

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q96JM4	LRIQ1_HUMAN	Homo sapiens	Leucine-rich repeat- and IQ domain-containing protein 1	32.252584	S1590	NaN	39894887;30620550	MDDDDAKLKAIEAELDKLSSISLEK EDIESDAKSETQSDSDTDSVELPES VLHCINIKNRSKAVEELIQDLEDT DILSCSYGAVSNNHMLRTRGLSTEY EESSEQLIKLSEIEKEEFMRSKTDC ATPDFVPEPSPHDLPMDEHVLPPDA DINFGYCEVEEKCROSFEEAWQEKQK ELEDKEKQTLKAQRDREEKQFQEEE EKRHCMWKQFKVEKKLENIQKQE QDKMNDELYKEEKIWKKEFKQHEE YIRNLHLQMEEEERTRFKDQOEKEK NSLLKQQNNAAVKIQAKYKAFVAYQ KYGPIIKEQIESKKRKAQEWKEKEAK IROKKEENRRKLEEEQRKEERKKQ KEEERKRREKEYEKKNIVKQEREQ LISKEKILREDASQQLIISALKKSGY NNKHLSEDISNDKGDIAKNLVDEN SKKQEDVLLWLVEESNMKENVDRQ TILKESIQVKLKESISSQTLADFKME EKNEENLAKKRCSEELVKQERKYENT DNKTELGNSDLKGNLKEQFPLQELK SDAQKEKIMKHVINENTGQKTOIIL GHNQEISEVKTNEEQKIKDNQQKKI QKVEKEEIQEQNGLLYKDKDTLVISV KQRSLSLTSENSKDVRENVLQEKI YSKSEIEENPKDNAWNNGIVFNNTT DTMINIEGKRNDQDYVLGRHAPCEG LSNYNAESSMVSKEVNSLKSEIRNIS EKCHENAPEPDSMTCCVSESTLLYS IEERRLAWIKSFKPWLEIFKQNOQK KIVRRKRPVKCPANMTPALDKLEILR CGPWDTLQQTVTTFQDLPGCVLST LAEACTNLQFLSLRRCGLTSLHSLSN CKKLYIDAQENHIEAIECENLENLC VLLNKNQTLHLGLDGCNTIQCLE LSYNKITRIGYSFFLEEKLVDNAGFC HHLGTSTSYLSLAQVWIPTGLCWSW IPITSLTKNSDCNFLSHLYWNCGLE SLKNLQQLLDHNLINTKGLCDTPT IVYLDCSHNHLTDVEGVENCGLLQI LKLOGNYLSELPLENLVLRELHL DDNSISTVEAFSSYWLPLLQNTISQ NSLTKIVPLFHFVSLEKLDVSHNCLS DLKSAIKWFDACYSHELSTLGNPLL QETNWRDSSLKVLPALRILNGNILN SNSESRTEEHNQLGSAGFLALCQSQ IREFNLLIENYITGKGDVFTLDTAEN LCHYFKKLMILSTEYRHAHERGDVTI TKKDESEAQKNHLAPTNSDSTLQNG VFYSCAREGEPDSDIPEKWMDSVS SHSPLSKSATCENMEGRHQEILVCO KREDSKASSIPTIRIPFKEVVMNSLL RNHQNIPESEKIMAAVVIQSYWRGY LMRRQTHFSTRLHTAATEGLPNSSI KNQTLKKGKRENIVNIRKQREKAAI LIQAVWKGFIKRLTATAEAIKNEE SDEEYREIDLDFIDEAALEEWLA LDSTRFPSQTLNLSNQLHWPKIPGN LKWDDTSFNLPSNPAQAWLCNDKE NLSSSEHTQFNRSSENKTSWTPES KTSRKSLLKSEKKEKISEEWGFKDIS TAQQMLKRAQKMKSKLKKKIDSTV RLALFKNNENKVSPLPKSPKMVQPRR DGYFEGIEEDPIHKDTTANEKLERN REYTYQWLHTQVGVHETTSRNMK CNHFLPELDPDLNNGRRVQLVARLV SREDTDLDLFSMTNGSALSUNREKK NQAHRHSAGSSSKLWFPSKLI	False	False	1.031	1.875	False	False	False	False	