

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A2BH40	ARI1A_MOUSE	Mus musculus	AT-rich interactive domain-containing protein 1A	15.818779	S697;S706;S710;T764	S56;S77;S234;T287;S302;S365;S384;S605;S697;S699;S703;S731;S765;S773;S1185;S1236;S1605;T1874;T1886;S1927;S1942	36852467	MAAQVAPAAASSLGNPPPPSELKK AEQQQREEAGGEAAAAAERGERMK AAAGQESEGPVAVGPPQPLKELQDG AESNGGGGGGAGSGGGPGAEPDL KNSNGNAGPRPALNNNLPEPPGGG GGGSSSSDGVGAPPHSAAAALPPP AYGFGQAYGRSPSAVAAAAAVFHQ QHGGQQSPGLAALQSGGGGLEPY AGPQQNSHDHGFPNHQYNSYYPNR SAYPPPPQAYALSSPRGGTPGSGAAA AAGSKPPSSSASASSSSSFAQQRF GAMGGGGPSAAGGGTPQPTATPTLN QLLTSPSSARGYQGYPGGDYGGGPQ DGGAGKGPADMASQCWGAAAAAAA AAAAVSGGAQQRSHHAPMSPGSSG GGGQPLARTQSSSPMDQMGKMRP QPYGGTNPYSQQQPPSGPQQGHG YPGQPYGSQTPQRYPMTMQGRAQS AMGSLSYAQQIPPYGQQGPSAYGQQ GQTPYYNQSPHPQQQPPYAQQPPS QTPHAQPSYQQPQTQQPQLQSSQP PYSQQPSQPPHQQSPTPYPSQQSTT QQHPQSQPPYSQPQAQSPYQQQQP QQPASSLSQQAAYPQPQPQQSQQT AYSQQRFPQPQELSQDSFGSQASSA PSMTSSKGGQEDMNLQSRPSSLP DLSGSIDDLPMTGEGALSPGVSTSGI SSSQGEQSNPAQSPFSPHTSPHLPGI RGPSPPVGPASVAQSRSGPLSPAA VPGNQMPRPPSGQSDSIMHPSMN QSSIAQDRGYMQRNPQMPQYTSPQ PGSALSPRQPSGGQMHSVGVGSYQQ NSMGSYGPQGSQYGPQGGYPRQPN YNALPNANYPNAGMAGSMNPMGA GGQMHHGQPGIPPYGTLPGRMAHA SMGNRPYGNMANMPPQVGS GMC PPPGMNRKTQESAVAMHVAANSI QNRPPGYPNMNQGGMMGTGPPYG QGINSMAGMINPQGPYPMGGTMA NNSAGMAASPEMMGLGDVCLTPAT KMNNKADGTPKTESKSKSSSSTTT NEKITKLYELGGEPERKMWVDLYLA FTEEKAMGMTNLPVAVGRKPLDLYRL YVSVKEIGGLTQVNKNKKWRELATN LNVGTSSSAASSLKKQYIQCLYAFEC KIERGEDPPPDIFAAADSKKSQPKIQ PPSPAGSGSMQGPQTPQSTSSSMAE GGDLKPPTPASTPHSQIPPLPGMSRS NSVGIQDAFPDGS DPTFQKRNSMTP NPGYQPSMNTSDMMGRMSYEPNK

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NNVDLILATPPFSRLEKLYSTMVRFL
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LAARAI VQKGSIGNLLGFLEDSLAA
TQFQQSQASLLHMQNPPFEPTSVD
MMRRAARALLALAKVDENHSEFTL
YESRLLDISV SPLMNSLVSQVICDVL
FLIGQS