

## Full-Length

## UBE2V1 (Human) Recombinant Protein (P01)

Catalog # H00007335-P01 Size 25 ug, 10 ug

## Applications



Specification	
Product Description	Human UBE2V1 full-length ORF ( AAH00468, 1 a.a 147 a.a.) recombinant protein with GST-tag at N-terminal.
Sequence	MAATTGSGVKVPRNFRLLEELEEGQKGVGDGTVSWGLEDDEDMTLTRWTGMIGPPRTIYENRIY SLKIECGPKYPEAPPFVRFVTKINMNGVNSSNGVVDPRAISVLAKWQNSYSIKVVLQELRRLMMS KENMKLPQPPEGQCYSN
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	41.91
Interspecies Antigen Sequence	Mouse (94); Rat (94)
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCI, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.
Storage Instruction	Store at -80°C. Aliquot to avoid repeated freezing and thawing.



Note

Best use within three months from the date of receipt of this protein.

## Applications

- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — UBE2V1	
Entrez GenelD	7335
GeneBank Accession#	BC000468
Protein Accession#	AAH00468
Gene Name	UBE2V1
Gene Alias	CIR1, CROC-1, CROC1, UBE2V, UEV-1, UEV1, UEV1A
Gene Description	ubiquitin-conjugating enzyme E2 variant 1
Omim ID	<u>602995</u>
Gene Ontology	Hyperlink
Gene Summary	Ubiquitin-conjugating E2 enzyme variant proteins constitute a distinct subfamily within the E2 prot ein family. They have sequence similarity to other ubiquitin-conjugating enzymes but lack the cons erved cysteine residue that is critical for the catalytic activity of E2s. The protein encoded by this g ene is located in the nucleus and can cause transcriptional activation of the human FOS proto-onc ogene. It is thought to be involved in the control of differentiation by altering cell cycle behavior. Mu Itiple alternatively spliced transcripts encoding different isoforms have been described for this gen e. A pseudogene has been identified which is also located on chromosome 20. Co-transcription of this gene and the neighboring upstream gene generates a rare transcript (Kua-UEV), which en codes a fusion protein comprised of sequence sharing identity with each individual gene product. [provided by RefSeq
Other Designations	DNA-binding protein OTTHUMP00000031795 OTTHUMP00000031796 OTTHUMP0000003998 0