

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
Q969V6	MRTFA_HUMAN	Homo sapiens	Myocardin-related transcription factor A	27.828308	S146;S295;T465;T475	S6;S124;S139;S156;T305;S310;S312;T313;S317;S320;S333;S385;S446;T447;S449;T450;S454;T456;S458;S482;S492;S507;S511;S685;S691;S695;S792;S807;S859	38665916;23301498;28657654;40307207;29351928;35254053	MPPLKSPAAFHEQRRSLERARTEDY LKRKIRSRPERSELVRMHILEETSAE PSLQAKQLKLRARLADDLNEKIAQ RPGPMELVEKNILPVESSLKEAIIIVG QVNYPKVADSSSFDESSDALSPQE PASHESQGSVPSPLEARVSEPLLSAT SASPTQVVSQPLMGRDSREMLFLAE QPPLPPPPPLPPLSLTNGTTIPTAKSTP TLIKSQPKSASEKSORSKKAKELKP KVKKLKYHQYIPDQKQDRGAPPMD SSYAKILQQQLFLQLQILNQQQQ HHNYQAILPAPPKSAGEALGSSGTTP VRSLSTTNSSSSSGAPGPCGLARON STSLTGKPGALPANLDDMKVAELKQ ELKLRSLPVSGTKTELIERLRAQDQ ISPVPGAPKAPAATSILHKAGEVVVA FPAARLSTGPALVAAGLAPAEEVVVAT VASSGVVKFGSTGSTPPVSPTPSERS LLSTGDENSTPGDTFGEMVTSPLTQ LTLQASPLQILVKEEGPRAGSCCLSP GGRAELEGRDKDQMLQEKDKQIEA LTRMLRQKQQLVERLKLQLEQEKRA QQPAPAPAPLGTVPVKQENSFSSCQL SQQLGPAHPFNPSLAAPATNHIDP CAVAPGPPSVVVKQEQALQPEPEPVPA PQLLLGPGQPSLIKGVAPPTLITDSTG THLVLTVTNKNADSPGLSSGSPQPF SSQPGSPAPAPSAQMDLEHPLOPLF GTPTSLLKKEPPGYEAMSQQPKQQ ENGSSSQMDDLFDILIQSGEISADF KEPPSLPGKEKPSPKTVCGSPLAAQP SPSAELPQAAPPPGSPSLPGRLEDF LESSTGLPLLTSGHDGPEPLSLIDDL HSQMLSSTAILDHPPSPMDTSELHF VPEPSTMGDLADGHLDSMDWLE LSSGGPVLSLAPLSTTAPSLFSTDFL DGHDLQLHWDSCSL	True	False	4.453	5.0	0.594	False