

Product datasheet for SC205886

KAT5 (NM 006388) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: KAT5 (NM 006388) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: KAT5

Synonyms: cPLA2; ESA1; HTATIP; HTATIP1; NEDFASB; PLIP; TIP; TIP60; ZC2HC5

ACCN: NM 006388

Insert Size: 453 bp

>SC205886 3'UTR clone of NM_006388 **Insert Sequence:**

The sequence shown below is from the reference sequence of NM_006388. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AAGGACTGGAGCAAGAGGGGGAAGTGGTGACCAGACACTGCCCACTGCAGTGCCAAGACGGCAGCAGGA CTGGGGCTGATAGCCCACCCCGCCCCACTGCAGCTCCCACAAAGCACTCTAAGGGAGATGGGGCTGAG GACAGCTCAAAAAGGAGAGGACAGGCCTGGCAGGGGCCCACTGGTGCCCAGCACCAAGGCGAGCTCCGG GCTCAGACCAACTCCAAGGTCAGCTGGCCACAGGCCCAGGCCTCCTCTGAAGCAGGGACCAGAGGGAGC CAGGCAGCTGTGTACAGTGAGAAGGGATCCGGATGGGGGAGCTCTGTACAGAGGGCTGGTGATTGTAAA

CCCCTATTGCCCCCGGCAATAAATTGTTTCTATATGCCA

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.

NM 006388.4 RefSeq:



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Summary: The protein encoded by this gene belongs to the MYST family of histone acetyl transferases

(HATs) and was originally isolated as an HIV-1 TAT-interactive protein. HATs play important roles in regulating chromatin remodeling, transcription and other nuclear processes by acetylating histone and nonhistone proteins. This protein is a histone acetylase that has a role in DNA repair and apoptosis and is thought to play an important role in signal transduction. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Jul

2008]

Locus ID: 10524

MW: 15.9