

Product datasheet for SC209106

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

SETD2 (NM_014159) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: SETD2 (NM_014159) Human 3' UTR Clone

Vector: pMirTarget (PS100062)

Symbol: SETD2

Synonyms: HBP231; HIF-1; HSPC069; HYPB; KMT3A; LLS; p231HBP; SET2

ACCN: NM_014159

Insert Size: 687 bp

Insert Sequence: >SC209106 3'UTR clone of NM_014159

The sequence shown below is from the reference sequence of NM_014159. 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.





SETD2 (NM_014159) Human 3' UTR Clone - SC209106

RefSeq: <u>NM 014159.7</u>

Summary: Huntington's disease (HD), a neurodegenerative disorder characterized by loss of striatal

neurons, is caused by an expansion of a polyglutamine tract in the HD protein huntingtin.

This gene encodes a protein belonging to a class of huntingtin interacting proteins

characterized by WW motifs. This protein is a histone methyltransferase that is specific for lysine-36 of histone H3, and methylation of this residue is associated with active chromatin. This protein also contains a novel transcriptional activation domain and has been found associated with hyperphosphorylated RNA polymerase II. [provided by RefSeq, Aug 2008]

Locus ID: 29072

MW: 25.7