

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	extracellular region
088990	ACTN3_MOUSE	Mus musculus	Alpha-actinin-3	25.019509	S369;T825	NaN	34418053	MMMVMQPEGLGAGEGPFSGGGGG EYMEQEEDWDRDLLLLPAWEKQQR KTFTAWCNSHLRKAGTQIENIEEDF RNLKLMLELVISGERLPRPKGK MRFHKIANVVKALDFIASKGVKLVSI GAEIIVDGNLKMILGMIWTHLRFI QDISVEETSAKEGLLLWCQRKTAPY RNVNVQNFHTSWKDLGALCALIHR HRPDLIDYAKLRKDDPIGNLNTAFEV AEKYLDIPKMLDAEDIVNTPKPDEKA IMTYVSCFYHAFAGAEQAETAANR KVLAVNQENEKLMEEYEKLAELLE WIRRTVPWLENRVGEPMSAMQRK LEDFRDYRRLHKPPRVQEKQLEIN FNTLQTKLRLSHRPAFMPSEGKLV DIANAWRGLEQVEKGYEDWLLSEIR RLQRLQHLAEKFQKASLHEAWTR GKEEMLNQHDYESASLQEVALLR RHEAFESDLAAHQDRVEHIAALAE LNELDYHEAASVNSRCQAICDQWD NLGTLTQKRRDALERMEKLETIDQ LQLEFARRAAPFNNWLDGAIEDLQD VWLVHSVEETQSLTAHEQFKATLP EADRREGAILGIQGEIQKICQTYGLR PKSGNPYITLSSQDINNKWDTVRKL VPSRDQTLQEEELARQVNERLRRQF AAQANAIGPWIQGKVEEVGRLAAGL AGSLEEQMAGLRQEQNIINYKSNI DRLEGDHQLLQESLVFDNKHTVYS MEHIRVGVWEQLLTSIARTINEVENQ VLTRDAKGLSQEQLNEFRASFNHFD RKRNGMMEPDDFRACLISMGYDLG EVEFARIMTMVDPNAAAGVVFQAFI DFMTRETAETDTAEQVVASFILAG DKNYITPEELRRELPAEQAEYCIRRM APYKGS GAPSGALDYVAFSSALYGES DL	False	False	3.421	1.712	1.731	1.581	0.794	3.705	1.354