

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	exti reg
Q8CCN5	BCAS3_MOUSE	Mus musculus	BCAS3 microtubule associated cell migration factor	46.058797	S797	S461;S480;S488;S838;S886;S898	22645316;40885482	MNETMATDSPRRPSRCTGGVVVVRP QAVTEQSYMESVVTFLODQVVPQAYS GSPLTEEKEKIVVWRFENADLNDTS RNLEFHELHSTGNEPPLLVMIGYSD GMQVWGPISGEAQELFVSRHGPVR AARILPAPQLGAQKCDNFAEKRP GVCKSIGSSGTTPPYCCVDLYSLRTG EMVKSQFKTPIYDLHCNKRILVVVL QEKIAAFDSCFTTKKFFVTSCYPCPG PNMNPALGSRWLYAENKLIRCHQ SRGGACGDNIQSYTATVLSAAKTLKS GLTMVGKVVTLTGLTPSGVTEDDV ALHCNSRRSPLVPGIITVIDTETVGE GOVLVSESDSDGIVAHFPAHEKPV CCMAFNSTGMLLVTTDTLGHDFHV FQILTHPWSSSQCAVHHLYTLHRGE TEAKVQDICFSDCRWVWVSTLRGT SHVFPINPYGGQPCVRTHMSPRVVN RMSRFQKSAGLEEIEQELTSKQGGGR CSPVPGLSSSPSGSPLHGKLTSDSY NNFTNNPNPNRPLSPLSMVVVTP AQIKQPMTLGTTTKRTGPLYFGAGCF SIKAPCKVKSPPOISPSKSMGGGFCV AAVFGTSRSWFANNAGLKREKQDS KQVVVESLYIISCYGTLVEHMIERPI STAPKISDDTPEIMTSPRASWTLVR TPQWNELOPPFNANHPLLLAEEAV QYYQLLAGSLPPGSPGPIRHGYSYD SLASDHSGQEDDEEWLSQVEIVHTHG PHRRLWMGPQFHFKTIQTSGQTTVI STSSSVLQSHGSDTPQPLDFDFTD DLDLNSLRQPVRSDPVSMPGSSRA VSDRRGVSTVTAASGTFDRSVTLLLE VCGSWPEGFGLRHMSSMEHSEEGLE RERLADAMAESPSRDVVGSGTELQR EGSIETLSNSSGSTSGSIPRNFQGYR SPLPTNESQPLSLFPTGFP	True	False	2.096	4.551	1.039	0.908	False	0.787	0.74