	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	Q148V8	FA83H_MOUSE	Mus musculus	Protein FAM83H	25.161187	NaN	S512;S513;S515;S522;S639;S660;T749;S752;S778;S806;S871;T873;S882;S893;S904;S915;T917;S926;T928;S937;S948;S959;S970;S977;S1035;S1041;S1057;T1072;S1080;S1098;S1177	34887587	MARRSQSSSQGDNPLAPGYLPPHYKEYYRLAVDALTEGGPEAYNRFLASEGAPDFLCPEELEHVSRHLQPPQYVAREPPEGTPSDVDMDGSSGTYWPVNSDQAVPELDLGWPLTFGFQGTEVTTLVQPPPPDSPSIKDEARRMIRSAQQVVAVVMDMFTDVDLLSEVLEAAARRVPVYILLDEMNAQHFLDMADKCRVNLHHVDFLRVRTVAGPTYYCRTGKSFKGHLKEKFLLVDCAVVMSGSYSFMWSFEKIHRSLAHVFQGELVSSFDEEFRILFAQSEPLVPSAGALARMDAYALAPYSGAGPLVGVPGVGAPTPFSFPKRAHLLFPPPREEGLGFPSFLDPDRHFLSAFRREELQRMPGGALEPHTGLRPLARPTEAGPFGELAGPRGFFQSRHLEMDAFKRHSYATPDGAGAVENFAAARQVSRQTFLSHGDDFRFQTSHFQRDQLYQQHYQWDPQFAPARPQGLFEKLRAGRPGFADPDDFALGAGHRFPELGADVHQRLEYVPSSASREVRHGSDPAFGPSPRGLEPSGASRPNLGQRFPCQATLRQGLDTASEAEPERRGGPEGRAGLRHWRLASYLSGCHGDGGEEGLPMEAEACEDEVLAPGGRDLLPSAFRTPAAFPAKGPKPGSGSGGGDSSEREGPEETSLAKQDSFRSRLNPLIQRSSRLRSSLIFASQAEGAVGTAAATTEKVQLMHKEQTVSETLGPSGEAVRSSASAKVAELLEKYKGPARDPGGAGGAVTSSSHSKAVVSQAWREEVVAPGGAGTERRSLESCLLDLRDSFAQQLHQEAERHPGAASLTAAQLLDTLGGTDRLPSRFLSAQGRSLSPQGRDSPPPEGLGTHQLPYSEPKGNPTPAYPERKGSPTPAYPERKGSPTPAYPERKGSPTQAYPERKGSPTSGFPNRRGSPTTGLMEQKGSPTSTYPDRRGSPVPPVPERRGSPVPPVPERRGSLTFAGESSKTGPTEEVSSGPMEVLRKGSLRLRQLLSPKNERRGEDEGSFPTPQENGQPESPRRPSLSRGDSTEAAAEERGSRVRLASATANALYSSNLRDDTKAILEQISAHGQKHRGVPAPGPAHSSPDVGRPTTAGDLAPDMSDKDKCSAIFRSDSLGTQGRLSRTLPGSAEERDRLLRRMESMRKEKRVYSRFEVFCKKDEAGSSGAGDNLADEDTRDSKMGKFVPKILGTFKSKK	None	None	None	None	None	None	None	None	None