

Product datasheet for MC227427

Rorb (NM_001289921) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Rorb (NM_001289921) Mouse Untagged Clone

Tag: Tag Free Symbol: Rorb

Synonyms: hstp; Nr1f2; Rorbeta; RZR-beta; RZRB

Mammalian Cell Neo

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >MC227427 representing NM_001289921

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ATGTCAAGAGATGCTGTAAAGTTCGGGAGGATGTCCAAGAAGCAGCGGGACAGCCTGTATGCTGAGGTGC AGAAGCATCAGCAAAGGCTGCAGGAGCAGCGGCAGCAGCAGAGTGGGGAGGCCGGAGGCCCTCGCCAGGGT GTACAGCAGCAGCATTAGCAATGGCCTCAGCAACCTGAACACCGAGACCGGCGCACATACGCCAACGGG CACGTCATTGACCTGCCCAAGTCCGAAGGTTATTACAGCATAGATTCCGGTCAGCCGTCTCCCGATCAGT ATTGCACAGAACATCATTAAGTCCCATTTGGAGACATGTCAGTACACCATGGAAGAACTCCATCAGCTGG CATGGCAGACCCACACCTACGAGGAAATCAAGGCGTATCAAAGCAAGTCCAGGGAGGCTCTGTGGCAGCA GTGTGCCATCCAGATCACCCATGCTATCCAGTACGTGGTGGAGTTCGCCAAGCGGATAACAGGCTTCATG GAGCTGTGTCAGAACGATCAGATCTTACTTCTGAAGTCAGGTTGCTTGGAAGTGGTTTTAGTGAGAATGT GTCGTGCCTTCAACCCATTAAACAACACTGTTCTGTTTGAAGGAAAATATGGAGGAATGCAAATGTTCAA AGCCTTAGGTTCGGATGACCTAGTGAATGAAGCATTTGACTTTGCGAAGAATCTGTGTTCCTTGCAGCTG ACTGAGGAAGAGTTGCTCTGTTCTCCTCTGTTGTTCTGATATCCCCAGACCGAGCCTGGCTGATCGAAC CAAGAAAAGTCCAGAAGCTTCAGGAAAAGATTTATTTTGCACTGCAACATGTGATTCAGAAGAACCACCT GGATGATGAGACCCTGGCAAAGTTAATAGCCAAGATACCAACTATCACGGCAGTCTGCAACTTGCATGGG GAGAAGCTGCAGGTATTTAAGCAGTCTCATCCAGACATAGTGAATACACTGTTTCCTCCATTGTACAAGG AGCTCTTTAATCCTGACTGTGCTGCGGTCTGCAAATGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATTACAAGGATGACGACGATAAGGTTTAA



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Restriction Sites: Sgfl-Mlul

ACCN: NM_001289921

Insert Size: 1158 bp

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).

OTI Annotation: Clone contains native stop codon, and expresses the complete ORF without any c-terminal

tag.

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 Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

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.

Note: Plasmids are not sterile. For experiments where strict sterility is required, filtration with

0.22um filter is required.

RefSeq: <u>NM 001289921.1</u>, <u>NP 001276850.1</u>

 RefSeq Size:
 8808 bp

 RefSeq ORF:
 1158 bp

 Locus ID:
 225998

 UniProt ID:
 Q8R1B8

 Cytogenetics:
 19 B

Gene Summary: The protein encoded by this gene is a member of the NR1 subfamily of nuclear hormone

receptors. It is a DNA-binding protein that can bind as a monomer or as a homodimer to hormone response elements upstream of several genes to enhance the expression of those

genes. The encoded protein has been shown to interact with NM23-2, a nucleoside

diphosphate kinase involved in organogenesis and differentiation, and to help regulate the expression of some genes involved in circadian rhythm. Three transcript variants encoding

different isoforms have been found for this gene. [provided by RefSeq, Feb 2014]

Transcript Variant: This variant (3) contains an alternate exon compared to variant 2. The resulting isoform (3) is shorter at the N-terminus compared to 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.