	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ex re
0	Q14721	KCNB1_HUMAN	Homo sapiens	Potassium voltage-gated channel subfamily B member 1	29.421095	NaN	S15;T128;S444;S457;S484;S496;S503;S519;S520;S541;S567;S590;S607;S656;S720;S772;S800;S805	28657654;26960196	MPAGMTKHGSRSTSSLPPEPMEIVR SKACSRRVRLNVGGLAHEVLWRTLD RLPRTRLGKLRDCNTHDSLLEVCDD YSLDDNEYFFDRHPGAFTSILNFYRT GRLHMMEEMCALSFSQELDYWGID EIYLESCCQARYHQKKEQMNEELKR EAETLREREGEEFDNTCCAEKRKKL WDLLEKPNSSVAAKILAIISIMFIVLS TIALSLNTLPELQSLDEFGQSTDNPQ LAHVEAVCIAWFTMEYLLRFLSSPK KWKFFKGPLNAIDLLAILPYYVTIFLT ESNKSVLQFQNVRRVVQIFRIMRILR ILKLARHSTGLQSLGFTLRRSYNELG LLILFLAMGIMIFSSLVFFAEKDEDD TKFKSIPASFWWATITMTTVGYGDIY PKTLLGKIVGGLCCIAGVLVIALPIPII VNNFSEFYKEQKRQEKAIKRREALE RAKRNGSIVSMNMKDAFARSIEMM DIVVEKNGENMGKKDKVQDNHLSP NKWKWTKRTLSETSSSKSFETKEQG SPEKARSSSSPQHLNVQQLEDMYN KMAKTQSQPILNTKESAAQSKPKEE LEMESIPSPVAPLPTRTEGVIDMRSM SSIDSFISCATDFPEATRFSHSPLTSL PSKTGGSTAPEVGWRGALGASGGRF VEANPSPDASQHSSFFIESPKSSMK TNNPLKLRALKVNFMEGDPSPLLPV LGMYHDPLRNRGSAAAAVAGLECAT LLDKAVLSPESSIYTTASAKTPPRSPE KHTAIAFNFEAGVHQYIDADTDDEG QLLYSVDSSPPKSLPGSTSPKFSTGT RSEKNHFESSPLPTSPKFLRQNCIYS TEALTGKGPSGQEKCKLENHISPDV RVLPGGGAHGSTRDQSI	False	True	2.232	2.542	2.013	2.182	1.583	5.0	2.