

Product datasheet for SC330699

UEVLD (NM 001261386) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: UEVLD (NM_001261386) Human Untagged Clone

Tag: Tag Free
Symbol: UEVLD

Synonyms: ATTP; UEV3

Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC330699 representing NM_001261386.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGGAGTTCGACTGCGAGGGCCTGAGACGGCTGCTTGGCAAGTACAAGTTCAGGGACCTAACTGTGGAA GAACTAAGGAATGTAAATGTATTTTTCCCACATTTCAAATATTCCATGGACACCCTATGTTTTTAAAGAT AGTTCTCAGAAAGACCTGCTGAATTTACTGGCACAATTCCTGTGATGTATCAGGGTGTTTCAGATACA AATTCAAAGACTGGGCAAATCATGAGAATAAAAACAGTCAATAAAATTACTGTGGTTGGAGGTGGAGAA CTCGGTATTGCCTGCACATTAGCCATTTCAGCCTC TCAGAAGGGACTAAAGGAGCCACGATGGACCTTGAAATCTTCAACCTTCCTAATGTGGAGATCAGCAAA GTGGAAATCATGACCTATGTAACATGGAAACTGAGTACATTTCCTGCAAATCGAGTGATCGGAATTGGA TGTAATCTGGATTCACAGAGATTACAGTATATTATTACAAATGTTTTGAAGGCACAGACTTCAGGCAAA

GAAGTATGGGTTATTGGCGAGCAAGGAGAAGACAAAGGGATATTA<mark>TGA</mark>

Restriction Sites: Sgfl-Mlul

ACCN: NM_001261386

Insert Size: 600 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).



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Reconstitution Method:

- 1. Centrifuge at 5,000xg for 5min.
- 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
- 3. Close the tube and incubate for 10 minutes at room temperature.
- 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
- 5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: <u>NM 001261386.1</u>

 RefSeq Size:
 1766 bp

 RefSeq ORF:
 600 bp

 Locus ID:
 55293

 UniProt ID:
 Q8IX04

 Cytogenetics:
 11p15.1

 MW:
 22.2 kDa

Gene Summary: Possible negative regulator of polyubiquitination.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (7) has multiple differences in the coding region, compared to variant 1. The resulting isoform (g) lacks two internal segments and has a shorter and distinct C-terminus, compared to isoform a. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.