

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8TF05	PP4R1_HUMAN	Homo sapiens	Serine/threonine-protein phosphatase 4 regulatory subunit 1	21.731224	NaN	S935	23301498	MADLSLLQEDLQEDADGFGVDDYS SESDVIIIPSAIDFVSDQDEMLTPLGRL DKYAAENIFNRQMVARSLLDTLRE VCDDERDCIAVLERISRLLADDSEPTV RAELMEQVPHIALFCOENRPSIPYAF SKFLLPIVVRYLADQNNQVRKTSQA ALLALLEQELIERFDVETKVCPLIEL TAPDSNDDVKTEAVAIMCKMAPMV GKDITERLILPRFCMCCDCRMFHV RKVCAANFGDICSVVGGQATEEMLL PRFFQLCSDNVWGVKKACAECFMA VSCATCQEIRRTKLSALFINLISDPSR WVRQAAFQSLGPFISTFANPSSSGQ YFKEESKSSEEMSVENKNRTRDQEA PEDVQVRPEDTPSDLVSNSSVILEN TMEDHAAEASGKPLGEISVPLDSSLL CTLSSESHQEAASNENDKKPGNYKS MLRPEVGTTSQDSALLDQELYNSFH FWRTPLEIDLDIELEQNSGGKPSPE GPEEESGVPSPSPNITMATRKELEE MIENLEPHIDDPDKAQUEVLSAAL RASSLDAHEETISIEKRSDLQDELDI NELPNCKINQEDSVPLISDAVENMD STLHYIHSDDLSNNSFSFDEERRT KVQDVVPQALLDQYLSMTDPSRAQT VDTEIAKHCAYSLPGVALTLGRQNW HCLRETYETLASDMQKVRRTLAFS IHELAVILGDQLTAADLVPIFNGFLK DLDEVIRIGVLKHLHDFLKLHIDKR REYLYLQEQFLVTDNSRNWRFRAL AEQLLLELYSPRDVYDYLRIALNL CADKVSSVRWISYKLVSEMVKLHA ATPPTFGVDLINELVENFGRCPKWS GRQAFVFCQTVIEDDCLPMDQFAV HLMPHLLTLANDRVPNVVLLAKTL RQTLLEKDYFLASASCHQEAVEQTI MALQMDRSDVKYFASIHIPASTKIS EDAMSTASSTY	None	None	None	None	None	None	None	None	

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