

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0 Q61830	MRC1_MOUSE	Mus musculus	Macrophage mannose receptor 1	28.83309	NaN	NaN	34289986	MRLLLLFATVSVIPVSVQLLDARQFLI YNEDHKRCVDALSASVQTATCNPE AESQKFRWVSDSQIMSVAFKLCGLGV PSKTDWASVTLYACDSKSEYQKWEC KNDTLFGIKGTELYFNYNRQEKNI KLYKSGSLWSRWKVYGTDDDLCSR GYEAMYSLLGNANGAVCAFPFKFEN KWYADCTSAGRSDGWLWCGITTDY DKDKLFGFCPLHFEGSERLWNKDP LTGILYQINSKALTWHQARASCKQ QNADLLSVTEIHEQMGLTGLTSSLSS GLWIGLNSLSVRSQWQWAGGSPFR YLNWLPGPSSEPGKSCVSLNPGKN AKWENLECVQKLGKYGKGNNTLN PFIIPSASDVPTGCPNQWVWPYAGHC YRIHREEKKIQYALQACRKEGGDLA SIHSIEEFDIFSQLGYEPNDELWIGL NDIKIQMYFEWSDGTPVFTFKWLP EPSHENNRQEDCVVMKGDGYWA DRACEQPLGYICKMVSQSHAVVPEG ADKGCGRKGRHGFYCYLIGSTLST FTDANHTCTNEKAYLTTVEDRYEQ FLTSLVGLRPEKYFWTGLSDVQKNG TFRWTVDEQVQFTHWNADMPGRK AGCVAMKTGVAGGLWDVLSCEEKA KFVCKHWAEGVTRPEPTTTPEPKC PENWGTTSKTSKMCFLYAKGKHEK KTWFESRDFCKAIGGELASIKSKDEQ QVIWRLITSSGSYHELFWLGLTYGSP SEGFTWSDGSPVSYENWAYGEPNN YONVEYCGELKGDPGMSWINDINCE HLNNWICQIQKGTLLPEPTPAPQD NPPVTADGWVYKYDYQYFSKEKET MDNARAFCKNFGDLATIKSESEKK FLWKYINKGGQSPYFIGMLISMDK KFIWMDGSKVDFVAWATGEPNFAN DDENCVTMYTNSGFWINDINCYPN NFCQRHNSINATAMPPTPTTGG CKEGWHLYKNCFKIFGFANEEKKS WQDARQACKGLKGNLVSINAQEQ AFVTYHMRDSTFNAWTGLNDINAE HMFLWTAGQGVHYTNWKGYPGG RRSSLSEADACVVVIGGNSREAGT WMDDTCDKQGYICQTQTDPSLPVS PTTTPKDGFTYKSSYSMLKLLP WHEAETYCKDHTSLASILDYPSNA FAWMKMHFPNVPWIALNSNLN EYTWDRWRVRYTNWGADEPKLKS ACVYMDVDGYWRTSYCNESFYFLCK KSDEIPATEPPQLPGKCPSEQTAWI PFYGHCHYFESSFTRSWGQASLECL RMGASLVSIETAAESSFLSYRVEPLK SKTNFWIGMFRNVEGKWLWLN PVSFVNWKTGDPGERNDCVLASS SGLWNNIHCSSYKGFICKMPKIIPV TTHSSITTKADQRKMDPQPKGSSKA AGVTVVLLVIGAGVAAFFYKRRH ALHIPQEATFENTLYFNSNLSPGTS TKDLMGNIEQNEHAI	False	True	2.803	3.206	2.757	2.298	1.511	4.794	3.338