

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	pl m
095163	ELP1_HUMAN	Homo sapiens	Elongator complex protein 1	21.026851	T1128;T1130	S471;S804;S867;S1171;S1174	36580660;39531497;23301498;38665916;28510447;34229054;30397120	MRNLKLFRTLEFRDIQPGNPQCFS LRTEQGTVLIGSEHGLIEVDPVSREV KNEVSLVAEGFLPEDGSGRIVGVQD LLDQESVCVATASGDVILCSLSTQQL ECVGSVASGIVMSWSWSPDQELVLLA TGQQTLLIMMTKDFEPILQQIHQDD FGESKFITVGVGRKETQFHGSEGRQ AAFQMQMHESALPWDDHRPQVTW RGDGGQFFAVSVVCPETGARKVRVW NREFALQSTSEPVAGLGPALAWKPS GSLIASTQDKPNQQDIVFFEKNGLL HGHFTLPLFKDEVKVNDLLWNADS SVLAVWLEDLQREESSIPKTCVQLW TVGNVHWYLLKQSLSFSTCGKSKIVS LMWDPVTPYRLHVLCOGWHYLAID WHWTTDRSVGDNSDLSNVAIDG NRVLVTVFRQTVVPPMCTYQLLFP HPVNQVTFLAHPQKSNDLAVLDAS NQISVYKCGDCPSADPTVKGAVGG SGFKVCLRTPHLEKRYKIQFENNED QDVNPLKGLLWIEEDVFLAVSHS EFSRPSVIHHLTAASSEMDEEHGQL NVSSAAVDGVIISLCCNSKTKSVVL QLADGQIFKYLWESPLAIKPKWNS GGFPVRFPPYCTQTELAMIGEEECVL GLTDRCRFFINDIEVASNITSFAVYD EFLLLTTHSHTCQCFCLRDAFRTL QAGLSSNHVSHGEVLKRVGSRIV TVVPQDTKLVLMQPRGNLEVVHHR ALVLAQIRKWLDKLMFKEAFECMRK LRINLNLIYDHNPKVFLGNVETFIQ IDSVNHINLFFTELKEEDVTKTMYP PVTSSVYLSRDPDGNKIDLVCDAMR AVMESINPHKYCLSILTSHVKKITPE LEIVLQVHLEQGNAPSDPDAVSAE EALKYLLHLVDVNELYDHSGLTYDF DLVLMVAEKSQKDPKEYLPFLNLT KMETNYQRFTIDKYLKRYEKAIGHLS KCGPEYFPECLNLIKDNLYNEALKL YSPSSQYQDISIAYGEHLMQEHMY EPAGLMFARCGAHEKALSFLTCGN WKQALCVAQLNFTKDQLVGLGRTL AGKLVEQRKHIDAAMVLEECAQDYE EAVLLLLEGAWEALRLVYKYNRL DIITNVKPSILEAQKNYMAFLDSQT ATFSRHKKRLLVVRELKEQAQAGL DDEVPHGQESDLFSETSSVVSSEM SGKYSHNSRISARSSKNRRKAERK KHSKKEGSPLEDLALLEALSEVVON TENLKDEVYHILKVLFLFEFDEQGR ELQKAFEDTLQLMERSLPEIWTLY QONSATPVLGPNSTANSIMASYQQQ KTSVPVLDALFIPPKINRRQWKLS LLD	True	False	4.687	4.565	1.639	1.094	0.53	1.