

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasm mem
Q9H6U6	BCAS3_HUMAN	Homo sapiens	BCAS3 microtubule associated cell migration factor	17.760791	NaN	S461;S480;S488;S838;S886;S898	34019948;39302247;30379171	MNEAMATDSPRRPSRCTGGVVVRP QAVTEQSYMESVVTFLQDVVPQAYS GTPLTEEEKIKIVWVRFENADLNDS RNLEFHEIHSTGNEPPLIMIGYSDG MQVWSIPISGEAQELFSVRHGPRAA RILPAPQFQAQKCDNF AEKRPLLGV CKSIGSSGTSPPYCCVDLYSLRTGEM VKSIOFKTPIYDLHCNKRILVVVLOE KIAAFDSCFTFKFFVTSCYPCPGPN MNPALGSRWLYAENKLRCHQSR GGACGDNIQSYTATVISAATLKSGL TMVGKVVTLTGLTPSGVTEDDAI HSNSRRSPLVPGIITVIDTETVGEQ VLVSESDSDGIVAHFPAHEKPVCC MAFNSTSGMLLVTTDTLGHDFHVFQI LTHPWSSSQCAVHHLYTLHRGETEA KVQDICFSDCRWVVVSTLRGTSHV FPINPYGGQPCVRTHMSPRVVNRMS RFQKSAGLEEIEQELTSKQGGRCSP VPGLS SSPSGSPLHGKLNQDSYNN FTNNNPGNRLSPLSLMVVMP QIKQPMTLGTITKRTGPLYFGAGCFS IKAPCKVKPPPQISPSKSMGGFCVA AIFGTSRSWFANNAGLKREKQSKQ VVVESLYIISCYGTLVEHMM EPRPLS TAPKISDDTPELMMTSPRASWTLVR TPOWNELQPPFNANHPLLLAADAV QYYQFLLAGLVPPGSPGPIRHSYD SLASDHSGQEDEEWLSQVEIVHTG PHRRLLWGPQFQFKTIHPSGQTTVI SSSSVLQSHGSPDTPQLLDFDFTD DLDLNSLRIQPVRSDPVSMPGSSRP VSDRRGVSTVIDAASGTFDRSVTLL VCGSWPEGFGLRHMSMEHTEEGL RERLADAMAESPSRDVVGSGTELQR EGSIETLSNSSGTSGISPRNFDGYR SPLPTNESQPLSLFPTGFP	True	False	2.246	5.0	1.172	1.058	0.747	0.871