

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q08189	TGM3_MOUSE	Mus musculus	Protein-glutamine gamma-glutamyltransferase E	26.222628	NaN	T111;T112	24788674	MSALQIQNVNWQVPMNRRHHTD KFSSQDSIVRRGQPWEIILVCNRSLE SGEDLNFIVSTGQPQSEARTKAVFS ISGRSTGGWNAALKANSNNLAI ASPVSAPIGLYTLSSVEISSRGRASSLK LGTFIGMLFNPWLQADDVFMSNHAE RQEYVEEDSGIHYVGSTNRIGMVGW NFGQFEEDILNISLSILDRSLNFRRD PVTDVARRNDPKYVCRVLSAMINGN DDNGVISGNWVSGNYTGGVDPRTWN GSVEILKNWKKSGFRPVQFGQCWV FAGTLNIVLRCLGVPSRVITNFNSA HDTDRNLSVDVYYDAMGNPLEKGS DSVWNFHVWNEGWFVRTDLGPTY NGWQVLDATPQERSQGVFQCGPAS VNAIKAGDVDRNFDMIFIFAEVNAD RITWIYNRRNNTQKQNSVDTHSIGK YISTKAVGSNSRMDVTDKYKYPEGS SEERQVHQKALDKLPNASFGATSS RNPEGEDKEPSISGKFKVTGILAVGK EVSLSLMLKNMTNDRKVTVMNMTA WTIVYNGTLVHEVWVKDSATISLDPE EEIQYVVKIAYSQYERYLKADNMIRIT AVCKVPDEAEVVVERDVILDNPALTL EVLEQAHVRKPVNVQMLFSNPLDQ PVNNCVLLVEGSLLRGLKIDVPSL RPKEKSRRIFEIFPTRSGTKQLLADF SCNKFFAIKAMPLIDVSE	False	True	2.994	2.131	1.818	0.681	0.658	5.0	1.609