

Product datasheet for **SC324589**

OGDH (NM_002541) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	OGDH (NM_002541) Human Untagged Clone
Tag:	Tag Free
Symbol:	OGDH
Synonyms:	AKGDH; E1k; KGD1; OGDC; OGDH2
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC (PS100020)
E. coli Selection:	Ampicillin (100 ug/mL)

Fully Sequenced ORF: >OriGene sequence for NM_002541.2
 CTGAGCCGGAGACAGGCAGTTGTGAAAACTTCAGGACAAAAATGTTTCATTTAAGGACT
 TGTGCTGCTAAGTTGAGACCATTGACGGCTTCCCAGACTGTTAAGACATTTTCACAAAAC
 AGACCAGCAGCAGCTAGGACATTTCAACAGATTGGTGCTATTCTGCACCTGTTGCTGCT
 GAGCCCTTTCTCAGTGGGACTAGTTCGAACTATGTGGAGGAGATGTAAGTGTGCTGGCTG
 GAAAAACCCAAAAGTGTACATAAGTCATGGGACATTTTTTTTCGCAACACGAATGCCGGA
 GCCCCACCGGGCACTGCCTACCAGAGTCCCCTTCCCCTGAGCCGAGGCTCCCTGGCTGCT
 GTGGCCCATGCACAGTCCCTGGTAGAAGCACAGCCCAACGTGGACAAGCTCGTGGAGGAC
 CACCTGGCAGTGCAGTCGCTCATCAGGGCATATCAGATACGAGGGCACCATGTAGCACAG
 CTGGACCCCTGGGGATTTGGATGCTGATCTGGACTCCTCCGTGCCCGCTGACATTATC
 TCATCCACAGACAACTGGGTTCTATGGCCTGGATGAGTCTGACCTCGACAAGGTCTTC
 CACTTGCCACCACCCTTTTCATCGGGGACAGGAATCAGCACTTCTCTGCGGGAGATC
 ATCCGTGGCTGGAGATGGCCTACTGCCAGCATATTGGGGTGGAGTTCATGTTTCAAT
 GACCTGGAGCAGTGCCAGTGGATCCGGCAGAAGTTTGAGACCCCTGGGATCATGCAGTTC
 ACAAATGAGGAGAAACGGACCCTGCTGGCCAGGCTTGTGCGGTCCACCAGGTTTGAGGAG
 TTCCTACAGCGGAAGTGGTCTCTGAGAAGCGCTTTGGTCTAGAAGGCTGCGAGGTAAGT
 ATCCCTGCCCTCAAGACCATCATTGACAAGTCTAGTGAGAATGGCGTGGACTACGTGATC
 ATGGGCATGCCACACAGAGGGCGGCTGAACGTGCTTCAAATGTCATCAGGAAGGAGCTG
 GAACAGATCTTCTGTCAATTCGATTCAAAGCTGGAGGCAGCTGATGAGGGCTCCGGAGAT
 GTGAAGTACCACCTGGGCATGTATCACCGCAGGATCAATCGTGTACCGACAGGAACATT
 ACCTTGTCTTGGTGGCCAACCCCTTCCCACCTTGAAGCCGCTGACCCCGTGGTGTGGGC
 AAGACCAAAGCCGAACAGTTTTACTGTGGCGACTGAAGGGAAAAAGGTGATGTCATC
 CTGTTGATGGGGATGCTGCATTTGCTGGCCAGGGCATTGTGTACGAGACCTTCCACCTC
 AGCGACCTGCCATCCTACACAACCTATGGCACCGTGCACGTGGTCAACAACCCAGATC
 GGCTTACCACCGACCCTCGGATGGCCCGCTCCTCCCCCTACCCACTGACGTGGCCCGA
 GTGGTGAATGCCCCATTTTCCACGTGAACTCAGATGACCCCGAGGCTGTCATGTACGTG
 TGCAAAGTGGCGCCGAGTGGAGGAGCACCTTCCACAAGGACGTGTTGTCGATTTGGTG



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TGTTACCGGCGCAACGGCCACAACGAGATGGATGAGCCCATGTTACGCGAGCCGCTCATG
TACAAGCAGATCCGCAAGCAGAAGCCTGTGTTACAGAAGTACGCTGAGCTGCTGGTGTGC
CAGGGTGTGGTCAACCAGCCTGAGTATGAGGAGGAAATTTCCAAGTATGATAAGATCTGT
GAGGAAGCTTTTGCCAGATCTAAAGATGAGAAGATCTTGACATTAAGCACTGGCTGGAC
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GAAAACCTTACTATTCATGGAGGGCTGAGCCGGATCTTGAAGACTCGTGGGAAATGGTG
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GAGGGCATCCACATTCGCTGAGCGGCCAGGACGTGGAGCGGGGCACATTCAGCCACCGC
CACCATGTGCTCCATGACCAGAATGTGGACAAGAGAACCCTGCATCCCCATGAACCATCTC
TGGCCCAATCAGGCCCTTATACTGTGTGCAACAGCTCACTGTCTGAGTACGGCGTGTG
GGCTTTGAGCTGGGCTTCGCCATGGCCAGTCTAATGCCCTGGTCTCTGGGAAGCCAA
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GCCAAGTGGGTGCGGCAGAATGGCATCGTGTGCTGCTGCCCCATGGCATGGAGGGCATG
GGTCCAGAACATTCCTCCGCCGCCAGAGCGGTTCTTGACAGATGTGCAACGATGACCCA
GATGTCCTGCCAGACCTTAAAGAAGCCAACTTCGACATCAATCAGCTATATGACTGCAAT
TGGGTTGTTGTCAACTGCTCCACTCCTGGCAACTTCTTCCACGTGCTACGACGCCAGATC
CTGCTGCCATTCGGAAGCCGTTAATTATCTTACCCCCAAATCCCTGTTGCGCCACCCC
GAGGCCAGATCCAGCTTTGATGAGATGCTTCCAGGAACCCACTTCAGCGGGTGTATCCCA
GAAGATGGCCCTGCAGCTCAGAACCAGAAAATGTCAAAGGCTTCTCTTCTGCACCGGC
AAAGTGTATTATGACCTACCCGGGAGCGCAAAGCACGCGACATGGTGGGGCAGGTGGCC
ATCACAAGGATTGACAGCTGTGCCATTCCTTTCCTGCTGCTGAGGAGGTGCAG
AAGTACCCCAATGCTGAGCTGGCTGGTCCAGGAGGAGCACAAGAACCAAGGCTACTAT
GACTACGTGAAGCCAAGACTTCGGACCACCATCAGCCGCCCAAGCCCGTCTGGTATGCC
GGCCGGGACCCAGCGGCTGCTCCAGCCACCGGCAACAAGAAGACCCACCTGACGGAGCTG
CAGCGCTCCTGGACACGGCCTTCGACCTGGACGTCTTCAAGAACTTCTCGTAGATGCTG
CCTAGGGTTGCTTGGGCCACTGCCCTCTCCACACCCATGACTGCCCTTGTCTTCAACT
AAAGAATAGTGCCTCAGCGCTGCCACACCACCGCCCTCTCGTGTGCCACCACCCCTC
CCTCTGCTCTCATAGGAGTTAGGCTGTCGTCCCTCCAGTGTGGTGGCCACAGGC
CACACGCTGCCAGGCTCTGCTGACTTCTGAGCAGTTTTCCAGGAGCCGGGGGAGCAG
GAGCAGGAAAGGTAGCCCCGAGGGATGTCCTTGGGAGGGGTGACTCTGGCCACAATC
CTCCCCACAGTCTACCCACTAGGATAGGAAGTGGGCTTGTGTGCTGGCTTCCGCTGT
CACCCAGCAAGGCACAGGCTCCTGTATTTGAGACTAGGATAGCTTCATCTTGAGCCTGAG
CCTTAGAATCTGTAGAGGAGCCTGGAGTCGGATCTAGCCATGGCTGGCAGAGGTTTCTAG
GGTGGGCCCCAGCCGTGGCGTGAAGTGAAGGATGACCCGGGGCAGCTGGCAGGAGAGGCC
TTGGCCTGACCTGGCACAGAAAGGGCAGCTTCACTCTCGAGTGTCCATTATCTGCTGT
TCCTTCGAGGGTTCAGGCTGTGTGTGGGGCCCAAGCATGCCCAACCCACCCCTCCTGGG
CCCAGGCAGCACCTGGAGCCACAGAGTCTGTGTGTAGCCAGGAAGCCCCGCTCAGGTAG
CCACCGCGGGGCACTGGCTGCTGTCTTGGTCTGTTAACCCCTCCACCTCCTCTCTTG
GACTCCCTCCCCACCCCAACCACTTTTCTTTCTCTTTAACCCAATGGAGACTTTCTGA
TGCATCGTTTTCTTGTGTGCCAAAGCAGGTCAGAAGAGGGAGAGGAGGGGCTGGGGGT
GAGGGGCCAGGCCATGGCCAAGGGGCCAGCTGCCCTCATTTTACTCTGACCTTACACA
GGGACAGATCTGATTTATTTATTTTGGTAAAAAAAAAAAAAAAAAGGAACAGAAACAATTT
GCATTGCATTGGCTTGACCCATAAACTAAGTTATATCCGTGAAAAAAAAAAAAAAAAAAAA

Restriction Sites:

Please inquire

ACCN:

NM_002541

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:	<u>NM_002541.2</u> , <u>NP_002532.2</u>
RefSeq Size:	4257 bp
RefSeq ORF:	3072 bp
Locus ID:	4967
UniProt ID:	<u>Q02218</u>
Cytogenetics:	7p13
Domains:	E1_dehydrog, transket_pyr
Protein Families:	Druggable Genome
Protein Pathways:	Citrate cycle (TCA cycle), Lysine degradation, Metabolic pathways, Tryptophan metabolism
Gene Summary:	<p>This gene encodes one subunit of the 2-oxoglutarate dehydrogenase complex. This complex catalyzes the overall conversion of 2-oxoglutarate (alpha-ketoglutarate) to succinyl-CoA and CO(2) during the Krebs cycle. The protein is located in the mitochondrial matrix and uses thiamine pyrophosphate as a cofactor. A congenital deficiency in 2-oxoglutarate dehydrogenase activity is believed to lead to hypotonia, metabolic acidosis, and hyperlactatemia. Alternative splicing results in multiple transcript variants encoding distinct isoforms.[provided by RefSeq, Sep 2009]</p> <p>Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1).</p>