

## OriGene Technologies, Inc.

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## Product datasheet for RC211739L3V

## RNF22 (TRIM3) (NM\_006458) Human Tagged ORF Clone Lentiviral Particle

## **Product data:**

Product Type:	Lentiviral Particles
Product Name:	RNF22 (TRIM3) (NM_006458) Human Tagged ORF Clone Lentiviral Particle
Symbol:	RNF22
Synonyms:	BERP; HAC1; RNF22; RNF97
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
Tag:	Myc-DDK
ACCN:	NM_006458
ORF Size:	2232 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC211739).
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 006458.2</u>
RefSeq Size:	3059 bp
RefSeq ORF:	2235 bp
Locus ID:	10612
UniProt ID:	<u>075382</u>
Cytogenetics:	11p15.4
Domains:	zf-B_box, NHL, Filamin, RING, BBC
MW:	80.6 kDa



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Gene Summary:The protein encoded by this gene is a member of the tripartite motif (TRIM) family, also called<br/>the 'RING-B-box-coiled-coil' (RBCC) subgroup of RING finger proteins. The TRIM motif includes<br/>three zinc-binding domains, a RING, a B-box type 1 and a B-box type 2, and a coiled-coil<br/>region. This protein localizes to cytoplasmic filaments. It is similar to a rat protein which is a<br/>specific partner for the tail domain of myosin V, a class of myosins which are involved in the<br/>targeted transport of organelles. The rat protein can also interact with alpha-actinin-4. Thus it<br/>is suggested that this human protein may play a role in myosin V-mediated cargo transport.<br/>Alternatively spliced transcript variants encoding the same isoform have been identified.<br/>[provided by RefSeq, Jul 2008]

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