	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q16821	PPR3A_HUMAN	Homo sapiens	Protein phosphatase 1 regulatory subunit 3A	22.903221	T927	S38;S42;S46;S49;T56;S65;S844	38253038	MEPSEVPSQISKDNFLEVPNLSDSL CEDEEVTFQPGFSPQPSRRGSDSSE DIYLDTPSSGTRRVSFADSFGFNLVS VKEFDCWELPSASTTFDLGTDIFHT EEYVLAPLFDLPSSKEDLMQQLQIQ KAILESTESLLGSTSIKGIIRVLNVSF EKLVYVRMSLDDWQTHYDILAEYVP NSCDGETDQFSFKIVLVPPYQKDGS KVEFCIRYETSVGTFWSNNNGTNYT FICQKKEQEPEPVKPWKEVPNRQIK GCLKVKSSKEESSVTSEENNFENPK NTDTYIPTIICSHEDKEDLEASNRNV KDVNREHDEHNEKELELMINQHLI RTRSTASRDERNTFSTDPVNFPNKA EGLEKKQIHGEICTDLFQRSLSPSSS AESSVKGDFYCNEKYSSGDDCTHQP SEETTSNMGEIKPSLGDTSSDELVQ LHTGSKEVLDDNANPAHGNGTVQIP CPSSDQLMAGNLNKKHEGGAKNIE VKDLGCLRRDFHSDTSACLKESTEE GSSKEDYYGNGKDDEEQRIYLGVNE KQRKNFQTILHDQERKMGNPKISVA GIGASNRDLATLLSEHTAIPTRAITAD VSHSPRTNLSWEEAVLTPEHHHLTS EGSALGGITGQVCSSRTGNVLRNDY LFQVEEKSGGINSEDQDNSPQHKQS WNVLESQGKSRENKTNITEHIKGQT DCEDVWGKRDNTRSLKATTEELFTC QETVCCELSSLADHGITEKAEAGTAY IIKTTSESTPESMSAREKAIIAKLPQE TARSDRPIEVKETAFDPHEGRNDDS HYTLCQRDTVGVIYDNDFEKESRLGI CNVRVDEMEKEETMSMYNPRKTH DREKCGTGNITSVEESSWVITEYQKA TSKLDLQLGMLPTDKTVFSENRDLR QVQELSKKTDSDAIVHSAFNSDTNR APQNSSPFSKHHTEISVSTNEQAIAV ENAVTTMASQPISTKSENICNSTREI QGIEKHPYPESKPEEVSRSSGIVTSG SRKERCIGQIFQTEEYSVEKSLGPMI LINKPLENMEEARHENEGLVSSGQS LYTSGEKESDSSASTSLPVEESQAQG NESLFSKYTNSKIPYFLLFLIFLITVY HYDLMIGLTFYVLSLSWLSWEEGRQ KESVKKK	False	False	2.902	3.009	2.403	2.155	1.169	2.369	2.156