

Product datasheet for SC330229

ROD1 (PTBP3) (NM_001244898) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: ROD1 (PTBP3) (NM_001244898) Human Untagged Clone
Tag: Tag Free
Symbol: PTBP3
Synonyms: ROD1
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC330229 representing NM_001244898.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

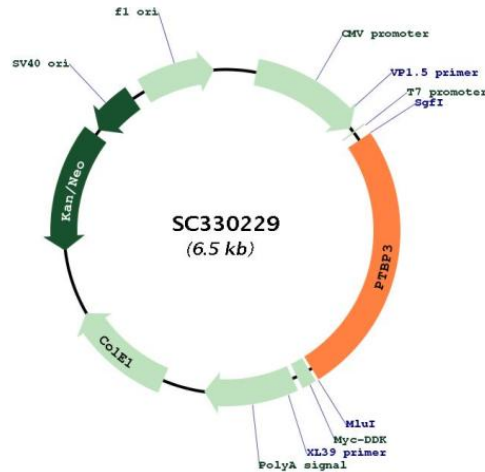
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Restriction Sites: Sgfl-Mlul



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Plasmid Map:


ACCN: NM_001244898

Insert Size: 1677 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).

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.

RefSeq: [NM_001244898.1](#)

RefSeq Size: 8009 bp

RefSeq ORF: 1677 bp

Locus ID: 9991

UniProt ID: [O95758](#)

Cytogenetics: 9q32

Protein Families: Druggable Genome

MW: 60.4 kDa

Gene Summary: The protein encoded by this gene binds RNA and is a regulator of cell differentiation. The encoded protein preferentially binds to poly(G) and poly(U) sequences in vitro. Several transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Oct 2011]
Transcript Variant: This variant (6) encodes the longest isoform (6). 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.