

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8K212	PACS1_MOUSE	Mus musculus	Phosphofurin acidic cluster sorting protein 1	9.108089	NaN	S28;T44;T249;S377;S379;S428;S493;T502;S517;S526;S527;S529;S532	33300544	MAERGGAGGGPGGSGGGSSQRGSG VAQSPQQPQQPPQPQQPTPPKLA QATSSSSSTSAAAAASSSSSTSTMA VAVASGSAPPGGPGPGRTPAPVQMN LYATWEVDRSSSSCVPRFLSLTLKKL VMLKEMDKDLNSVVIKLVKQSKRI LRSNEIILPASGLVETELQLTFSLQYP HFLKRDANKLQIMLQRRKRYKNRTI LGYKTLAVGLINMAEVMQHPNEGAL VLGLHSNVKDVSVPAEIKIYSLSSQ PIDHEGKSKLSDRSPDIDNYSEEEE ESFSSEQEGSDDPLHGQDLFYEDD LRKVKKTRRKLSTSAITRQPNIKQK FVALLKRFKVSDEVGFGLEHVSREQI REVEEDLDELYDSLEMYPNSDSGPE MEETESILSTPKPKLKPFFEGMSQSS SQTEIGSLNSKGLGKDTTSPMELA ALEKVKSTWIKNQDDSLTETDTLEIT DQDMFGDVSTSLVVPEKVKTPMKSS KADLQGSASPSKVEGHTTPRQKRST PLKERQLSKPLSERTNSSDSERSPDL GHSTQIPRKVVYDQLNQILVSDAALP ENVILVNTTDWQGGQYVAELLQDQRK PVVCTCSTVEVQAVLSALLTRIQRYS NCNSSMPRPVKVAAVGSQSYLSSIL RFFVKSLASKTPDWLGYMRFLIPLG SHPVAKYLGVSYSSTFLDSAWR DLFSRSEPPVSEPLDVVGRVMQYVN GATTHQLPVAEAMLTCTRHKFPDED SYQKFIPFIGVVKVGLVEDSPSTAGD GDDSPVSLTVPSTSPSSSGLSRDA TATPPSSPSMSSALAVGSPNSPYGD VIGLQVDYWLGHGERRREGDKRD ASSKNTLKSFRSVQVSRLPHAGEA QLSGTMAMTVVTKEKNKKVPTIFLS KKPREKEVDSKSQVIEGSRILCSAK QQQTMLRVSIDGVEWSDIKFFQLAA QWPTHVKHFPVGLFSGSKTT