

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ex re
Q99798	ACON_HUMAN	Homo sapiens	Aconitate hydratase, mitochondrial	20.21434	S193	S559;S670	29351928;32119511;34725712;28510447;38665916	MAPYSLLVTRLQKALGVROYHVASV LCQRAKVAMSHFEPNEYIHYDLLEK NINIVRKRLNRPLTLSEKIVYGHLLD PASQEIERGKSYLRRLRDRVAMQDA TAQMAMLQFISGLSKVAVPSTIHC DHLIEAQVGGGEKDLRRAKDINQEVY NFLATAGAKYGVGFWKP GSGIIHQII LENYAYPGVLLIGTDSHTPNGGGLG GICIGVGGADAVDMAGIPWELKCP KVIGVKLTGSLSGWSSPKDVILKVAG ILTVKGGTGAIVEYHGPGVDSISCTG MATICNMGAEIGATTSVFPYNHRMK KYLKGTGREDIANLADEFKDHLPD PGCHYDQLIEINLSELKPHINGPFTP DLAHPVAEVGKVAEKEGWPLDIRVG LIGSCTNSSYEDMGRSAAVAKQALA HGLKCKSQFTITPGSEQIRATIERDG YAQILRDLGGIVLANACGPCIGQWD RKDIKKGEKNTIVTSYNNRNFTRND ANPETHAFVTSPEIVTALAIAAGTLKF NPETDYLGTGDKKFRLEAPDADEL PKGEFDPGQDTYQHPPKDSSGQHV DVSPTSQRLQLEPFDKWDGKDLED LQILIKVKGCTTDHISAAGPWLKFR GHLDNISNNLLIGAINIENGKANSVR NAVTFEFGVPTDARYYKKGIRVW VIGDENYEGSSREHAALPRHLGG RAITKSFARIHETNLKKQGLLPLTFA DPADYNKIHPVDFKLTIQGLKDFTPGK PLKCIKHFNQTETILLNHTFNETQ IEWFRAGSALNRMKELQQ	True	False	4.197	2.446	4.904	2.227	1.613	2.344	2.