

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region							
070373	XIRP1_MOUSE	Mus musculus	Xin actin-binding repeat-containing protein 1	25.629171	NaN	S205;S208;S213;S295;S332;S353	36288343	MADAQMVAPTPTIQMRTEEDLSLP HPSAFEGFLPPPPKETFSKFQQQRQ ASELRRLYKHHPELRRNLEEAFAE DLAEVLGSEEPTEGDVQCMRWIFEN WRLDAIGDHERPAAREPVSGGNVQA TSRKFEEGSGFTNSSDQEPEGLRPSG GDVQAARQMFETKPLDALRGQEEA TQTTMREPAATGDVQTRKLFETR LDRLGSRPSIQEQSPLERSEIQELK GDVKKTVKLFQTEPLCAIQDAEGTIH EVKAACREEQSNVRSARWLFETR PLDAFNQDPSQVRVIRGISLEEGALP DVSATRWIFETQPLDAIREIEVDEKD FQPSFDLIPPGPDVQHQRHLFETCSL DTLKERETEAEVPPKEEVIPGDVRS TLWLFETKPLDAFRDQVQVGHQRV GHQEGEGLVTECLPSNGTSVLPVLSQ GVPQNDGLKGDVKTFFKNLFETPLD SIGQGEPSAYGNINRGQNTDSAEQS QGSADPVYAMQDSRQGLHALTSVSR EQVGGDVQGYKWMFETQPLDTLG RSPSTIDVVRGITRQEVVAGDVGTTR WLFETQPLEMIHQEQEQKPEEEEG KGGPGPPPELPPKGDVQTRWLFET YPMSELAEKRESEVTPVSKAETQS CTWMFQPSLNPAGSGEQHLQTS QVPAGDRQTRHVFETESLPSNQS SGRKPVRVYCSRVEIPSGQVSRQKEVF QALEAGKKEVPETTINLGSIPTGSVH KFTWLFENCMPGSLAAESIRGDNLQ EEQPKGSAGHGTPEROETAARTLR TLHATPGILHHGGILMEARGPGLC LAKYVLPSPGQGRPYRKEELVCGEL PRIVRQVLRRTDQGLLVQEDTA GQLQLHPLMLPGDPGNIEDMDP ELQQLLACGLGVSVSKTGLVMQETG QGLVALTAYSQPLTSRAPERSVQ LLASCIDKDLHSLHSLRWEPTDTP SSGPATEESQVRVPTESIHHVTPDST MEMGQLRISGSTPCPPPSRAAGKVV LPNGKPVAQAPLQEARKKRDISHAG QKGAASGRPEASPLGSGAPDLQEA MQNLRRLATAEAQSLHQVLSRHPQ GSDPVATSMQVQDVLQASTPATGVT QGSIRPVAGSEARIPAVPRKLL	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None