

Product datasheet for **SC311142**

PPAN-P2RY11 (NM_001040664) Human Untagged Clone

Product data:

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| Product Type: | Expression Plasmids |
| Product Name: | PPAN-P2RY11 (NM_001040664) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | PPAN-P2RY11 |
| Synonyms: | BXDC3; P2RY11; P2Y11; PPAN; Ssf-1; SSF1 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | pCMV6-Entry (PS100001) |
| E. coli Selection: | Kanamycin (25 ug/mL) |



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Fully Sequenced ORF: >SC311142 representing NM_001040664.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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GCTCGTTTAGTGAACCGTCAGAATTTTGTAAACGACTCACTATAGGGCGGCCGGGAATTCGTGACTG
GATCCGGTACCGAGGAGATCTGCCGCCCGGATCGCC
ATGGGACAGTCAGGGAGGTCCTCCGGCACCAGAAGCGCGCCCGCCAGGCGCAGCTCCGCAACCTCGAG
GCCTATGCCCGCGAACCCGCACTCGTTTCGTGTTACGCGAGGCTGCACGGGTGCAACATCCGGCAGCTC
AGCCTGGACGTGCGGCGGGTCATGGAGCCGCTCACTGCCAGCCGTCTGCAGGTTCTGAAGAAGAAGTCTG
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ACCAATGTCTACTTTAAGCTGATGCGCCTCCAGGAGGCCACCTTGACCTTCCAGGTCAAGAAGTAC
TCGCTGGTGGTGTGTTCTCCTCACTGCGCCGGCACCGCATGCACGAGCAGCAGTTTGGCCACCCA
CCCCTCCTGGTACTCAACAGCTTTGGCCCCATGGTATGCATGTGAAGCTCATGGCCACCATGTTCCAG
AACCTGTTCCCTCCATCAACGTGCACAAGGTGAACCTGAACACCATCAAGCGTGCCTCCTCATCGAC
TACAACCCGACTCCAGGAGCTGGACTCCGCCACTATAGCATCAAAGTTGTTCTGTGGGCGCGAGT
CGCGGGATGAAGAAGCTGCTCCAGGAGAAGTCCCAACATGAGCCGCTGCAGGACATCAGCGAGCTG
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CCTCAGGCTGTCGCTGGCCGTGGCAACATGCGGGCCAGCAGAGTGCAGTGCAGGCTCACCAGATCGGC
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AACGTGGATGCTCGGGCGCTGGAGCACCCGCTGCCCGAGCTTTGCAGACATAGCCAGGCCACAGCA
GCCCTGGAGCTGGGGCCCTACGTGGGCTACCAGGTGATGCGGGCCCTCATGCCCTGGCCTTCTGTGTC
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GACAGCTGGAACCCAGAGGACGCCAAGAGCACTGGCCAAGCCCTGCCCTCAATGCCACAGCCGCCCT
AAACCGTCAGAGCCCCAGTCCCCTGAGCTGAGCCAATGA
ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCTGGAT
TACAAGGATGACGACGATAAGGTTAAACGGCCGGC
  
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Restriction Sites: SgfI-MluI
 ACCN: NM_001040664
 Insert Size: 2385 bp

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| 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: | <ol style="list-style-type: none">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_001040664.2 |
| RefSeq Size: | 3208 bp |
| RefSeq ORF: | 2385 bp |
| Locus ID: | 692312 |
| UniProt ID: | Q9NQ55 |
| Cytogenetics: | 19p13.2 |
| Protein Families: | Stem cell - Pluripotency, Transmembrane |
| MW: | 87.9 kDa |
| Gene Summary: | <p>This locus represents naturally occurring read-through transcription between the adjacent PPAN and P2RY11 genes. Alternative splicing results in two transcript variants, one of which encodes a fusion protein that shares sequence identity with each individual gene product. This transcript is found to be ubiquitously expressed and is up-regulated by agents inducing granulocytic differentiation. However, its functional significance in vivo remains unclear. [provided by RefSeq, Nov 2010]</p> <p>Transcript Variant: This variant (1) represents the shorter transcript but encodes the longer isoform (1), which is a fusion of the PPAN and P2RY11 products. 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.</p> |