

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9UJX6	ANC2_HUMAN	Homo sapiens	Anaphase-promoting complex subunit 2	34.455997	NaN	S218;S314;S470;S534;S697;T810	40914422	MAAAVVVAEGDSDSRPQELLVAV NTVSTGLVPPAALGLVSSRTSGAVPP KEEELRAAVEVLRGHLHSVLEEFW VEVLQNDLQANISPEFWNAISQCEN SADEPQCLLLLLDAFGLLESRLDPYL RSLELLEKWTRLGLLMGTGAQGLRE EVHTMLRGVLFSTPRTFQEMIQRL YCCFLRVYMQSKRKGEGGTDPELEG ELDSRYARRRYRLLQSPLCAGC SSD KQCCWCRQALEQFHQLSQVLHRLS LLERSAEAVTTTLHQVTRERMEDR CRGEYERSFLREFHKWIERVVGWLG KVFLQDGPAPASPEAGNLRWRRC HVQRFYRIYASLRIEELFSIVRDFPD SRPAIEDLKYCLERTDQRQQLLVSLK AALETRLLHPGVNTCDIITLYISAIKA LRVLDPSMVILEVACEPIRRYLRTRE DTVVRQIVAGLTGDSGTDGLAVELS KTDPASLETGQDSEDDSGEPEDWVP DPVDADPGKSSSKRRSSDIISLLVSIY GSKDLFINEYRSLADRLRHQFSFSP EREIRNVELLKLRFGEAPMHFCEVM LKDMADSRINANIREEDEKRPAAE QPPFGVYAVLSSEFWPPFKDEKLEV PEDIRAALAEAYCKKYEQLKAMRTLS WKHTLGLVTMDVELADRTL SVAVTP VQAVILLYFDQASWTLEELSKAVK MPVALLRRRMSVWLQQGVLREPP GTFVIEEERPDQDRDNMVLIDSDE SDSGMASQADQKEEELLLFWTYIQA MLTNLESLSLDRIYNMLRMFVVTGP ALAEIDLQELQGYLQKKVRDQQLVY SAGVYRLPKNCS	True	False	4.414	5.0	1.296	0.842	False	0.971	0.773