

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	p
Q02241	KIF23_HUMAN	Homo sapiens	Kinesin-like protein KIF23	19.658599	T758	S155;S160;S605;S682;S684;T738;S814;S867;S902;S911;T927	40596516;34019948;40914422;30379171	MKSARAKTPRKPTVKKGSQTNLKDP VGVYCRVRPLGFPDQECIEVINNTT VOLHTPEGYRLNRNGDYKETQYSFK QVFGTHTTQKELFDVVANPLVNDLI HGKNGLLFTYGVGTSGKTHMTGSP GEGGLLPRCLDMIFNSIGSFQAKRY VFKSNDRNSMDIQCEVDALLERQKR EAMPNPKTSSSKRQVDFEADMIV QEFCKAEEVDEDSVYGVFVSYIEIYN NYIYDLLEVPFDPKPKPPQSKLLRE DKNHNMYVAGCTEVEVKSTEEAFE VFWRGQKRRRIANTHLNRESSRSHS VFNIKLVQAPLDADGDNLQEKIQI TISQSLVLDLAGSERTNRTRAEGNRL REAGNIQSLMTRLTCMDVIRENO MYGTNKMVPYRDSKLTFLFKNYFD GEGKVRMIVCVNPKAEDYEENLQV MRFAEVTQEEVEVARPVDKAICGLTP GRRYRNQPRGPVGNPEPLVDVVLQS FPPLPSCIEILDINDEQTLPRLIEALEK RHNLRQMMIDEFNKQSNAFKALLQ EFDNAVLSKENHMQGLNEKEKMI SGQKLEIERLEKKNKTLEYKIEILEKT TTIYEEDKRNLDQEELETQNKLQRO FSDKRRLEARLQGMVTETTMKWEK ECERRVAAKQLEMQNKLVWKDEKL KQLKAVTEPKTEKPERPSRERDREK VTQRSVSPSPVPLSSNIAQISNGQQ LMSQQLHRRSNSCSSISVASCISE WEQKIPTYNTPLKVTSIARRRQVEPG QSKTCIVSDRRRGMWTEGREVVPT FRNEIEIEEDHCGRLLFPDQNPPI RLRHRRRSRAGDRWVDHQPASNMQ TETVMOPHVPHAITVSVANEKALAK CEKYMLTHQELASDGEIETKLIKDI YKTRGGGQSVQFTDIETLKQESPNG SRKRRSSTVAPAPDGAESEWTDVE TRCSVAVEMRAGSQLGPGYQHHAQ PKRKKP	True	False	4.638	5.0	3.636	1.434	1.502	2.