

## OriGene Technologies, Inc.

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## Product datasheet for RC223522L3V

## UBE2J2 (NM\_058167) Human Tagged ORF Clone Lentiviral Particle

## **Product data:**

Product Type:	Lentiviral Particles
Product Name:	UBE2J2 (NM_058167) Human Tagged ORF Clone Lentiviral Particle
Symbol:	UBE2J2
Synonyms:	NCUBE-2; NCUBE2; PRO2121
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
Tag:	Myc-DDK
ACCN:	NM_058167
ORF Size:	777 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC223522).
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. <u>More info</u>
OTI Annotation:	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
RefSeq:	<u>NM 058167.2</u>
RefSeq Size:	2267 bp
RefSeq ORF:	780 bp
Locus ID:	118424
UniProt ID:	<u>Q8N2K1</u>
Cytogenetics:	1p36.33
Domains:	UBCc
Protein Families:	Transmembrane



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<b>GRIGENE</b> UBE2J2 (NM_058167) Human Tagged ORF Clone Lentiviral Particle – RC223522L3V	
Protein Pathways:	Parkinson's disease, Ubiquitin mediated proteolysis
MW:	28.7 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. This enzyme is located in the membrane of the endoplasmic reticulum. Multiple alternatively spliced transcript variants have been found for this gene, but the full-length nature of some variants has not been defined. [provided by RefSeq, Jul 2008]

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