

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
Q7Z2W7	TRPM8_HUMAN	Homo sapiens	Transient receptor potential cation channel subfamily M member 8	24.917859	NaN	NaN	28411811	MSFRAARLSMRNRRNDTLDSTRTL YSSASRSTDLSYSESDLVNFIQANFK KRECVFFTKDSKATENVCKCGYAQS QHMEGTQINQSEKWNYKKHTKEFP TDAFGDIOFETLGGKGYIRLSCDTD AEILYELLTQHWHLKTPNLVISVTGG AKNFALKPRMRKIFSRLYIAQSKGA WILTGGETHYGLMKYIGEVVRDNTISR SSEENIVAIGIAAWGMVSNRDTLIRN CDAEGYFLAQYLMDDFTRDPLYILD NNHTLHLLVDNGCHGHPVVEAKLR NQLEKYISERTIQDSNYGGKPIVCF QGGGKETLKAINTSIKNKIPCVVVEG SGQIADVIASLVEVEDALTSSAVKEK LVRFLPRTVSRLPEEETESWIKWLKE ILECSHLLTVIKMEEAGDEIVSNAISY ALYKAFSTSEQDKDNWNGOLKLLLE WNQLDLANDEIFTNDRRWESADLQ EVMFTALIKDRPKFVRLFLENGLNL RKFLTHDVLTELFNHFSTLVYRNL QIAKNSYNDALLTFVWKLVANFRRG FRKEDRNGRDEMDELHDSVSPITRH PLQALFIWAILQNKKELSKVIWEQTR GCTLAALGASKLLKTLAKVKNNDINAA GESEELANEYETRAVELFTECYSSDE DLAEQLLVYSCEAWGGSNCLELAVE ATDQHFIAPGVQNFSLKQWYGEIS RDTKNWKIILCFIIPLVGCGFVSFRK KPVDKHKKLLWYVAFFTSFVVFVS WNVVFIYAFLLLFAYVLLMDFHSVP HPPELVLYSLVFLFCDEVQWYVN GVNYFTDLWNVMDTLGLFYFIAGIV FRLHSSNKSSLYSGRVIFCLDYIIFTL RLIHIFTVSRNLGPKHIMLQRMILIDVF FFLFLFAVWMVAFGVARQGILRQNE QRWRWIFRSVIYEPYLAMFGQVPSD VDGTTYDFAHCTFTGNESKPLCVEL DEHNLPRFPEWITIPLVCIYMLSTNI LLVNLVAMFGYTVGTQVQENNDQV WKFQRYFLVQEYCSRLNIPFPFIVFA YFYMVVKCKFKCCCKEKNMESSVC CFKNEDNETLAWEGVMKENYLVKI NTKANDTSEEMRHRFRQLDTKLNLD LKGLLKEIANKIK	False	True	2.26	2.496	2.205	5.0	1.208	5.0	2.438