

Recombinant Human IDH2, His-tagged

Cat. No. IDH2-28080TH **Lot. No.** (See product label)

SPECIFICATION

Product Overview	Recombinant fragment, corresponding to amino acids 217-442 of Human IDH2 with an N terminal His tag; MWt 26 kDa.
Species	Human
Source	E.coli
ProteinLength	217-442 a.a.
Description	Isocitrate dehydrogenases catalyze the oxidative decarboxylation of isocitrate to 2-oxoglutarate. These enzymes belong to two distinct subclasses, one of which utilizes NAD(+) as the electron acceptor and the other NADP(+). Five isocitrate dehydrogenases have been reported: three NAD(+)-dependent isocitrate dehydrogenases, which localize to the mitochondrial matrix, and two NADP(+)-dependent isocitrate dehydrogenases, one of which is mitochondrial and the other predominantly cytosolic. Each NADP(+)-dependent isozyme is a homodimer. The protein encoded by this gene is the NADP(+)-dependent isocitrate dehydrogenase found in the mitochondria. It plays a role in intermediary metabolism and energy production. This protein may tightly associate or interact with the pyruvate dehydrogenase complex.
Conjugation	HIS
Form	Lyophilised:Reconstitute with 87 µl aqua dest.

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Storage buffer	Preservative: None Constituents: 0.5% Trehalose, 6M Urea, 100mM Sodium phosphate, 10mM Sodium chloride, pH 4.5
Storage	Shipped at 4°C. Upon delivery aliquot and store at -80oC. Avoid freeze / thaw cycles.
Sequences of amino acids	VGMGMYNTDESISGFAHSCFQYAIQKKWPLYMSTKNTILK AYDGRFKDIFQEIFDKH YKTD FDKNKI WYEHRLIDDMV AQVLKSSGGFVWACKNYDGDVQSDILAQGFGLG LMTS VLVCPDGKTIEAEAAHGTVTRHYREHQKGRPTSTNPIA SIFAWTRGLEHRGK LDGNQDLIRFAQMLEKVCVETVESGA MTKDLAGCIHGLSNVKLNEHFLNTTDFLDTI K
Sequence Similarities	Belongs to the isocitrate and isopropylmalate dehydrogenases family.
GENE INFORMATION	
Gene Name	IDH2 isocitrate dehydrogenase 2 (NADP+), mitochondrial [Homo sapiens]
Official Symbol	IDH2
Synonyms	IDH2; isocitrate dehydrogenase 2 (NADP+), mitochondrial; isocitrate dehydrogenase [NADP], mitochondrial;
Gene ID	3418
mRNA Refseq	NM_002168
Protein Refseq	NP_002159
MIM	147650

Uniprot ID	P48735
Chromosome Location	15q21-qter
Pathway	Citrate cycle (TCA cycle), organism-specific biosystem; Citrate cycle (TCA cycle), conserved biosystem; Citrate cycle, first carbon oxidation, oxaloacetate => 2-oxoglutarate, organism-specific biosystem; Citrate cycle, first carbon oxidation, oxaloacetate =>
Function	NAD binding; isocitrate dehydrogenase (NADP+) activity; isocitrate dehydrogenase (NADP+) activity; magnesium ion binding; oxidoreductase activity;

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