	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	extracel region
0	Q3V3Q7	PACS2_MOUSE	Mus musculus	Phosphofurin acidic cluster sorting protein 2	32.157358	T393	S361;S387;S424;S662;S665	35822049;40021952	MAERGRLGLPGAPGALNTPVPMNL FATWEVDGSSPSCVPRLCSLTLKKL AVLRELEKELLSVVIAVKMQYPHFLK REGNKLQIMLQRRKRYKNRTILGYK TLAAGSINMAEVMQHPSEGGQVLSL CSSIKEASVKVAEIWIVSLSSQPIDHE DSAMQAGPKTKSTDNYSEEEYESFS SEQEASDDAVQGQDLDEDDFDVGK PKKQRRSIVRTTSMTRQQNFKQKVV ALLRRFKVSEEVLDSEQDPAEHVPE VEEDLDLLYDTLDVENPSDSGPDMD DDDSVLSTPKPKLRPYFEGLSHSSSQ TEIGSIHSARSHREPPSPADVPEKTR SLGGKQQLSDSVSDTVALSAAVPRE PSGQPEDSPEAETSTLDVFTEKLPPS GRIIKTESLVIPSTRSESKPAGRRGRS TSLKERQPARPQNERANSLDNERCP DTRSQLQIPRKTVYDQLNHILISDDQ LPENIILVNTSDWQGQFLSDVLQKH TLPVVCTCSAADVQAAFSTIVSRIQR YCNCNSQPPTPVKIAVAGAQHYLSAI LRLFVEQLSHKTPDWLGYMRFLIIPL GSHPVARYLGSVDYRYNNFFQDLA WRDLFNKLEAQSSVQDTPDIVSRIT QYISGANCAHQLPIAEAMLTYKQKSP DEESSQRFIPFVGVVKVGIVEPSSAT SGDSDDAAPSSSSILSSTPPSASTSPA AKEASPTPPSSPSVSGGLSSPSQGVG AELMGLQVDYWTAAQPADRKRDAE KKDMPTTKNTLKCTFRSLQVSRLPS SGEAAATPTMSMTVVTKEKNKKVM FLPKKTKDKEVESKSQCIEGISRLICT AKHQQNMLRVLIDGVECSDVKFFQL AAQWSSHVKHFPICIFGHSKATF	False	False	0.375	2.531	3.0	3.0	False	False	False