

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membr
Q9BZ71	PITM3_HUMAN	Homo sapiens	Membrane-associated phosphatidylinositol transfer protein 3	22.903221	S750;S756;T782	S30;S31;S109;S295;S298;S321;S343;S495;S612;S907;S928;S946	38253038	MAKAGRAGGPPPGGGAPWHLRNVL SDSVESDDEFFDAREEMAEGKNAI LIGMSQWNSNDLVEQIETMGKLEDE HQEGGTAPCTSSILQEKQRELYRVSL RRQRFPAQGSIEIHEDSEEGCPQRS CKTHVLLLVHGGNILDGTAGDPSC KAADIHTFSSVLEKVTTRAHFPAALGH ILIKFVPCPAICSEAFSLVSHLNPYSH DEGCLSSSQDHVPLAALPLLAISSPQ YQDAVATVIERANQVYREFLKSSDGI GFSGQVCLIGDCVGLLAFDAICYSA GPSGDSPASSSRKGSISSTQDTPVAV EEDCSLASSKRLSKSNIDISSGLEDE EPKRPLPRKQSDSSTYDCEAITQHH AFLSSIHSSVLKDESETPAAGGPQLP EVS LGRFDVSDFFLFGSPLGLVLA MRRTVLPGLDGFQVRPACSQVYSFF HCADPSASRLEPLLEPKFHLVPPVSV PRYQRFPLGDGQSLLLADALHTHSP LFLEGSRRDSPPLLDAPASPPQASRF QRPGRRMSEGSSESSSSDMSMA PVGASRITAKWWGSKRIDYALYCPD VLTAFPTVALPHLFHASYWESTDVV AFILRQVMRYESVNIKESARLDPAA SPANPREKWLRKRTQVKLRNVTAN HRANDVIAAEDGPQLVGRFMYGPL DMVALTGEKVDILVMAEPSGRWV HLDTEITNSSGRITVNPVPRRLGCVG VYPVKMVVVRGDQTCAMSYLTVLPRG MECVVFSIDGSFAASVSIMGSDPKV RPGAVDVVRHWQDLGYMILYITGRP DMQKQRVVSWSLSQHNFPGMIFFS DGLVHDPLRQKAIFLRNLMQECFIKI SAAYGSTKDISVYVSLGLPASQIFIVG RPTKKYQTCCQFLSEGYAAHLAAL ASHRSRPKKNNSRMILRKGSFGLHA QPEFLRKRNLRRRTMSVQQDPPAA NPKPERAQSQPESDKDHERPLPALS WARGPPKFESVP	True	False	4.536	3.29	1.316	1.36	1.154	1.835