

Product datasheet for SC211109

OriGene Technologies, Inc.

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Amino terminal enhancer of split (AES) (NM_001130) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: Amino terminal enhancer of split (AES) (NM_001130) Human 3' UTR Clone

Symbol: Amino terminal enhancer of split

Synonyms: AES; AES-1; AES-2; ESP1; GRG; Grg-5; GRG5

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_001130

Insert Size: 939 bp

Insert Sequence: >SC211109 3'UTR clone of NM_001130

The sequence shown below is from the reference sequence of NM_001130. The complete

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

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

 ${\tt TGGGCTTAGCTGTGCCTTTCCAATAAAGATGTGAGAAGCTTC}$

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul





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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.

RefSeq: <u>NM 001130.6</u>

Summary: The protein encoded by this gene is similar in sequence to the amino terminus of Drosophila

enhancer of split groucho, a protein involved in neurogenesis during embryonic

development. The encoded protein, which belongs to the groucho/TLE family of proteins, can

function as a homooligomer or as a heteroologimer with other family members to

dominantly repress the expression of other family member genes. Three transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Jul 2008]

Locus ID: 166

MW: 34.6