	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasm mem
0	O35954	PITM1_MOUSE	Mus musculus	Membrane-associated phosphatidylinositol transfer protein 1	35.498311	NaN	T59;T282;T287;S300;S304;S319;S326;S329;S342;S345;S346;S373;S382;S593;S600;S621;S895;S1236	34678516;22645316	MLIKEYHILLPMSLDEYQVAQLYMIQ KKSREESSGEGSGVEILANRPYTDGP GGNGQYTHKVYHVGSHIPGWFRAL LPKAALQVEEESWNAYPYTRTRYTC PFVEKFSIEIETYYLPDGGQQPNVFN LSGAERRQRIVDTIDIVRDAVAPGEY KAEEDPRLYRSAKTGRGPLADDWAR TAAQTGPLMCAYKLCKVEFRYWGM QAKIEQFIHDVGLRRVMLRAHRQA WCWQDEWIELSMADIRALEEETAR MLAQRMAKCNTGSEGPEAQTPGKS STEARPGTSTAGTPDGPEAPPGPDAS PDASFGKQWSSSSRSSYSSQHGGGV SPQSLSEWRMQNIARDSENSSEEEF FDAHEGFSDSDEVFPKEMTKWNSN DFIDAFASPTEVEGVPDPTVMATKGI EDGARAPRDSEGLDGAGDLVVEACS VHALFLILHSGSILDSGPGDTNSKQA DVQTLSTAFEAVTRVHFPEALGHVA LRLVPCPPICAAAYALVSNLSPYSHD GDSLSRSQDHIPLAALPLLATSSSRY QGAVATVIARTNQAYAAFLRSSEGTG FCGQVVLIGDGVGGILGFDALCHSA SAGPGSRGSSRRGSMNNEMLSPEV GPVRDPLADGVEVLGRASPEPSALPA QRTFSDMANPDPDGSQNSLQVAST ATSSGEPRRASTASCPPASSEAPDGP TNAARLDFKVSGFFLFGSPLGLVLAL RKTVMPALEVAQLRPACEQIYNLFH AADPCASRLEPLLAPKFQAIAPLAVP RYQKFPLGDGSSLLLADTLQTHSSLF LEELEMMVPSTPTSASGAFWKGSEL GNEPASQTAAPSTTSEVVKILDRWW GNKRIDYSLYCPEALTAFPTVTLPHL FHASYWESADVVAFILRQVIEKERPQ LTECEEPSIYSPAFPREKWQRKRTQV KIRNVTSNHRASDTVVCEGRPQVLN GRFMYGPLDVVTLTGEKVDVYVMTQ PLSGKWIHFGTEVTNSSGRLTFPVPS ERALGIGVYPVRMVVRGDHTYAECC LTVVSRGTEAVVFSIDGSFTASVSIM GSDPKVRAGAVDVVRHWQDSGYLIV YVTGRPDMQKHRVVAWLSQHNFPH GVVSFCDGLTHDPLRQKAMFLQSLV QEVELNIVAGYGSPKDVAVYAALGLS PSQTYIVGRAVRKLQAQCQFLSDGYV AHLGQLEAGSHSHAPSGPPRAALAK SSYAVAAPVDFLRKQSQLLRSRGPS QVDREGPGTPPTTLARGKTRSISLKL DSEE	None	None	None	None	None	None	None	None