

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	end reti
P59759	MRTFB_MOUSE	Mus musculus	Myocardin-related transcription factor B	69.970158	S207;S209;T214;S225;T234;T237;S245;T638;S757;S838;T858;T922;T932	S66;S531;S535;S537;S913	22517741;22645316;16452088;40885482;36852467;36084651;22826440;35822049;40021952;30059200;40997131;30016717	MIDSSKKQPGFPEILTAEDFEPFKE KECLEGSNQKSLKEVLQLRLQQRRT REQLVDQGMPPPKSPAAPFHEQIKSL ERARTENFLKHKIRSRPDRSELVRM HILEETFAEPLQATQMKLKRARLA DDLNEKIAQRPGPMELVEKNILPVD SSVKEAIIIGVVKEDYPHTHGFEFDFE DSSDALSPDQPASQESQSGSAASPSE PKVSASPPPVTASTPAQFTSVSPA EFLKTPLTADQPPTTRSTAPVLPNTV SSAKSGPMLVKQSHPKNPNDKHRS KKCKDPKPRVKKLKYHQYIPPNQKG EKSEPMDSNYARLLQQQLFLQL QILSQQQQQQQQHYNYQTILPAPI KTDKNSSSGSNSSSSMPARRPGP LPSSLDLKVSELKTELKLRGLPVSG TKPDLIERLKPQEVTSNNLATGSIVA VSSATIVTSNPEVTVALPVTTLHNAV TSSVSTFKADLALPATSSVPHVENAH SPLPISPSPEQSSLSTDDTNMTDTF TEIMTMMSPSQLLCSPLRVVSHDD SLSPSSSTLSTLELDAAEKDRKLQEK EKQIEELKRKLEQEQKLVVLMQML EVEKRGQQRPPDPQSDPPHPFNFS DPKHGSGSSIKDEASLPDCSSPQQ PITVPGHSGQPISTGSQLVAKKTV VVKQEVMAQAEQQNVVSQFYLLS QGQPALVAQPAQLTTOTTQLLLPV SIQGSNVTSQLPVGSLQLQTPAQG RVQAQPHVAAATQVPAAALPSALTS ALPQKQEAFFQHVGLGQPQVRRKFT NSAPNTVLQYQRQPGPTNQPFVSK TSNPALQSRAPLAPLQNGPSLASKP SSPPPPQFVVQHSLEFATPITKTKDP PRYEEAIKQTRSTQPALPEVSSVHSQ QMDLDFDILIKSGEISFPIKEEPSIS KMKPVTASITMPVNTVSRPPQV QIAPPVSLEPVNSLASLENQLEAFL DGTLPASATDTGPLQNSSEDRESFSLI EDLQNDLLSHSSMLYQSHSPMETS EAQLVSGTPCLSLDSDSNLDNME WLDITMPTTSSGLTPLSTTAPSMFSA DFLDPQDLPLPWD	True	False	1.316	4.434	1.299	1.01