

Product datasheet for SC329840

PIG3 (TP53I3) (NM_001206802) Human Untagged Clone

Product data:

Insert Size:

Product Type: Expression Plasmids Product Name: PIG3 (TP53I3) (NM_001206802) Human Untagged Clone Tag: Tag Free Symbol: PIG3 PIG3 Synonyms: Mammalian Cell Neomycin Selection: Vector: pCMV6-Entry (PS100001) E. coli Selection: Kanamycin (25 ug/mL) **Fully Sequenced ORF:** >SC329840 representing NM_001206802. Blue=Insert sequence Red=Cloning site Green=Tag(s) GCTCGTTTAGTGAACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTG GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC ATGTTAGCCGTGCACTTTGACAAGCCGGGAGGACCGGAAAACCTCTACGTGAAGGAGGTGGCCAAGCCG AGCCCGGGGGGAGGGTGAAGTCCTCCTGAAGGTGGCGGCCAGCGCCCTGAACCGGGCGGACTTAATGCAG AGACAAGGCCAGTATGACCCACCTCCAGGAGCCAGCAACATTTTGGGACTTGAGGCATCTGGACATGTG GCAGAGCTGGGGCCTGGCTGCCAGGGACACTGGAAGATCGGGGGACACAGCCATGGCTCTGCTCCCCGGT GGGGGCCAGGCTCAGTACGTCACTGTCCCCGAAGGGCTCCTCATGCCTATCCCAGAGGGATTGACCCTG ACCCAGGCTGCAGCCATCCCAGAGGCCTGGCTCACCGCCTTCCAGCTGTTACATCTTGTGGGAAATGTT CAGGCTGGAGACTATGTGCTAATCCATGCAGGACTGAGTGGTGTGGGCACAGCTGCTATCCAACTCACC CGGATGGCTGGAGCTATTCCTCTGGTCACAGCTGGCTCCCAGAAGAAGCTTCAAATGGCAGAAAAGCTT GGAGCAGCTGCTGGATTCAATTACAAAAAAGAGGATTTCTCTGAAGCAACGCTGAAATTCACCAAAGTA CAAGCAAATGCTGGTGAATGCTTTCACGGAGCAAATTCTGCCTCACTTCTCCACGGAGGGCCCCCAACG TCTGCTGCCGGTTCTGGACAGAATCTACCCAGTGACCGAAATCCAGGAGGCCCATAA ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC **Restriction Sites:** Sgfl-Mlul ACCN: NM 001206802

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747 bp

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ORIGENE PIG3 (T	P53I3) (NM_001206802) Human Untagged Clone – SC329840
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:	 Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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:	<u>NM 001206802.2</u>
RefSeq Size:	961 bp
RefSeq ORF:	747 bp
Locus ID:	9540
UniProt ID:	<u>Q53FA7</u>
Cytogenetics:	2p23.3
Protein Families:	Druggable Genome
Protein Pathways:	p53 signaling pathway
MW:	25.4 kDa
Gene Summary:	The protein encoded by this gene is similar to oxidoreductases, which are enzymes involved in cellular responses to oxidative stresses and irradiation. This gene is induced by the tumor suppressor p53 and is thought to be involved in p53-mediated cell death. It contains a p53

in cellular responses to oxidative stresses and irradiation. This gene is induced by the tumor suppressor p53 and is thought to be involved in p53-mediated cell death. It contains a p53 consensus binding site in its promoter region and a downstream pentanucleotide microsatellite sequence. P53 has been shown to transcriptionally activate this gene by interacting with the downstream pentanucleotide microsatellite sequence. The microsatellite is polymorphic, with a varying number of pentanucleotide repeats directly correlated with the extent of transcriptional activation by p53. It has been suggested that the microsatellite polymorphism may be associated with differential susceptibility to cancer. Alternatively spliced transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, May 2011]

Transcript Variant: This variant (3) lacks a coding exon in the 3' region, compared to variant 1. The resulting isoform (2) has a shorter and distinct C-terminus, compared to isoform 1. This variant lacks the 5'-most non-coding exon because of alternate splicing, the extension at the 5' end is uncertain.

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