	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	extracellular region
0	O43525	KCNQ3_HUMAN	Homo sapiens	Potassium voltage-gated channel subfamily KQT member 3	35.076234	T655	T81;T246	28657654;25303527	MGLKARRAAGAAGGGGDGGGGGGG AANPAGGDAAAAGDEERKVGLAPGD VEQVTLALGAGADKDGTLLLEGGGR DEGQRRTPQGIGLLAKTPLSRPVKR NNAKYRRIQTLIYDALERPRGWALLY HALVFLIVLGCLILAVLTTFKEYETVS GDWLLLLETFAIFIFGAEFALRIWAA GCCCRYKGWRGRLKFARKPLCMLDI FVLIASVPVVAVGNQGNVLATSLRSL RFLQILRMLRMDRRGGTWKLLGSAI CAHSKELITAWYIGFLTLILSSFLVYL VEKDVPEVDAQGEEMKEEFETYADA LWWGLITLATIGYGDKTPKTWEGRLI AATFSLIGVSFFALPAGILGSGLALKV QEQHRQKHFEKRRKPAAELIQAAW RYYATNPNRIDLVATWRFYESVVSFP FFRKEQLEAASSQKLGLLDRVRLSN PRGSNTKGKLFTPLNVDAIEESPSKE PKPVGLNNKERFRTAFRMKAYAFW QSSEDAGTGDPMAEDRGYGNDFPIE DMIPTLKAAIRAVRILQFRLYKKKFK ETLRPYDVKDVIEQYSAGHLDMLSRI KYLQTRIDMIFTPGPPSTPKHKKSQK GSAFTFPSQQSPRNEPYVARPSTSEI EDQSMMGKFVKVERQVQDMGKKL DFLVDMHMQHMERLQVQVTEYYPT KGTSSPAEAEKKEDNRYSDLKTIICN YSETGPPEPPYSFHQVTIDKVSPYGF FAHDPVNLPRGGPSSGKVQATPPSS ATTYVERPTVLPILTLLDSRVSCHSQ ADLQGPYSDRISPRQRRSITRDSDTP LSLMSVNHEELERSPSGFSISQDRD DYVFGPNGGSSWMREKRYLAEGET DTDTDPFTPSGSMPLSSTGDGISDSV WTPSNKPI	False	True	1.67	3.283	2.906	1.475	0.954	5.0	1.91