	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	extra regio
0	Q9QXL1	KI21B_MOUSE	Mus musculus	Kinesin-like protein KIF21B	28.898226	T287	S580;T583;S1150;S1168;S1217;T1239;S1243	34105348;30059200	MAGQGDCCVKVAVRIRPQLSKEKIE GCHICTSVTPGEPQVLLGKDKAFTYD FVFDLDTWQEQIYSTCVSKLIEGCFE GYNATVLAYGQTGAGKTYTMGTGFD TVTSEEEQGIIPRAIAHLFRGIDERKR RAQEKGVTGPEFKVSAQFLELYNEEI LDLFDSTRDPDARHRRSNIKIHEDA NGGIYTTGVTSRLINSQEELIQCLKQ GALSRTTASTQMNVQSSRSHAIFTIH LCQMRVCAQPDLVNETVTGLPDGA APTGTEYETLTAKFHFVDLAGSERLK RTGATGERAKEGISINCGLLALGNVI SALGDQSKKVVHVPYRDSKLTRLLQ DSLGGNSQTIMIACVSPSDRDFMET LNTLKYANRARNIKNKVVVNQDKTS QQISALRAEIARLQMELMEYKAGKR VIGEDGTEGYSDLFRENAMLQKENG ALRLRVKAMQEAIDAINNRVTQLMS QEANLLLAKAGDGNEAIGALIQNYIR EIEELRTKLLESEAMNESLRRSLSRA SARNPYSLGASPAGPAFGGSPATSM EDASEVIRKAKQDLERLKKKEVRQR RKSPEKEAFKKRAKLQAENSEETDE NEAEEEEEERDESGCEEEEGREDE DEDSGSEESLVDSDSDPEEKEVNFQ ADLADLTCEIEIKQKLIDELENSQRR LQTLKHQYEEKLILLQNKIRDTQLER DRVLQNLSTMECYTEEKANKIKADY EKRLREMNRDLQKLQAAQKEHARL LKNQSRYERELKKLQAEVAEMKKAK VALMKQMREEQQRRRLVETKRNRE IAQLKKEQRRQEFQIRALESQKRQQ EIVLRRKTQEVSALRRLAKPMSERVA GRVGLKPPNMDSGAEVSASTTSSEA ESGARSVSSIVRQWNRKIDHFLGDR PTATVNGGRPARKKFQKKGASQSFS KAARLKWQSLERRIIDIVMQRMTIV NLEADMERLIKKREELFLLQEALRR KREHLQAESPEEEKGLQELAEEIEVL AANIDYINDSITDCQATIVQLEETKEE LDSTDTSVVISSCSLAEARLLLDNFL KASIDKGLQVAQKEAQIRLLEGRLRQ TDMTGSSQNHLLLDALREKAEAHPE LQALIYNVQHENGYASTDEEVSEFS EGSFSQSFTMKGSTSHDDFKFKGEP KLSAQMKAVSAECLGPPLDSSTKNIT KSLASLVEIKEDGVGFSIRDPYYRDK VSRTVSLPTRGSTFPRQSRGATDTSP LTRRKSYDRGQPIRSTDMGFTPPSSP PTRPRNDRNVFSRLTSNQSQGSALD KSDDSDSSLSEVLRGIITPIGGAKGA RTAPLQCISMAEGHTKPILCLDATDE LLFTGSKDRSCKMWNLVTGQEIAAL KGHPNNVVSIKYCSHSGLVFSVSSS YIKVWDIRDSAKCIRTLTSSGQVISG DACIATSTRAITSAQGEHQINQMALS PSGSMLYVASGNAVRIWELNRFQPI GKLTGHIGPVMCLTVTQTSNQHDLV VTGSKDHYVKMFQLGDCVTGTIGPT HNFEPPHYDGIECLAIQGDILFSGSR DNGIKKWDLDQQELIQQIPNAHKD WVCALAFVPGRPMLLSACRAGFIKV WNVDNFTPIGEIKGHDSPINAICTNS KHIFTASSDCRVKLWNYVPGLTPCL PRRVLAIKGRAPPCPDLPPPPLTLPIL PFPVFPPPRSELLLHVT	None	None	None	None	None	None	None	None	None