

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
Q9WU40	MAN1_MOUSE	Mus musculus	Inner nuclear membrane protein Man1	38.939326	S28;S296;S298;S311	S28;S136;S137;S140;S261;S263;S287;S412;S787;T893;S921	40997131;36852467	MAAATAAAAPQQLSDEELFSQLRRY GLSPGPVTESTRPVLYKKLKKLREEE QQQQQQQQQQHRRAGGRGNKTRN SNNNTATAMGGRPGSGDLAYLRS PAGLGRLSASAAESPVAGGSGGAAA VPAAGSKVLLGFSSDESDEASPRE QAGGGGGGARRDRAALQYRGLRA PPAPPAAGEVTGGHPGERRKPWSW WGARRPAGPEPQPPAAGSDGAAEDA DEELADGEDRDPEAEPLWASRAV NGSRLLPYSSCREHYSDSEEEEEEG EEDGDVAPARQVLKDDSLARHRPRR SHSKPFSALTAKSGGSRQETSVOGG GALAMNDRAAAGSLDRSRNLEEA AAEPGGGGGGCGDPVDSIPRYRA GAKKLAPLLSPPSPDGDSTLESPTGP LLKTNHIGGGAFGVDSPGLYANSL PPGATAAAAPGTLRINHANHTGSNH TYLKTAYGKPKLCEPEEELLQQFKRE EVSPTGSFSAHYLSMFLTAACLFFL ILGLTYLGMRTGVPEDGGLKNPFD ETFGKIQESEKNLLMSTLYKLDHRL AQIAGDHECGSSSRMLSVQEAAY LKNLGPEDVFNLSLLWIFKNGKD VGIRCVGYGPEEDLTNITDVQFLQST RPQMFWCRFRRAFITVTHRLLLLC LGVVLCVALRYMRYRWTKEEETR QMYDMVKIIVLVRSHNEACQETKD LQPYMPLPHVRDSLIPQDRKKMKK VWDRAVDFLAANESRVRTETRRVG GADFLVWRWIQPSASCDKTLVIPSK VWQGQAFHLDRRNSPPNSLTPCLKI RNMFDPVMEIGDHWHLAIQEAILEK CSDNDGIVHIAVDRNSREGCVYVKC LSPEYAGKAFKALHGSWFDGKLVTV KYLRLDRYHHRFPQALTCNTPLKPA NKHMNLSHLRLRGLANSQGS	True	False	1.506	5.0	1.075	1.874	1.359	3.3