

Product datasheet for SC329665

UBE2H (NM_001202498) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: UBE2H (NM_001202498) Human Untagged Clone

Tag: Tag Free
Symbol: UBE2H

Synonyms: E2-20K; GID3; UBC8; UBCH; UBCH2

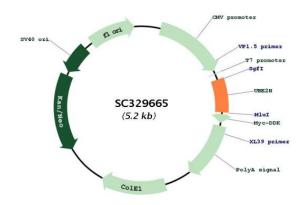
Vector: pCMV6-Entry (PS100001)

Fully Sequenced ORF: >SC329665 representing NM_001202498.

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

Restriction Sites: Sgfl-Mlul

Plasmid Map:



ACCN: NM 001202498

Insert Size: 342 bp



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com

UBE2H (NM_001202498) Human Untagged Clone - SC329665

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).

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 001202498.1</u>

 RefSeq Size:
 4778 bp

 RefSeq ORF:
 342 bp

 Locus ID:
 7328

 UniProt ID:
 P62256

Protein Pathways: Ubiquitin mediated proteolysis

7q32.2

MW: 12.9 kDa

Cytogenetics:

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. The encoded protein sequence is 100% identical to the mouse homolog and 98% identical to the frog and zebrafish homologs. Three alternatively spliced transcript variants have been found for this gene and they encode distinct isoforms.

[provided by RefSeq, Feb 2011]

Transcript Variant: This variant (3) has an alternate splice site in the 5' region, resulting in a downstream AUG start codon, as compared to variant 1. The resulting isoform (3) has a shorter N-terminus, as compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record

were based on transcript alignments.