

Product datasheet for **SC300041**

PAX2 (NM_000278) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	PAX2 (NM_000278) Human Untagged Clone
Tag:	Tag Free
Symbol:	PAX2
Synonyms:	FSGS7; PAPRS
Mammalian Cell Selection:	Neomycin
Vector:	<u>PCMV6-Neo</u>
E. coli Selection:	Ampicillin (100 ug/mL)
Fully Sequenced ORF:	>OriGene ORF within SC300041 sequence for NM_000278 edited (data generated by NextGen Sequencing)

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ATGGATATGCACTGCAAAGCAGACCCCTTCTCCGCGATGCACNNNGGCACGGGGGTGTG
AACCCAGCTCGGGGGGGTGTGGTGAACGCGCCGCCCTACCCGACGTGGTGAGGCAGCGC
ATCGTGGAGCTGGCCACCAGGGTGTGCGGCCCTGTGACATCTCCCGGCAGCTGCGGGTC
AGCCACGGCTGTGTGACAAAATCCTGGGCAGGTACTACGAGACCGGCAGCATCAAGCCG
GGTGTGATCGGTGGCTCCAAGCCAAAAGTGGCGACGCCAAAAGTGGTGGACAAGATTGCT
GAATACAAACGACAGAACCCGACTATGTTCCGCTGGGAGATTGAGACCGGCTCCTGGCC
GAGGGCATCTGTGACAATGACACAGTGGCCAGCGTCTCTCCATCAACAGAATCATCCGG
ACCAAAGTTCAGCAGCCTTTCCACCCAACGCCGATGGGGCTGGGACAGGAGTGACCGCC
CCTGGCCACACCATTGTTCCAGCACGGCCTCCCCTCCTGTTTCCAGCGCCTCCAATGAC
CCAGTGGGATCCTACTCCATCAATGGGATCCTGGGGATTCTCGCTCCAATGGTGAGAAG
AGGAAACGTGATGAAGATGTGTCTGAGGGCTCAGTCCCCAATGGAGATCCAGAGTGGT
GTGGACAGTTTGCGAAGCACTTGCAGCTGACACCTTACCCAGCAGCAGCTGGAAGCT
TTGGATCGGGTCTTTGAGCGTCTTCCCTACCCTGACGTCTTCCAGGCATCAGAGCACATC
AAATCAGAACAGGGGAACGAGTACTCCCTCCAGCCCTGACCCCTGGGCTTGATGAAGTC
AAGTCGAGTCTATCTGCATCCACCAACCCTGAGCTGGGCAGCAACGTGTGAGGCACACAG
ACATACCCAGTTGTGACTGGTGTGACATGGCGAGCACCCTCTGCCTGGTTACCCCTCCT
CACGTGCCCCCACTGGCCAGGGAAGCTACCCACCTCCACCCTGGCAGGAATGGTGCCT
GGGAGCGAGTTCTCCGGCAACCCGTACAGCCACCCCAAGTACACGCGCTACAACGAGGCT
TGGAGATTGAGCAACCCCGCCTTACTAAGTTCCCTTATTATTATAGTGCCGCCCCCGG
TCCGCCCTGCCGCTGCTGCCGCTGCCTATGACCGCCACTAG

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Clone variation with respect to NM_000278.3
44 c=>n;45 a=>n;46 g=>n

Restriction Sites: Please inquire



[View online »](#)

ACCN:	NM_000278
Insert Size:	1300 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:	The open reading frame of this TrueClone was fully sequenced and found to differ from the protein associated to this reference by three amino acids.
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_000278.2 , NP_000269.2
RefSeq Size:	4207 bp
RefSeq ORF:	1182 bp
Locus ID:	5076
UniProt ID:	Q02962
Cytogenetics:	10q24.31
Protein Families:	Druggable Genome
Gene Summary:	<p>PAX2 encodes paired box gene 2, one of many human homologues of the <i>Drosophila melanogaster</i> gene <i>prd</i>. The central feature of this transcription factor gene family is the conserved DNA-binding paired box domain. PAX2 is believed to be a target of transcriptional suppression by the tumor suppressor gene WT1. Mutations within PAX2 have been shown to result in optic nerve colobomas and renal hypoplasia. Alternative splicing of this gene results in multiple transcript variants. [provided by RefSeq, Dec 2014]</p> <p>Transcript Variant: This variant (b) lacks an alternate in-frame exon and uses an alternate splice site in the 3' coding region, compared to variant e. This results in a protein (isoform b) with a shorter, distinct C-terminus compared to isoform e. 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. CCDS Note: The coding region has been updated to add an additional glycine residue in the protein C-terminal region, supported by the available transcript and conservation data.</p>