

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O54988	SLK_MOUSE	Mus musculus	STE20-like serine/threonine-protein kinase	23.2672	NaN	S14;T183;S189;S340;S341;S344;S347;S348;S354;S372;S543;S561;S566;S643;S647;S666;S775;S777;T812;S816;T1095	22645316;25153642;36288343;28135057	MSFFNFRKIFKLGSEKKKKQYEHVK RDLNPPEEFWEIIGELGDGAFGKVVYK AQNKETNVLAAAKVIDTKSEEELED YMVEIDILASCDHPNIVKLLDAFYFE NNLWILIEFCAGGAVDAVMLELERP LTESQIQVVCQQTLEALNYLHDNKII HRDLKAGNIFLFTLDGDIKLADFGVSA KNTRTIQRRDSFIGTPYWMapevVM CETSKDRPYDYKADVWSLGITLIEMA EIEPPHHELNPmrvLLKIAKSEPPTL AOPSKWSSNFKDFLRKCLEKNVDA RWTTSQLLQHPFVTVDSNKPVRELI AEAKAEVTEEVEDGKEEDEEEEAEN ALPIPANKRASSDLSIASSEEDKLSQ NACILESVSERTEQSTSEDKFSNKIL NEKPTTDGPEKAVDEHASDVNLETG AELNDQTVGIHENGREKKRPKLENL PDTQDQQTVDVNSVSEENENNRVT LETNTDCLKPEEDRNKENQETLESK LIQSEEINDTHIQTMDLVSQETGEKE ADFQAVDNEVGLTKEEQEKLKGDG TAQKVITSDRSSEVGTDEALDDTQKA AELSKAAQSGEGDEALAPTQTLAEK PTEGPEAGGAEIEPPGGERVEDKQP EQQPAVCEAEGQLTSTSETTRATLE QPETDEVEQVSESNIEELERLVVTG AEARALGSEGEAAATEVDLERKENA QKVPVKAESQAPAASQPSEPHVLP SININSETTENKEEMGALPKPETILP PEPEHEKGNDDSGTGSTVENSSGD LNLSSISFLSKAKDSGSVSLQETRRQ KKTLKKTRKFIVDGVESVTTSKIVT DSDSKTEELRFLRRQELRELRLLQK EEQRAQQQLNGKLQQQREQIFRRFE QEMLSKKRQYDQEIENLEKQKQTI ERLEQEHTNRLRDEAKRIKGEQEKE LSKFQNVLKNRKEVMNEVEKAPR ELRRELTkrrKEELAQSQAHAQEQEF VQKQQQLDGSLLKIIQQQKAELANI ERECLNnkQQLMRAREAAIWELEE RHLQEKHQLLkQQLKDQYFMQRHQ LLKRHEKETEQMQRYNQRLIEELKN RQTQERARLPKIQRSEAKTRMAMFK KSLRINSTATPDQDREKIKQFAAQEE KRQKNERMAHQKHESQMRDLQL QCEANVRELHQLQNEKCHLLVEHE TQKLKELDEEHSQELKEWREKLRPR KKTLEEEFARKLQEQEVFFKMTGES ECLNPSAQSRISKFYPIPTLHSTGS