

Product Data Sheet

CISD1 Protein, Human (His)

Cat. No.:	HY-P76262
Synonyms:	CDGSH iron-sulfur domain-containing protein 1; MitoNEET; C10orf70; ZCD1
Species:	Human
Source:	E. coli
Accession:	Q9NZ45/NP_060934.1 (K32-T108)
Gene ID:	55847
Molecular Weight:	Approximately 14 kDa.

PROPERTIES	
AA Sequence	KRFYVKDHRN KAMINLHIQK DNPKIVHAFD MEDLGDKAVY CRCWRSKKFP FCDGAHTKHN EETGDNVGPL IIKKKET
Appearance	Lyophilized powder
Formulation	Lyophilized from sterile PBS, pH 7.4.
Endotoxin Level	<1 EU/µg, determined by LAL method.
Reconsititution	It is not recommended to reconstitute to a concentration less than 100 μg/mL in ddH ₂ O. For long term storage it is recommended to add a carrier protein (0.1% BSA, 5% HSA, 10% FBS or 5% Trehalose).
Storage & Stability	Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage.
Shipping	Room temperature in continental US; may vary elsewhere.

DESCRIPTION

Background	The CISD1 protein functions as an L-cysteine transaminase, facilitating the reversible transfer of the amino group from L- cysteine to the alpha-keto acid 2-oxoglutarate. This enzymatic process results in the formation of 2-oxo-3- sulfanylpropanoate and L-glutamate. The catalytic cycle is orchestrated in the presence of the pyridoxal 5'-phosphate (PLP) cofactor, which initiates transamination by forming an internal aldimine with the epsilon-amino group of the active site Lys- 55 residue on the enzyme (PLP-enzyme aldimine). This internal aldimine is subsequently displaced by the formation of an external aldimine with the substrate amino group (PLP-L-cysteine aldimine). The external aldimine undergoes
	deprotonation, leading to the formation of a carbanion intermediate. In the presence of 2-oxoglutarate, this intermediate regenerates PLP, ultimately yielding the final products 2-oxo-3-sulfanylpropanoate and L-glutamate. The active site lysine residue is implicated in controlling proton transfer in the carbanion intermediate, while PLP stabilizes the carbanion structure through electron delocalization, a phenomenon known as the electron sink effect. Additionally, CISD1 plays a

crucial role in regulating the maximal capacity for electron transport and oxidative phosphorylation and may be involved in iron-sulfur cluster shuttling and/or redox reactions. Notably, it can transfer the [2Fe-2S] cluster to an apo-acceptor protein, particularly under oxidative stress conditions, suggesting a role as a redox sensor that regulates mitochondrial iron-sulfur cluster assembly and iron trafficking.

Caution: Product has not been fully validated for medical applications. For research use only.

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