

Product datasheet for RC202106

UBE2E1 (NM_003341) Human Tagged ORF Clone

Product data:

Product Type: Expression Plasmids

Product Name: UBE2E1 (NM_003341) Human Tagged ORF Clone

Tag: Myc-DDK
Symbol: UBE2E1
Synonyms: UBCH6
Mammalian Cell Neomycin

Selection:

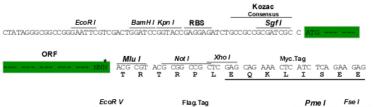
Vector: pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)Restriction Sites: Sgfl-Mlul

Cloning Scheme:

Cloning sites used for ORF Shuttling:





GAT CTG GCA GCA ANT GAT ATC CTG GAT TAC AGG GAT GAC GAC GAT AAG GTT TAA AACGGCCGGCCC

D L A A N D I L L D Y K D D D D K V stop

ACCN: NM_003341

ORF Size: 579 bp



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^{*} The last codon before the Stop codon of the ORF

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OTI Disclaimer: 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.

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).

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 003341.5</u>

 RefSeq Size:
 1845 bp

 RefSeq ORF:
 582 bp

 Locus ID:
 7324

 UniProt ID:
 P51965

Cytogenetics: 3p24.2 Domains: UBCc

Protein Pathways: Ubiquitin mediated proteolysis

MW: 21.4 kDa

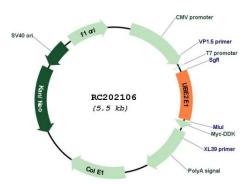
Gene Summary: The modification of proteins with ubiquitin is an important cellular mechanism for targeting

abnormal or short-lived proteins for degradation. Ubiquitination involves at least three classes of enzymes: ubiquitin-activating enzymes, or E1s, ubiquitin-conjugating enzymes, or E2s, and ubiquitin-protein ligases, or E3s. This gene encodes a member of the E2 ubiquitin-conjugating enzyme family. Three alternatively spliced transcript variants encoding distinct

isoforms have been found for this gene. [provided by RefSeq, Jan 2011]



Product images:



Circular map for RC202106