



# Human ACO2 blocking peptide (CDBP0298)

This product is for research use only and is not intended for diagnostic use.

## PRODUCT INFORMATION

<b>Product Overview</b>	Blocking/Immunizing peptide for anti-Aconitase 2 antibody
<b>Antigen Description</b>	The protein encoded by this gene belongs to the aconitase/IPM isomerase family. It is an enzyme that catalyzes the interconversion of citrate to isocitrate via cis-aconitate in the second step of the TCA cycle. This protein is encoded in the nucleus and functions in the mitochondrion. It was found to be one of the mitochondrial matrix proteins that are preferentially degraded by the serine protease 15(PRSS15), also known as Lon protease, after oxidative modification. [provided by RefSeq, Jul 2008]
<b>Species</b>	Human
<b>Conjugate</b>	Unconjugated
<b>Applications</b>	Apuri, BL, ELISA
<b>Format</b>	Lyophilized powder
<b>Size</b>	100 µg
<b>Preservative</b>	None
<b>Storage</b>	Shipped at ambient temperature, store at -20°C.

## GENE INFORMATION

<b>Gene Name</b>	<a href="#">ACO2 aconitase 2, mitochondrial [ Homo sapiens (human) ]</a>
<b>Official Symbol</b>	ACO2
<b>Synonyms</b>	ACO2; aconitase 2, mitochondrial; ICRD; ACONM; aconitate hydratase, mitochondrial; citrate hydro-lyase;

<b>Entrez Gene ID</b>	<a href="#">50</a>
<b>mRNA Refseq</b>	<a href="#">NM_001098.2</a>
<b>Protein Refseq</b>	<a href="#">NP_001089.1</a>
<b>UniProt ID</b>	Q99798
<b>Chromosome Location</b>	22q13.2
<b>Pathway</b>	2-Oxocarboxylic acid metabolism, organism-specific biosystem; 2-Oxocarboxylic acid metabolism, conserved biosystem; Biosynthesis of amino acids, organism-specific biosystem; Biosynthesis of amino acids, conserved biosystem; Carbon metabolism, organism-specific biosystem; Carbon metabolism, conserved biosystem; Citrate cycle (TCA cycle), organism-specific biosystem; Citrate cycle (TCA cycle), conserved biosystem; Citrate cycle (TCA cycle, Krebs cycle), organism-specific biosystem; Citrate cycle (
<b>Function</b>	3 iron, 4 sulfur cluster binding; 4 iron, 4 sulfur cluster binding; aconitate hydratase activity; iron ion binding;