encoded protein is required for proper protein ubiquitination function of APC/C and for the interaction of APC/C with certain transcription coactivators. Multiple transcript variants encoding different isoforms

have been found for this gene. [provided by RefSeq, Nov 2008]

Locus ID: 51434 MW: 23.3