

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q8N4C8	MINK1_HUMAN	Homo sapiens	Misshapen-like kinase 1	33.867134	T672	S324;S326;S641;S701;S754;S763;S777;S778;S782;T891	39894887;28657654	MGDPAPARSLDDIDL.SALRDPAGIFE LVEVVGNGTYGQVYKGRHVKTGQLA AIKVM DVTEDEEEIKQEINMLKKYS HHRNIATYYGAFIKKSPGNDQLW LVMEFCGAGSVTDLVKNTKGNALKE DCIAYICREILRGLAHLHAHKVIHRDI KGNQVLLTENAEVKLVDFGVSQAQLD RTVGRRNTFIGTPYWM APEVIACDE NPDATYDYRSDIWSLGITAIEMAEGA PPLCDMHPMRALFLIPRNPPLKLS KKWSKFFIDFIDTCLIKTYLSRPTEQ LLKFFPIRDQPTERQVRIQLKDHIDR SRKKRGEKEETEYEYSGSEEDDSH GEEGEPSSIMNVPGESTLRREFLRL QQENKSNSEALKQQOQLQQOQRD PEAHIKHLLHQRRRIEEQKEERRR VEEQRREREQRKLQEKEQQRRL DMQALRREERRQAEREQEYKRKQ LEEQRQSERLQRQLQEHAYLKSLO QQQQQQQLKQOQQQLLPGDRKPL YHYGRGMNPADKPAWAREVEERTR MNKQNSPLAKSKPGSTGPEPPIPO ASPGPPGPLSQTPPMQRPVPEQEGP HKSVAHRVPLKPYAAPVPRSLSQ DQPTRNLAAFPASHDPDPAIPAPTAT PSARGAVIRQNSDPTSEGPGSPNPP AWVRPDNEAPPKVPQRTSSIATALN TSGAGGSRPAQAVRARPRNSAWQI YLORRAERCTPKPPGPPAQP GPPN ASSNPDLRRSDPGWERSDVL PASH GHLPQAGSLERNRVGVSSKPDSSPV LSPGNKAKPDDHRSRGRPADFVLL KERTLDEAPRPPKAMDYSSSEEV ESSEDDDEEGEGGPAEGSRDTPGGR SDGDTDSVSTMVVDVEEITGTQPP YGGGT MVVQRTPEERNLLHADS GYTNLPDVVQPSHSP TENS KGQSP SKDGGSDYQSRGLVKAPGKSSFTMF VDLGIYQPGSGSIPITALVGGEGT RLDQLQYDVRKGSVNVNPTNTRA HSETPEIRKYKRFNSEILCAALWGV NLLVGTENGLMLLDRSGQKVYGLI GRRRFQOMDVLEGLNLLITISGKR KLRVYLSWLRNKILHNDPEVEKKQ GWTTVGDMEGCGHYRVVYKERIKFL VIALKSSVEVYAWAPKPYHKFMAFK SFADLPHRPLLDLTVEEGRQLKVIY GSSAGFHAVD VDSGNSYDIYIPVHIQ SQITPHAIIFLPNTDGMEMLLCYEDE GVVNTYGRHKD VVLQW GEMPTSV AYICSNQIMGWGEKAIEIRSVETGHL DGVFMHKRAQRLKFLCERNDKVFF ASVRS GSSQVYFMTLNRNCIMNW	True	True	4.391	2.953	False	False	5.0	False