

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	ex re	
0	P53597	SUCA_HUMAN	Homo sapiens	Succinate-CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial	23.565205	S316	NaN	34229054;40596516;38665916;40914422;28510447;29351928	MTATLAAAADIATMVSGSSGLAAARLLSRSFLLPQNGIRHCSYTASRQHLYVDKNTKICQGFQKQGTFFHSQQAL EYGTKLVGGTTPGKGGQTHLGLPVFNTVKEAKEQTGATASVIYVPPFAAA AINEAIEAEIPLVVCITEGIPQDDMVRVKHKLLRQEKTRRLIGPNCVGINPGECKIGIMPGHIHKKGRIGIVSRSGTLTYEAVHQTTQVGLGQSLCVGIGGDPFN GTDFIDCLEIFLNDSETEGIIIGEIGG NAEENAAEFLKQHNSGPNKPVVSF IAGLTAPPGRRMGHAGAIAGGKGA KEKISALQSAGVVVSMSPAQLGTTIY KEFEKRKML	True	False	2.822	1.948	4.852	1.725	0.755	2.977	2.0