

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P11024	NNTM_BOVIN	Bos taurus	NAD(P) transhydrogenase, mitochondrial	27.604395	NaN	NaN	28100784	MANLLKTVVTGCSCPFLSNLGSCKV LPGKKNFLRTFHTHRILWCSAPVKP GIPYKQLTVGVPKEIFQNEKRVALSP AGVQALVKQGFNVVVEGAGEASKF SDDHYRAAGAQQGAKVLDLVLV KVRAPMLNPTLGVHEADLLKTSGTL ISFIYPAQNPDLLNKLSKRKTTVLAM DQVPRVTIAQGYDALSSMANIAGYK AVVLAANHFGRFFTGQITAAGKVPP AKILIVGGVAGLASAGAASKMGAIIV RGFDTRAAALEQFKSLGAEPLVDL KESGEGQGGYAKEMSKEFIEAEMKL FAQQCKEVDILISTALIPGKKAPILFN KEMIESMKEGSVVVDLAAEAGGNFE TTKPGELYVHKGIHIGYTDLP SRMA TQASTLYSNNITKLLKAISPDKDNFY FEVKDDDFDGTMGHVIRGTVMKD GQVIFPAPTPKNIPQGAPVKQKTVAE LEAEKAATITPFRKTMTSASVYTAGL TGILGLGIAAPNLAFSQMVTTFLAG IVGYHTVWGVTPALHSPMLSVTNAI SGLTAVGGLVLMGGHLYPSTTSQGL AALATFISSVNIAGGFLVTQRM LDM FKRPTDPPPEYNYLYLLPAGTFVGGYL ASLYSGYNIEQIMYLGSGLCCV GALA GLSTQGTARLGNALGMIGVAGGLAA TLGGLKPCPELLAQMSGAMALG GTI GLTIAKRIQISDLPQVAAFHSLVGLA AVLTCIAEYIIEYPHFATDAAANLTKI VAYLGTIYGGVTFSGSLVAYGKLOGI LKSAPLLPGRHLLNAGLLAGSVGGI IPFMMDPSFTTGITCLGSVSALSAVM GVTLTAAGGADMPVITVLNSYSG WALCAEGFLLNNLLTIVGALIGSSG AILSYIMCVAMNRSANLILGGYGT STAGGKPMIEISGTHTEINLDNAIDMI REANSIIITPGYGLCAAKAQYPIADLV KMLSEQGGKVRFGIHPVAGRMPGQ LNVLLAEAGVPYDIVLEMDEINHDF PDTDLVLVIGANDTVNSAAQEDPNSI IAGMPVLEVWKSQVIVMKRSLGVG YAAVDNPIFYKPNAMLLGDAKKT DALQAKVRESYQK	None	None	None	None	None	None	None		

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