

Product datasheet for SC206117

OriGene Technologies, Inc.

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Carboxypeptidase B2 (CPB2) (NM_016413) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Carboxypeptidase B2 (CPB2) (NM 016413) Human 3' UTR Clone

Symbol: Carboxypeptidase B2

Synonyms: CPU; PCPB; TAFI

Mammalian Cell

Neomycin

Selection:

Vector: pMirTarget (PS100062)

ACCN: NM_016413

Insert Size: 483 bp

Insert Sequence: >SC206117 3'UTR clone of NM_016413

The sequence shown below is from the reference sequence of NM_016413. 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).

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.





Carboxypeptidase B2 (CPB2) (NM_016413) Human 3' UTR Clone - SC206117

RefSeq: <u>NM 016413.3</u>

Summary: Carboxypeptidases are enzymes that hydrolyze C-terminal peptide bonds. The

carboxypeptidase family includes metallo-, serine, and cysteine carboxypeptidases. According to their substrate specificity, these enzymes are referred to as carboxypeptidase A (cleaving aliphatic residues) or carboxypeptidase B (cleaving basic amino residues). The protein encoded by this gene is activated by trypsin and acts on carboxypeptidase B substrates. After thrombin activation, the mature protein downregulates fibrinolysis. Polymorphisms have been described for this gene and its promoter region. Alternate splicing results in multiple

transcript variants. [provided by RefSeq, Jun 2013]

Locus ID: 1361 **MW:** 18.5