

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	extrac regior
Q702N8	XIRP1_HUMAN	Homo sapiens	Xin actin-binding repeat-containing protein 1	27.15334	T1137;S1174	S205;S208;S213;S295;S332	31637018;35254053	MADTQTQVAPTPTMRMATAEDLPL PPPPALEDLPLPPPKESFSKFHQQRQ ASELRRLYRHHPELKRKNLAEAVAE DLAEVLGSEEPTEGDVQCMRWIFEN WRLDAIGEHHERPAAKEPVLCCGDVQA TSRKFEEGSFANSTDQEPTRPQPGG GDVRAARWLFETKPLDELTAQAKEL EATVREPAASGDVQGRMLFETRPL DRLGSRPSLQEQSPLELRSEIQELKG DVKKTVKLFQTEPLCAIQDAEGAIHE VKAACREEQSNVRSARWLFETRPL LDAINQDPSQVRVIRGISLEEGARPD VSATRWFETQPLDAIREILVDEKDF QSPDLIPPGPDVQQQHLFETRAL DTLKGDEEAGAEAPPKEEVVPGDVR STLWLFETKPLDAFRDKVQVGHQR VDPQDGEHLSSDSSSALPFSQASAP QRDELKGDVKTFRKLFETLPLDSIG QGEVLAHGSPSREEGTDASAGQAQGI GSPVYAMQDSKGRHLHALTSVSRQI VGGDVQGYRWMFETQPLDQLGRSP STIDVVRGITRQEVVAGDVGTARWLF ETOPLEMIHQEQEQERQKEEGKSQ GDPOEAPPKGDVQTRWLFETCPM SELAEKQGEVTDPTAKAEAQSCW MFKPQVDRPVGSREQLQVQVQV AGERQTDHRVFEFTEPLQASGRPCGR RPVRYCSRVEIPSGQVSRQKEVFQAL EAGKKEEQEPRVIAGSIPAGSVHKFT WLFENCPMGSLAAESIQGGNLEEQ PMSPSGNRMQESQETAAGTLRRL HATPGILHHGGILMEARGPGELCLA KYVLSGTGGHPYIRKEELVSGELPR IICQVLRDPDQVQGLLVQEDPTGQL QLKPLRLPTPGSSGNIEDMDPELQQ LLACGLGTSVARTGLVMQETEQGLV ALTAYSQPRLTSKASERSVVOLLAS CIDKGDLSGLHSLRWEPADPSPVP ASEGAQSLHPTESIHHVPLDPSMG MGHLRASGATPCPPQAIKAVPLAG EAAAPALQNTKEQEDSHSGQKGM AVLGKSEGATTPPGPGAPDLLAAM QSLRMATAEAQSLHQVNLKHKQG PTPTATSNPIQDGLRKAGATQSNIRP GGGSDPRIPAAPRKVSREEQALPRGL PGGWVTIQDGIYTAHPVRTFDPPGG VQLSQREPOSRHRETALSVQAPRPL QGGPGQSTGPGREEPGGCTQMAWG PPGKAMAEEVCPGGLOAAETLKTAP LGRHILASGPAAGASPHPHNAFVP PPPTLPAAVTGPDFPAGAHRAEDSIQ QASEPLKDLLHSHSSPAGQRTPPG SQTKTPLDPTMPPKKKQLPPKPA HLTQSHPPQRLPKPLPSFSFSSEVG QREHQGERDTAIPQPAKVPTTVDO GHIPLARCPGSHSQPSLQHGLSTTA PRPTKNQATGSNAQSSEPPKLNALN HDPTSPQWGPGPSGEQPMEGSHQG APESPDSLQRNQLQGLLNQVQAL EKEAASSVDVQALRRLEAVPQLGG AAPQAPAAHQKPEASVEQAFGELTR VSTEVAQLKEQTLARLLDIEEAVHKA LSSMSSLQPEASARGHFQPPKDHS AHKISVTVSSARPSGSGQEVGGQT AVKNQAKVECHTEAQSQVKIRNHTE ARGHTASTAPSTRRQETSREYLCPPR VLPSSRDSPPSPTFISIQSATRKPLET PSFKGNPDVSVKSTQLAQDIGOALL HQKGVODKTGKDKITQCSVQPEPAP PSASPLPRGWQKSVLELQTGPGSSQ HYGAMRTVTEQYEEVDQFGNTVLM SSTTVTEQAEPPRNPGSHLGLHASP LLRQFLHSPAGFSSDLTEAETVQVS CSYSQAAQ	None	None	None	None	None	None	None		