	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	A6NFQ2	TCAF2_HUMAN	Homo sapiens	TRPM8 channel-associated factor 2	26.21016	T305	NaN	29351928	MATIAAAAFEALMDGVTCWDVPRGP IPSELLLIGEAAFPVMVNDKGQVLIA ASSYGRGRLVVVSHEGYLSHTGLAP FLLNAVSWLCPCPGAPVGVHPSLAP LVNILQDAGLEAQVKPEPGEPLGVYC INAYNDTLTATLIQFVKHGGGLLIGG QAWYWASQHGPDKVLSRFPGNKVT SVAGVYFTDTYGDRDRFKVSKKVPKI PLHVRYGEDVRQDQQQLLEGISELD IRTGGVPSQLLVHGALAFPLGLDASL NCFLAAAHYGRGRVVLAAHECLLCA PKMGPFLLNAVRWLARGQTGKVGV NTNLKDLCPLLSEHGLQCSLEPHLN SDLCVYCCKAYSDKEAKQLQEFVAE GGGLLIGGQAWWWASQNPGHCPLA GFPGNIILNCFGLSILPQTLKAGCFP VPTPEMRSYHFRKALSQFQAILNHE NGNLEKSCLAKLRVDGAAFLQIPAE GVPAYISLHRLLRKMLRGSGLPAVSR ENPVASDSYEAAVLSLATGLAHSGT DCSQLAQGLGTWTCSSSLYPSKHPI TVEINGINPGNNDCWVSTGLYLLEG QNAEVSLSEAAASAGLRVQIGCHTD DLTKARKLSRAPVVTHQCWMDRTE RSVSCLWGGLLYVIVPKGSQLGPVPV TIRGAVPAPYYKLGKTSLEEWKRQM QENLAPWGELATDNIILTVPTTNLQA LKDPEPVLRLWDEMMQAVARLAAE PFPFRRPERIVADVQISAGWMHSGY PIMCHLESVKEIINEMDMRSRGVW GPIHELGHNQQRHGWEFPPHTTEA TCNLWSVYVHETVLGIPRAQAHEAL SPPERERRIKAHLGKGAPLCDWNV WTALETYLQLQEAFGWEPFTQLFAE YQTLSHLPKDNTGRMNLWVKKFSE KVKKNLVPFFEAWGWPIQKEVADSL ASLPEWQENPMQVYLRARK	None	None	None	None	None	None	None	None	None