

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
088588	PACS1_RAT	Rattus norvegicus	Phosphofurin acidic cluster sorting protein 1	12.481722	NaN	S28;T44;T249;S377;S379;S428;S493;T502;S517;S526;S527;S529;S532	18683930	MAERGGAGGGPGGAGGGSSQRGSG VAQSPQQPPQPSQPQPTPPKLA QATSSSSSTSAAAASSSSSTSTMSMA VAVASGSAPPGGPGPGRTPAPVQMN LYATWEVDRSSSSCVPRFLSLTLKKL VMLKEMDKDLNSVVIKLVKQSGKRI LRSNEIILPASGLVETELQLTFSLQYP HFLKRDANKLQIMLQRRKRYKNRTI LGYKTLAVGLINMAEVMQHPNEGAL VLGLHSNVKDVSVPAEIKIYSLSSQ PIDHEGKSKLSDRSPDIDNYSEEEE ESFSSEQEGSDDPLHGQDLFYEDD LRKVKKTRRKLSTSAITRQPNIKQK FVALLKRFKVSDEVGFGLHVSREQI REVEEDLDELYDSLEMYPNSDSGPE MEETESILSTPKPKLKPFFEGMSQSS SQTEIGSLNSKGLGKDTTSPMELA ALEKVKSTWIKNQDDSLTETDTLEIT DQDMFGDASTSLVVPEKVKTPMKSS KADLQGSASPSKVEGHTTPRQKRST PLKERQLSKPLSERTNSSDSERSPDL GHSTQIPRKVVYDQLNQILVSDAALP ENVILVNTTDWQGGYVAELLQDQRK PVVCTCSTVEVQAVLSALLTRIQRYS NCNSSMPRPVKVAAVGSQSYLSSIL RFFVKSLASKTPDWLGHMRFLIVPL GSHPVAKYLGSDSRYSSTFLDSAW RDLFSRSEPPVSEPLDVVGRVMQYV NGATTTTHQLPVAEAMLTCTRHKFPDE DSYQKFIPFIGVVKVGLVEDSPSTAG DGDDSPVSLTVPSTSPSSSGLSRD ATATPPSSPSMSSALAVGSPNSPYG DVIGLQVDYWLGHGERRREGDKR DASSKNTLKSFRSVQVSRPLPHAGE AQLSGTMAMTVVTKEKNKKVPTIFL SKKPREKEVDSKSQVIEGISRLICSAK QQQTMLRVSIDGVEWSDIKFFQLAA QWPTHVKHFPVGLFSGSKPT