

## OriGene Technologies, Inc.

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## Product datasheet for SC335426

## Retinoic Acid Receptor beta (RARB) (NM\_001290276) Human Untagged Clone

## **Product data:**

Product Type:	Expression Plasmids
Product Name:	Retinoic Acid Receptor beta (RARB) (NM_001290276) Human Untagged Clone
Tag:	Tag Free
Symbol:	Retinoic Acid Receptor beta
Synonyms:	HAP; MCOPS12; NR1B2; RARbeta1; RRB2
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
runy sequenceu Okr.	Blue=Insert sequence Red=Cloning site Green=Tag(s) GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC ATGATTTACACTTGTCACCGAGATAAGAACTGTGTTATTAATAAAGTCACCAGGAATCGATGCCAATAC TGTCGACTCCAGAAGTGCTTTGAAGTGGGAATGTCCAAAGAATCTGTCAGGAATGACAGGAACAAGAAA AAGAAGGAGACTTCGAAGCAAGAATGCACAGAGAGCTATGAAATGACAGCTGAGTTGGACGATCTCACA GAGAAGATCCGAAAAGCTCACCAGGAAATGCACAGAGAGCTATGAAATGACAGCTGAGTTGGACGATCTCACA GAGAAGATCCGAAAAGCTCACCAGGAAACTTTCCCTTCACTCTGCCAGCTGGGTAAATACACCACGAAT TCCAGTGCTGACCATCGAGTCCGACTGGACCTGGGCCTCTGGGACAAATTCAGTGAACTGGCCACCAAG TGCATTATTAAGATCGTGGAGTTTGCTAAACGTCTGCCTGGTTTCACTGGCTTGACCATCGCAGACCAA ATTACCCTGCTGAAGGCCGCCTGCCTGGACATCCTGATTCTTAGAATTGCACCAGGTATACCCCAGAA CAAGACACCATGACTTTCTCAGACGGCCTTACCCTAAATCGAACTCAGATGCACAATGCTGGATTTGGT CCTCTGACTGACCTTGTGTTCACCTTTGCAACCAGCTCCTGCCTTTGGAAATGCACAAGCTGGACACAGAAACA GGCCTTCTCAGTGCCATCTGCTTAATCTGTGGAGACCGCCAGGACCTTGAGGAACCGACAAAAGTAGAT
	AAGCTACAAGAACCATTGCTGGAAGCACTAAAAATTTATATCAGAAAAAGACGACCCAGCAAGCCTCAC ATGTTTCCAAAGATCTTAATGAAAATCACAGATCTCCGTAGCATCAGTGCTAAAGGTGCAGAGCGTGTA ATTACCTTGAAAATGGAAATTCCTGGATCAATGCCACCTCTCATTCAAGAAATGCTGGAGAATTCTGAA GGACATGAAACCCTTGACCCCAAGTTCAAGTGGGAACACAGCAGGAGCACAGTCCTAGCATCTCACCCAGC TCAGTGGAAAACAGTGGGGTCAGTCAGTCACCACTCGTGCAATAA ACGCGTACGCGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC
Restriction Sites:	Sgfl-Mlul
ACCN:	NM_001290276
Insert Size:	1011 bp



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	Retinoic Acid Receptor beta (RARB) (NM_001290276) Human Untagged Clone – SC335426
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 M	<ul> <li>ethod: 1. Centrifuge at 5,000xg for 5min.</li> <li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>3. Close the tube and incubate for 10 minutes at room temperature.</li> <li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>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.</li> </ul>
RefSeq:	<u>NM 001290276.1</u>
RefSeq Size:	3222 bp
RefSeq ORF:	1011 bp
Locus ID:	5915
UniProt ID:	<u>P10826</u>
Cytogenetics:	3p24.2
Protein Families:	Druggable Genome, Nuclear Hormone Receptor, Transcription Factors
Protein Pathways	: Non-small cell lung cancer, Pathways in cancer, Small cell lung cancer
MW:	37.9 kDa
Gene Summary:	This gene encodes retinoic acid receptor beta, a member of the thyroid-steroid hormone receptor superfamily of nuclear transcriptional regulators. This receptor localizes to the cytoplasm and to subnuclear compartments. It binds retinoic acid, the biologically active form of vitamin A which mediates cellular signalling in embryonic morphogenesis, cell growth and differentiation. It is thought that this protein limits growth of many cell types by regulating gene expression. The gene was first identified in a hepatocellular carcinoma where it flanks a hepatitis B virus integration site. Alternate promoter usage and differential splicing result in multiple transcript variants. [provided by RefSeq, Mar 2014] Transcript Variant: This variant (1, also known as beta-2) is a predominant transcript. This variant can initiate translation from an upstream AUG site and also from a downstream, in- frame AUG site (PMID: 12118004). The isoform (2, also known as beta', formerly, beta-4) represented in this RefSeq is derived from the downstream AUG start codon, and has a

shorter N-terminus, compared to isoform 1. Three variants in this gene encode the same isoform 2. 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.

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