



# Human ACO1 blocking peptide (CDBP0296)

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-ACO1/Aconitase 1 antibody
<b>Antigen Description</b>	The protein encoded by this gene is a bifunctional, cytosolic protein that functions as an essential enzyme in the TCA cycle and interacts with mRNA to control the levels of iron inside cells. When cellular iron levels are high, this protein binds to a 4Fe-4S cluster and functions as an aconitase. Aconitases are iron-sulfur proteins that function to catalyze the conversion of citrate to isocitrate. When cellular iron levels are low, the protein binds to iron-responsive elements (IREs), which are stem-loop structures found in the 5' UTR of ferritin mRNA, and in the 3' UTR of transferrin receptor mRNA. When the protein binds to IRE, it results in repression of translation of ferritin mRNA, and inhibition of degradation of the otherwise rapidly degraded transferrin receptor mRNA. The encoded protein has been identified as a moonlighting protein based on its ability to perform mechanistically distinct functions. Alternative splicing results in multiple transcript variants [provided by RefSeq, Jan 2014]
<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

**Gene Name** [ACO1 aconitase 1, soluble \[ Homo sapiens \(human\) \]](#)

<b>Official Symbol</b>	ACO1
<b>Synonyms</b>	ACO1; aconitase 1, soluble; IRP1; ACONS; HEL60; IREB1; IREBP; IREBP1; cytoplasmic aconitate hydratase; IRE-BP 1; citrate hydro-lyase; iron regulatory protein 1; ferritin repressor protein; epididymis luminal protein 60; aconitate hydratase, cytoplasmic; iron-responsive element binding protein 1; iron-responsive element-binding protein 1;
<b>Entrez Gene ID</b>	<a href="#">48</a>
<b>mRNA Refseq</b>	<a href="#">NM_001278352.1</a>
<b>Protein Refseq</b>	<a href="#">NP_001265281.1</a>
<b>UniProt ID</b>	P21399
<b>Chromosome Location</b>	9p21.1
<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>	4 iron, 4 sulfur cluster binding; RNA binding; RNA binding; aconitate hydratase activity; iron-responsive element binding; metal ion binding; protein binding;