	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparat
0	P97434	MPRIP_MOUSE	Mus musculus	Myosin phosphatase Rho-interacting protein	43.165225	NaN	S193;S219;S221;S225;S227;S266;S270;S289;S292;T295;S364;S492;S617;T645;S799;S976;S992;S1013;S1015;S1015;S979	22517741;30031227;34887587;22645316	MSAAKENPCRKFQANIFNKSKCQN CFKPRESHLLNDEDLTQAKPIYGGW LLLAPDGTDFDNPVHRSRKWQRRF FILYEHGLLRYALDEMPTTLPQGTIN MNQCTDVVDGEARTGQKFSLCILTP DKEHFIRAETKEIISGWLEMLMVYP RTNKQNQKKKRKVEPPTPQEPGPAK MAVTSSSGGSSGSSSSIPSAEKVPTT KSTLWQEEMRAKDQPDGTSLSPAQ SPSQSQPPAACTPREPGLESKEDEST ISGDRVDGGRKVRVESGYFSLEKAK QDLRAEEQLPPLLSPPSPSTPHSRRS QVIEKFEALDIEKAEHMETNMLILTT PSSDTRQGRSERRAIPRKRDFASEAP TAPLSDACPLSPHRRAKSLDRRSTES SMTPDLLNFKKGWLTKQYEDGQWK KHWFVLADQSLRYYRDSVAEEAADL DGEINLSTCYDVTEYPVQRNYGFQIH TKEGEFTLSAMTSGIRRNWIQTIMK HVLPASAPDVTSSLPEGKNKSTSFET CSRSTEKQEAEPGEPDPEQKKSRAR ERRREGRSKTFDWAEFRPIQQALAQ ERASAVGSSDSGDPGCLEAEPGELE RERARRREERRKRFGMLDTIDGPG MEDTALRMDIDRSPGLLGTPDLKTQ NVHVEIEQRWHQVETTPLREEKQVP IAPLHLSLEDRSERLSTHELTSLLEK ELEQSQKEASDLLEQNRLLQDQLRV ALGREQSAREGYVLQATCERGFAAM EETHQKKIEDLQRQHQRELEKLREE KDRLLAEETAATISAIEAMKNAHREE MERELEKSQRSQISSINSDIEALRRQ YLEELQSVQRELEVLSEQYSQKCLE NAHLAQALEAERQALRQCQRENQE LNAHNQELNNRLAAEITRLRTLLTG DGGGESTGLPLTQGKDAYELEVLLR VKESEIQYLKQEISSLKDELQTALRD KKYASDKYKDIYTELSIAKAKADCDIS RLKEQLKAATEALGEKSPEGTTVSGY DIMKSKSNPDFLKKDRSCVTRQLRN IRSKSVIEQVSWDN	None	None	None	None	None	None	None