

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9JM52	MINK1_MOUSE	Mus musculus	Misshapen-like kinase 1	18.754664	NaN	S324;S326;S644;S720;S729;S745;S746;S750;T867;S702	22645316;33300544	MGDPAPARSLDDIDLALRDPAGIFE LVEVVGNGTYGQVYKGRHVKTGQLA AIKVM DVTEDEEEEIKQEINMLKKYS HHRNIATYYGAFIKKSPPGNDDQLW LVMEFCGAGSVTDLVKNTKGNALKE DCIAYICREILRGLAHLHAHKVIHRDI KGQNVLLTENAEVKLVDFGVSAQLD RTVGRRNTFIGTPYWMapeVIACDE NPDATYDYRSDIWSLGITAIEMAEGA PPLCDMHPMRALFLIPRNPpPRLKS KKWSKKFTDFIDTCLIKTYLSRPpTE QLLKFPFIRDQPTerQVRIQLKDHD RSRKKRGEKEETEYEYSGSEEDDS HGEEGEPSSIMNVPGESTLRREFLR LQQENKSNSEALKQQQQLOQQQQR DPEAHIKHLLHQRQRRIEEQKEERR RVEEQQRREERQRKLQEKEQQRRL EDMQALRREEERRQAEREQeYKRK QLEEQRQSERLQRQLQQEHAYLKSL QQQQQQQQQLKQQQQQQQILPGDR KPLYHYGRGINPADKPAWAREVEER ARMNKQQNSPLAKAKPSSAGPEPPI SQASPPPGPLSQTPPMQRpVEPQE GPHKSLVAHRVPLKPYAAPVPRSQS LQDQPTRNLAAFPASHDPDPAAVPT PTATPSARGAVIRQNSDPTSEGPGPS PNPPSWVRPDNEAPPKVPQRTSSIA TALNTSGAGGSRPAQAVRASNPDLR RSDPGWERSDSVLPASHGHLPOAG SLERNRNRVGASTKLDSSPVLSPGN KAKPEDHRSRGRPASyKRAIGEDF VLLKERTLDEAPKPPKAMDYSSSS EEVESSEEEEEEGDGEPSEGSRDTP GGRSDGDTDSVSTMVVDVVEISGT QPSYGGGTmVVQRTPEEERSLLLAD SNGYTNLPDVVQPSHSPTENSKGQS PPTKDGGSDYQSRGLVKAPGKSSFT MFVDLGIYQPGGSGDTIPITALVGGE GGRLDQLQFDVRKGSVVNVNPTNT RAHSETPEIRKYKKRFNSEILCAALW GVNLLVGTENGLMLLDRSGQGVY GLIGRRRFQQMDVLEGLNLLITISGK RNKLRVYYLSWLRNKILHNDPEVEK KQGWTTVGDMEGCGHYRVVKYERI KFLVIALKNSVEVYAWAPKPYHKFM AFKSFADLPHRPLLVDLTVEEGQRL KVIYGSSAGFHAVDSDGNSYDIYIP VHIQSQITPHAIIFLPNTDGMEMLLC YEDEGVYVNTYGRiIKDVVLQWGEM PTSVAYICSNQIMGWGEKAIEIRSVE

