	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion
0	Q3UHJ0	AAK1_MOUSE	Mus musculus	AP2-associated protein kinase 1	53.614683	T354;T359;T360;T395;S406;S414;S416;T441;T445;T448;S572;T578;S635;S648;T670;T740	S14;T234;S235;T354;T389;T441;T604;S616;T618;S621;S622;S635;S648;T651;S729;S844;S935;S936	22517741;22645316;33300544;40885482;36731797;36288343;35822049;40021952;34678516;39627609	MKKFFDSRREQGSSGLGSGSSGGG GSSSGLGSGYIGRVFGIGRQQVTVDE VLAEGGFALVFLVRTSNGVKCALKR MFVNNEHDLQVCKREIQIMRDLSG HKNIVGYIDSSINNVSSGDVWEVLIL MDFCRGGQVVNLMNQRLQTGFTE NEVLQIFCDTCEAVARLHQCKTPIIH RDLKVENILLHDRGHYVLCDFGSAT NKFQNPQAEGVNAVEDEIKKYTTLS YRAPEMVNLYSGKIITTKADIWALGC LLYKLCYFTLPFGESQVAICDGSFTIP DNSRYSQDMHCLIRYMLEPDPDKR PDIYQVSYFSFKLLKKECPVPNVQNS PIPAKLPEPVKASEAAVKKTQPKARL TDPIPTTETSIAPRQRPKAGQTQPNP GILPIQPALTPRKRATVQPLPQAAGP SNQPGLLPSVSQPKAQATPSQPLQS SQPKQPQAPPTPQQTPATQTQGLPT QAQATPQHQQQHLLKQQQQQQQQP QQPTAPPQPAGTFYQQQQQQQQQQ AQTQQFQAVHPAAQQPVTAQFPVGS QGGAQQQLMQNFYHQQQQQQQQQ QQLMAQQAALQQKTAVVVPQSQAQ PATAPQAAAAQEPGQIQAPVRQQPK VQTTPPPTIQGQKVGSLTPPSSPKTQ RAGHRRILSDVTHSAVFGVPASKST QLLQAAAAEASLNKSKSATTTPSGSP RTSQQNVSNASEGSTWNPFDDDNF SKLTAEELLNKDFAKLGEGKLPEKL GGSAESLIPGFQPTQGDAFTTPSFSA GTAEKRKGGQAVDSGIPLLSVSDPFI PLQVPDAPEKLIEGLKSPDTSLLLPD LLPMTDPFGSTSDAVIDKADVAVESL IPGLEPPVAQRLPSQTESVTSNRTDS LTGEDSLLDCSLLSNPTAGLLEEFAP IALSAPTHKAAEDSNLISGFGVAEGS EKVAEDEFDPIPVLITKNTQGGHSR NSSGSSESSLPNLARSLLLVDQLIDL	False	False	3.968	3.057	1.38