

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8K3F6	KCNQ3_MOUSE	Mus musculus	Potassium voltage-gated channel subfamily KQT member 3	25.775171	NaN	T82	34678516	MGLKARRAAGAAGGGGEGGGGGG GAANPAGGDSAVAGDEERKVLAPG DVEQVTLALGAGADKDGTLLEGGG REEGQRRTPQGIGLLAKTPLSRPVKR NNAKYRRIQTLIYDALERPRGWALLY HALVFLIVLGLLAVLTTTFKEYETVS GDWLLLETFAIIFFGAEFALRIWAA GCCCRYKGWGRGLKFARKPLCMLDI FVLIASVPVAVGNQGNVLTSLRSL RFLQILRMLRMDRRGGTWKLLGSAI CAHSKELITAWYIGFLTLLSSFLVYL VEKDVPEMDAQGEEMKEEFETYAD ALWWGLITLATIGYGDKTPKTWEGR LIAATFSLIGVSFFALPAGILGSLAL KVQEQHRQKHFEKRRKPAELIQAA WRYATNPRLDLVATWRFYESVVS FPFRKEQLEAAASQKLGLLDRVRL SNPRGSNTKGKLFPLNVDIAEESPS KEPKPVGLNNKERFRTAFRMKAYAF WQSSEDAGTGDPMAEDRGYGNDFL IEDMIPTLKAIRAVRILQFRLYKKKF KETLRPYDVKDVEQYSAGHLDMLS RIKYLQTRIDMIFTPGPPSTPKHKKS QKGSFTYPSQSPRNEPYVARAAT SETEDQSMGKFVKVERQVHDMG KKLDFLVDMMHMOHMERLQVHVTE YYPTKGASSPAEGEKEDNRYSDLK TIICNYSETGPPDPPYSFHQVPIDRVG PYGFFAHDVPKLRGGPSSTKAQAN LPSSGSTYAEERPTVLPILTLDDSCVSY HSQTELGQPSYDHSRQRRSITRDS DTPLSLMSVNHHEELERSPSGFSISQ DRDDYVFGPSGGSSWMREKRYLAE GETD'TDTPFTPSGSMPSSTGDGI SDSIWTPSNKPT	False	True	1.354	1.584	2.786	1.881	1.426	4.594	2.25

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