

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane
Q8K212	PACS1_MOUSE	Mus musculus	Phosphofurin acidic cluster sorting protein 1	37.975063	S772	S28;T44;T249;S377;S379;S428;S493;T502;S517;S526;S527;S529;S532	33300544;40885482	MAERGGAGGGPGGGSSQKRSQ VAQSPQQPQQPPQPPQPTPKLA QATSSSSSTSAAAAASSSSSTSTMA VAVAGSAPPGGPGGRTPAPVQMN LYATWEVDRSSSSCVPRFLSLTKKL VMLKEMDKDLNSVVIIVKLGSKRI LRSNEIILPASGLVETELQLTFSLQYP HFLKRDANKLQIMLQRRKRYKNRTI LGYKTAVGLINMAEVMQHPNEGAL VLGLHSNVKDVSVPAEIKIYSLSSQ PIDHEGKSKLSDRSPDIDNYSEEEE ESFSSEQEGSDDPLHGQDLFYEDD LRKVKTRRRLTSTSAITRQPNKOK FVALLKRFKVSDEVGFLEHVSREQI REVEEDLDELVDSEMYNPSDSGPE MEETESILSTPKPKLPFFEGMSQSS SQTEIGSLNSKGLGKDTTSPMELA ALEKVKSTWIKNQDDSLTETDTLEIT DQDMFGDVSTSLVVPEKVKTPMKSS KADLQGSASPSKVEGHTPRQKRST PLKERQLSKPLSERTNSSDSERSPDL GHSTQIPRKVVYDQLNQLVSDAALP ENVILVNTTDWQGQYVAELLQDQRK PVVCTCSTVEVQAVLSALLTRIORYC NCNSSMPRPVKVAAVGSQSYLSSIL RFFVKLSLASKTPDWLGYMRFLIPLG SHPVAKYLGVSVDVSRYSSTFLDSAWR DLFSRSEPPVSEPLDVVGRVMQYVN GATTHQLPVAEAMLTCTRHKFPDED SYQKFIPIFGVVKVGLVEDSPSTAGD GDDSPVSLTVPSTSPSSSGLSRDA TATPPSSPSMSSALAVGSPNSPYGD VIGLQVDYWLGHGERRREGDKRD ASSKNTLKSVMFRSVQVSRPLPHAGEA QLSGTMAMTVVTKEKNKKVPTIFLS KKPREKEVDKSKQVIEGISRLICSAK QQQTMRLRVSIDGVEWSDIKFFQLAA QWPTHVKHFFPVGLFSGSKTT	False	False	1.731	3.44	1.019	1.345	3.868	1.689

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