

# Product datasheet for RC402774

## p21 Ras (HRAS) (NM\_005343) Human Mutant ORF Clone

### **Product data:**

Product Type:	Mutant ORF Clones
Product Name:	p21 Ras (HRAS) (NM_005343) Human Mutant ORF Clone
Mutation Description:	A146V
Affected Codon#:	146
Affected NT#:	437
Nucleotide Mutation:	HRAS Mutant (A146V), Myc-DDK-tagged ORF clone of Homo sapiens v-Ha-ras Harvey rat sarcoma viral oncogene homolog (HRAS), transcript variant 1 as transfection-ready DNA
Effect:	Cosello syndrome
Symbol:	HRAS
Synonyms:	C-BAS/HAS; C-H-RAS; C-HA-RAS1; CTLO; H-RASIDX; HAMSV; HRAS1; p21ras; RASH1
E. coli Selection:	Kanamycin (25 ug/mL)
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
Tag:	Myc-DDK
ACCN:	NM_005343
ORF Size:	567 bp
<b>Restriction Sites:</b>	Sgfl-Mlul

#### OriGene Technologies, Inc.

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	p21 Ras (HRAS) (NM_005343) Human Mutant ORF Clone – RC402774
ORF Nucleotide Sequence:	<pre>&gt;RC402774 representing NM_005343 Red=Cloning site Blue=ORF Green=Tags(s)</pre>
	TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC GCC <mark>GCGATCGC</mark> C
	ATGACGGAATATAAGCTGGTGGTGGTGGGGGCGCCGGCGGTGTGGGCAAGAGTGCGCTGACCATCCAGCTGA TCCAGAACCATTTTGTGGACGAATACGACCCCACTATAGAGGATTCCTACCGGAAGCAGGTGGTCATTGA TGGGGAGACGTGCCTGTTGGACATCCTGGATACCGCCGGCCAGGAGGAGTACAGCGCCATGCGGGACCAG TACATGCGCACCGGGGAGGGCTTCCTGTGTGTGTTTGCCATCAACAACACCAAGTCTTTTGAGGACATCC ACCAGTACAGGGAGCAGATCAAACGGGTGAAGGACTCGGATGACGTGCCCATGGTGGTGGGGAACAA GTGTGACCTGGCTGCACGACTGTGGGAATCTCGGCAGGCTCAGGACCTCGCCCGAAGCTACGGCATCCC TACATCGAGACCTCGGTCAAGACCCGGCAGGGAGTGGAGGATGCCTTCTACACGTTGGTGGGGAACCA GGCAGCACAAGCTGCGGCTGAAGACCCCGCAGGGAGTGGAGGATGCCTTCTACACGTTGGTGCGTGAGATCC GGCAGCACAAGCTGCGGAAGCTGAACCCTCCTGATGAGAGTGGCCCCGGCTGCATGAGCTGCAAGTGTGT GCTCTC
	<b>AGCGGACCG</b> ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC TGGATTACAAGGATGACGACGA TAAG <b>GTTTAA</b>
Protein Sequence:	e: >RC402774 representing NM_005343 Red=Cloning site Green=Tags(s)
	MTEYKLVVVGAGGVGKSALTIQLIQNHFVDEYDPTIEDSYRKQVVIDGETCLLDILDTAGQEEYSAMRDQ YMRTGEGFLCVFAINNTKSFEDIHQYREQIKRVKDSDDVPMVLVGNKCDLAARTVESRQAQDLARSYGIP YIETSVKTRQGVEDAFYTLVREIRQHKLRKLNPPDESGPGCMSCKCVLS
	SGPTRTRRLEQKLISEEDLAANDILDYKDDDDKV
<b>Restriction Sites:</b>	Sgfl-Mlul
Cloning Scheme:	Cloning sites used for ORF Shuttling: Sgf I ORF Mlu I GCGATCGC C ATG NIRN ACG CGT
	Kozac         Consensus         EcoRI       BamHI Kpn I       RBS       SgfI         CTATASGEGGEGGGAATTOGTOGACTGGATGGGGAACTGGGGGGAACTGGGCGGGGAATTOGTOGGACTGGGGGGGGGG
	ORF <u>Miu i Noti Xhol</u> Myc.Tag ACG CGT ACG CGG CCC CTC GAG CAA CTC ATC TCA GAA GAG T R T R P L E Q K L I S E E

 EcoR V
 Flag.Tag
 Pme I
 Fse I

 GAT CTG GCA GCA AAT GAT ATC CTG GAT TAC AAG GAT GAC GAC GAC GAC GAT AAG GTT TAA ACGGCCGGGCC
 D
 L
 A
 N
 D
 I
 L
 D
 Y
 K
 D
 D
 D
 K
 V
 stop

\* The last codon before the Stop codon of the ORF

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OTI Disclaimer:	Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <u>custsupport@origene.com</u> or by calling 301.340.3188 option 3 for pricing and delivery.
	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.
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).
RefSeq:	<u>NP 005334</u>
RefSeq Size:	567 bp
RefSeq ORF:	570 bp
Locus ID:	3265
Cytogenetics:	11p15.5
Protein Families:	Druggable Genome
Protein Pathway	S: Acute myeloid leukemia, Axon guidance, B cell receptor signaling pathway, Bladder cancer, Chemokine signaling pathway, Chronic myeloid leukemia, Endocytosis, Endometrial cancer, ErbB signaling pathway, Fc epsilon RI signaling pathway, Focal adhesion, Gap junction, Glioma, GnRH signaling pathway, Insulin signaling pathway, Long-term depression, Long-term potentiation, MAPK signaling pathway, Melanogenesis, Melanoma, Natural killer cell mediated cytotoxicity, Neurotrophin signaling pathway, Non-small cell lung cancer, Pathways in cancer, Prostate cancer, Regulation of actin cytoskeleton, Renal cell carcinoma, T cell receptor signaling pathway, Thyroid cancer, Tight junction, VEGF signaling pathway
MW:	20.8 kDa

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#### CRIGENE p21 Ras (HRAS) (NM\_005343) Human Mutant ORF Clone – RC402774

Gene Summary:This gene belongs to the Ras oncogene family, whose members are related to the<br/>transforming genes of mammalian sarcoma retroviruses. The products encoded by these<br/>genes function in signal transduction pathways. These proteins can bind GTP and GDP, and<br/>they have intrinsic GTPase activity. This protein undergoes a continuous cycle of de- and re-<br/>palmitoylation, which regulates its rapid exchange between the plasma membrane and the<br/>Golgi apparatus. Mutations in this gene cause Costello syndrome, a disease characterized by<br/>increased growth at the prenatal stage, growth deficiency at the postnatal stage,<br/>predisposition to tumor formation, cognitive disability, skin and musculoskeletal<br/>abnormalities, distinctive facial appearance and cardiovascular abnormalities. Defects in this<br/>gene are implicated in a variety of cancers, including bladder cancer, follicular thyroid cancer,<br/>and oral squamous cell carcinoma. Multiple transcript variants, which encode different<br/>isoforms, have been identified for this gene. [provided by RefSeq, Jul 2008]

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