	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ext reg
0	Q61941	NNTM_MOUSE	Mus musculus	NAD(P) transhydrogenase, mitochondrial	30.328585	NaN	NaN	37453647;39927985;36288343	MAHLLKTVVAGCSCPFLSNLGSSKV LPGKRDFVRTLRTHQALWCKSPVKP GIPYKQLTVGVPKEIFQNEKRVALSP AGVQALVKQGFNVVVESGAGEASKF PDDLYRAAGAQIQGMKEVLASDLVV KVRAPMVNPTLGAHEADFLKPSGTL ISFIYPAQNPDLLNKLSERKTTVLAM DQVPRVTIAQGYDALSSMANISGYKA VVLAANHFGRFFTGQITAAGKVPPA KILIVGGGVAGLASAGAAKSMGAVV RGFDTRAAALEQFKSLGAEPLEVDL KESGEGQGGYAKEMSKEFIEAEMKL FAQQCKEVDILISTALIPGKKAPVLFS KEMIESMKEGSVVVDLAAEAGGNFE TTKPGELYVHKGITHIGYTDLPSRMA TQASTLYSNNITKLLKAISPDKDNFH FEVKDDFDFGTMSHVIRGTVVMKD GKVIFPAPTPKNIPEEAPVKPKTVAEL EAEKAGTVSMYTKTLTTASVYSAGLT GMLGLGIVAPNVAFSQMVTTFGLAG IIGYHTVWGVTPALHSPLMSVTNAIS GLTAVGGLALMGGHFYPSTTSQSLA ALATFISSVNIAGGFLVTQRMLDMF KRPTDPPEYNYLYLLPGGTFVGGYLA ALYGGYNIEEIMYLGSGLCCVGALG GLSTQGTARLGNALGMIGVAGGLAA TLGGLKPDPQLLAQMSGAMAMGGT IGLTIAKRIQISDLPQLVAAFHSLVGL AAVLTCMAEYIVEYPHFAMDATSNF TKIVAYLGTYIGGVTFSGSLVAYGKLQ GILKSAPLLLPGRHALNAGLLAASVG GIIPFMADPSFTTGITCLGSVSGLSTL MGVTLTAAIGGADMPVVITVLNSYS GWALCAEGFLLNNNLLTIVGALIGSS GAILSYIMCVAMNRSLANVILGGYGT TSTAGGKPMEISGTHTEINLDNAVE MIREANSIVITPGYGLCAAKAQYPIAD LVKMLTEQGKKVRFGIHPVAGRMPG QLNVLLAEAGVPYDIVLEMDEINSDF PDTDLVLVIGANDTVNSAAQEDPNSI IAGMPVLEVWKSKQVIVMKRSLGVG YAAVDNPIFYKPNTAMLLGDAKKTC DALQAKVRESYQK	True	False	False	False	4.334	False	False	2.906	Fal