

Product datasheet for RC200629L4V

OriGene Technologies, Inc.

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UBE3A (NM_130838) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: UBE3A (NM_130838) Human Tagged ORF Clone Lentiviral Particle

Symbol: UBE3A

Synonyms: ANCR; AS; E6-AP; EPVE6AP; HPVE6A

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_130838

ORF Size: 2556 bp

ORF Nucleotide

Sequence:
OTI Disclaimer:

The ORF insert of this clone is exactly the same as(RC200629).

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeg: NM 130838.1

 RefSeq Size:
 4491 bp

 RefSeq ORF:
 2559 bp

 Locus ID:
 7337

 UniProt ID:
 Q05086

 Cytogenetics:
 15q11.2

Domains: HECT

Protein Families: Druggable Genome





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Protein Pathways: Ubiquitin mediated proteolysis

MW: 97.8 kDa

Gene Summary: This gene encodes an E3 ubiquitin-protein ligase, part of the ubiquitin protein degradation

system. This imprinted gene is maternally expressed in brain and biallelically expressed in other tissues. Maternally inherited deletion of this gene causes Angelman Syndrome, characterized by severe motor and intellectual retardation, ataxia, hypotonia, epilepsy, absence of speech, and characteristic facies. The protein also interacts with the E6 protein of human papillomavirus types 16 and 18, resulting in ubiquitination and proteolysis of tumor protein p53. Alternative splicing of this gene results in three transcript variants encoding three isoforms with different N-termini. Additional transcript variants have been described,

but their full length nature has not been determined. [provided by RefSeq, Jul 2008]