

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3UHJ0	AAK1_MOUSE	Mus musculus	AP2-associated protein kinase 1	29.484366	T360;S572	S14;T234;S235;T354;T389;T441;T604;S616;T618;S621;S622;S635;S648;T651;S729;S844;S935;S936	34678516;36288343;35822049;22645316;22517741;33300544	MKKFFDSRREQSSGLGSGSSGGG GSSSGLGSGYIGRVFGIGRQQVTVDE VLAEGGFALVFLVRTSNGVKCALKR MFVNNNEHDLQVCKREIQIMRDLG HKNIVGYIDSSINNVSSGDVWEVLIL MDFCRGGQVVNLMNQRLQTFTE NEVLQIFCDTCEAVARLHQCKTPIIH RDLKVENILLHDRGHYVLCDFGSAT NKFQNPQAEGVNAVEDEIKKYTTLS YRAPEMVNLYSGKIITTKADIWALGC LLYKLCYFTLPFGESQVAICDGSFTIP DNSRYSQDMHCLIRYMLEPDPDKR PDIYQVSYFSFKLLKKECPVNVQNS PIPAKLPEPVKASEAAVKKTQPKARL TDDPIPTTETSIAPRQRPKAGQTQPNP GILPIQPALTPRKRATVQPLPQAAGP SNQPGLLPSVSQPKAQATPSQPLQS SQPKQPQAPPTPQQT PATQTQGLPT QAQATPQHQQHLLKQQQQQQQQQ QQPTAPPQAGTFYQQQQQQQQQQ AQTOQFQAVHPAAQQPVTAQFPVGS QGGAAQQQLMQNFYHQQQQQQQQQ QQLMAQQAALQKKTAVVVPQSQAQ PATAPQAAAAQEPGQIQAPVRQPK VQTTTPPTIQGQKVGSLTPPSSPKTQ RAGHRRILSDVTHSAVFGVPASKST QLLQAAAAEASLNKSKSATTTPSGSP RTSQQNVSNASEGSTWNPFDNDF SKLTAEELLNKDFAKLGEGKLEPEL GGSAESLIPGFQPTQGDFTTSPFSA GTAEKRRGGQAVDSGIPLLSVSDPFI PLQVPDAPEKLEGLKSPDTSLLLPD LLPMTDPFGSTSDAVIDKADVAVESL IPGLEPPVAQRLPSQTESVTSNRTDS LTGEDSLLDCSLLSNPTAGLLEEFAP IALSAPTHKAAEDSNLISGFGVAEGS EKVAEDEFDPIPVLITKNTQGGHSR NSSGSSESSLPNLARSLLLVDQLIDL