	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q924C5	ALPK3_MOUSE	Mus musculus	Alpha-protein kinase 3	25.629171	NaN	S229;S1199	36288343	MGSRRAAGRGWGLGGRAGAGGDSE DDGPVWTPGPASRSYLLSVRPEASL SSNRLSHPSSGRSTFCSIIAQLTEET QPLFETTLKSRAVSEDSDVRFTCIVT GYPEPEVTWYKDDIELDRYCGLPKYE ITHQGNRHTLQLYRCQEEDAAIYQA SARNTKGIVSCSGVLEVGTMTEYKIH QRWFAKLKRKAAAKMREIEQSWKH GKEASGEADTLLRKISPDRFQRKRRL SGVEEAVLSTPVREMEEGSSAAWQE GETESAQHPGLGLINSFAPGEAPTN GEPAPENGEDEERGLLTYICEVMEL GPQNSPPKESGAKKKRKDEESKPGE QKLELEKAEGSQCSSENVVPSTDKP NSSRREKSTDTQPAQTQPRGRVARG PGIESTRKTASVLGIQDKVQDVPAPA PAPVPAPALAPAPVPVPAPTPVPSRS SEQVYFSLKDMFMETTRAGRSQEEE KPPPPSTRVAGESPPGKTPVKSRLEK VPMVSSQPTSSMVPPPIKPLNRKRF APPKSKVESTTTSLSSQTSESMAQSL GKALPSASTQVPTPPARRRHGTRDS PLQGQTSHKTPGEALESPATVAPTKS ANSSSDTVSVDHDSSGNQGATEPM DTETQEDGRTLVDGRTGSRKKTHTD GKLQVDGRTQGDGAQDRAHASPRT QAGEKAPTDVVTQGSERPQSDRSSW KNLVTQRRVDMQVGQMQAGERWQ QDPGDARIQEEEKETQSAAGSIPVAF ETQSEQLSMASLSSLPGALKGSPSG CPRESQAIECFEKSTEAPCVQERSDL MLRSEEAAFRSHEDGLLGPPSGNRT YPTQLPPEGHSEHLGGQTHQRSEQE DSLSQCPKKEQPQEPLHVGLSGGHS TGLSQEVPAMPSLPGTGLTSSLQEE LPGTAASLHTNTDVPLPSRDQDFPS SAPTLQLGPGSPTQSHPPEAMATSS EGACAKEPNVDGRSSGTRSCDPGLI DSLKNYLLLLLKLSSPETSEARAESQ EVADTGGLTSSSTLVPTMEVAGLSP RTSRRILERVENNHLVQSAQTLLLSP CTSRRLTGLLDREVQAGQQALAAAQ CSRGPCPTPLTIPAIVVGEEGSAGED SEERTSQESDKKGLLGEVEGHTVES RTQEPCQEEAMPGEALTGLPAATPE ELALGARRKRFLPKVRAGSDGEANK AEERESPTVSPRGPRKGLTPGSPGTP GRERRSPTQARKASMLEVPGAEEEP ATGDLVSRSKDSGLDSEPAVDEGKQ EALAKQRKAKDLLKAPQVIRKIRVEQ FPDSSGSLKLWCQFFNIVSDSVLTW AKDQHPVGEVNRSAGDEGPAALAIV QASPTDCGVYRCTIQNEHGSASTDF CLSPEVLSGFISREEGEVGEEIEMTP MVFAKGLADSGCWGDKLFGRLVSE ELRGGGHGLQKASRAKVIYGLEPIFE SGRTCIIKVSSLLVFGPSSETSLLGRN YDVTIQGCKIQNMSREYCKIFAAEAR AASGFGEVPEIIPLYLIYRPANNIPYA TLEEDLGKPLQTYCSRQWGCAGAPA AASSSEALQKCQTFQHWLYQWTNG SFLVTDLTGADWKMTDVQIATKLRG YQGLKESCFPALLDQFASSHQCNTY CDMLGLKPLKGPEAAHPQAKAKGSK SPSAGRKGSQLSPQPQKKGLPSPQG SRKSAPSSRATLQASQAATVQLLGQ PPVQDGSSKAQSMR	None	None	None	None	None	None	None	None	None