

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P38266	AIM3_YEAST	Saccharomyces cerevisiae	Altered inheritance of mitochondria protein 3	9.365641	NaN	S57;S58;S64;S476;T729;T861	33229814	MGFWENNKDSITSGLKSAAGKYGYQ GTYVAKTGYKASKKHYNNKARRE RKSGKKNSSDEEYDSEDEMEYERKP TDIRSLKDPKSFPPPLKPGQKTYTG QQQQQMPNGQASYAFQAYQGQPG AGSTEQSQAQPQYNQYPQQQLQQ GVMPQQQLQQGVVPPQPPPIYGEQ VPPYGSNSNATSYSQSLPQQNQPNNA IPSQVSLNSASQOSTGFVSNLQYG TQSSNPAPSPSFQNGLQCHQPPQYV SHGSTNLGQSQFPSSGQQQQPTTQF GQQVLPSPAQPQQQQGQPLPPPRG QVILPAPGEPLSNGFGQQQQQQQQ QQQPLNQNNALLPQMNVEGVSGM AAVQPVYQGAMSSTTNMQDSNPSY GASPMQGGPPVGGQPPVPVRMQPQ PPQPMQQGNIYPIEPLDSTGSTPHF EVTDFDPDAPAPKPKIDIPTVDVSSLP PPPTHDRGAVVHQEPAPSGKIQPN TTSSAASLPKHSRTTTADNERNNSG NKENDESTSKSSILGHYDVDVNIMP PPKPFRLGLDVPSEHTTKNAPERA VPILPPRNNVEPPPPSRGNFERTES VLSTNAANVQEDPISNFLPPPFPFRH TETKQNQNSKASPVEMKGEVLPGH PSEEDRNVEPSLVPQSKPQSQSQFR RAHMETQPIQNFQPPPKPFRRSQSS NSSDSSYTIDGPEANHGRRGRIAK HHDGDEYNPKSENSTENGLGDAP NSFIRKRAPTPPAPSRSEKLHEGTIT SEVDSSKDANKYEKSIPPVTSSIAQ QSTKKAPPVVKPKPRNFSLKANAY PKELTREATGQDEVLSITNELSHIK LRKTNVNLEKLGSKKVKDSSPVPS DLDEKYVSASGSITPPRPPSRSSPK KVPVVPKKNLKKKPPVVPKPKK LLKSLEPRPIEMERAYSGDISAADDN LNPFERYKRNVPQEDDRLHKLK