

## Product datasheet for **SC128054**

### PPP2R2D (NM\_018461) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	PPP2R2D (NM_018461) Human Untagged Clone
Tag:	Tag Free
Symbol:	PPP2R2D
Synonyms:	B55D; B55delta; MDS026
Vector:	<u>pCMV6-XL4</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Cell Selection:	None
Fully Sequenced ORF:	>NCBI ORF sequence for NM_018461, the custom clone sequence may differ by one or more nucleotides

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ATGGCAGGAGCCGGAGGCGGGCTGCCCGGGCGGCAACGACTTCCAGTGGTCTTCTCGCAGGTCA
AGGGGGCCATCGACGAGGACGTGGCCGAAGCGGACATCATTTCCACCGTTGAGTTAATTACTCTGGAGA
TCTTCTTGCAACAGGAGACAAGGGCGGCAGAGTTGTTATTTTTTCAGCGTGAACAAGAGAATAAAAGCCG
CCTCATTCTAGGGGAGAATAAATGTTTACAGCACCTTTCAAAGTCATGAACCGGAGTTTGACTATTTGA
AAAGTCTAGAAATTGAGGAAAAAATTAATAAAATTAGGTGGTTACCACAACAGAATGCTGCTCATTTTCT
ACTGTCTACAAATGATAAACTATAAAATTATGAAAAATAAGTGAACGGGATAAAAGAGCAGAAGGTTAT
AACCTGAAAGACGAAGATGGAAGACTTCGAGACCCATTTAGGATCACGGCGCTACGGGTCCCAATATTGA
AGCCCATGGATCTTATGGTAGAAGCGAGTCCACGGCGAATTTTTGCAAAATGCTCACACATATCATATAAA
TTCCATTTTCAGTAAATAGTGATCATGAAACATATCTTTCTGCAGATGACCTGAGAATTAATTTATGGCAC
TTAGAAATCACAGATAGAAGCTTTAACATCGTGGACATCAAGCCTGCTAACATGGAGGAGCTGACCGAAG
TCATCACTGCAGCCGAGTTCCACCCGCACCAGTGAACGTGTTTCGCTACAGCAGTAGCAAAGGGACCAT
CCGCTGTGTGACATGCGCTCCTCGGCCCTGTGCGACAGACACTCCAAGTTTTTTGAAGAGCCTGAAGAT
CCCAGCAGTAGGTCCTTCTCAGAAAATAATTTTCATCCATATCCGATGTAATAATTCAGTCATAGTGGC
GGTACATGATGACCAGAGACTACCTGTGCGTGAAGGTGTGGGACCTCAACATGGAGAGCAGGCCGGTGGGA
GACCCACAGGTCCACGAGTACCTGCGCAGCAAGCTCTGCTCTCTATGAGAACGACTGCATCTTTGAC
AAGTTTGTGAGTGTGCTGGAACGGTTCGGATAGCGCCATCATGACCGGGTCTATAAACAATTCTTCAGGA
TGTTTGTAGAGACACGCGGAGGGATGTGACCCTGGAGGCTCGAGAGAGAGCAGCAAACCGCGGCCAG
CCTCAAACCCCGAAGGTGTGTACGGGGGTAAAGCGGAGAAAGACGAGATCAGTGTGGACAGTCTGGAC
TTCAACAAGAAGATCCTGCACACAGCCTGGCACCCCGTGACAATGTCATTGCCGTGGCTGCCACCAATA
ACTTGACATATCCAGGACAAAATCAACTAG
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<b>5' Read Nucleotide Sequence:</b>	<p>&gt;OriGene 5' read for NM_018461 unedited            ACGACTACTATAGGGCGGCCGGAATTCGCACGAGGCCCGCGCGGCGGCGCGGGCG            GCGCCGGCGGTGGTGCGGCCCGGGGGCTGAGCGCTCGGCTGCAGCGGCGGAGGCCGT            CTCCTGGTCTGCCCGGTCCCCGCCGCTCCCGCCCGGCTGCCATGGCAGGAGCCGGA            GCGGGCGGCTGCCCCGCGGGCGCAACGACTTCCAGTGGTGCTTCTCGCAGGTCAAGGGG            GCCATCGACGAGGACGTGGCCGAAGCGGACATATTTCCACCGTTGAGTTTAATTACTCT            GGAGATCTTTCTTGAACAGGAGACAAGGGCGGCAGAGTTGTTATTTTTCCAGCGTGAACAA            GAGAATAAAAGCCGCCCTCATTCTAGGGGAGAATAAATGTTTACAGCACCTTTCAAAGT            CATGAACCGGAGTTTGACTATTTGAAAAGTCTAGAAAATTGAGGAAAAAATTAATAAAATT            AGGTGGTTACCACAACAGAATGCTGCTCATTCTACTGTCTACAAATGATAAACTATA            AAATTATGAAAAAAGTGAACGGGATAAAAGAGCAGAAGTTAATCTGAAAGACGAA            GATGGAAGACTTCGAGACCCATTTAGGATCACGGCGCTACGGGTCCAATATTGAAGCCC            ATGGATCTTATGGTAGAAGCGAGTCCACGGCGAATTTTTGCAAATGCTCACACATATCAT            ATAAATCCATTTAGTAAATAGTGATCATGAAACATATCTTCTGCAAATGACCTGAGA            AATTAATTATGGCACTTAGAAAACACAGATAGAAGCTTTAACATCGGGGACATCAAGCCT            GCTAACATGGAGGAGCTACCGAAGTCATCACTGCAGCCGGAGTTCAACCGGACCAAGTGC            CACGTGTTCTGCTACAGCAAAGCAAAGT</p>
<b>3' Read Nucleotide Sequence:</b>	<p>&gt;Forward primer walk for NM_018461 unedited            NAAACAGCGAGCACGCGGCCCGCCCTCAACTCGGTAGTGTGTACGGGGGTAAGCGGNAG            GAAANACGAGCATCAGTGTGGACAGTCTGGACTTCAACAAAAAGATCCTGCACACAGCCT            GGCACCCCGTGGACAATGTCATTGCCGTGGCTGCCACCAATAACTTGTACATATTCAGG            AAAAACTCAACTAGAGACGCGAACACACAGGAGAAAGCCCGCTCCGCTGGAGGCCCGGTG            TGGTTCGCCCTCGGCGAGGCGGAGACAGGCGCTGCTGCTCACGTGGAGACGCTCTCGAA            GCAGAGTTGACGGACACTGCTCCAAAAGCGTGGGGGTATGTGTGCNNCATTTCTGTNTC            NNNNCTGCTNNTCCGTGNNTGGTCTTCTTCCACCCCCCCACCACGGGAGAAGACCCC            CAGAAACGACAACCCCCAGCAGACAAACCACCACCCTAATTACCCCCCCCCAACAA            CCGCACGAACCAATTACACCGGTGCCAAACACCCCCCGCACAAATACCTTAACAGCAC            GCCACGCTAAACAGAAGCGAACTCCACACATCAACACACCCCAACCCACCCCCCCAC            TCGCCACGAATATGTCACCTTTTCCACCCGACCAAGAGAGCATATAACCCACCGGAGC            GGCTATACCCCCCGGCACATGCCGAACCGACACCAACCGCCAGTTTTTCAACACG            AGTTCCTCCCTTGTGACACACCTTATATTTGAAATCCAAACACGAAGCAGCATGACCGCG            GACACGTTAGCCGANCCACAATAAAAACACGCCTATGCACCCTAGTATCAAAACCCACCG            AAATACGATGGATAACACGGGTTCCACAAACAATAGACCAACCGGTAATAAAAAAT</p>
<b>Restriction Sites:</b>	NotI-NotI
<b>ACCN:</b>	NM_018461
<b>Insert Size:</b>	4700 bp
<b>OTI Disclaimer:</b>	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).
<b>Components:</b>	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).

<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>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></ol>
<b>RefSeq:</b>	<a href="#">NM_018461.2</a> , <a href="#">NP_060931.2</a>
<b>RefSeq Size:</b>	2105 bp
<b>RefSeq ORF:</b>	1362 bp
<b>Locus ID:</b>	55844
<b>UniProt ID:</b>	<a href="#">Q66LE6</a>
<b>Cytogenetics:</b>	10q26.3
<b>Protein Families:</b>	Druggable Genome, Phosphatase
<b>Protein Pathways:</b>	Tight junction
<b>Gene Summary:</b>	<p>B regulatory subunit of protein phosphatase 2A (PP2A) that plays a key role in cell cycle by controlling mitosis entry and exit. The activity of PP2A complexes containing PPP2R2D (PR55-delta) fluctuate during the cell cycle: the activity is high in interphase and low in mitosis. During mitosis, activity of PP2A is inhibited via interaction with phosphorylated ENSA and ARPP19 inhibitors. Within the PP2A complexes, the B regulatory subunits modulate substrate selectivity and catalytic activity, and also may direct the localization of the catalytic enzyme to a particular subcellular compartment (By similarity).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (1) encodes the longer isoform (a).</p>