	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regio
0	P35609	ACTN2_HUMAN	Homo sapiens	Alpha-actinin-2	27.792899	S431;T435	T237	30379171;35008409;23271734;29351928	MNQIEPGVQYNYVYDEDEYMIQEEE WDRDLLLDPAWEKQQRKTFTAWCN SHLRKAGTQIENIEEDFRNGLKLML LLEVISGERLPKPDRGKMRFHKIAN VNKALDYIASKGVKLVSIGAEEIVDG NVKMTLGMIWTIILRFAIQDISVEET SAKEGLLLWCQRKTAPYRNVNIQNF HTSWKDGLGLCALIHRHRPDLIDYS KLNKDDPIGNINLAMEIAEKHLDIPK MLDAEDIVNTPKPDERAIMTYVSCF YHAFAGAEQAETAANRICKVLAVNQ ENERLMEEYERLASELLEWIRRTIP WLENRTPEKTMQAMQKKLEDFRDY RRKHKPPKVQEKCQLEINFNTLQTK LRISNRPAFMPSEGKMVSDIAGAWQ RLEQAEKGYEEWLLNEIRRLERLEH LAEKFRQKASTHETWAYGKEQILLQ KDYESASLTEVRALLRKHEAFESDLA AHQDRVEQIAAIAQELNELDYHDAV NVNDRCQKICDQWDRLGTLTQKRR EALERMEKLLETIDQLHLEFAKRAA PFNNWMEGAMEDLQDMFIVHSIEE IQSLITAHEQFKATLPEADGERQSIM AIQNEVEKVIQSYNIRISSSNPYSTVT MDELRTKWDKVKQLVPIRDQSLQEE LARQHANERLRRQFAAQANAIGPWI QNKMEEIARSSIQITGALEDQMNQL KQYEHNIINYKNNIDKLEGDHQLIQ EALVFDNKHTNYTMEHIRVGWELLL TTIARTINEVETQILTRDAKGITQEQ MNEFRASFNHFDRRKNGLMDHED FRACLISMGYDLGEAEFARIMTLVDP NGQGTVTFQSFIDFMTRETADTDTA EQVIASFRILASDKPYILAEELRRELP PDQAQYCIKRMPAYSGPGSVPGALD YAAFSSALYGESDL	True	False	4.696	3.632	2.438	1.929	0.992	3.319	4.353