

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|-----------------|--------------|---------------|-----------------------|-------------------------------------|---|
| P35609 | ACTN2_HUMAN | Homo sapiens | Alpha-actinin-2 | 15.309242 | S431;T435 | T237 | 30379171;23271734;29351928;35008409 | MNQIEPGVQYNYVYDEDEYMIQEEE WDRDLLDPAWEKQQRKTFWAWCN SHLRKAGTQIENIEEDFRNGLKML LLEVISGERLPKPDGRKMRFHKIAN VNKALDYIASKGVKLVSIGAEEIVDG NVKMTLGMWITILRFAIQDISVEET SAKEGLLLWCQRKTAPYRNVNIQNF HTSWKDGLGLCALIHRHRPDLIDYS KLNKDDPIGNINLAMEIAEKHLDIPK MLDAEDIVNTPKPDERAIMTYVSCF YHAFAGAEQAETAANRICKVLAVNQ ENERLMEEYERLASELLEWIRRTIP WLENRTPEKTMQAMQKKLEDFRDY RRKHKPPKVQEKQLEINFNTLQTK LRISNRPAFMPSEGKMSVSDIAGAWQ RLEQAEEKGYEELLNEIRRLERLEH LAEFRQKASTHETWAYGKEQILLQ KDYESASLTEVRALLRKHEAFESDLA AHQDRVEQIAAIAQELNELDYHDAV NVNDRQCQKICDQWDRLGTLTQKRR EALERMEKLETTIDQLHLEFAKRAA PFNNWMEGAMEDLQDMFIVHSIEE IQSLITAHEQFKATLPEADGERQSIM AIQNEVEKVIQSYNIRISSNPYSTVT MDELRTKWDKVKQLVPIRDQSLQEE LARQHANERLRRQFAAQANAIGPWI QNKMEIARSSIQTGALEDQMNQL KQYEHNIINYKNNIDKLEGDHQLIQ EALVFDNKHTNYTMEHIRVGVWELL TTIARTINEVETQILTRDAKGITQEQ MNEFRASFNHFDRRKNGLMDHED FRACLISMGYDLGAEAFARIMTLVDP NGQGTVTFQSFIDFMTRETADTDTA EQVIASFRILASDKPYILAEELRREL PDQAQYCIKRMPAYSGPGSVPGALD YAAFSSALYGESDL |

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