

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	extracellular region	
0	P13086	SUCA_RAT	Rattus norvegicus	Succinate--CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial	23.439218	NaN	NaN	38843836	MTAAVVAAAATATMVGSSGLAAAR LLSRTFLLQQNGIRHGSYTASRKNY IDKNTKVICQGFTGKQGTFSQQAL EYGTKLVGGTTPGKGGKKHLGLPVF NTVKEAKEKTGATASVIYVPPFAAA AINEAIDAEIPLVVCITEGIPQQDMVR VKHKLTRQGKTRLIGPNCPGIINPGE CKIGIMPGHIIKKGRIGIVSRSGTLTY EAVHQTTQVGLGQSLCIGIGDPFN GTNFDCLDVFLKDPATEGIVLIGEIG GHAEENAAEFLKEHNSGPKAPVVS FIAGITAPPGRRMGHAGAIAGGKGG AKEKISALQSAGVIVSMSPAQLGTC MYKEFEKRKML	True	False	1.445	1.29	5.0	1.255	False	0.88	1.446