

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q96GW7	PGCB_HUMAN	Homo sapiens	Brevican core protein	27.086674	S125	S418	28657654	MAQLFLPLLAALVLAQAPAAALADVL EGDSSSEDRAFRVRIAGDAPLQGVLG GALTIPCHVHYLRPPPSRRAVLGSPR VKWTFLSRGREAEVLVARGVRVKVN EAYRFRVALPAYPASLTDVSLALSEL RPNDSGIYRCEVQHIDDSSDAVEV KVKGVVFLYREGSARYAFSFGAQE ACARIGAHIAPEQLYAAYLGGYEQC DAGWLSDQTVRYPIQTPREACYGDM DGFPGVRNYGVVDPDDLVDVYCYAE DLNGELFLGDPPEKLTLEEARAYCQ ERGAEIATTGQLYAAWDGGLDHCSP GWLADGSVRYPIVTPSQRGGGLPG VKTLFLFPNQTGFPNKHSRNFVYCF RDSAQPSAIPASNPASNPASDGLEA IVTVTETLEELQLPQEATESESRGAIY SIPIMEDGGGGSTPEDPAEAPRTLL EFETQSMVPPTGFSEEEGKALEEEE KYDEEEKEEEEEVEDEALWA WPSELSSPGPEASLPTEPAAQEEELS QAPARAVLQPGASPLPDGESEASRP PRVHGPPPTETLPTPRERNLASEPST LVEAREVGEATGGPELGGVPRGESE ETGSSEGAPSLPATRAPEGTRELEA PSEDNSGRTAPAGTSVQAQPVLPTD SASRGGVAVVPASGDCVPSCHNGG TCLEEEEGVRCLCLPGYGGDLCDVG LRFENPGWDAFQGACYKHFSTRRS WEEAETQCRMYGAHLASISTPEEQD FINNRYREYQWIGLNDRTIEGDFLW SDGVPLLYENWNPGQDSYFLSGEN CVVMVWHDQGGQWSDVPCNYHLSY TCKMGLVSCGPPPELPLAQVFRPR LRYEVDTVLRYRCREGLAQRNLPLIR CQENGRWEAPQISCVRRRPARALHP EEDPEGRQGRLLGRWKALLIPSSP MPGP	False	True	2.552	2.8	2.572	1.548	4.242	3.478	4.702