

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac 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	20.267722	S316	NaN	34229054;29351928;28510447;38665916	MTATLAAAADIATMVSGSSGLAAAR LLSRSFLLPQNGIRHCSYTASRQHLY VDKNTKICQGFYKQGTFFHSQQAL EYGTKLVGGTTPGKGGQTHLGLPVF NTVKEAKEQTGATASVIYVPPFAAA AINEAIEAEIPLVVCITEGIPQDDMVR VKHKLLRQEKTRLIGPNCVGINPGE CKIGIMPGHIHKKGRIGIVSRSGTLTY EAVHQTTQVGLGQSLCVGIGGDPFN GTDFFIDCLEIFLNDSETEGIIIGEIGG NAEENAAEFLKQHNSGPNKPVVSF IAGLTAPPGRRMGHAGAIAGGKGGGA KEKISALQSAGVVVSMSPAQLGTTIY KEFEKRKML	True	False	3.694	1.949	4.851	1.722	0.758	2.978	2.0