

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8N4C8	MINK1_HUMAN	Homo sapiens	Misshapen-like kinase 1	11.460591	T672	S324;S326;S641;S701;S754;S763;S777;S778;S782;T891	28657654	MGDPAPARSLDDIDLSALRDPAGIFE LVEVVGNGTYGQVYKGRHVKTGQLA AIKVM DVTEDEEEEIKQEINMLKKYS HHRNIATYYGAFIKKSPPGNDQQLW LVMEFCGAGSVTDLVKNTKGNALKE DCIAYICREILRGLAHLHAHKVIHRDI KGQNVLLTENA EVKLVD FGVSAQLD RTVGRRNTFIGTPYWMAPEVIACDE NPDATYDYRS DIWSL GITAIEMAEGA PPLCDMHPMRALFLIPRNPPRLKS KKWSKKFIDFIDTCLIKTYLSRPPT EQ LLKFPFIRDQPTERQVRIQLKDHIDR SRKKRGEKEETEY EYSGSEEEEDDSH GEEGEPSSIMNVPGESTLRREFLRL QQENKSNSEALKQQQQQLQQQQQRD PEAHIKHLLHQRRRIEEQKEERRR VEEQQRREREQRKLQEKEQRRLE DMQALRREEERRQAEREQEYKRKQ LEEQRQSERLQRQLQQEHAYLKS LQ QQQQQQQLQKQQQQQLLPGDRKPL YHYGRGMNPADKPAWAREVEERTR MNKQQNSPLAKSKPGSTGPEPPIQ ASPGPPGPLSQTPPMQRPVEPQEGP HKSLVAHRVPLKPYAAPVPRSQSLQ DQPTRNLAAFPASHDPDPAIPAPTAT PSARGAVIRQNSDPTSEGPGSPNPP AWVRPDNEAPPKVPQRTSSIATALN TSGAGGSRPAQAVRARPRSNSAWQI YLQRR AERGTPKPPGPPAQPPGPPN ASSNPDLRRSDPGWERSDSVLPASH GHL PQAGSLERNRVGVSSKPDSSPV LSPGNKAKPDDHRSRPGRPAD FVLL KERTLDEAPRPPKAMDYSSSSEEV ESSEDEEEEGEGGPAEGSRDTPGGR SDGDTDSVSTMVVHDVEEITGTQPP YGGGTMVVQRTPEEERNLLHADSN GYTNLPDVVQPSHSPTENSKGQSPP SKDGSGDYQSRGLVKAPGKSSFTMF VDLGIYQPGGSGDSIPITALVGGEGT RLDQLQYDVRKGSVVNVNPTNTRA HSETPEIRKYKKRFNSEILCAALWGV NLLVGTENGLMLLDRSGQKVG LI GRRRFQQMDVLEGLNLLITISGKR N KLRVYYSWLRNKILHNDPEVEKKQ GWTTVGDMEGCGHYRVVKYERIKFL VIALKSSVEVYAWAPKPYHKFMAFK SFADLPHRPLLVDLTVEEGQRLKVIY GSSAGFHAVDVDSGNSYDIYIPVHIQ SQITPHAIIFLPNTDGMEMLLCYEDE

