

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	ext req
0 Q9WUM5	SUCA_MOUSE	Mus musculus	Succinate--CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial	29.036547	NaN	NaN	37507081;37453647;33300544;36288343;39927985	MTATVAAAAATATMVSSSSGLAAAR LLSRTFLLQQNGIRHGSYASRKH IDKNTKIICQGFQKQGFHSSQQA YGTCLVGGTTPGKGGQKHLGLPV TVKEAKEKTGATASVIVPPFAAA NEAIDAEIPLVVCITEGIPQQDM KHRLTRQGTTRLIGPNCVGPVNG CKIGIMPGHIHKKGRIGVSRSGT EAVHOTTOVGLGQSLCIGIGDFF GTDFFIDCLEVFLNDPATEGIIIG GHAEENAAAFKEHNSGPKAPVVS FIAGITAPPGRRMGHAGAIAGGK AKEKISALQSAGVVVMSPAQLGT YKEFEKRKML	True	False	False	False	4.573	False	False	False	False