

Product datasheet for SC327494

PRKAR1B (NM 001164758) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: PRKAR1B (NM_001164758) Human Untagged Clone

Tag: Tag Free

Symbol: PRKAR1B

Synonyms: PRKAR1

Mammalian Cell Neomycin

Selection:

Vector:

pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC327494 representing NM_001164758.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGGCCTCCCGCCCGCCTGCCCCTCGGAGGAGGAGGAGGAGCCTGAAGGGCTGTGAGCTGTACGTGCAG CTGCACGGGATCCAGCAGGTCCTCAAAGACTGTATCGTCCACCTCTGCATCTCCAAGCCCGAACGCCCC AAGTCAAACTCACAGTCGGACTCCCATGATGAGGAGGTGTCGCCCACCCCCCGAACCCTGTGGTGAAG GCCCGCCGCCGAGGAGGAGGCGTGAGTGCCGAGGTGTACACCGAGGAGGACGCCGTGTCCTACGTCAGG AAGGTGATTCCCAAGGACTACAAAACCATGACTGCGCTGGCCAAGGCCATCTCCAAGAACGTGCTCTTC GAGACTGTTATACAGCAAGGGAATGAAGGAGACAACTTCTATGTCGTTGATCAAGGGGAAGTGGATGTG GGCACCCCAGGGCTGCGACCGTGAAAGCCAAGACGGACCTCAAGCTCTGGGGGATCGACCGGGACAGC TACCGGCGCATCCTTATGGGCAGCACGCTGAGGAAACGCAAGATGTACGAGGAGTTCCTCAGCAAGGTC TCCATCCTAGAGTCCCTGGAGAAGTGGGAGCGTCTGACCGTGGCGGATGCGCTGGAGCCCGTCCAGTTT GAAGATGGAGAAAATTGTGGTCCAGGGAGAGCCTGGGGACGACTTTTACATCATCACGGAGGGCACC GCGTCCGTGCTGCAGCGCCGGTCCCCCAATGAGGAGTACGTGGAGGTGGGGCGCCTGGGACCCTCTGAC TACTTCGGGGAGATTGCACTGCTGAACCGGCCCCGGGCCACTGTCGTGGCCCGGGGGCCCCTC AAGTGTGTGAAGCTGGACCGGCCCCGCTTCGAGCGTGTGCTGGGGCCCTGCTCTGAGATCCTCAAGAGG

AACATTCAGCGTTACAACAGCTTCATCTCCCTCACCGTCTGA

 ${\color{blue} \textbf{ACGCGTACGCGGCCGCTC} \textbf{GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT} }$

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul



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PRKAR1B (NM_001164758) Human Untagged Clone - SC327494

ACCN: NM 001164758

Insert Size: 1146 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: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

> into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA

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.

NM 001164758.1 RefSeq:

RefSeq Size: 2488 bp RefSeq ORF: 1146 bp 5575 Locus ID: **UniProt ID:** P31321 7p22.3

Cytogenetics:

Protein Families: Druggable Genome

Protein Pathways: Apoptosis, Insulin signaling pathway

MW:

The protein encoded by this gene is a regulatory subunit of cyclic AMP-dependent protein **Gene Summary:**

> kinase A (PKA), which is involved in the signaling pathway of the second messenger cAMP. Two regulatory and two catalytic subunits form the PKA holoenzyme, disbands after cAMP binding. The holoenzyme is involved in many cellular events, including ion transport, metabolism, and transcription. Several transcript variants encoding the same protein have

been found for this gene. [provided by RefSeq, Aug 2015]

Transcript Variant: This variant (3) differs in the 5' UTR compared to variant 1. All of the variants encode the same protein. Sequence Note: This RefSeg 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.