

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	e. ri
Q7TPR4	ACTN1_MOUSE	Mus musculus	Alpha-actinin-1	36.545784	NaN	S6;T12;S471;S677;S890	21540332;33300544;37453647;28528544;27301331	MDHYDSQQTNDYMQPEEDWDRDL LLDPAWEKQQRKTFTAWCNSHLRK AGTQIENIEEDFRDGLKMLLLEVIS GERLAKPERGKMRVHKISNVNKALD FIASKGVKLVSIGAEIIVDGNVKMTL GMIWTIILRFIAIQDISVEETSAKEGLL LWCQRKTAPYKNVNIQNQFHISWKD GLGFCALIHRRPELIDYGKLRKDDP LTNLTAFDVAERFLDIPKMLDAEDI VGTARPDEKAIMTYVSSFYHAFSGA QKAETAANRICKVLAVNQENEQLM EDYEKLASDLEWIRRTIPWLENRVP ENTMHAMQOKLEDFRDYRRLHKPP KVQEKQLEINFNTLQTKRLSNRP AFMPSEGRMVSDINNAWGCLEQAE KGYEEWLLNEIRRLERLDHLAEKFR QKASIHEAWTDGKEAMLKQDYET ATLSEIKALLKKHEAFESDLAAHQDR VEQIAAIAQELNELDYDSSPSVNARC QKICDQWDNLGALTQKRREALERTE KLETTIDQLYLEYAKRAAPFNWME GAMEDLQDTFIVHTIEEQGLTTAHE QFKATLPDADKERLAILGIHNEVSKI VQYHVN MAGTNPYTTITPQEINGK WDHVRQLVPRRDQALTEEHARQQH NERLRQFGAQANVIGPWIQTKMEE IGRISIEMHGTLEDQLSHLROYEKSI VNYKPKIDQLECDHQLQEQALIFDNK HTNRYNMEHIRVGEQLLTTIARTIN EVENQILTRDAKGISQEQMNEFRAS FNHFDRDHSGLGPEEFKACLISLG YDIGNDPQGEAEFARIMSIVDPNRL GVVTFQAFIDFMSRETADTDTADQV MASFKILAGDKNYITEDELRRRELPPD QAEYCIARMAPYAGPDSVPGALDYM SFSTALYGESDL	False	True	3.408	2.937	1.725	1.566	1.206	4.416	1.