

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|--|--------------|---------------|-------------------------------|----------|---|
| Q8CCN5 | BCAS3_MOUSE | Mus musculus | BCAS3 microtubule associated cell migration factor | 16.122943 | NaN | S461;S480;S488;S838;S886;S898 | 22645316 | MNETMATDSPRRPSRCTGGVVVRP QAVTEQSYMESVVTFLQDVVPQAYS GSPLTEEKEKIVWRFENADLNDS RNLEFHELHSTGNEPPLVMIGYSD GMQVWGIPISGEAQELFSVRHG PVR AARILPAPQLGAQKCDNFAEKRP GVCKSIGSSGTPPYCCVDLYSLRTG EMVKSIQFKTPIYDLHCNKRLVVVL QEKIAAFDSCFTFKKFVTSYPCPG PNMNPALGSRWLAYAENKLIRCHQ SRGGACGDNIQSYTATVLSAAKTLKS GLTMVGKVVTQLTGTLPSTGVEDDV ALHCNSRRSPLVPGIITVIDTETVGE GQVLVSESDSDGIVAHFPAHEKPV CCMAFNTSGMLLVTTDTLGHDFHV FQILTHPWSSSQCAVHHLTYTLHRGE TEAKVQDICFSDCRWVWVSTLRGT SHVFPINPYGGQPCVRTHMSPRVVN RMSRFQKSAGLEEIEQELTSKQGGGR CSPV PGLSSSPSGSPLHGKLTSDSY NNFTNNNPGNPRLSPLPSLMVVTPL AQIKQPMTLGTITKRTGPYLFGAGCF SIKAPCKVKSPQISPSKSMGGEFCV AAVFGTSRSWFANNAGLKREKDQS KQVVVESLYIISCYGTLVEHMIERPI STAPKISDDTPLEIMTSPRASWTLVR TPQWNELOPPFNANHPHLLAAEAV QYYQLLLAGSLPPGSPGPITRHGSYD SLASDHSGQEDEEWLSQVEIVTHTG PHRRLWMGPQFHFKTIQTSGQTTVI STSSSVLQSHGPSDTPQPLDFD TD DLDLNSLRIQPVRSDPVSM PGSSRA VSDRRGVSTVTDAAAGTFDRSVTLLE VCGSWPEGFGLRHMSSMEHSEEGL RERLADAMAESPRDVVGSGETELQR EGSIETLSNSSGSTSGSIPRNF DGYR SPLPTNESQPLSLFPTGFP |