

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	extra regic
Q02241	KIF23_HUMAN	Homo sapiens	Kinesin-like protein KIF23	14.702989	T758	S155;S160;S605;S682;S684;T738;S814;S867;S902;S911;T927	30379171;34019948	MKSARAKTPRKPTVKKGSQTNLKDP VGVCYCRVRLGFPDQECCIEVINNTT VOLHTPEGYRLNRNGDYKETQYSFK QVFGTHHTQKELFDVVANPLVNDLI HGKNGLLFTYGVTVSGKTHMTGSP GEGGLLPRCLDMIFNSIGSFQAKRY VFKSNDRNSMDIQCEVDALLERQKR EAMPNPKTSSSKRQVDPEFADMIV QEFCKAEVDEDSVYGVFVSYIEIYN NYIYDLLEEVFFDPIKPKPPQSKLLRE DKNHNMYVAGCTEVEVKSTEEAFE VFWRGQKRRRIANTHLNRESSRSHS VFNIKLVQAPLDADGDNVLQEKEQI TISQLSLVDLAGSERTNRTRAEGNRL REAGNINQSLMTRLRTCMDVLRNO MYGTNKMVVPYRDSKLTHLFKNYFD GEGKVRMIVCVNPKAEDYEENLQV MRFAEVTQEVEVARPVDKAICGLTP GRRYRNQPRGPVGNELVTDVVVLOS FPPLPSCIELDINDEQTLPRLIEALEK RHNLRQMMIDEFNKQSNAFKALLO EFDNAVLSKENHMQGKLNKEKEMI SGQKLEIERLEKKNKTLEYKIEIELEKT TTHYEDKRNLQEQLETONQKLRQ FSDKRRLEARLQGMVTETTMKWEK ECERRVAAKQLEMQNKLWVKDEKL KQLKAVTEPKTEKPERPSRERDREK VTQRSVSPSPVPLSSNYIAQISNGQQ LMSQPQLHRRSNSCSSISVASCISE WEQKIPTYNTPLKVTSIARRRQEPG QSKTCIVSDRRRGMWYTEGREVVPT FRNEIEIEDHCGRLLFQPDQNAPPI RLRHRRSRVAGDRWVDHQPASNMQ TETVMQPHVPHAITVSVANEKALAK CEKYMLTHQELASDGEIETKLIKGD YKTRGGGQSVQFTDIETLKQESPNG SRKRRSSTVAPAQPDGAESEWTDVE TRCSVAVEMRAGSQLGPGYQHHAQ PKRKKP	True	False	4.638	5.0	1.597	1.44	1.507	2.071	1.772

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